

**MECHANISM AND GENETICS OF GRAIN MOLD
RESISTANCE IN SORGHUM**

THESIS

Submitted to the
Dr. Panjabrao Deshmukh Krishi Vidyapeeth, Akola
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IN
AGRICULTURE
(PLANT PATHOLOGY)**

BY
GULAB DAULATRAO AGARKAR

DEPARTMENT OF PLANT PATHOLOGY
POST GRADUATE INSTITUTE,

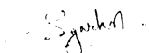
**Dr. PANJABRAO DESHMUKH
KRISHI VIDYAPEETH, AKOLA (M. S.)
INDIA**

DECLARATION OF STUDENT

I hereby declare that the experimental work and its interpretation of thesis entitled "Mechanism and genetics of grain mold resistance in sorghum" or part there of has not been submitted for any other degree or diploma of any University nor the data have been derived from any thesis/ publication of any University or Scientific Organization. The sources of materials used and all assistance received during the course of investigation have been duly acknowledged.

Akola

Dated :


G.D. AGARKAR
Enrolment No. AGNG/1/78/69

CERTIFICATE

This is to certify that the thesis entitled "**Mechanism and genetics of grain mold resistance in sorghum**" submitted in partial fulfillment of the requirements for the degree of Doctor of Philosophy (Ph. D.) in Agriculture of Dr. Panjabrao Deshmukh Krishi Vidyapeeth, Akola is a record of bonafide research work carried out by **Shri. Gulab Daulatrao Agarkar**, under my guidance and supervision. The subject of the thesis has been approved by the Students Advisory Committee.

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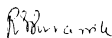


Dr. R. B. Somani

Chairman,

Advisory Committee

Courtersigned



Associate Dean,

Post Graduate Institute

Dr. Panjabrao Deshmukh Krishi

Vidyapeeth, Akola

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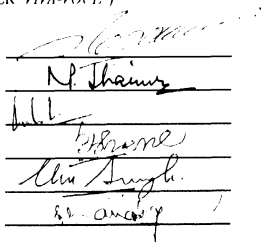
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External Member (Dr. K. H. Jadhav)



Handwritten signatures of the members of the Students Advisory Committee and the External Examiner, each on a line.

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Place : Akola

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ABBREVIATIONS USED

Abbreviated forms/symbol	Terms/unit
1	2
AFP	Antifungal protein
am	Ante meridian
CD	Criticle difference
CE	Catechin equivalent
Cl	<i>Curvularia lunata</i>
cm	Centimeter(s)
cv(s)	Cultivars
dSm ⁻¹	desi Simen
DAA	Days after anthesis
DAF	Days after flowering
ed.	editor(s)
e.g.,	for example
et al.	and others
°C	Degree centigrade
F ₁	First filial generation
F ₂	Second filial generation
f.sp.	Formae specialis
FAA	Formalin Acetic-acid Alcohol
F.p.	<i>Fusarium pallidoroseum</i>
g	gram(s)
gca	general combining ability
ha	hectare
h	hour(s)
pH	H ion conc.
H ₁	Heterosis (over mid parent)
H ₂	Heterobeltiosis (over better parent)
i.e.	that is
<i>in vitro</i>	in glass
kg/cm ⁻²	Kilogram per centimeter square
L	Linieus
l	litre
min	minutes
mol wt	molecular weight
max	maximum
min	minimum
M	molar
ml	millilitre
mg	milligram
µm	micrometer
nm	nanometer (10 ⁻⁹ meter)

N	normal
no	number
PA	Phenolic acid
PC	Phenolic compound
p.m.	post meridian
%	Percent
sca	specific combining ability
SE	standard error
sp.	species plural
vol	volume
v/v	volume/volume
var	variety
wt	weight
vivavoce	orally
vs.	against
viz.	Namely
%	per cent
+	plus minus
<	is less than
≥	is more than or equal than
=	is equal to plus or minus
x	is multiplied by

CHAPTER – I

INTRODUCTION

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CHAPTER – I

INTRODUCTION

Sorghum (*Sorghum bicolor* (L.) Moench) is the fifth major cereal crop in the world after wheat (*Triticum* spp.), rice (*Oryzae* spp.), maize (*Zea mays*) and barley (*Hordeum vulgre*). It is cultivated widely throughout tropical, sub-tropical and temperate regions within latitudes of 45°N and 45°S and mainly in Africa, Asia, North and South America.

The countries of Semi Arid Tropics (SAT) accounts 83% (about 36 million ha) of the total world area sown (43 million ha) under sorghum. In India the crop is grown on 12.5 million ha which accounts for 35% of the sorghum area in the SAT. The SAT countries produce on an average about 63% of world sorghum, of which Indian contribution accounts for 21% (FAO, 1995).

In India, sorghum is grown in areas receiving 500 to 1000 mm annual precipitation with temperature ranging between 26 and 32°C. Plain and plateau below 1000 m elevation offer excellent scope for successful cultivation of the crop in two seasons viz., 'Kharif' (June to October) as rainfed crop and 'Rabi' (October to February) with protective irrigation constituting 60 and 40 percent cultivation respectively. The main areas of sorghum cultivation in India are in states of Maharashtra, Karnataka, Madhya Pradesh, Andhra Pradesh, and Rajasthan. It is mostly used as staple food, feed and forage in the states of Maharashtra, Northern Karnataka and part of Madhya Pradesh.

Sorghums in the tropics, have to sustain a hostile environment where unreliable rainfall, poor soils, pests, diseases and parasitic weeds all constantly exert a harsh selection pressure. The traditional cultivars are photo-sensitive and when rains cease earlier, there is poor grain filling which results in low yields. Therefore, to have cultivars of short duration, photo in-sensitive and good grain filling and drought tolerant is the primary objective of breeders. Several early maturing and high yielding hybrids have been developed which have replaced the traditional varieties to a large extent. However, these cultivars, lack the inherent grain mold escape mechanism of

the local sorghums and the grains that mature during the wet weather are always vulnerable to infection by several fungi. The grain mold problem has achieved a greater significance and because of this, control of grain mold has become a major activity in many sorghum improvement programs.

Definitions of 'grain mold' (GM) found in recent literature appear to fit into one of the two general concepts of fungal- related grain deterioration. The first concept describes a condition resulting from fungal infection and colonization of grain, occurring any time between anthesis and harvest. Here GM can be broadly defined as a fungal component of pre-harvest grain deterioration, involving numerous fungal species interacting in different ways with the plant (i.e. parasitically and/or saprophytically).

The second concept restricts the definition of GM to a condition caused by infection and colonization of spikelet tissues prior to grain maturity. In this limited definition, few fungi are thought to be involved. The magnitude of field fungi that colonize grain after physiological maturity are not part of GM *per-se*, but rather constitute a component of weathering, or general post harvest grain deterioration.

On practical level, the two concepts are similar. Early and late infections in first concept can be seen as analogous to the GM and weathering of second concept (Forbes *et al.*, 1992). Various terms (e.g. grain molds, seed molds, grain deterioration, grain weathering, head molds and head blight) have been used in literature to describe the association between deteriorated grain of sorghum and numerous fungal species (Castor, 1981; Williams and Rao, 1981). Fungal related grain deterioration, whether occurring before or after grain maturity, can cause economic losses in several ways viz, a) moldy and discolored pericarp, b) a soft and chalky endosperm, c) decreased grain filling and size (low yields), d) reduced germination, e) presence of mycotoxins, f) decreased dry matter, density and test weight, g) altered composition of grain, h) low storability, i) low nutritive value, j) low acceptability by the consumers and farmers and k) low market price.

The mold may develop in sorghum inflorescence at any stage from the young inflorescence to the mature head, provided that climatic conditions are suitably

humid. Generally, it seems that wet weather following flowering is necessary for grain mold development and longer the wet period greater the mold development (Rao and Williams, 1977). Dry weather during flowering and grain development followed by wet weather near maturity will not promote such serious mold as when the wet weather occurs from the time of flowering onward.

Few fungi infect sorghum spikelet tissues during early stages of grain development. These are (in approximate order of importance) *Fusarium moniliforme* Sheld., *Curvularia lunata* (Wakker) Boedijn, *Fusarium pallidoroseum* (Cooke) Sacc. (*F. semitectum* Berg., and Rave.), and *Phoma sorghina* (Sacc.). *F. moniliforme* and *C. lunata*, and they are significant worldwide (Castor, 1981; Frederiksen *et al.* 1982; Williams and Rao, 1981; Bandyopadhyay, 1986). Several fungi belonging to about 21 genera have been reported so far with sorghum grain by various workers.

In Vidarbha region of Maharashtra, the most common grain mold fungi in relative order of frequencies are *C. lunata* (40-60%), *F. moniliforme* (15-20%), *F. pallidoroseum*, *Drechslera* spp., *Phoma sorghina*, *Alternaria* spp. and *Aspergillus* spp. (5-30%) (Anon., 1984).

C. lunata and *F. moniliforme* secrete amylases, cellulases and pectinases resulting in disintegration of endosperm and germ tissue. These fungi also interfere with carbohydrate translocation to developing kernels causing reduction in size and weight of the kernels, causing physical, physiological and chemical changes. Endosperm of molded grain appears chalky because of partial hydrolysis of starch and protein. Molded grain may be contaminated with mycotoxins and present health hazards to consumers (Castor and Frederiksen, 1980).

In general, avoidance and/or sowing GM resistant cultivars are the only practical and economical methods for control of GM in sorghum. Chemical control of GM is usually impractical and too costly.

Several characteristics contribute resistance to field deterioration of grain viz., loose heads, seed completely enclosed in glumes (Murty, 1975), colored grain with high tannins and presence of pigmented testa (Harris and Burns, 1973). However, some white-grained varieties and relatively small glumes and even a

cultivar with compact panicle have been reported to be less susceptible (Williams and Rao, 1978). Thin mesocarp, rate of water absorption and conductivity of seed leachates (Glueck *et al.*, 1977), hardness of seed (Rana *et al.*, 1978), endosperm texture, pericarp thickness, surface wax and grain integrity (Glueck and Rooney, 1977), flavan-4-ols and hardness of grain (Jambunathan *et al.*, 1991) and phenolic acid (Hahn and Rooney, 1986) are known to influence GM resistance. Presence of phenolic compounds (PC) and phenolic acid (PA) in mature caryopsis of sorghum in wet or dry environment (Waniska *et al.*, 1989), stele layer and its thickness and electrical conductivity of grain leachates (Somani, 1992) are also reported to play roles in resistance. It is apparent that several factors, independently or in combination, contribute to GM resistance. The most important are tannins, flavan-4-ols, and phenolic acids, type of proteins and grain hardness.

The recently improved early-maturing cultivars have higher harvest index and give stable and high yields under favorable environment, but when they flower fill grains and often mature in wet weather, it results in:

- i) increased susceptibility to parasitic and saprophytic fungi that destroy the grain,
- ii) loss of seed viability and sprouting on the panicle, and
- iii) poor food quality.

Since farmers preferences depend upon the consumption value of the grain and its market price, grain deterioration problem becomes crucial for the extension and adoption of high yielding cultivars.

The high yielding, white-grained cultivars developed recently do not possess sufficient levels of resistance to grain molds. Colored-grained sorghum germplasm lines with high level of mold resistance have been used for developing white-grained genotypes with good levels of resistance in elite materials (varieties) of good grain quality and yield. Mold resistance in white-grained types is associated with grain hardness, while that of brown-grained types with either high tannins or flavan-4-ols or grain hardness. Cultivars with a combination of these factors are highly resistant. It appears that flavan-4-ols is not produced in white-grained types,

either mold-resistant or mold-susceptible. At present the only factor known to be responsible for mold resistance in white-grained cultivars is grain hardness. Intensification of efforts to breed mold resistance into high-yielding cultivars is in order, so that farmers can efficiently grow mold-free sorghums (Mukuru, 1992). Due to grain mold problem, seed production of the hybrids that are grown in Maharashtra state is being taken up in southern States. The Maharashtra State Government is thus emphasizing priority to grain mold research in the state.

In view of the above, and the recommendations made at the International workshop held at Harare, Zimbabwe in March, 1988 grain mold has been identified as number one problem of sorghum in India. The present investigation, undertaken at Sorghum Research Unit, Dr. Panjabrao Deshmukh Krishi Vidyapeeth (Dr PDKV), Akola (M.S.) during 1994-1997 and at International Crops Research Institute for the Semi-Arid Tropics (ICRISAT), Patancheru, Andhra Pradesh during 1995-1997 had the following objectives.

1. To determine the infection sites and colonization of *F.moniliforme*, *F. pallidorseum* and *C. lunata* at different stages of host maturity.
2. To understand mold resistance heritability using a simple 10 x 10 diallel among resistant and susceptible lines.
3. To study the physical characters viz. glume color, glume covering, grain hardness, electrical conductivity of grain leachates, endosperm texture, pericarp color and thickness, presence of testa layer in relation to mechanisms of resistance to grain mold.
4. To study the biochemical parameters viz., i) proteins ii) soluble sugar iii) tannins iv) flavan-4-ols v) protein fractions viz., albumin and globulin, prolamins, cross-link prolamins, glutelin-like, glutelin and residues in relation to biochemical mechanism of host resistance.

CHAPTER – II

REVIEW OF LITERATURE

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CHAPTER II

REVIEW OF LITERATURE

2.1 Predisposition

It is well recognized that predisposition of sorghum panicles to wet and humid weather from flowering to grain maturity period favors infection by grain mold fungi . (Tarr ,1962; Balasubramanian, 1977; Gray *et al.* 1971; Koteswara Rao and Poornachandrudu, 1977; Williams and Rao, 1978; Gangadharan *et al.*, 1978) reported that the wet weather and heavy rainfall stimulate the development of molds at all the stages from the emergence of ears to the ripening of grains, wet weather conditions have a greater bearing on the extent of moldiness. A highly significant correlation exist between the percent of mold grains in earhead and rainfall.

Siddiqui and Khan (1973) reported that grain maturity and not the flowering stage, must coincide with rains for grain mold development, other factors being normal. Under favorable high moisture conditions mature sorghum grains are invaded by *Alternaria*, *Cladosporium*, *Phoma* and *Fusarium semitectum* (Seitz *et al.*, 1975).

Dry weather during flowering and grain development followed by wet weather at maturity does not promote serious mold of wet weather that continues from the time of flowering onwards (Williams and Rao, 1978).

Screening of sorghum lines for grain mold resistance under field conditions has successfully been done by spraying panicles with water (Anahosur, 1983; Deshmukh, 1989) or by providing sprinkler irrigation on rainfree days (Butler and Bandyopadhyay, 1990).

A good level of grain mold development was recorded at Hyderabad, Coimbatore, Dharwad and Baraut in years when rainfall was high during flowering to grain maturity period (Indira *et al.*, 1991). Somani (1992) also highlighted the importance of rainfall in grain mold development.

2.2 Screening/Identification of grain mold resistance

Screening methods used by different workers and sources of resistant/tolerant to grain mold are summarised below:

Reference	Lines	Remarks
1	2	3
Kulkarni et al. (1975)	Selections H-142, H-143, and H-145 from a cross CSV-4 (resistant) x H-112 (susceptible)	-
Glueck (1977)	E-35-1, IS-2327 and IS-2328 (less susceptible)	relatively small glumes
Gangadharan et al. (1976)	K-3 (resistant), CSV-4 (CS-3541), CO-21, CO-22, SPV-34, SPV-181 and IS-3880 (moderately resistant)	-
Anahosur and Patil (1982)	IS-14332, IS-3443 and IS-2328	-
Rao and Rao (1982)	IS(s)-14332 (completely free), 2328, E-35-1, M(s)-38776, 6090, 90324, 61743, 62467, 62522 and 64083 (resistant), SPV-104 and CSH-1 (susceptible). SPV-126, M-90894, E-35-1, SPV(s)-315, 312, 371 and 311 (resistant), 138, SPV-247, CS-3541, CSV-8R, and M-90253 (moderately resistant)	- Field & laboratory ratings

Contd..

1	2	3
Rao et al. (1984)	CS-3541 (resistant) CSV-5, SPV(s)-35,81,102,126,141 and 249 (moderately resistant) IS(s)-3927,9327,9333 and IS-9530 (resistant)	ICRISAT genetic stock
Chandrasekaran et al. (1985)	SPV(s)-126,346,544,617 and 679 and IS(s) 6265, 8283 and 14332 (promising)	TGMR, disease intensity and germination
Shrotria et al. (1986)	IS(s) 4006,5959,6047,6335,7237, 8131,2930,13798,13804,13598 and PAB-105	Field and laboratory test.
Stenhouse et al. (1990)	IS(s)-9470,15119 (moderately resistant) IS(s)9470, 25077,23585,1815,10696, 10942 and 20884 (resistant)	Male sterility
Anahosur (1992)	SPV(s)-126,312,346,351,386,472, 2219B,DMS-1B,MR(s)-750,849, IS(s)-3443,3547,14332,10892, 14372,14380,22995, and 24996 (resistant)	Screening methods, field grade, TGMR, ger- mination, no loss in 100 grain mass

Contd..

Contd..

1	2	3
	IS(s)-14375,14380,24995, 24996 and 108922	Brown sorghums
Rao et al. (1995)	IS(s)-7173,23773,23783 and 34219 (resistant)	<i>in vitro</i> with <i>F.monili-</i> <i>forme</i> <i>C.lunata</i> <i>F.pallido-</i> <i>roseum</i>

2.3 Effect of grain mold on seed weight

Infection by *F. moniliforme* and *C. lunata* has been reported to interfere with carbohydrate translocation to developing kernels, and thus causing reduction in size and weight of seed (Bhatnagar, 1971; Gray *et al.*, 1971; Mathur *et al.*, 1975; Castor and Frederiksen, 1977).

Significant grain weight losses (40-70%) due to infection by several grain mold fungi have been reported (Gray *et al.*, 1971; Sundaram *et al.*, 1972; Glueck and Rooney, 1976; Castor, 1977; Singh and Agrawal, 1989).

Forbes, *et al.* (1989) compared severity with loss in grain weight, the standard deviation of grain weight, grain density, electrolyte leachate, percentage germination and visual appraisal of moldy, off colored or smaller grain. Grain density and grain weight were less closely associated with severity.

Somani (1992) recorded reduction in test weight by 53,51,24 and 9% in combine inoculations (*F. moniliforme* and *C. lunata*), 46,40,20 and 8% due to *C. lunata* and 42,25,7 and 5% due to *F. moniliforme* when inoculated at 10,20,30 and 40 DAF in 296-B cultivar. Martinez *et al.* (1994) reported decrease in test weight and percentage of seed germination and molding was higher in the white cultivars than in red or brown ones.

2.4 Effect of grain hardness on mold development

Plant breeders and technologists have used the terms "hardness" or "vitreousness" to describe the endosperm textural characteristics of sorghum grain. Although a substantial number of reports have dealt with techniques to measure hardness of wheat grains (Obuchowski and Bushuk, 1980) only limited data available on hardness of sorghum grain (Rooney and Sullins, 1969; Maxson *et al.*, 1971).

Grain hardness in sorghum is contributed by several factors, such as grain shape and size, thickness of pericarp, the adherence of pericarp to endosperm and starch-protein interactions (Greenwell and Schofield, 1986). Abdelrahman and Hosney (1984) reported that cross-link prolamin may be advantageous as it confers hardness to grain.

Rana *et al.* (1977) reported that tan plant type having grains with lower water absorption capacity and higher grain hardness could contribute to mold resistant cultivars. Glueck and Rooney (1980) opined that corneous endosperm texture and more epicuticular wax contributed to increased weathering resistance. Rana *et al.* (1984) reported hardness and rate of absorption of water are predominant additive character offering resistance to grain deterioration.

Mukuru (1988,1992) reported that mold resistance in the white-grain advanced selection was associated with grain hardness. Resistance in red-grain types was associated with flavan-4-ols and grain hardness, while that of brown grain types was associated with either high tannins or flavan-4-ols or grain hardness. Cultivars with combination of these factors are highly resistant. At present the only factor known to be responsible for mold resistance in white-grained cultivars is grain hardness.

In white sorghum without test, grain hardness contributes positively to mold resistance (Bandyopadhyay, 1988; Stenhouse *et al.*, 1990).

Ratios of hardness to grain mass in white-red and brown grained sorghums suggested, that resistance in white and red grains might be due to hardness. Correlations between hardness, grain mass and threshed grain mold ratings (TGMR) generally supported the conclusions (Reddy *et al.* 1991). It was further pointed out that, the flavan-4-ols and hardness were inherited as dominant traits and these together were responsible for resistance in red grained hybrids. Mold resistance in white grained sorghum genotypes without testa could be attributed to their having harder grains than mold susceptible genotypes. Association of both flavan-4-ols and grain hardness with resistance to grain mold is reported. Grain hardness is governed by prolamin content.

Kumari *et al.* (1992) reported that hard grains showed less incidence of grain molds than soft grains during development. Microscopic examination showed more intense deposition of protein bodies in hard than in the soft grains. The presence of fungal hyphae in the endosperm of soft grains and pitted starch granules was clearly visible microscopically. Extract of immature and mature hard and soft endosperm were inhibitory to *F. moniliforme* growth. These inhibitors were heat labile and non-dialyzable indicating that protein factors may be involved. The activities of inhibitors to serine proteases were comparatively higher in endosperm of hard grain during development. The endosperm of hard grain contained more protein and prolamin than that of soft grains.

Mukuru (1992) suggests that for sorghum lines to be mold-resistant their grain must remain hard and vitreous in the field under wet and warm conditions - the ideal environment for grain mold development. Sorghum grains with hard and vitreous endosperm are also known to be least susceptible to storage weevils. However, it was reported that grains with a harder, more vitreous endosperm were less digestible than those having soft endosperm.

2.5 Endosperm texture in relation to grain mold

The relative proportion of the corneous to floury endosperm within a sorghum kernel is often referred to as endosperm texture. Texture can be determined by a visual examination of longitudinal half kernel. The rating ranges from 1 to 5, where 1 means very little floury endosperm (<20%) almost completely corneous and 5 rating means essentially all floury (>80%) endosperm. Munck (1981) reported that determination of percent of soft (or 'floury') portion in kernels of sorghum is another measure of hardness.

Kirleis *et al.* (1984) developed a method for measuring endosperm texture by quantitatively determining the corneous and floury areas of sectioned sorghum grain using a light microscope. The corneous endosperm characteristics are not necessary for resistance to weathering; however, if all other things were equal, a line with more corneous grain would resist deterioration more than a floury endosperm line, because of the more dense structure and organization (Clark *et al.*, 1973; Ellis, 1972, 1975; Garud, 1992).

Glueck *et al.* (1977) and Glueck and Rooney (1980) used several lines that had consistently ranked high among the most resistant to weathering at all locations in several years of testing, and concluded that grains with more corneous endosperms were more likely to resist deterioration than floury endosperm lines. However, Mansuetus (1990) reported that endosperm texture had no effect on grain mold score, but in all cases the presence of the testa decreased mold incidence.

Somani (1992) reported that corneousness of endosperm contributes towards resistance in white grain genotypes. He used iodine vaporization method. Hardness grades were attributed on the basis of lower absorption corneous endosperm do not take much color.

2.6 Electrical conductivity of grain leachates and grain mold development

Glueck *et al.* (1977) suggest several possible mechanisms for resistance to grain deterioration, including rate of water absorption and conductivity of seed leachates and these tests may be useful as preliminary screening method. Glueck and Rooney (1980) reported that cultivars with rapid rate of water uptake exhibited less resistance

to weathering. Composition of leachates from these cultivars was richer in nutrients. A thicker mesocarp and softer endosperm texture usually corresponded to increased water absorption and richer leachates. Forbes (1986) reported that automatic measuring of seed leachates and its correlation with seed germination could become an efficient technique for studying the effect of GM severity on viability. Grain mold severity, caused by *F. moniliforme* was positively correlated with electrolyte leachates (Forbes et al., 1989).

Somani (1992) also reported that electrical conductivity of seed leachates was more in susceptible cultivars.

2.7 Glume color and glume covering

Murty (1975) noticed *Curvularia* infection on the portion of seed not covered by glume and concluded that open heads with seeds completely enclosed in long papery glumes are relatively resistant to field deterioration. Gangadharan et al. (1978) noticed that fully enclosed grain by glume and loose heads are the factors associated with resistance. Glueck and Rooney (1980) reported that panicle shape, glume characters, wet season avoidance, seed size contribute to increased resistance to grain molding. Longer glumes are considered protective to the grain, but only if the glumes do not trap water. Narayana and Prasad (1980) suggested besides permeability of grain, hard grain, papery pericarp as reported by various workers, the glume characters, viz. glume permeability, tight/loose attachment of glume to seed were also other possible factors and structural and chemical composition of seed also contribute to resistance. Studies of Mansuetus, et al. (1988) showed that disease incidence was negatively associated with glume cover ($r=-0.56$), glume length (-0.56) and glume area (-0.62) at the boot stage. Somani (1992) also reported that covered kernel with compact glumes showed less grain molds.

2.8 Pericarp/Mesocarp

Swanson and Hunter (1936) pointed out that the great discrepancy between laboratory and field germination of better known sorghum varieties was due to the relative thickness of starchy layer of cells located in mesocarp. When seeds soaked in water for 2 hours, the varieties having thick mesocarp (70-80 μm) absorbed about 33% more

moisture than the varieties which had thin mesocarp (20-50 μm). Glueck and Rooney (1977) found that water enters through the pericarp, especially through hilum area. Long enveloping glumes does not necessarily protect the grain from weathering, open glume types are probably more vulnerable to sprouting on the head, pigmentation in pericarp and glume is not necessarily related to resistance, though it may impart slight degree of resistance. They also reported that thin mesocarp sorghum withstand weathering better than those with thick mesocarp and that quantity of surface wax was probably not a factor since the most susceptible lines had adequate or greater quantities of surface wax when compared to resistant lines. However, alteration in the distribution of wax on the surface of pericarp may affect water uptake by the grain. A thin pericarp on sorghum caryopsis normally corresponds to less weathering. Glueck and Rooney (1980) reported that at physiological maturity, colonies of fungi was observed in starchy mesocarp and the cross and tube cells of the pericarp in all cultivars. Since fungal colonies were observed inside the pericarp are readily hydrolyzed by saprophytic fungi. Hence, a thick mesocarp that contains starch and protein support more fungal colonies than sorghum with thin mesocarp. The mesocarp is thin when the Z gene is dominant (Z-) and thick when the gene is recessive (zz). Also most of free phenolic compounds are located in pericarp and the adjoining testa layer. These apparently bioactive compounds would be diluted with starch etc, in a thick pericarp.

Narayana and Prasad (1980) suggested besides permeability of grain, hard grain, papery pericarp as reported by various workers, the glume characters, viz. glume permeability, tight / loose attachment of glume to seed were also other possible factors and structural and chemical composition of seed also contribute to resistance.

Castor (1981) from his histopathological studies reported that mesocarp provides an ideal environment for early colonization and a jumping - off point for fungi to continue the deterioration on the grain after the infection has taken place. Miller (1981) noticed that thick mesocarp types are susceptible to grain mold. Bandyopadhyay *et al.* (1988) when screened 26564 selected accessions from the world collection of sorghum germplasm in field during the 1980-85 rainy seasons

found that all resistant accessions except one, had colored pericarp. Waniska *et al.* (1992) observed that sorghum cultivars that exhibit some resistance to deterioration have thin pericarp besides corneous endosperm and specific phenolics. The red pericarp trait also conferred grain mold resistance to certain extent. The effect of a red pericarp was enhanced by the presence of a intensifier gene. The effect of both pigmented testa and red pericarp were additive. Mesocarp thickness did not play a significant role in grain mold resistance.

2.9 Loss of seed viability and germination

Tarr (1962) reported that *Aspergillus*, *Fusarium* and *Rhizoctonia* species were responsible for poor emergence of sorghum seedlings, since these fungi destroy starchy endosperm of seed and thus deprive young seedling of it's food. Arif and Ahmed (1969) found that *Fusarium* spp. were the most harmful, followed by *Aspergillus*, *Penicillium* and *Helminthosporium* in reducing germination. Narasimhan and Rangaswamy (1969) found a viability reduction of 40 to 80% when healthy sorghum seeds were treated with mold isolates. Bhatnagar (1971) and Castor (1977) noticed that fungus infected seeds often exhibited a reduction in germination and emergence, which caused poor stand in the farmers fields.

Tripathi (1974) obtained 56% germination with moldy sorghum grains, whereas apparently clean grains gave 76% germination. Further, they reported reduced germination by (42%) due to *Colletotrichum graminicola* followed by *C. lunata* (40%), *F. moniliforme* (37%), *Phoma insidiosa* (26%), *Penicillium* spp. (23%) and *Aspergillus flavus* (18%). Castor (1977) reported reduction of germination from 95 to 77% with grain harvested from *Fusarium* spp. inoculated heads. Rao and Williams (1977) obtained viability loss upto 100% in sorghum grains with severe *Fusarium* and *Curvularia* infection. Denis and Girard (1977) and Castor and Frederiksen (1980) considered, loss in viability to be so important part of grain mold, that they recommend a germination test as part of standard evaluation for identification of grain mold resistance. Munghate (1980) recorded 0.75 to 25.25% and 0.75 to 5.30% loss in germination due to *C. lunata* and *F. moniliforme*, respectively. Bhale and Khare (1982) observed pre- and post-emergence mortality due to seed borne *C. lunata* and

F. moniliforme in water agar seedling symptom test method. Vidyasekaran (1983) and Granja and Zambolim (1984) recorded low germination of sorghum seeds due to severe infection of *F. moniliforme*.

Forbes (1986) stated that severity appears to be more closely associated with seed viability than with yield, two measures of severity (ergosterol concentration and propagules of *F. moniliforme* g-1 seed tissue) were highly correlated with percent germination than with seed mass or grain density. Automatic measuring of seed leachates and correlation with germination could become an efficient technique for studying the effect of GM severity on viability. Deshmukh (1989) reported that *C. lunata*, *F. moniliforme* and *Exserohilum halodes* caused considerable reduction in germination. Singh and Agrawal (1989) noted that seedling from infected seeds were less vigorous than those from healthy seeds. Wu and Cheng (1990) reported that the incidence of *C. lunata* in blotter test was negatively correlated with seed vigour. The incidence of three pathogens (*C. lunata*, *F. moniliforme* and *P. sorghina*) in blotter test was significantly correlated with abnormal seedlings in seedling evaluation, seedling growth and accelerated aging tests.

2.10 Seed mycoflora

Seed mycoflora of sorghum varies from region to region. Frequently occurring fungi associated with moldy sorghum grain mold complex reported by different workers throughout the world is summarised below:

Location	Fungi detected	Infection	
		frequency(%)	Reference
1	2	3	4
Georgia	<i>Colletotrichum graminicola</i> ,	-	Luttrell
	<i>Curvularia</i> spp., <i>Fusarium</i> spp.,		(1950)
	and <i>Penicillium</i> spp.		

Contd..

Contd..

1	2	3	4
Kansas, USA	<i>Alternaria tenuis</i> , <i>Curvularia</i> spp., <i>Fusarium</i> spp. and <i>Aspergillus</i> spp.	-	Swarup (1955)
India	<i>Curvularia</i> spp., <i>F. moniliforme</i> , and <i>F. oxysporum</i> ,	-	Mathur et al. (1967)
Edembergh	<i>Ascochyta sorghi</i> <i>F. moniliforme</i> , <i>C. lunata</i> , <i>C. graminicola</i> , <i>Drechslera</i> spp., and <i>Peronosclerospora sorghi</i> .	-	Noble and Richardson (1968)
Maharashtra	<i>Aspergillus niger</i> , <i>C. lunata</i> , <i>F. moniliforme</i> , <i>F. semitectum</i> , <i>Helminthosporium</i> <i>rostratum</i> and <i>H. tetramera</i> <i>F. moniliforme</i>	-	Bhagwat and Pedgaonkar (1973)
	<i>C. lunata</i>	42%	Tripathi (1974)
	<i>F. moniliforme</i>	19%	
	<i>A. flavus</i> -	15%	
	<i>Phoma insidiosa</i>	13%	

Contd..

Contd..

1	2	3	4
India	<i>F. moniliforme</i> and <i>F. semitectum</i> <i>C. lunata</i> <i>A. tenuis</i> <i>F. moniliforme</i> <i>F. roseum</i> <i>A. niger</i> <i>F. oxysporum</i>	- 35% 31.2% 10.0% 12.5% 3.1% 2.5%	Mathur et al. (1975) Khare et al. (1976)
Madhya Pradesh	<i>C. lunata</i> , <i>Alternaria</i> spp., <i>Fusarium</i> spp. and <i>Verticillium</i> spp.	-	Sharma et al. (1976)
USA	<i>Fusarium</i> and <i>Curvularia</i>	-	Castor 7)
USA	<i>C. lunata</i> and <i>F. moniliforme</i>	-	Castor and Frederiksen (1977)
India	<i>Curvularia</i> , <i>Fusarium</i> and <i>Phoma</i>	-	Reddy and Reddy(1977)
USA	<i>Alternaria</i> spp., <i>C. lunata</i> <i>F. moniliforme</i> , <i>Fusarium</i> spp., <i>H. tetramera</i> and <i>Phoma</i> spp.	-	Castor and Frederiksen (1980)
India	<i>Curvularia</i> spp and <i>Fusarium</i> spp.	-	Williams and Rao (1978)

Contd..

Contd..

1	2	3	4
Brazil	<i>Alternaria alternata</i> and <i>Curvularia</i> spp.	-	Pinherio et al. (1979)
Karnataka	<i>A. alternata</i> , <i>C. lunata</i> <i>Drechslera specifera</i> , <i>F.</i> <i>moniliforme</i> , <i>F. semitectum</i> , <i>Phoma sorghina</i> and <i>Trichothecium roseum</i>	-	Anahosur and Hegde (1980)
USA	<i>Gibberella fujikuroi</i>	-	Castor and Frederiksen (1981)
India	<i>Curvularia</i> spp and <i>Fusarium</i> spp.		Williams and Rao (1981)
Madhya Pradesh	<i>C. lunata</i>	2-49%	Bhale and Khare (1982)
	<i>F. moniliforme</i>	4-30%	
	<i>P. insidiosa</i>	2-5%	
Philippines	<i>F. moniliforme</i> and <i>C. lunata</i>	-	Dayan and Dalmacio (1982)
Taiwan	<i>F. moniliforme</i> , <i>C. lunata</i> , and <i>Phoma sorghina</i>	-	Wu (1983)
India	<i>F. moniliforme</i> , <i>F. oxysporum</i> , <i>F. semitectum</i> and <i>F. solani</i>	-	Gopinath and Shetty (1985)

Contd..

Contd..

1	2	3	4
India	<i>C. lunata</i>	27.7%	Singh and Agrawal (1987)
	<i>F. moniliforme</i>	9.4%	
	<i>P. sorghina</i>	8.4%	
	<i>C. lunata</i> , <i>Drechslera sorghicola</i>	-	Gupta and Singh (1988)
	<i>F. moniliforme</i> and <i>Phoma</i> spp.	-	
China	<i>F. moniliforme</i> and <i>F. oxysporum</i>		Liang and Bai (1988)
Vidarbha	<i>C. lunata</i>	62%	Deshmukh (1989)
	<i>F. moniliforme</i>	26%	
India	<i>C. lunata</i> , <i>F. moniliforme</i> and	-	Singh and Agrawal (1989)
	<i>Phoma sorghina</i>		
Taiwan	<i>C. lunata</i> , <i>Drechslera maydis</i> and <i>F. moniliforme</i>	-	Wu and Cheng (1990)
Thailand	<i>C. lunata</i> , <i>Fusarium</i> spp.,	-	Boon-Long (1992)
	<i>Colletotrichum</i> spp. and		
			Contd..

Contd..

1	2	3	4
	<i>Phoma</i> spp.		
Central America Caribbean Basin	<i>F. moniliforme</i> and <i>C. lunata</i>	-	Wall and Meckenstock (1992)
Vidarbha	<i>C. lunata</i> and <i>F. moniliforme</i>	44.6% 40.8%	Somani (1992)

Biochemical parameters

2.11 Proteins (protein fractions)

Sorghum proteins varied in their properties and amino acid composition. It has also been reported that protein content has been negatively related to lysine content. (Virupaksha and Sastry, 1968). Deosthale *et al.* (1972) reported that sorghum genotypes, growing environment and nitrogen fertilization influenced the protein content in sorghum. Gupta and Gupta (1974) suggested that the quality of protein depends on amino acid composition and proportion of various classes of proteins. They also observed that in sorghum levels of prolamin and glutelin progressively increased from early milky stage to maturity stage. There was progressive increase in the fraction IV (glutelin like) from 7 to 35 days of maturity period, which then stabilized until maturity. Insoluble protein contents in the residues were high during the first 7 days and decreased until 21 days, with very little change thereafter.

Jambunathan *et al.* (1975) and Guiragossian *et al.* (1978) reported distribution of nitrogen in different fractions of proteins from the grains of IS-11167 and IS-11758. Glueck *et al.* (1977) reported that the proteins are hydrolyzed and partially used in the synthesis of fungal protein and remain in the moldy grain. Guiragossian *et al.* (1978) observed that mutation in P-721 sorghum decreased quantity of kafirin (Prolamin) with an increase in the albumin-globulin fraction. Rooney and Miller (1982) reported that the

matrix protein is comprised mainly of **glutelins** (alkali-soluble proteins) and **prolamins**. (**alcohol-soluble protein**). The prolamin, exists in small spheres called protein bodies. Physical and chemical characteristics of the grain from different sorghum varieties influenced solubility's and chemical scores of protein fractions (Neucere and Sumrell, 1979). Nwasike *et al.* (1979) observed that, in general cross-link prolamin is high in sorghum, as compared to pearl millet and corn. Jambunathan *et al.* (1983) reported that protein content in 146 grain samples obtained from the ICRISAT breeding program, lysine, normal sorghum grains and their progenies, ranged from 7.1 to 19.1%.

Subramanian *et al.* (1983) reported 8.8 to 13.2% with mean value 10.56% protein content in 18 brown sorghum genotypes. They also reported protein content in 8 grain samples comprising land races, hybrids and local cultivars, ranged from 6.8 to 19.6%. Also observed that protein content of developing grain was high 7 and 14 days after anthesis, and a decline was observed toward maturity. Cross link prolamin may be advantageous as it confers hardness to the grain (Abdelrahman and Hosene, 1984). Subramanian and Jambunathan (1984) estimated protein content in sorghum germplasm accessions that varied from 4.4% to 21.1% with mean value of 11.4%. Van Scoyoc *et al.* (1988) reported that the percentage distribution of fraction I, expressed as percent of total nitrogen, declined rapidly and steadily in developing sorghum grain. However, when expressed as N content/endosperm, the patterns of nitrogen per endosperm for fraction I were very different from those expressed as a percentage of total protein. Prolamin synthesis was low during first seven days after anthesis (DAA), reached its maximum from 14 to 28 days and declined thereafter. Synthesis of cross-linked prolamin steadily increased until maturity. This suggests that prolamin was synthesized in different proportion during different stages of grain development. Onset of active synthesis of glutelin was observed from 21 days after anthesis, with little change until maturity.

Subramanian ^A_L *et al.* (1990) studied the distribution of protein fractions in grains revealed that, fraction I, comprising albumin – globulin including non-protein nitrogen, and fraction V (glutelin) together constitute about 41.55% of the protein in eight sorghum cultivars. Variation in fraction II (prolamin) fraction III (cross-linked prolamin) contents was observed among the cultivars. To elucidate the pattern of synthesis of protein fraction in

grain, studies were made at different grain maturity periods, using one cultivar. Fraction I, synthesis was initiated at seven DAA. Prolamin increased from 14 to 28 days and declined toward maturity. Glutelin did not change beyond 14 days and declined toward maturity. Glutelin did not change beyond 14 day after anthesis until maturity.

Indira *et al.* (1991) noticed that grain hardness is generally governed by prolamin content. Desai *et al.* (1994) reported grain protein content from 8.94% to 12.38%, averaging 10.63% in 13 sorghum cultivars comprises of land race hybrids, local cultivars. Kumari and Chandrashekar (1992) reported the endosperm of hard grains contained more protein and prolamin than that of soft grains. Somani *et al.* (1993) in analysis of grain revealed that the crude protein content was slightly reduced in the discolored grain. The amino acid spectrum was also changed in these grains and it is suggested that this may be due by hydrolysis of weakened protein matrix.

Seetharaman *et al.* (1994) reported the presence of antifungal proteins in several sorghum cultivars using antibodies raised against zeamatin, barley chitinase and bean chitinase. Seetharaman *et al.* (1996) reported changes in sorghum caryopsis antifungal proteins (AFP) in different tissues and during development imbibition, and germination, Somatin, chitinase and glucanase levels increased during caryopsis development and ^{highest} at physiological maturity (30 DAA). Ribosome inactivating protein levels were higher at 15 DAA and decreased subsequently. Somatin and chitinase levels were significantly different between sorghum cultivars. Somatin content at physiological maturity correlated with mold rating ($R(2) = 0.65$). Seed AFPs were present in endosperm and migrated toward the extension of caryopsis upon imbibition. AFPs leached out of immature seeds but were retained in the pericarp of mature seeds. Levels of these proteins also changed significantly during seed germination and were present in the shoots of germinating caryopsis.

2.12 Soluble sugars

Edwards and Curtis (1943) analysed 26 sorghum samples for soluble sugars and the content varied from 0.81 to 1.59%. Sorghum seed with sugary endosperms have been reported to contain at least twice the quantity of sugars that normal seed contain (Karper and Quinby, 1963).

Glueck *et al.* (1977) reported that in deteriorated grain the soluble carbohydrates are usually decreased as they are used to provide energy for the growth and development of the fungi. Neucere and Sumrell (1980) reported that free sugar content in five varieties of sorghum varied from 2.34 to 6.01%. Subramanian *et al.* (1980) reported the total sugar content of the 10 sorghum cultivars varied from 1.30 to 5.19%. The high lysine Ethiopian lines, IS-11163 and IS-11758 had higher sugar content i.e. 5.19 and 4.43 % respectively. They also estimated that sucrose is the predominant sugar in the sorghum grain. The proportion of sucrose ranged from 68.7 to 82.7% of soluble sugars in the sorghum cultivars. Subramanian *et al.* (1983) reported the soluble sugar content of the 18 brown sorghum genotypes range^d from 1.1. to 2.5%.

2.13 Tannins and Flavan-4-ols

Swain and Bate-Smith (1962) defined tannins as phenolic compounds (PC) having molecular weights between 500 and 3000. Depending on their molecular structure, tannins are customarily divided into hydrolyzable and condensed tannins. Hydrolyzable tannins are complex molecules containing ester type linkages, which yield on hydrolysis, a sugar and phenol residue consisting ester type linkages. These consist of either gallic or ellagic acid, which is dimer of the former. Condensed tannins are formed by the polymerization of molecular units having the general structure of flavanoids, the most important of which are flavan-3-ols (catechin) and flavan-3, 4-diols (leucoanthocyanidins). Relatively little is known about the occurrence and distribution of proanthocyanidins and leucoanthocyanidins in sorghum tissues. As a result of injury or physiological stress, sorghum leaf tissue frequently develop red coloration due to anthocyanidins. Leucoanthocyanidins could be precursors of these pigments. Harris (1969) reported that brown-seeded hybrids have higher tannins levels than red or yellow seeded hybrids. Weinges *et al.* (1969) and Watterson and Butler (1983) noticed that certain monomeric flavanols, such as flavan-3,4 -diols and flavan-4-ols can give rise to anthocyanidins and therefore these can be distinguished from the oligomeric flavan-3-ols by the name "leucoanthocyanidins".

Maxson *et al.* (1972) stated that kernels with a testa or a coloured pericarp should be tested for tannins and those with values lower than 0.05 catechin equivalents/g of sorghum be selected. McMillan *et al.* (1972) mentioned that tannins present in the grains of

certain varieties resist bird depredation. Bullard and Elias (1980) reported that resistance to bird depredation is a complex phenomenon that may be associated with non-tannin polyphenols, as well as tannins.

Harris and Burns (1973) reported that sorghum seed tannins content was strongly and negatively correlated with pre harvested seed molding indices. Brown seed with high tannin contents and the presence of pigmented testa are relatively resistant to field deterioration (Ellis, 1972; Murty, 1975). High tannin sorghums tend to be less digestible and nutritionally inferior to sorghums in which tannin is absent or present at low levels (Maxson *et al.*, 1973; Jambunathan and Mertz, 1973; Mabbayard and Tipton, 1975) reported that pericarp color may not be a reliable indicator for tannin concentration.

Price *et al.* (1979) and Hagerman and Butler (1981) noticed that tannins binds certain proteins very strongly and thereby diminish the digestibility and nutritional value of high tannin sorghum grains. Most cultivars with pigmented testa containing polymer of flavan-3-ols (tannins) resist weathering. (Glueck and Rooney, 1980; Hahn *et al.*, 1983; Bandyopadhyay *et al.*, 1988). Hagerman and Butler (1980) noticed tannin associated proteins consists of three major components, two of which are high molecular weight prolamins and one of these was quite rich in proline. Butler (1981,1982) reported that the most important factor controlling the affinity of protein for sorghum tannin, was the amount of proline that it contains. The affinity of protein for tannin can be predicated with reasonable accuracy from its proline content. Also reported that flavan-4-ols monomers may contribute to bird repellency of high-tannin sorghums. Rooney and Miller (1981) pointed that phenolic compounds in sorghum caryopsis improve resistance to birds, insects and molds as well as pre harvest germination.

Hahn *et al.* (1983) separated PA and identified eight acids from sorghum extracts by HPLC, their concentration and whether bound or free differed with cultivar. Sorghum grain resistant to fungal attack contained both a greater variety and larger amounts of identified PA and unidentified compounds. Resistant cultivars had more PA in free form. Subramanian *et al.* (1983) reported tannin content (catechin equivalents) in 18 brown sorghum genotypes which ranged from 0.13 to 7.22 CE %. They reported that variation in tannin content was much larger than variation in the other constituents. Detailed polyphenol

analysis on selected genotypes indicated that some lines had insignificant levels of condensed tannins, that none of them was a group II sorghum and that the levels of flavan-4-ols were relatively high. Cultivars with red pericarp (with or without pigmented testa) containing significant levels of flavan-4-ols exhibit resistance to weathering. Mukuru (1992) found that cultivars with white pericarp without pigmented testa do not contain significant levels of flavan-4-ols, yet some of these cultivars exhibit resistance to weathering. Apparently, flavan-4-ols and related compounds are involved in some way with resistance to grain weathering. Hahn *et al.* (1984) concluded that phenolic compounds i.e. phenolic acids, flavanoides, anthocyanidins and tannins are located primarily in the pericarp and testa layers of the sorghum caryopsis. Ring (1984) detected phenolic compounds in leaves and glumes of sorghum. Jambunathan *et al.* (1986) analysed polyphenol concentrations in grain, leaf and callus tissues of mold susceptible and mold resistant sorghum cultivars and reported that the level of flavan-4-ols were two to three-fold higher in mold-resistant cultivars than in mold-susceptible cultivars. Doherty *et al.* (1987) and Forbes (1986) reported that free PC and tannin contents in caryopses increase significantly during development, reaching maximum levels 7-18 DAA. At maximum, the levels of free PC and tannins were two to eight times higher than those observed in mature grain. PC and tannins are apparently being bound to cellular tissues, and therefore are not extractable for analysis. The high level of free PC and tannins occurred during the period of early invasion and colonization of fungi of the caryopsis. Hence, it is likely that these compounds are involved in the resistant mechanisms of grain weathering. Bandyopadhyay *et al.* (1988) reported 24 grain mold resistant accessions with colored pericarp had negligible amounts of tannin (less than 1.0 CE%) and 14 of the 24 lacked the testa layer. The range of tannin content in the resistant accessions was 0.1-10.7 CE%. Mansuetus *et al.* (1988) observed that resistant cultivars had higher free PC content in their glumes and mature caryopses, and showed greater increase in these compounds in response to infection than susceptible cultivars. Measurements of parahydroxybenzoic, coumaric, vanillic and gentisic acid contents indicated that phenolic-bound acids provide the plant with back-up defence to *F. moniliforme* when free PA have been depleted from the caryopses.

Forbes *et al.* (1989) reported that resistant cultivars also respond more quickly to fungal invasion via increased levels of PC and pigmentation of spikelet tissues than do susceptible cultivars. Waniska *et al.* (1989) quantified PC and PA in mature caryopses of sorghum grown in wet or dry environments. Seventeen cultivars varying in pericarp color and presence of pigmented testa exhibited different degree of resistance to molding in wet environment. Sorghum caryopses with white pericarp had lower free PC contents (14 micro g/ caryopses) than those with red pericarp (41 micro g / caryopses) when grown in the dry environment. This difference diminished under humid conditions. Cultivars with a pigmented testa were more resistant to grain mold, had higher free PC content (151 micro g/ caryopses), and had softer endosperm texture than cultivars without a pigmented testa. In cultivars without pigmented testa, higher free PC and free PA contents especially free P-coumaric, ferulic and caffeic acids, were observed in mold susceptible cultivars. A scatter plot of free PC vs. free P-coumaric acids indicated that mold susceptibility was related to higher levels of P-coumaric acid, regardless of environment. Jambunathan *et al.* (1990) analysed methanol and acidified-methanol extracts of grains harvested at different developmental stages for flavan-4-ols and reported that the concentration of flavan-4-ols in mold-resistant grains were at least 2-fold higher than in mold-susceptible grains in both extract at or after 30 days flowering (DAF). Concentration of flavan-4-ols in mature grains could, therefore, be an indicator of their potential resistance or susceptibility to grain mold, and this method could be an important tool in screening sorghum cultivars for such characteristics.

Jambunathan and Kherdekar (1991) reported that methanol and acidified methanol extract of leaves of mold resistant accessions contained at least 3-fold higher concentrations of flavan-4-ols than susceptible accessions at 56, 63 and 70 DAF. The concentration of flavan-4-ols was monitored in the flag leaves of mold resistant accession that had no testa at 77, 84, 91 and 98 DAF, and it decreased sharply at or after 70 DAF. The estimation of concentration of flavan-4-ols in sorghum leaves, therefore offer scope for screening sorghum accessions for their grain mold resistance. Jambunathan *et al.* (1991) reported that ergosterol concentration increased with increasing DAF in the mold susceptible accessions and was 10-fold higher in grains collected at 50 DAF than in the corresponding mold-resistant accessions. It is suggested that ergosterol concentration could be used to

assess the magnitude of mold damage in sorghum grains. The correlation coefficient between ergosterol and flavan-4-ols concentration was significant ($P < 0.01$) and negative in colored mold-susceptible and mold-resistant accessions that did not have testa, but no significant correlation was observed in white mold resistant and mold-susceptible sorghum. It was concluded that there could be another genetic factor or mechanism besides flavan-4-ols associated with mold resistance in white grained sorghums. Mukuru (1992) reported mold resistance in the white-grained types was associated with grain hardness, while that of brown-grained types was associated with either high-tannin or flavan-4-ols or grain hardness. and also reported that flavan-4-ols is not produced in white-grained types, either mold-resistant or mold-susceptible.

Resistant cultivars respond more quickly than the susceptible cultivars to fungal invasion via increased levels of phenolic compounds in glume tissues. Extensive deterioration does not occur before physiological maturity because the developing grain contains 3 to 10 times the level of specific phenolic compounds of mature grain. In sorghum grown under wet conditions, grain of resistant cultivars contains lower levels of free phenolic compounds at maturity compared with grain of susceptible cultivars, hence phenolic compounds content in grain is a predictor of the sorghum cultivars level of resistance to grain molding Waniska et al. (1992). Martinez et al. (1994) evaluated 9 sorghum cultivars for resistance to *F. moniliforme* [*G. fujikuroi*] in field experiment in Argentina. Three cultivars (SC-630-11E, MF-5107 and MF-5097) were resistant; these cultivars had a higher concentration of flavan-4-ols than the white pericarp cultivars (BTX623, ICSB-34, BArg 34 and Bvar) and less tannin than the control brown pericarp cultivars (B-1509 and MF-5194). Melakeberhan et al. (1996) evaluated 10 sorghum genotypes with differences in phenolic compound concentration and grain mold resistant over three crop seasons (1989, 1990 and 1992) to assess changes in phenolic compounds during seed development and how these changes influence grain molding. Flavan-4-ols concentration were high and similar for both the mold resistant and mold-susceptible genotypes at early stages of seed development. In susceptible genotypes, the flavan-4-ols concentration dropped by 67% between third and last sampling dates compared with a 20% decline for the resistant genotypes in the same period. The results also showed that highest incidence of seed infection by fungi occurred between 25

and 35 days after anthesis. *Alternaria*, *Fusarium* (especially *F. moniliforme*) *Cladosporium*, and *Epicoccum* species were the major fungi isolated from the seeds. Menkir *et al.* (1996) identified sorghum accessions with high level of grain mold resistance, originating from diverse geographical areas and belonging to different botanical races. Resistance to grain mold in these sorghums was strongly associated with high concentration of phenolic compounds (apigeninidin, flavan-4-ols and tannin), kernel hardness and pericarp color. Each of these kernel properties contributed to grain mold resistance differently in white, red and brown pericarp sorghum accessions, respectively.

2.14 Presence of testa layer and resistance to grain mold

Ellis (1972) studied the morphological characters indicating grain mold resistance and reported that pigmented testa was the most influential seed characteristic affecting weathering resistance in the field. Furthermore, within a given genetic background and when the pigmented testa was absent, lines with red or lemon yellow were more resistant than lines with white pericarp to grain mold. The ability of pigmented testa to resist grain mold development is attributed to its high tannin content.

Maxson *et al.* (1972) described the level of tannins and their effect on nutritional value as related to the presence or absence of the testa and spreader in sorghum. Cummings and Axtell (1973) proposed a scheme to classify sorghum in groups I, II and III based on chemical analysis and redefined by Price and Butler (1977). Group I does not have testa, group II has a testa (B_1-B_2-SS), and group III has a testa and spreader (B_1-B_2-S-). Harris and Burns (1973) and Murty (1975) reported that brown seed with high tannin content and the presence of pigmented testa are resistant to field deterioration. Glueck and Rooney (1980) identified that sorghum lines high in tannin content and having testa layer were more resistant to grain mold fungi. The presence or absence of testa layer is controlled by the B_1 and B_2 genes. When the complementary B_1 and B_2 genes are dominant (B_1-B_2-), testa pigmentation is present and when either or both genes are homozygous recessive ($b_1b_1B_2-$, $b_1b_1b_2b_2$), pigmented testa is controlled by another gene (Tp) in which brown is dominant to purple. The spreader gene (S) allows the brown colour of pigmented testa to be present in epicarp ($S-$).

Bandyopadhyay (1986) pointed out that IS-14384 has no testa and no tannin and is promising against grain mold (ICRISAT, 1986). Cultivars with red pericarp (with or without a pigmented testa) containing significant level of flavan-4-ols exhibit resistance to weathering (Jambunathan *et al.*, 1986; Mukuru, 1992).

Jimnez and Valleja (1986) reported that tolerant entries to *C. lunata* were those which had a pigmented testa and floury endosperm. In the presence of testa, texture was not an important trait but in the absence of testa comeousness or floury endosperm showed maximum tolerance than those with intermediate texture. Waniska *et al.* (1989) reported that, PC and PA were quantified in mature caryopses of sorghum grown in wet or dry environments. Seventeen cultivars varying in pericarp color and presence of a pigmented testa exhibited different degree of resistance to molding in the wet environment. Mansuetus (1990) correlated mold incidence with the presence of testa. Eighty percent of the selection were brown grained with testa and with TGMR of 2 or less. Twelve percent were red or white without testa and TGMR of 3 or less indicating the importance of testa in imparting grain mold resistance (Stenhouse *et al.*, 1990). The effect of both pigmented testa and a red pericarp were additive, mesocarp thickness did not play significant role in grain mold resistance.

2.15 Artificial inoculation of sorghum with fungi at different stages of grain development

Castor (1977) inoculated *F. moniliforme*, *F. semitectum*, *C. lunata* *C. protruberata*, *Alternaria* spp. and *Helminthosporium* spp. at various times after flowering. The *Fusarium* and *Curvularia* isolates were the principal pathogens causing discoloration and reduction in seed viability. Seeds from the heads inoculated with *Fusarium* at flowering had the highest proportion of split pericarps. Rao and Williams (1977) reported high levels of grain mold in sorghum, when heads inoculated at anthesis with conidial and mycelial suspensions of *F. moniliforme*, *F. semitectum* and *C. lunata*. Castor and Frederiksen (1980) reported that inoculation (2.2×10^3 conidia/ml) at anthesis or within 2 to 3 days of anthesis resulted in greatest damage, suggesting that floral tissues (glumes, stigmas, styles etc.) are most susceptible at flowering and becomes less susceptible thereafter. They inoculated sorghum lines with *C. lunata* at anthesis and noticed that infection occurs in glumes, lemma, palea and

lodicles in 5 days after anthesis. Colonization of pedicel tissues resulted in kernel abortion or in reduced kernel filling. Narayana and Prasad (1982) inoculated *F. moniliforme* and *C. lunata* singly and in combination in equal proportion at preflowering and post flowering periods. *F. moniliforme* was observed to be more infective at flowering and *C. lunata* at soft dough stage. Combine inoculation was more effective from anthesis to dough stage. Deshpande *et al.* (1985) noticed that artificial inoculation with *C. lunata* not only reduced seed germination and seedling vigour index but increased abnormal seed germination than with *Fusarium*. Bandyopadhyay and Mughogho (1988a) inoculated *F. moniliforme*, *F. pallidoroseum* and *C. lunata* on flowering panicles of six genotypes which were watered with sprinkler on the days of no rainfall from flowering to grain maturity and harvest. They reported that TGMF of susceptible and resistant genotypes were significantly greater and germination of susceptible genotypes lower in plot with sprinkler irrigation than in plot receiving only rainfall.

Singh *et al.* (1993) inoculated with a mixed spore suspension of *F. moniliforme*, *F. pallidoroseum* and *C. lunata* at 50% anthesis on 14 accessions. Genotypes IS 7173, IS 23773, IS 23783 and IS 34219 were completely resistant to mold upto 55 days after inoculation and a further six ^(IS-7326, IS-4943, IS-5326, IS-4011, IS-5292 and IS-27741) developed moderate to low mold growth.

Somani (1992) inoculated *F. moniliforme* and *C. lunata* at 10, 20, 30 and 40 DAF and observed that if the infection takes place at flowering, no grain development takes place and germination increases linearly with the advancement of grain development. *F. moniliforme* infection is usually at or earlier to soft dough stage where as that of *C. lunata* is from dough stage onwards.

2.16 Histopathology of infection by molding fungi

Mathur, *et al.* (1975) noticed embryonic infection by *F. moniliforme* and *F. semitectum*. They could locate deep seated infection of *Alternaria tenuis*, *C. lunata* and *Drechslera sorghicola*.

Castor (1977) observed that the scutellum was often partially degraded in seed from *Fusarium* inoculated heads and suggested that the fungus could destroy the embryo indirectly by interfering with translocation from the endosperm to the embryo during germination. All the grain from susceptible *Fusarium* – inoculated heads contained *Fusarium* in the endosperm and 78% of grain from *Curvularia* – inoculated heads had that fungus in the

endosperm. Rao and Williams (1977) noticed that inoculations at anthesis or within 2-3 days of anthesis resulted in invasion in greatest tissues. That indicates that floral tissues (Glumes, stigmas style etc.) are most susceptible at flowering and becomes less susceptible there after. Abdullah and Dadhum (1978) observed embryonic infection by *Aspergillus fumigatus* and *Gibberella fujikuroi* in sorghum. Castor and Frederiksen (1980) compared seeds from heads inoculated with *F. moniliforme*, *F. semitectum* and *C. lunata* and observed that *F. semitectum* produced minimum discoloration and no degradation of endosperm and germ tissues. *C. lunata* resulted in partial degradation of the endosperm and appeared to progress slowly inward from the pericarp. *F. moniliforme* appeared to colonize the region where hilum, scutellum and endosperm fuses. They also reported that there are some evidences suggesting that *F. moniliforme* growth over the kernel surface occurs after complete colonization of the endosperm tissues. *F. moniliforme* was found initially beneath the glume, around the germ and hilum where as *Curvularia* produced mold on portions of the seed not covered by glumes. Blakely and Castor (1979) reported that fungi enter through hylar and stylar ends and they further stated that the embryo region which is under the hylar areas was damaged the most implying that the hylar area is the most readily accessible to fungi. Castor (1979) isolated fungi from different grain components and noticed that embryo contain less fungi than any other tissue.

Castor (1981) studied histopathology of grain mold and provided that infection take place by relatively few species of fungi during anthesis. Also a number of different tissues appear to be involved in resistance to colonization. Since the infection takes place at such an early stage the presence of a testa probably has little effect on initial colonization. Mesocarp provide an ideal environment for early colonization. Spikelet tissues, including sterile lemma, palea, lodicules, anthers, and filaments are sites of infection at anthesis with *C. lunata* and *F. moniliforme*. The ovary which develops into kernel, is not colonized until it expands between 5 and 10 days after anthesis. Forbes (1986) and Bandyopadhyay (1986) noticed initial infection by *F. moniliforme* on a susceptible cultivar at the apical end on spikelet tissues; lemma, palea, glumes, filaments and senescing styles. Fungus mycelium advances basipetally, either by colonizing spikelet tissues or by growing in voids between these tissue. Early colonization of glumes (3-4 days following inoculation) was found to be

very heavy and caused little cellular disruption or pigmentation in the host. Within 5 days of inoculation, mycelium can be seen in all parts of spikelets, with the denser growth around the ovary base. Lodicules appears to be severe as an important source and always surrounded by dense fungal growth, but extensive colonization of lodicule tissue *per se* has been questioned.

In the next stages of invasion, a dense mycelial mat progresses acropetally between the aleurone layer and the pericarp. Subsequent invasion of the endosperm, embryonic tissues and pericarp originates from this peripheral mat. When environmental conditions are favourable, mycelial growth pushes through the pericarp, producing a white or pink fungal mass which can completely cover the grain.

Infection by *C. lunata* differs from that of *F. moniliforme*, *C. lunata* can infect the apical part of the ovary wall from the colonized lemma, palea, lodicules, filaments, pollen grains and decaying styles, within 5-10 days mycelium penetrates the pericarp and ramifies through the cross and tube cells. Colonization does not usually continue directly into the endosperm, but rather through the placental sac, which can also lead to invasion of the embryo. Gopinath *et al.* (1987) noticed 43% embryonic infection of *F. moniliforme*, while *F. semitectum* and *F. solani* colonized the embryonic tissue in 8 and 5% seed, respectively.

Singh *et al.* (1988) could detect infection by *F. moniliforme* and *C. lunata* after anthesis, while that of *Phoma sorghina*, 8 DAA. Singh and Agarwal (1989) detected *C. lunata* and *F. moniliforme* infection in embryonic tissue and *Phoma sorghina* in the aleurone layer only.

Deshmukh and Raut (1993) microscopically examined the seeds infected by *C. lunata* (*Cochliobolus lunatus*) showed a clear distribution of pathogen in pericarp, seed-coat, aleurone layer, endosperm and embryo. The scutellum showed heavy colonization by mycelium and mycelia were also present in coleoptile, coleorrhiza, and embryonic axis.

2.17 Inheritance/Genetics of resistance

The grain mold problem is so complex that it is almost certain that grain mold resistance is the results of the additive effects of many genes affecting several plant characteristics (Murty *et al.*, 1980). Rana *et al.* (1978) found that water absorption capacity and seed hardness, two factors that have an effect on susceptibility are governed by additive genes. Glueck and

Rooney (1980) reported the inheritance of testa is controlled by complementary B₁ and B₂ testa genes. Dabholkar and Baghel (1980) made 7×7 diallel cross amongst sorghum lines resistant and susceptible to grain mold caused by *Curvularia* and observed both gca and sca components of variance were significant for reaction to grain mold. However, gca was larger in magnitude suggesting predominance of additive genetic variance. Dabholkar and Baghel (1982) reported additive gene action to be predominant in the inheritance of tannins in grain. Dabholkar and Baghel (1983) crossed two varieties of sorghum highly susceptible to grain mold, two moderately resistance and three less susceptible in a diallel fashion excluding reciprocals. Grain mold was developed by artificially inoculating earhead with spore suspension of *C. lunata*. Both additive and non-additive genetic components of variance determined the expression of mold reaction. Genes for resistance were recessive to genes for susceptibility. SPV-29 and 2219 B appeared to possess dominant genes for resistance. Narayana and Prasad (1983) studied a diallel set of crosses involving resistant and susceptible lines to mold for heterosis, combining ability and genetic variation. Additive gene action was found to be predominant in the inheritance of *Fusarium* molds.

Kataria *et al.* (1990) studied mode of resistance to grain molds, largely caused by *C. lunata* (*Cochliobolus lunatus*) and *F. moniliforme* (*Gibberella fujikuroi*) in F₁, F₂ and backcross generations from 3 crosses involving 2 resistant and 2 susceptible lines. Analysis of the results indicated that additive effects were significant in all crosses except IS402 × IS 10892, but were generally smaller in magnitude than the dominance effects. Duplicate type interaction were also important in most crosses.

Esele *et al.* (1993) used four parental cultivars with distinct characteristics and gene markers for caryopsis traits as a base population to generate F₁, F₂ and BC₁ populations. These populations were evaluated for resistance to grain mold. The presence of pigmented testa (B₁-B₂-), a red pericarp (R-Y-) a thin mesocarp (Z-) and an intensifier gene (I-) were all dominantly inherited. A pigmented testa was the single most important trait conferring grain mold resistance. The effect of red pericarp was enhanced by the presence of the intensifier gene. The effect of both a pigmented testa and a red pericarp were additive. Mesocarp thickness did not play a significant role in grain mold resistance. Shivanⁿ₁ *et al.* (1994) indicated that the genetic constitution of the susceptible genotypes is identical, while the

resistant genotypes differ in their genetic constitution. Inheritance of grain mold resistance was governed by 4 independently segregating genes, 2 with complementary interactions and the other 2 with additive interactions. Ghorade (1995) reported predominant additive gene action for characters like 100 seed weight, hardness, grain density, water absorption rate, conductivity of grain leachate, germination, fungal load of *Fusarium* spp., *Curvularia* spp. and other mold spp. in F_1 and F_2 diallel progenies.

2.18 Heterosis

Hybrid vigour was first noticed by Kolreuter (1763) in the interspecies crosses of *Nicotiana* and was studied thereafter by numerous workers. The first clear approach to the concept of heterosis was made by Shull (1904) and he coined the term "Heterosis" to denote the increased vigour, size, fruit fullness, speed of development, resistance to disease and insect-pests or to the climatic adversities. Heterosis (H_1) is expressed over the mid parent value and heterobeltiosis (H_2) is expressed over the superior parents. Heterosis could also serve more practical purpose in providing the breeders a mean of increasing disease resistance in resistance breeding programs.

The terms "heterosis" and "hybrid vigour" are synonymous and it has been suggested by Whaley (1952) that a more precise usage would be to term the superiority of the hybrids as hybrid vigour and to refer the mechanisms by which the superiority is developed as heterosis. Heterosis is the genetic expression of the beneficial effects of hybridization. The presence of heterosis in sorghum has been reported by many workers (Aggarwal and Chavan, 1957; Quinby, 1963; Kambal and Webster, 1966; Rao, 1970a; Tripathi *et al.* 1976; Goyal and Joshi, 1976; Laosuan and Atkins, 1977).

Giriraj and Goud (1981) studied the diallel analysis of eight varieties and reported heterosis to the extent of 28.08% for total plant height. Saradaman (1981) studied the heterosis and found that, in general, all hybrids showed heterosis for plant height and panicle length.

Kanaka (1982) found that heterosis for grain yield ranged from -11.36 to 74.72% and 1000 grain weight from -16.47 to 69.25%.

Kulkarni and Shinde (1983) studied the F_1 generation of seven variety diallel of sorghum and heterosis was found to be very high (65.5%) for grain yield, high for plant height (40.4%) and low for days to 50% bloom (8.0%). Swamlata and Rana (1988) studied six newly developed ms lines and crossed with few restorers and they reported that heterosis over ms line varied from 109 to 147% for yield characters. Over restorers, heterosis was 84% for grain yield and 12% for biological yield.

Ghorade (1995) studied 11x11 diallel cross excluding reciprocals and observed highest estimates of percentage of heterobeltiosis upto - 10.38% for days to 50 % flowering, 64.70% for plant weight, -41.02% for glume coverage, 19.61% for 100 grain weight, 6.29% for grain hardness, 2.76% for germination, -5.44% for conductivity of grain leachates, -5.69% for fungal load of *Fusarium* spp., -12.88% for *Curvularia* spp. and -8.91 for other fungal spp.

2.19 Combining ability

Combining ability can be defined as the relative ability of a biotype to transmit desirable performance to its crosses. It provides a means of understanding the nature of gene action in yield heterosis and is desirable tool to study and compare the performance of a line in hybrid combinations.

According to Sprague and Tatum (1942) the general combining ability denotes the average performance of line in a hybrid combination, while specific combining ability designates those cases in which certain combinations do relatively better or worst than would be expected on the basis of average performance of the lines involved. Srihari and Nagur (1980) observed that the variance due to general combining ability (gca) and specific combining ability (sca) were almost equal with respect to the character like grain weight of ear indicating the presence of both additive and non additive genetic variance of this character. Highest heterosis (121.34%) has been shown by combination (Swarna x CS-3541) associated with highest sca effects (26.38) even though its parents recorded low gca effects for grain weight of ear, indicating that non additive gene action is operating predominantly for expression of this character. Sanghi and Monapora (1981) observed predominance of non additive gene action for plant height. Bijapure *et al.* (1980) observed very high gca under rainfed condition. Indi and Goud (1981) found predominance of additive genetic effects for

plant height and dominance effects predominated for most of the traits. Similarly, over dominance was observed for grain yield and panicle weight. Patidar and Dabholkar (1981) found influence of additive gene effects in inheritance of 1000 seed weight and grain yield in sorghum. However, sca effects were significant for grain yield and number of grains/ear.

Borikar and Bhale (1982) reported more pronounced estimates of sca variances for all characters indicating the preponderance of non additive gene action for yield and yield components. The gca was also considerable for days to flowering and plant height. Kanaka (1982) observed predominance of additive gene action for plant height. He also reported additive and non additive gene actions for days to 50% flowering and panicle length. Karale *et al.* (1984) recorded highly significant gca and sca values for all the characters in *Kharif* and *Rabi* seasons. The gca values were of higher magnitudes than sca variances for days to flowering, plant height, panicle length and 1000 seed weight indicating the importance of additive and additive x additive gene action.

Palaniswamy and Subramanian (1986) reported additive gene action for plant height, grain yield, whereas, non additive gene action for days to bloom and peduncle length. Patil and Thombre (1986) reported the characters plant height, days to 50% flowering, 1000 seed weight under the control of additive gene effects while panicle weight and grain yield under non additive gene action. Chandrashekharappa (1987) reported predominance of additive gene action for plant height, days to 50% flowering (DTF) while non additive gene action for 1000 seed weight. Nimbalkar and Bapat (1987) noticed additive and non additive gene action for DTF and plant height. Shekar *et al.* (1987) recorded 1000 grain weight, days to 50% flowering and plant height predominantly under the control of additive whereas grain yield under non additive gene action. Hugar *et al.* (1988) reported non additive gene action more important for DTF, plant height, grain yield and 1000 grain weight. Patil (1990) reported that the sca effects were more stable than gca for grain yield and 1000 seed weight. Amsalu and Bapat (1990) recorded the additive gene action for DTF.

2.20 Correlation studies

It is a known fact that some plant and grain characters are associated with yield and grain deterioration. If the association is assessed, it is possible to select plants based on yield and grain mold resistance traits. Several workers have found out the characters association in sorghum. Hardness and density correlated inversely with milling yield, and the endosperm texture was negatively correlated with hardness, test weight and kernel density (Maxson *et al.*, 1971). Gangadharan *et al.* (1978) reported a highly significant correlation between the per cent of mold grain in ear head and rainfall. Patel *et al.* (1980) observed positive significant correlation between DTF with 1000 seed weight. Similar results were found by Patil and Thombre (1985). They also observed that the plant height and DTF expressed positive strong correlation. However, Reddy and Nagur (1981) recorded negative correlation. The percentage for kernel floating was correlated with percentage vitreousness, grain hardness determined as a work required for grinding, breaking strength at individual kernel (Hallgren and Murty, 1983). Mansuetus *et al.* (1988) observed that disease incidence was negatively associated with glume cover, glume length and glume area at boot stage.

Forbes *et al.* (1989) reported highly significant correlation ($r=0.94$) amongst two estimates of severity (Ergosterol concentration and colony forming units of *F. moniliforme*). Also reported colony forming units of *F. moniliforme* was positively associated with electrolyte leachate ($r=0.98$), visual assessment as percentage of kernels moldy ($r=0.93$), negative associated with percent germination ($r=-0.98$) and negative association between electrolyte leachate and percentage germination ($r=-0.95$). Indira *et al.* (1991) reported high correlation ($r=0.88$) between September rainfall and mean disease score at four locations. Jambunathan *et al.* (1991) reported significant ($P<0.01$) correlation coefficient between ergosterol and flavan-4-ols concentration and negative in coloured mold-susceptible and mold-resistant accessions that did not have testa, but no significant correlation in white mold-resistant and mold susceptible sorghum. Jambunathan *et al.* (1992) reported negative correlation with ergosterol concentration and grain hardness (Stenvert method). Mukuru (1992) reported significant correlation ($r=-0.66$, $P = 0.01$) between flavan-4-ols and mold resistance. Ghorade (1995) reported that grain hardness exhibited significant positive association with grain density, germination, grain yield, 100 seed weight, and negative

association with water absorption rate and fungal load of all types of fungi. Also reported that germination exhibited positive significant association with 100 grain weight, grain hardness, grain density and negative significant association with conductivity of grain leachate, water absorption rate and all three types of fungal load.

CHAPTER – III

MATERIALS AND METHODS

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CHAPTER III

MATERIALS AND METHODS

3.1 Materials

The experimental materials comprised of 10 parental lines of sorghum (*Sorghum bicolor* (L.) Moench) with different characteristic viz., good grain quality, earliness, adaptation, yield, varying mold reactions, presence of testa, mesocarp thickness and grain hardness. These included three resistant (IS-2284, IS-6335 and IS-9471); five moderately resistant elite (SPV-946, SPV-1201, SRT-26B, GJ-35-15-15 and ICSB-101 B), and two susceptible (SPV-104 and AKms-14B) lines. The pedigree and characteristics of these lines are summarised below:

Characteristics							
Line	Pedigree	Pl ht		TGM R	Testa	Seed	Source
		DTF	cm			hardness kg/cm ²	
SPV-1201	PVK-400×IS-3443-12	73	244	3.0	A	8.67	Dr.PDKV
ICSB-101 B	(Ind.Syn.89-1×RS/R 20-682)-5-1-3	72	167	3.0	A	6.94	ICRISAT
AKms-14B	(MR-807×BTX-678) ×AKms-2B	66	140	4.5	A	6.95	Dr.PDKV
SRT-26B	(M-211×ICSB-27)	70	179	3.0	A	6.56	Dr.PDKV
GJ-35-15-15	(2077A×M-25)× Marvan	78	237	3.0	A	8.11	Dr.PDKV
SPV-946	(SPV-475×SPV-462)	77	264	3.0	A	7.25	Dr.PDKV
SPV-104	(SPV-148×SPV-512)	80	179	4.5	A	6.08	Dr.PDKV
IS-2284	(Haak Doom) Q-2-2-37	70	283	2.0	P	5.91	ICRISAT
IS-6335	(purbi gooseneck banwar)	66	255	2.0	P	6.68	ICRISAT
IS-9471	(South Africa)	72	286	2.0	P	6.10	ICRISAT

A = absent, P = present.

3.2. Methods

3.2.1 Location

The experiments were conducted at two locations viz., Sorghum Research Unit, Dr. PDKV, Akola, latitude 20° 42' N and longitude 77° 02' E with an altitude of 281 m, soil vertisol medium to deep black and ICRISAT, Patancheru. latitude 17°32' N and longitude 78° 16' E with an altitude of 545 m soil-vertisol medium to deep black, 26 km northwest of Hyderabad.

3.2.2 Predisposition

The meteorological data on precipitation number of rainy days, relative humidity and temperature during rainy season 1995, 1996, were collected from Sorghum Research Unit, Dr. PDKV, Akola (Table 1). Since testing at ICRISAT was done under controlled conditions, the meteorological data were been considered Data from 23rd meteorological week to 45 meteorological week the period from flowering to maturity of sorghum crop at both the centres was taken in to consideration

3.2.3 Experimental layout

The experiments were conducted in a completely randomized block design with two replications. Fifty-five treatments (45 test crosses + 10 parents) were included in the experiment. The plot size was double row 3-m long, with row spaced at 45 cm and plants at 10 cm. The trial was sown on 27th June 1995 at Akola and harvested on 26th October 1995.

The experiment was repeated at Akola during the rainy season, 1996. It was sown on 22nd July and harvested on 20th November.

The experiment was also conducted at ICRISAT during the rainy season 1996. It was sown on 17th June and harvested on 21 Oct. The plot size was at double row 4 m long, row spaced at 75 cm and plants within row at 10 cm. At this location overhead sprinkler irrigation was provided from the onset of flowering to grain maturity i.e. black layer formation, to harvest about 2 week after physiological maturity. The plots were sprinkled for 1 hr in the morning if it did not rain the previous night and the same morning and for an additional 1 hr in the evening if it did not rain throughout day. Experimental field with sprinkler irrigation and aerial view of

the experimental field at Patancheru (Plate 1). The seeds of all these trials were used for studying physical, pathological and biochemical characters.

3.2.4 Data recording

Five plants were randomly selected from each replication of a treatment for recording data in F_1 crosses and parents. Fifteen plants were randomly selected per replication in each treatment for recording data in F_2 progenies. Data were recorded on agronomic traits, physical, pathological and biochemical characters. All above characteristics of the kernel of parents and crosses obtained from 1995 trial at Akola and 1996 trials at both locations, whereas, biochemical characteristics could only be studied for 1995 harvest at Akola. All characters were studied using 10×10 diallel analysis, except protein fractions, grain color, glume color pre-treatment of seed and presence of testa.

3.2.4.1 100-seed weight

Hundred kernels weight (g) of each line under various experiments was always recorded after compositing the threshed grain and sun-drying to about 12% moisture, using 'ADCO' precision electronic balance (model AD 200 C) unless specifically mentioned otherwise. All the values reported are means of two observations.

3.2.4.2 Grain hardness

A manually operated 'Kiya' hardness tester (Kiya Seisakusho Ltd., Japan) with bar-type probe (0.5 cm diameter) was used with a maximum permissible load of 20 kg for the determination of hardness of single grain.

Grains were equilibrated to a moisture content $6.5 \pm 1.0\%$ before hardness determination was made. Twenty randomly selected grains from each replications were tested for hardness, mean was calculated and expressed in kg/cm^2 .

3.2.4.3 Endosperm texture

Endosperm texture was determined by visual examination of longitudinal cut half kernels. The rating ranged from 1 to 5 (1=0-20% almost completely corneous and 5 = 81-100% floury). Ten seeds were soaked in water overnight and were cut into two halves to the long axis from the hilar region and endosperm texture was determined

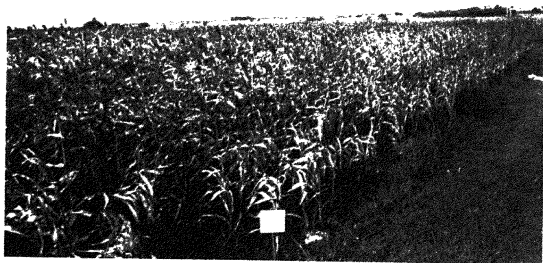
Plate 1 : Experimental field at Patancheru

A. Sprinkler irrigation

B. Aerial view of experimental field



A. Sprinkler irrigation



B. Aerial view of experimental field

Plate 1. Experimental field at Patancheru

using hand lens. The observations were recorded for each replication and mean was calculated.

3.2.4.4 Electrical conductivity of grain leachates

Electrical conductivity of grain leachates was measured using the method of Hendricks and Taylorson (1976) with some modification. Three g seed sample of each line were washed twice with glass distilled water (gdw) and placed in clean 'Coming', glass test tubes and 12 ml of gdw was dispensed to each tube. After four hr of soaking at 25°C the supernatants were used for measuring the electrical conductivity separately. 'YSI model 32' conductance meter of scientific division, Yellow Spring Instrument Co. INC., was used. The electrical conductivity was expressed in dSm^{-1} . The test was run twice.

3.2.4.5 Day's to 50% flowering

The number of days required for flowering of 50% of the plants in a plot from the date of sowing was recorded.

3.2.4.6 Plant height

Plant height (cm) was measured from the ground to the tip of ear head at physiological maturity.

3.2.4.7 Cob length

Cob length (cm) was measured from base to the tip of ear head.

3.2.4.8 Glume (grain) covering

Observation on grain covering i.e. amount of grain covered by the glume for all the lines under study were recorded. The characters were described as follows and shown in (Plate 2A):

1= 0 % grain covered

2=25% grain covered

3=50% grain covered

4=75% grain covered

5=Grain fully covered

6=Glume longer than grain.

Plate 2 : A. Glume covering

- 1 0% grain covering
- 2 25% grain covering
- 3 50% grain covering
- 4 75% grain covering
- 5 100% grain covering
- 6 > 100% grain covering

Plate 2 : B. Infected earheads

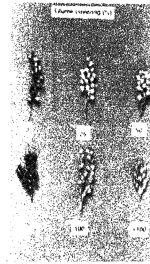
- 1 *F. moniliforme*
- 2 *F. pallidoroseum*
- 3 *C. lunata*
- 4 *Phoma sorghina*

Plate 2 : C. Infected seeds

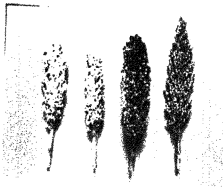
- 1 *F. moniliforme*
- 2 *F. pallidoroseum*
- 3 *C. lunata*

Plate 2 : D. Glume color

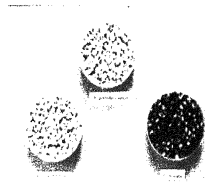
- 1 Tan
- 2 Red
- 3 Purple



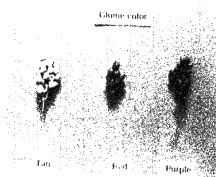
A



B



C



D

Plate 2. Glume covering, infected earhead, infected seed and glume color

3.2.4.9 Pericarp (Mesocarp)

Kernels were soaked in sterile water for 16 hr. The kernels were then removed and surfaces blot-dried. Free hand sections along the short axis with a new razor blade were taken. The sections of each line were selected and placed on clean glass slide separately. Two drop of stain (Mac Grunwald's) used by Scheuring and Rooney (1979) were placed over the sections. After 1 min, when the stain was evaporated the slide was held in slanting position and few drop of ethyl alcohol (95%) were added with the help of dropper for removing excess stain. After the evaporation of ethyl alcohol, the sections were mounted in glycerol with coverslip.

'Filar micrometer' was calibrated using slide micrometer. Sections were observed under low power (10 x objectives and 15 x Filar micrometer) and with help of screw drum, measurement of thickness, in micrometer (μm) of the pericarps were taken and the average of five observations was recorded. Observations for the presence of testa layer (dark brown/red layer adjacent to aleurone layer) were also recorded to confirm the results of bleach test.

3.2.4.10 Threshed grain mold rating (TGMR)

Panicles were harvested 14 days after physiological maturity, threshed and bulked was evaluated for grain mold severity. This is an estimate of the percentage of molded grain surface area and was recorded on a 1-5 scale, where :

- 1= no mold visible on grain surface,
- 2= scant superficial mold growth and 1-10% grain surface covered by mold,
- 3= moderate mold growth and 11-25% of grain surface molded,
- 4= considerable mold growth and 26-50% of grain surface molded, and
- 5= extensive mold growth and >50% grain surface molded.

3.2.4.11 Seed germination

The Ragdoll's (rolled paper towel) method (ISTA, 1966) and standard blotter plate method of (ISTA, 1976) were used for germination studies. In rolled paper towel, 100 seeds were placed at equidistant on two layers of moist paper towel (48x25 cm; 55-58 sheet/kg). Seeded paper towel were incubated for 10 days at room temperature (i.e. $27^{\circ}\text{C} \pm 2^{\circ}\text{C}$). The rolled paper were then unrolled after removing rubber bands gently

upper cover paper was removed and germinated seeds was counted. Unless otherwise mentioned the test was based on 400 seeds/ genotypes.

In standard blotter plate method grain germination was determined by incubating 100 grains from each genotypes in nine cm diameter Petri-plates (25 grains per dish) lined with wet filter paper for four days at $27^{\circ}\text{C} \pm 2$.

3.2.4.12 Detection of grain mold fungi

To detect seed borne mycoflora, standard blotter plate method (SBM) of ISTA (1976) was used throughout the study. Twenty five seeds of each line was placed at equidistant on sterile three layer blotting paper discs presoaked in sterile water in each plate. The seeded plates were kept for incubation at $27^{\circ}\text{C} \pm 2^{\circ}\text{C}$ under diurnal cycle for seven days. Fungi on incubated seeds were detected by stereoscopic binocular microscope with side illumination system. Research microscope was used for detailed observations species were identified by comparing the descriptions given in IMI description plates of fungi and bacteria and from book "Dematiaceous hyphomycetes" by Ellis (1971). Infected earhead and seeds by different mold fungi are shown in (Plate 2B and C).

Observations on colony count of *Fusarium moniliforme*, *Fusarium pallidoroseum* and *Curvularia lunata* separately and for other fungi were recorded in each plate on germinated and ungerminated seeds. The infection score were recorded on a 1-5 scale where:

- 1 = no infection
- 2 = up to 10% seed covered by fungal colony.
- 3 = up to 25% seed covered by fungal colony
- 4 = up to 50% seed covered by fungal colony.
- 5 = 50 % seed covered by fungal colony.

3.2.5 Biochemical characters

3.2.5.1 Proteins and protein fractions

Fifty five grain samples (10 parents and 45 F_1) grown during rainy season 1995 at Akola were used in the study. The grain samples were dried at 37°C for 48 hr and ground in Udy cyclone mill (UD Corporation, Boulder, Co.) to pass through a 0.4 mm

The meal was defatted in Soxhlet apparatus, using n-hexane and used for further analysis. All values reported are means of duplicate analysis.

3.2.5.1.1 Protein

The total nitrogen in the meal was determined by using Technicon auto analyser (TAA) described by Singh and Jambunathan (1980), which involves the conversion of organic nitrogen into ammonia by digesting the sample with sulphuric acid using block digester (BD 40). Ammonical nitrogen reacts with sodium phenate in the presence of sodium hypochlorite to form indo-phenol blue complex. This color complex is measured at 660 nm.

Reagents :-

Acid mixture. Five parts (v/v) orthophosphoric acid in 100 parts of sulphuric acid .

Kjel tabs auto. Each tablet contains 1.5 g K_2SO_4 and 7.5 mg selenium.

Alkaline sodium potassium tartrate. Dissolve 75 g sodium hydroxide and 50g sodium potassium tartrate in about 900ml distilled water, cool dilute to 1l.

Alkaline phenol. Mixed 138 ml of phenol (88%) with 500ml of 5N NaOH in an ice bath ; dilute to 1l with distilled water .

Sodium hypochlorite (NaOCl). Dilute, if necessary, commercially available bleach to get 5% NaOCl .

Ammonium sulphate standards. Dissolve 4.717g oven dried ammonium sulphate in 1000 ml distilled water (1000 ppm N stock solution). Take stock solution to give 10,20,30,40 and 50 ppm N into a 1000 ml volumetric flask and dilute to 1l by adding 4% sulphuric acid.

One hundred mg defatted flour was transferred to a Technicon digestion tube (75ml). In each set of 40 tubes one blank, one check, two random samples selected from preceding set, and 36 regular (unknown) samples were taken and 3 ml of acid mixture and 1 Kjel tab to each tube were added. Then the set was heated in Block digester at 375°C, and maintained for 1 to 1.5 hr for digestion, digest were cooled and dissolved with minimum amount of water. Volume was made up to 75 ml, and the solution was mixed thoroughly after putting the stopper. The aliquot

from each tube were transferred in to Technicon sample cup and analysed it using TAA.

Nitrogen in % was calculated by using formula

$$N\% = \frac{\text{Sample peak height} \times 75 \times 100}{1.8 (\text{slope}) \times 1000 \times \text{sample weight (mg)}}$$

Where, 5 is the made up volume and 1.8 is the net division on chart paper for 10 ppm N. The crude protein was calculated by using the factor ($N\% \times 6.25$).

3.2.5.1.2 Protein fractionation

Protein fractionation of all the parents (10) and F_1 crosses (20) obtained at Akola during 1995 season was carried out. For protein fractionation, crosses were selected in such a way that each cross had one resistant (IS-9471), one elite (SPV-946) and one susceptible (SPV-104) line. Landry and Moreaux (1970) method was used to obtain fractions of protein. The residue after fraction V was treated with 0.1 N sodium hydroxide solution and protein soluble in the alkali was referred to as fraction VI or residual protein. The detailed protein fractions I-VI were as follows :

Fraction	Protein class	Solvent system (reagents)	Extraction time (min)
I	Albumin, globulin free peptide	0.5 M NaCl	60,30,30
II	Prolamin	70% isopropanol	30,30
III	Cross link prolamin	70% isopropanol with 0.6% 2 ME ^a	30,30
IV	Glutelinlike	Borate buffer (pH ¹⁰)	30,30,15
V	Glutelin	Borate buffer (pH ¹⁰) 06% 2ME ^a with 0.5% SDS ^b	30,30,15,
VI	Residual protein	0.1 N NaOH	30,30,15

^a = 2 Mercapto-ethanol ^b = Sodium dodecyl sulphate.

Reagents :

0.5M NaCl , Dissolve 29.2 g Sodium chloride in water and make upto 1l, **70% Isopropanol**, Mix 700 ml isopropanol with 300 ml water, **Isopropanol-2-mercapto-ethanol**, Mix 700 ml isopropanol and 6 ml 2-mercaptoethanol with 294 ml of water, **Borate buffer (pH ¹⁰)**, Dissolve 6.2 g boric acid, 29.2 g sodium chloride and 6 ml 2 mercapto ethanol in water, Adjust pH to 10.0 and finally make up to 1l, **Detergent-borate buffer (pH¹⁰)**, Dissolve 62. g boric acid 29.2 g sodium chloride, 6 ml 2 – mercapto ethanol and 5 g sodium dedocyl sulphate in water. Adjust pH to 10 and make to 1 l with water, **0.1 N NaOH**, Dissolve 4 g NaOH in water make up to 1 l with water.

Extraction

- Defatted flour (1g) was put in to 50 ml screw cap centrifuge tube to which 20 ml 0.5 m sodium chloride solution were added and shaken for 1 hr. After centrifuge at 2500 rpm for 15 min the clear supernatant were collected in to 50 ml volumetric flask. Extraction repeated twice with 15 and 10 ml of the solvent for 30 min each, the aliquots were pooled and made upto 50 ml and filtered through Whatman No.1 filter paper (Fraction I). The extract were stored at 4°C.
- To the residue 20 ml 70% isopropanol were added and shaken for 30 min, the contents centrifuged, the extraction repeated twice of the residue with 15 and 10 ml solvent. The aliquots were pooled and made upto 50 ml and filter (Fraction II).
- Proceed similarly with the residue by using 70% isopropanol containing 0.6% ME (Fraction III).
- To the residue 20 ml borate buffer were added and shaken for 30 min, the contents centrifuged, and supernatant collected. Re-extracted the residue twice with borate buffer for 30 and 15 min and pooled the supernatant and volume were made upto 50 ml with borate buffer (Fraction IV).

- The residue were extracted with 20ml borate buffer containing 0.5% sodium dodecyl sulphate for 30 min and re-extracted twice referred above. The extract were pooled and volume made to 50 ml with borate buffer containing SDS (Fraction V).
- The residue were treated with 20 ml. 0.1 N sodium hydroxide solution (Fraction VI).

Estimation :

Ten ml of fraction (I to VI) were pipetted out in to Technicon digestion tube and few boiling chips were added to it. One hundred mg original flour (taken for protein fractionation) were added into Technicon digestion tube and 3 ml concentration sulphuric acid containing 5% orthophosphoric acid, digestion tablet containing 1.5g potassium sulphate and 7.5 mg selenium were added into each tube. Both flour sample and fractions were digested separately at 380°C for 1 hr. The digest were cooled for 5 min, then dissolved it in water (minimum quantity) and made upto 75 ml and mixed well.

Estimated N content as given in protein. Calculated mg nitrogen N in the 10 ml aliquot and expressed the 'N' at each fraction as % of the flour 'N'.

$$\text{Nitrogen in the fraction} = \frac{\text{mg N in 50 ml of the fraction}}{\text{mg N present in flour}} \times \frac{100\%}{100\%}$$

3.2.5.2 Soluble sugars

Soluble sugars were extracted with hot aqueous – ethyl alcohol and the sugars on treatment with phenol sulphuric acid produces a stable and sensitive golden yellow color (Dubois, *et al.*, 1956).

80% ethyl alcohol (ethanol). 800 ml of ethanol in 1 l. volumetric flask and volume was make with distilled water.

5% phenol. Dissolved 5 g phenol in water and made up to 100 ml with water.

96% sulphuric acid (v/v). Use 96% sulphuric acid, specific gravity 1.84, diluted according to the purity.

Glucose (w/v) standard. (Stock = 1000 mg/ 1000 ml.) Dissolved 1000 mg of glucose in water and made up to 1 l.

Working standard. Pipette out 10 ml of stock standard in to a 100 ml volumetric flask and made up volume to 100 ml (the final concentration will be 100 µg/ml)

Defatted flour (100 mg) was placed in to a boiling tube to which 30 ml of hot 80% ethanol was added and shaken on a vortex mixture. After settling the material for 20-30 min, the supernatant was filtered through Whatman No. 41 filter paper in a beaker. For complete extraction of sugars the above step was repeated for 3-4 times. The extract was then placed on hot sand bath until the ethanol is evaporated.

Ten ml water were added to dissolve the contents and transferred in to a 100 ml volumetric flask, the contents in the beaker were washed 2-3 times and added to volumetric flask and volume was made to 100 ml with water one ml aliquot and 1 ml water as blank were taken into test tubes separately and 1 ml 5% phenol was added in each tube and shaken, then 5 ml 96% sulphuric acid was added and shaken vigorously on a vortex mixture and then test tubes were cooled in water.

The absorbance of the golden yellow were read at 490 nm against the blank. The standard were run with different concentration (i.e. 10, 20, 30, 40 and 50 µg of glucose standard) from the working standard, keeping the volume to 1 ml with water, added reagents 1 ml 5% phenol and 5 ml 96% sulphuric acid.

Total soluble sugar (%)=

$$\frac{\text{Conc. of standard } (\mu\text{g})}{\text{Absorbance of standard}} \times \frac{\text{Absorbance for 1 ml sample extract}}{\text{Absorbance of standard}} \times \frac{1 \text{ (Conversion of g)}}{1,000,000} \times \frac{100 \text{ ml (vol. make up)}}{0.1\text{g (Sample weight)}} \times 100$$

3.2.5.3 Tannins

Price et al. (1978) Vanillin – HCL method was used for estimating tannins. Vanillin with tannin on acid catalyzation results in formation of an color complex which absorbs maximally at 500 nm. The reaction is specific for leucoanthocyanidins and

catechins although, anthocynidin and dihydrochiones may interfere to some extent, expressed as catechin equivalents (CE).

Reagents

- **8% hydrochloric acid in methanol (v/v).** Mixed 8 ml. conc. HCL in methanol and made upto 100 ml.
- **1% Vanillin.** Dissolved 1 g vanillin (Sigma) in methanol and made upto 100 ml.
- **Vanillin – Hydrochloric acid reagent.** Mixed equal volume solution 1 and solution 2 before use.
- **Hydrochloric acid in methanol (4%) (v/v).** Mixed 4 ml conc. HCL in 100 ml. methanol.
- **1% Hydrochloric acid in methanol (v/v).** Mixed 1 ml conc. HCL. with 99 ml. methanol.
- **Standard solution.** Stock solution, 1 mg/ml of catechin (Sigma) in methanol (0.1 ml) is equivalent to 10 µg catechin/ml). (stock solution can be stored for long time in stopper bottles in cold). Pipette out 0.1, 0.2, 0.4, 0.6, 0.8 and 1 ml. of catechin standard and made up the volume to 1 ml with methanol.

Extraction

Defatted flour (0.5g) was taken in centrifuge tube to which 10 ml acid methanol (1%) was added in each tube and shaken for 20 min in shaker and centrifuged for 10 min and aliquot was transferred to 20 ml volumetric flask. Again 5 ml acid methanol (1%) was added and shaken for 20 min and centrifuged for 10 min and aliquot was added to first extraction and the volume was made to 20 ml, and mixed well.

Estimation

One ml of extract was pipetted out in to test tubes, to which 5 ml freshly prepared Vanillin-HCL reagent was added slowly to the extract and to the catechin standard of different concentrations. Individual samples blank were prepared by adding 5 ml of 4% HCL in methanol to 1 ml aliquot. The absorbance were read at 500 nm against reagent blank.

Catechin equivalent (CE) in % were calculated by using formula.

$$\text{CE \%} = \frac{(\text{mg catechin/ml})}{\text{vol. of extract taken (1ml)}} \times \frac{\text{Vol. made up (20 ml)}}{\text{wt. of sample (500 mg)}} \times 100$$

3.2.5.4 Flavan-4-ols

The anthocyanidins are determined by ionizing the middle ring of flavonoides by acid, yielding a pink color. The intensity of pink color is directly proportional to the concentration of flavan-4-ols. In case of catechin – like compounds the cyanidins are first obtained by boiling, which consequently get ionized to give pink color.

Grain samples were extracted with methanol and the phenolic compounds were then adsorbed in polyvinyl pyrrolidone (PVP) layers. The PVP was subsequently cleaned and treated with acid to ionize the flavanoid ring, if any. The results in all cases were expressed as A_{550g}^{-1} , (Butler, 1982).

Reagents

Butanol, Hydrochloric acid, Acetic acid, 0.1 N acetic acid, Diluted 5.71 ml acetic acid to 1 l with water, 300 ml butanol were taken in a 500 ml separating funnel to which 150 ml water were added shaken vigorously and kept overnight on stand. The top layer removed (water saturated butanol) and mixed with HCl in a bottle in 70 : 30 ratio, Water saturated butanol were mixed with methanol and N/10 acetic acid in ratio 70:15:15, which is to be used for sample blank, Methanol, Methanol –HCL 1% : Mixed 1 ml concentrated HCL in methanol and make upto 100 ml with methanol.

Extraction

Defatted flour (200 mg) was taken in screw cap test tubes to which five ml methanol was added, then tubes were placed in Stuart tube rotator (TR-2) and mixed for 1 hr. The mix was centrifuged and the supernatant was collected in a vial. The above steps were repeated using the residue and all the extracts was pulled in the above vial. This is referred to as Methanol extract.

To the above residue five ml methanol HCL (reagent 8) was added and above steps were repeated; again residue was re-extracted with additional five ml. of methanol HCL and pool the extract which was used for the estimation of Acid methanol extract.

0.5 ml methanol and acid methanol extract was separately taken in test tubes to which seven ml (reagent, 5) water saturated butanol with HCL was added. Blank was prepared by using (reagent 6). All the test tubes were placed in tube rotator for one hr and absorbency was read at 550 nm. All the results were calculated as A₅₅₀^{g⁻¹} dry weight.

3.2.6.1 Glume color

Glume is the outer most protective covering of the developing kernels. Glume colors were described as below and shown in (Plate 2D).

W =white, S =sienna (yellow), M =mahogany (brown), R =red, P =purple, B =black, G =grey, O=other.

3.2.6.2 Grain color

Parental lines and F₁ crosses were screened for grain color. Which was visually assessed and designated as

W =white, Y=yellow, R=red, Br=brown, Bu= buff and O=other

3.2.6.3 Presence of testa layer

Bleach test. For studying presence of testa layer Kofoid *et al.* (1976) method was followed. Parental lines and hybrids (F₁'s) were screened for the presence of testa.

Fifty kernels of individual line were placed separately in test tube to which 10 ml of reagent solution containing one part of KOH pellets and five parts of 6% sodium hypochlorite (a house hold bleach) was added. The tubes were swirled for few minutes and then placed on water bath maintained at 70°C for 10 min and shaken occasionally. After 10 min of bleaching the contents of the tubes were poured on to the strainer and rinsed thoroughly with running tap water separately. Excess of water was taken off and the sample damped in 2-3 folds of a paper towel. When the bulk of

water was absorbed, kernels were transferred to fresh towel and examined. Kernels with testa layer turned black while without testa layer, remained light in color.

3.2.7 Pre-treatment of seed

Seed were pre-treated with 0.1% mercuric chloride solution for two min followed by washings in three changes of sterile water. Then these surface sterilized seeds were plated on pre-moistened three layered blotting paper in sterile petri-plates for detection of internally seed borne fungi. Pre-treated seeds were also tested for germination following the method described earlier.

Twenty five seeds pre-treated with 0.1% mercuric chloride were placed in each petri-plates, replicated three times and incubated at $27^{\circ}\text{C} \pm 2^{\circ}\text{C}$ under diurnal cycle for seven days, for two replications and for two centres. Observations on fungal counts, infection score on germinated and ungerminated seeds were recorded as per method described earlier.

$$\text{Increased}(\%) = \frac{\text{count in treated} - \text{count in untreated}}{\text{count in treated}} \times 100$$

$$\text{Reduction}(\%) = \frac{\text{count in untreated} - \text{count in treated}}{\text{count in untreated}} \times 100$$

$$\text{Increased over location}(\%) = \frac{\text{maximum mean value} - \text{minimum mean value}}{\text{minimum mean value}} \times 100$$

3.2.8 Histopathology of Infection

Histopathology of infection and colonization of major molding fungi *F. moniliforme*, *F. pallidoroseum* and *C. lunata*, at different stages of host maturity i.e. at anthesis, 10,20,30 and 40 DAF were investigated. An experiment was conducted during the rainy season 1996 at Akola.

Two susceptible lines AKms-14B and SPV-104 were selected for the purpose. Five plots (2.75 x 3 m) for each genotypes were randomly distributed. All recommended package of practices were followed.

The inoculum were separately multiplied on autoclave sorghum grains in 250 ml Erlenmeyer flasks incubated for 7-10 days at 30°C. Grain showing profuse sporulation of the pathogen were removed from the flasks, washed in a glass jar with distilled water, and strained through double layered cheese cloth. Spores and the mycelial fragments (cfu) in the resulting suspension were counted with a haemocytometer and appropriately diluted with distilled water so as to obtain 1×10^6 cfu/ml.

Thirty cobs of each genotypes were selected randomly from individual plot at flowering of which 10 cobs were sprayed with inoculum of three major mold fungi, separately. Spraying was done carefully to cover the entire panicle to runoff taking due care to protect other cobs from spray. Inoculated cobs were tagged and covered with brown paper bags individually. The other plots were sprayed with fungicidal mixture of Thiram 0.2% + Carbendazim 0.07%.

Similarly, cobs were selected from 2nd set and plots of these lines and inoculations were done following the methodology described above at 10 DAF. The inoculated cobs were covered with brown paper bags and the bags of previously inoculated cobs were replaced with perforated paper bags. Unbagged, uninoculated and other plots were sprayed with Thiram 0.2% + Carbendazim 0.07%. This way 3rd, 4th and 5th set of plots were inoculated at 20,30, and 40 DAF, respectively.

Randomly 15 artificially inoculated kernels of each set for each pathogen were picked up and preserved in FAA (Formalin-acetic acid-alcohol) solution. Microtomy and microscopy were taken up at ICRISAT in 1997 to study the process of infection and histopathology.

3.2.9 Microtomy and microscopy

For microtomy of paraffin wax sectioning method described by Gopinath *et al.* (1987) was used on slight modifications. Different steps are as under.

Killing, Fixing and Hardening, Dehydration, Clearing, Embedding/section cutting, Staining.

Procedure

- **As the samples were preserved in FAA solution, killing of host (grain) tissue is not necessary.**
- **Samples preserved in FAA were removed, thoroughly washed with water and soaked in water for 3-4 hr.**
- **The seeds were softened by boiling in water for 5 min and fixed in acetic acid-alcohol (1:3) solution for 24 hr. Then the seeds were washed in running water for 30 min-1 h before next procedure.**
- **All the samples were dehydrated with series of tertiary butyl alcohol i.e. 30,50, 60, 70,80,90 and 100% concentrations, for 30-45 min in each concentration**
- **The samples were then placed in xylene, a solvent of paraffin wax for 1 hr called clearing (dealcoholization).**
- **After clearing, the seed samples were infiltrated with paraffin wax, till it become free from xylene. This step was repeated 2-3 times, till all the xylene was removed from melted paraffin wax. Thereafter all the paraffin wax was replaced with fresh paraffin and kept for 24 hr at 57°C in oven. At this stage samples becomes ready for embedding.**
- **The samples were then embedded in paraffin wax in plastic mould, oriented as desired and labeled.**
- **The embedded blocks were removed from plastic mould and put in ice cold water for fine crystal texture and softened by immersing the blocks in 1% aqueous solution of sodium lauryl sulphate for 24 h.**
- **Then blocks were washed with water and transferred to a mixture of glycerin and acetic acid (1:1) for 7 days.**
- **Serial microtome sections were cut (15-20 μ m thick), dewaxed and stained in Pianzeze III b stain (Dhinara and Sinclair, 1995) and finally mounted in DPX mountant for observation under compound microscope.**
- **For fluorescence microscopy the sections were stained with fungiflour/celluliflour after dewaxing and observed under fluorescence microscope.**

3.2.10 Statistical analyses

The data were analysed as per standard method suggested by Panse and Sukhatme (1954). For 10x10 diallel, two sets of diallel viz. parents + F_1 crosses and parents + F_2 progenies were analysed for all characters as per the method 2, model I of Griffings (1956 b) as further extended by Singh (1973a, 1973 b).

3.2.10.1 Heterosis

Heterosis (H_1) expressed over mid parental values, heterobeltiosis (H_2) expressed over better parent; were estimated by using the following method of Laosuwan and Atkins (1977).

$$\text{Heterosis } (H_1) = \frac{\overline{F_1} - \overline{MP}}{\overline{MP}} \times 100$$

$$\text{Where, } \overline{MP} = \frac{\overline{P_1} + \overline{P_2}}{2}$$

$$\text{eterobeltiosis } (H_2) = \frac{\overline{F_1} - \overline{BP}}{\overline{BP}} \times 100$$

Where, $\overline{F_1}$ = Mean of hybrid (F_1)

$\overline{P_1}$ = Mean of parent 1

$\overline{P_2}$ = Mean of parent 2

\overline{MP} = Mid parental value

\overline{BP} = Mean of better parent

(Better parent changes as per the characters)

Test of significance, however, were calculated for the numerators of the expression i.e. $H_1 = \overline{F_1} - \overline{MP}$ and $H_2 = \overline{F_1} - \overline{BP}$.

3.2.10.2 Combining ability analysis

Combining ability analysis was based on plot means. It was carried out for all the characters, using method 2 model 1 of Griffings (1956 b) and further extended by Singh (1973 a, 1973 b).

The mathematical model for combining ability analysis (Model - 1) was assumed as :

$$X_{ijk} = \mu + g_i + g_j + S_{ij} + e_{ijk}$$

$$i, j = 1 \text{-----} p$$

$$k = 1 \text{-----} b$$

Where,

p = number of parents,

b = number of replications,

μ = population mean,

g_i = gca effect of i th parent,

g_j = gca effect of j th parent,

S_{ij} = sca effect of cross between i th and j th parents.

e_{ijk} = mean error effect.

The restrictions imposed to the model are :

$$\sum_i g_i = 0$$

and

$$\sum_j S_{ij} + S_{ii} = 0 \quad (\text{for each } i).$$

Computation of gca and sca effects

The various effects were estimated as follows :

$$\text{Population mean } \mu = \frac{2}{p(p-1)} X_{..}$$

gca effects of i th parent =

$$\hat{g}_i = \frac{1}{p+2} [(X_{i.} + X_{ii}) - \frac{2}{p} X_{..}]$$

and sca effects of ij th cross =

$$\hat{S}_{ij} = x_{ij} - \frac{1}{p+2} (X_{ii} + X_{ii} + X_{.j} + X_{ij}) + \frac{2}{(p+1)(p+2)} X_{..}$$

Where,

p = number of parents,

$X_{i.}$ = total of array involving i th parent,

X_{ii} = mean of i th parent,

X_{ij} = mean value of ij th cross,

$X_{..}$ = total of all the treatments in the table without

reciprocals i.e. $\sum_i \sum_j X_{ij}$ such that $i \leq j$

3.2.10.3 Correlation coefficient

In order to study the degree of association between different characters contributing to grain mold development, simple correlation coefficients were worked out from the respective variances and co-variances as per the formulae suggested by Hays *et al.* (1955).

$$\text{Correlation between } ij = \frac{\text{covariance of } ij}{\sqrt{\text{variance of } i \times \text{variance of } j}}$$

Significance of correlation coefficient was determined from table of correlation coefficient of 5 and 1% level of significance (Fisher, 1958). The 'r' values were compared against $n-2$ degree of freedom.

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RESULTS

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RESULTS

4.1 Analysis of variance

4.1.1 Analysis of variance (Pooled F_1) for physical, agronomic, and pathological traits, Akola and Patancheru, 1996

The pooled analysis of variance over two locations for parents and F_1 crosses is presented in (Table 2). Data revealed that the two locations, were significantly different from each other for all the characters, except plant height and mesocarp thickness. This is indicative of the fact that the two locations were ecologically different and genotypes were tested under sufficiently diverse environments.

The variance due to treatments-parents, crosses and parents vs crosses were highly significant for all the 15 characters studied, except infection by *F. pallidoroseum* and other fungi in parents and glume covering, and infection by *F. pallidoroseum* in parents vs. crosses. However, variances due to treatments \times locations, parents \times locations and crosses \times locations interactions were significant for all characters except cob length and glume covering in treatments \times locations, endosperm texture, cob length, glume covering, and infection by *F. moniliforme*, *F. pallidoroseum*, *C. lunata* and other fungi. in parents \times locations and grain hardness, cob length and glume covering in crosses \times locations interaction. Variance due to parents vs crosses \times locations interaction was observed to be significant for the trait grain hardness, electrical conductivity, DTF, TGM and germination.

4.1.2 Analysis of variance (Pooled F_2), for physical, agronomic and pathological traits, Akola and Patancheru, 1996

The pooled analysis of variance over two locations for parents and F_2 progenies for 11 characters is presented in (Table 3). Data revealed that the two locations significantly differ from each other for all the characters except cob length and glume covering.

The variance due to treatments, parents, crosses and parents vs. crosses were highly significant for all the 11 characters studied, except infection by *F. pallidoroseum* and other fungi for treatments, parents and crosses and plant height in parents vs. crosses. However variance due to treatments \times locations interactions were significant for all characters except glume covering and infection by *F. moniliforme*. For parents \times (locations) interactions only grain hardness, electrical conductivity and TGMR were significant and for progenies \times locations interactions grain hardness, electrical conductivity, plant height, cob length, TGMR and germination were significant. However, for parents vs progenies \times locations interactions significantly differed from each other for all the characters except glume covering, and infection by *F. moniliforme* and *C. lunata*.

4.1.3 Analysis of variance for biochemical characters of parental lines and F_1 at Akola, 1995

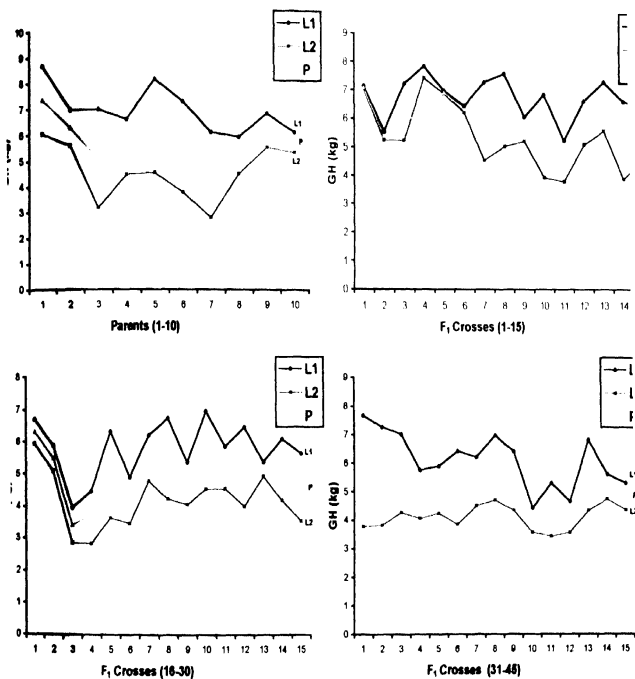
For biochemical characters viz. protein, soluble sugars, tannins and flavan-4-ols were studied from the samples collected at Akola (rainy season, 1995). The analysis of variance is presented in (Table 4).

The treatment differences were highly significant for all the traits studied. Further partitioning of treatment variance into components viz., parents, crosses and parents vs. crosses revealed that all the three components differed significantly among themselves for all the characters. Replication differences were non-significant for all the traits studied.

It may thus be concluded that the parental lines included in this investigation possessed sufficient variability for the characters studied.

4.2. Mean performance, heterosis and heterobeltiosis, Akola and Patancheru, 1996

The mean performance of parental lines, F_1 crosses, F_2 progenies, heterosis (H_1) and heterobeltiosis (H_2) were computed as per the procedure given in chapter III. The character wise results are presented in (Tables 5 to 20).



L1 = Akola L2 = ICRISAT P = Pooled

Figure 1. Grain hardness of parents and F₁ crosses at Akola and Patancheru, 1996

1 SPV-1201	1 SPV-1201 x ICSB-101B	16 ICSB-101B x IS-6335	31 GJ-35-15-15 x SPV-946
2 ICSB-101B	2 SPV-1201 x AKms-14B	17 ICSB-101B x IS-9471	32 GJ-35-15-15 x SPV-104
3 AKms-14B	3 SPV-1201 x SRT-26B	18 AKms-14B x SRT-26B	33 GJ-35-15-15 x IS-2284
4 SRT-26B	4 SPV-1201 x GJ-35-15-15	19 AKms-14B x GJ-35-15-15	34 GJ-35-15-15 x IS-6335
5 GJ-35-15-15	5 SPV-1201 x SPV-946	20 AKms-14B x SPV-946	35 GJ-35-15-15 x IS-9471
6 SPV-946	6 SPV-1201 x SPV-104	21 AKms-14B x SPV-104	36 SPV-946 x SPV-104
7 SPV-104	7 SPV-1201 x IS-2284	22 AKms-14B x IS-2284	37 SPV-946 x IS-2284
8 IS-2284	8 SPV-1201 x IS-6335	23 AKms-14B x IS-6335	38 SPV-946 x IS-6335
9 IS-6335	9 SPV-1201 x IS-9471	24 AKms-14B x IS-9471	39 SPV-946 x IS-9471
10 IS-9471	10 ICSB-101B x AKms-14B	25 SRT-26B x GJ-35-15-15	40 SPV-104 x IS-2284
	11 ICSB-101B x SRT-26B	26 SRT-26B x SPV-946	41 SPV-104 x IS-6335
	12 ICSB-101B x GJ-35-15-15	27 SRT-26B x SPV-104	42 SPV-104 x IS-9471
	13 ICSB-101B x SPV-946	28 SRT-26B x IS-2284	43 IS-2284 x IS-6335
	14 ICSB-101B x SPV-104	29 SRT-26B x IS-6335	44 IS-2284 x IS-9471
	15 ICSB-101B x IS-2284	30 SRT-26B x IS-9471	45 IS-6335 x IS-9471

4.2.1 100-grain weight

Data presented in (Table 5) revealed that for parental lines, the range of variation for 100-grain weight was from 1.69 to 2.90 g at Akola, whereas it was from 1.36 to 2.54 g at Patancheru. However, for crosses the range was from 1.76 to 3.41g and 1.52 to 2.98 g at Akola and Patancheru, respectively. Among the parental lines, SPV-104 recorded the highest 100-grain weight both at Akola (2.9 g) and Patancheru (2.54 g). Among the crosses, the highest grain weight at Akola and in pooled data exhibited by the cross SPV-1201 \times SPV-104 (3.41 g) and (2.94 g), respectively, whereas at Patancheru ICSB-101B \times SPV-104 (2.58 g) recorded the highest 100 grain weight. Among the crosses, SPV-1201 \times SPV-104 was significantly superior at Akola. Across the two locations, SPV-104 recorded the highest 100-grain weight (2.72 g), followed by SPV-1201 (2.66 g) and ICSB-101 B(2.06 g).

Out of 45 crosses four exhibited significant positive heterosis (Table 12). The positive heterosis of highest magnitude (23.72%) was observed in AKms-14B \times IS-6335 followed by SRT 26B \times IS-6335 (22.94%) and AKms- 14 B \times IS-9471 (20.95%)

As regards heterobeltiosis none of the crosses exhibited significant positive heterobeltiosis, however, the positive heterobeltiosis ranged between 18.75 (SRT-26B \times IS-6335) to 0.18 % (ICSB 101B \times SPV-104). Promising crosses exhibiting positive heterobeltiosis appear were SRT-26B \times IS 6335 (18.35%). SRT-26B \times IS-9471 (18.29%) and AKms-14B \times IS-9471 (16.41%).

4.2.2 Grain hardness

That the range of variation for grain hardness for parental lines was from 5.91 to 8.67 kg/cm² at Akola, and 2.78 to 6.03 kg/cm² at Patancheru (Table 5, Fig. 1). However, in pooled data the range of variation was 4.43 to 7.35 kg/cm². The parental line SPV-1201 recorded the highest grain hardness at Akola 8.67 kg/cm² and Patancheru 6.03 kg/cm² as well as in pooled data 7.35 kg/cm². For crosses the range was from 3.96 to 8.71 kg/cm² and 2.82 to 7.43 kg/cm² at Akola and Patancheru, respectively. The

highest (7.85, 7.43 and 7.64 kg/cm²) grain hardness was recorded for cross SPV-1201 × GJ-35-15-15 at both locations and in pooled data, respectively.

In F₂ progenies (Table 8) the range of variation for grain hardness was from 3.25 to 7.31 kg/cm² and 2.13 to 6.26 kg/cm² at Akola and Patancheru, respectively. However, when pooled over locations, the grain hardness ranged between 3.26 and 5.93 kg/cm². Maximum grain hardness was recorded in F₂ progenies of SPV-1201 × IS 9471 (7.31 kg/cm²) at Akola, whereas F₂ progenies of SPV 946 × IS 2284 (6.26 kg/cm²) and 5.93 kg/cm² exhibited maximum grain hardness at Patancheru and pooled data, respectively.

In F₁ diallel set, heterosis ranged from 0.40 to 11.84% over mid parental value (Table 12). The cross SPV 1201 × GJ 35-15-15 (11.84 %) recorded the highest heterosis, but the least heterosis was recorded in SPV 946 × IS 2284 (0.40%). None of the crosses exhibited significant positive heterosis, however, three crosses recorded significant negative heterosis.

Four crosses exhibited positive heterobeltiosis but it was not significant. The highest positive heterobeltiosis was exhibited in AKms-1201 × IS 2284 (5.49%) followed by SPV-1201 × GJ-35-15-15 (3.95%).

In F₂ diallel set only one F₂ progeny SPV-946 × IS-2284 (7.98 %) exhibited positive heterosis but, it was non-significant (Table 15).

4.2.3 Endosperm texture

It is seen from the (Table 5), that for parental lines the range of variation for endosperm texture was from 22.40 to 56.95 % at Akola, whereas, at Patancheru the same ranged between 25.90 to 59.15%. However, for crosses the range was from 21.15 to 74.00% and 25.55 to 66.60% at Akola and Patancheru respectively. The corneous endosperm texture at both locations and average over locations was exhibited by parental line GJ 35-15-15 (22.40, 25.9 and 24.15%), respectively. Among crosses corneous endosperm texture at Akola was exhibited by the cross SPV-1201 × GJ 35-15-15 (21.15 %), whereas, at Patancheru location ICSB-101 B × GJ 35-15-15 (25.55%) exhibited the corneous endosperm texture. Cross SPV-1201 × GJ 35-

15-15 (25.13%) exhibited lowest endosperm texture when pooled over locations. ^oCorneus, intermediate and floury endosperm textures are depicted in Plate 3.

In F_1 diallel only three crosses exhibited negative heterosis (Table 12). Highest negative heterosis was noted by cross SPV-1201 \times GJ 35-15-15 (-6.07%) followed by IS-2284 \times IS-6335 (-3.07%). Sixteen crosses exhibited significant positive heterosis. Highest significant positive heterosis was recorded in cross AKms-14 B \times SPV-946 (63.01%) followed by AKms-14B \times SRT-26B (52.31%).

Whereas, 21 crosses indicated negative heterobeltiosis, none of them is significant. Highest negative heterobeltiosis was indicated by crosses SPV-946 \times IS 6335 (-19.63%) followed by cross SPV-946 \times IS 2284 (-19.43%). Only four crosses recorded significant positive heterobeltiosis. Cross AKms-14B \times SPV-946 recorded highest significant positive heterobeltiosis (58.01%).

4.2.4 Electrical conductivity

It is observed from the (Table 5, Fig. 2) the range of variation for electrical conductivity for parental line was from 105.0 to 300.50 dSm^{-1} at Akola and 214.00 to 846.50 dSm^{-1} at Patancheru. However, in pooled data the range of variation was 167.25 to 553.25 dSm^{-1} . The parental line IS-9471 (105.00 dSm^{-1}) exhibited lowest electrical conductivity at Akola and IS-6335 (214.00 dSm^{-1}) at Patancheru, while in pooled data parental line IS-2284 (167.25 dSm^{-1}) exhibited lowest value. For crosses the range was 90.00 to 237.50 dSm^{-1} and 82.00 to 485 dSm^{-1} at Akola and Patancheru, respectively. Among crosses lowest electrical conductivity at Akola was exhibited by the cross IS-6335 \times IS 9471 (90.00 dSm^{-1}) and at Patancheru and pooled over locations SPV 946 \times IS 9471 exhibited lowest electrical conductivity (82.00 and 102.25 dSm^{-1} , respectively).

In F_2 progenies (Table 8) the mean values ranged from 155.00 dSm^{-1} (IS 6335 \times IS-9471) to 553.00 dSm^{-1} (SRT-26B \times SPV-104) at Akola; whereas, it was 158.00 dSm^{-1} (SRT-26B \times IS -9471) to 685.00 dSm^{-1} (SPV-1201 \times GJ 35-15-15) at Patancheru. However, when pooled over locations, the electrical conductivity ranged

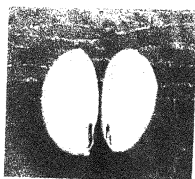
Plate : 3 Endosperm texture of sorghum midian longitudinal half
kernels of sorghum showing corneous, intermediate and
floury endosperm

The rating for the kernels are

A-1,B-2,C-3,D-3,E-3,F-4,G-5 and H-5 (3-3X)

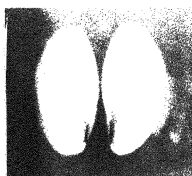


A

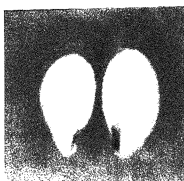


B

Corneous



C

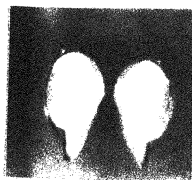


D



E

Intermediate



F



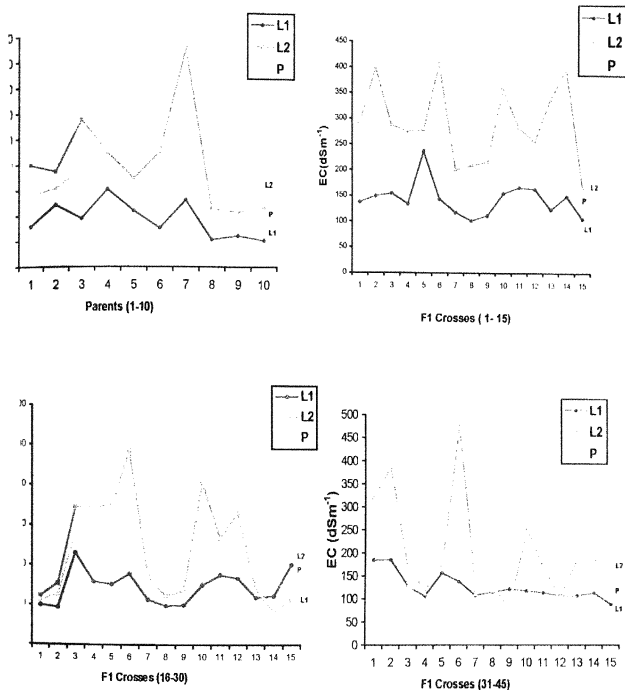
G

Floury



H

Plate 3. Endosperm texture



L1 = Akola L2 = Patancheru, P = Pooled

Figure 2. Electrical conductivity of parents and F_1 crosses at Akola and Patancheru, 1996

PV-1201	1 SPV-1201 x ICSB-101B	16 ICSB-101B x IS-6335	31 GJ-35-15-15 x SPV-946
SB-101B	2 SPV-1201 x AKms-14B	17 ICSB-101B x IS-9471	32 GJ-35-15-15 x SPV-104
Kms-14B	3 SPV-1201 x SRT-26B	18 AKms-14B x SRT-26B	33 GJ-35-15-15 x IS-2284
SRT-26B	4 SPV-1201 x GJ-35-15-15	19 AKms-14B x GJ-35-15-15	34 GJ-35-15-15 x IS-6335
J-35-15-15	5 SPV-1201 x SPV-946	20 AKms-14B x SPV-946	35 GJ-35-15-15 x IS-9471
SPV-946	6 SPV-1201 x SPV-104	21 AKms-14B x SPV-104	36 SPV-946 x SPV-104
SPV-104	7 SPV-1201 x IS-2284	22 AKms-14B x IS-2284	37 SPV-946 x IS-2284
-2284	8 SPV-1201 x IS-6335	23 AKms-14B x IS-6335	38 SPV-946 x IS-6335
-6335	9 SPV-1201 x IS-9471	24 AKms-14B x IS-9471	39 SPV-946 x IS-9471
-9471	10 ICSB-101B x AKms-14B	25 SRT-26B x GJ-35-15-15	40 SPV-104 x IS-2284
	11 ICSB-101B x SRT-26B	26 SRT-26B x SPV-946	41 SPV-104 x IS-6335
	12 ICSB-101B x GJ-35-15-15	27 SRT-26B x SPV-104	42 SPV-104 x IS-9471
	13 ICSB-101B x SPV-946	28 SRT-26B x IS-2284	43 IS-2284 x IS-6335
	14 ICSB-101B x SPV-104	29 SRT-26B x IS-6335	44 IS-2284 x IS-9471
	15 ICSB-101B x IS-2284	30 SRT-26B x IS-9471	45 IS-6335 x IS-9471

between 165.50 and 589.50 dSm^{-1} and minimum electrical conductivity recorded in progeny IS-6335 \times IS 9471 (165.50 dSm^{-1})

All 45 crosses indicated negative heterosis out of which 30 crosses exhibited significant negative heterosis (Table 12). Highest significant negative heterosis was recorded by cross SPV-104 \times IS 9471 (-70.69 %) followed by cross SRT-26B \times IS 6335 (-61.63 %).

As regards heterobeltiosis all 45 crosses exhibited negative heterobeltiosis, of which 38 indicated significant negative heterobeltiosis. The range in negative direction varied from -10.55 to -80.89%. The highest significant negative heterobeltiosis was exhibited by cross SPV-104 \times IS-9471 (-80.89 %) followed by cross SPV-104 \times IS 6335 (-74.33 %).

In F_2 diallel set (Table 15) nine progenies exhibited negative heterosis, of which three progenies exhibited significant negative heterosis. The highest negative heterosis exhibited by progeny SRT-26B \times IS-9471 (-33.09 %) followed by AKms- 14 B \times IS-6335 (-30.99 %) and SPV-104 \times IS-9471 (-19.68%). Whereas, 29 progenies exhibited negative heterobeltiosis of which 13 progenies exhibited significant negative heterobeltiosis. The highest negative heterobeltiosis was exhibited by SRT- 26B \times IS-9471 (-51.44 %) followed by AKms-14 B \times IS 6335 (-50.23 %).

4.2.5 Days to 50% flowering

The range for days to 50% flowering, among parental lines was from 66.00 to 80.00 and 59.00 to 79.00 days at Akola and Patancheru, respectively (Table 5). The early flowering was exhibited by IS-6335 (66.00 days) at Akola, (59.00 days) Patancheru and (62.50 days) in pooled data. The crosses ranged between 59.50 and 76.00 days and 59.00 and 85.50 days at Akola and Patancheru, respectively. The earliest flowering crosses was SRT-26B \times IS-6335 (59.00 and 59.75 days) at Patancheru and in pooled data, respectively.

The range of desirable heterosis was from - 0.17 to- 17.59 % (Table 12) 18 out of 45 crosses exhibited significant negative heterosis, highest recorded by

cross SPV-104 \times IS 9471 (-17.59 %) followed by SPV-104 \times IS-6335 (-13.32%). Only five crosses exhibited significant positive heterosis.

Heterobeltiosis in desired direction was exhibited by 36 crosses, while positive and significant heterobeltiosis was exhibited by only one cross. The cross SPV-104 \times IS 6335 recorded highest magnitude of heterobeltiosis (-22.04 %) followed by cross SPV-104 \times IS -9471 (-21.41 %). Both crosses were also significantly superior to other crosses for heterosis in negative direction.

4.2.6 Plant height

It is observed from the (Table 6), that the range of variation was from 136 cm to 283 cm at Akola, and 140 cm to 285.50 cm at Patancheru, however, in pooled data the range of variation was 138 cm to 273.75 cm. The parental line AKms-14B exhibited least height (136, 140, 138 cm) at Akola, Patancheru and in pooled data, respectively. For cross the dwarfs cross was ICSB-101B \times AKms-14B (183 cm) at both locations and pooled data, however tallest cross was SPV-1201 \times IS-9471 (343m 340 and 341.50 cm) at Akola, Patancheru and in pooled data, respectively.

In F_2 progenies (Table 8) the range was from 157.90 to 260.00 cm at Akola and from 184.15 to 318.25 cm at Patancheru. The least height was exhibited by progeny ICSB-101B \times AKms-14 B (157.90, 184.15 and 171.02 cm) at Akola, Patancheru and pooled data, respectively.

Significant positive heterosis was observed in (Table 13) 41 crosses and maximum heterosis (56.81 %) was observed in SPV-1201 \times AKms-14B followed by SPV-1201 \times ICSB 101B (54.40 %). Only three crosses exhibited negative heterosis but, none of the crosses exhibited significant. Highest negative heterosis was observed in SPV-1201 \times SPV-946 (-3.93 %).

Only seven crosses exhibited heterobeltiosis in negative direction and out of them, only three crosses were significant. They were SRT-26 \times GJ 35-15-15 (-19.04 %), AKms- 14B \times GJ-35-15-15 (-12.02 %) and SPV-1201 \times SPV-946 (-10.78 %).

In F_2 diallel set (Table 15) 18 progenies exhibited heterosis in negative direction and none of them was significant. The highest negative heterosis was exhibited by SPV-1201 \times SPV-104 (-12.55 %) followed by SPV-1201 \times ICSB-101B (-10.06 %).

As regards to heterobeltiosis, 35 progenies exhibited negative heterobeltiosis and seven progenies exhibited significant negative heterobeltiosis. The range was from -23.32 to -1.21 %. The highest negative heterobeltiosis was exhibited in SPV-1201 \times ICSB-101B (-23.32 %) followed by SRT-26B \times GJ-35-15-15 (-21.61 %).

4.2.7 Cob length

Mean values for this character varied from 21.20 cm to 26.20 cm and from 17.00 to 28.90 cm in case of parental lines at Akola and Patancheru, respectively. (Table 6), whereas in pooled data the range was 19.10 cm to 27.55 cm. Parental line ICSB-101B recorded maximum ear head length (26.20, 28.90 and 27.55 cm) at Akola, Patancheru and pooled data, respectively. Among crosses the range was 20.40. (SPV-1201 \times SPV-104) to 29.70 cm (GJ-35-15-15 \times IS-6335) at Akola, 21.00 (AKms-14 B \times SRT-26B) to 32.10 cm (ICSB 101B \times IS-9471) at Patancheru and 20.95 cm (SPV-1201 \times SPV-104) to 30.80 cm (ICSB-101B \times IS-9471) in pooled data.

In F_2 progenies (Table 8) mean values ranged from 18.95 (SPV-1201 \times SPV-104) to 30.05 cm (ICSB-101 B \times AKms-14 B) at Akola, whereas, it was 20.00 (SPV-946 \times SPV-104) to 29.85 cm in (SPV-946 \times IS-6335) at Patancheru. However, in pooled data, the cob length ranged between 21.25 to 28.90 cm and the maximum cob length recorded in progeny (ICSB-101 B \times AKms-14 B).

In F_1 diallel set (Table 13) the estimates for heterosis ranged from 0.10 to 22.40% in positive direction and from -2.42 to -4.23 % in negative direction. Five crosses were significant in positive direction. The highest heterosis % in positive direction was recorded by cross SPV-104 \times IS-6335 (22.40 %) followed by AKms-14B \times GJ-35-15-15 (19.79 %).

Only one cross AKms-14B \times GJ-35-15-15 (19.54%) exhibited significant heterobeltiosis in positive direction. This cross was also significantly superior to other crosses for heterosis in positive direction.

In F_2 diallel set (Table 15) 42 progenies exhibited positive heterosis of which only three progenies exhibited significant positive heterosis, the highest positive heterosis was exhibited in GJ-35-15-15 \times IS 2284 (27.54 %) followed by SPV-946 \times IS-6335 (17.26%) and GJ-35-15-15 \times IS-9471 (14.92%). Whereas, 25 progenies exhibited positive heterobeltiosis, of which only one progeny GJ-35-15-15 \times IS-2284 (25.60 %) exhibited significant positive heterobeltiosis.

4.2.8 Glume covering

The range for glume covering for parental lines was exhibited from 50 to 75% at Akola, Patancheru and in pooled data (Table 6). Parental lines ICSB-101B and IS-6335 recorded maximum 75% glume covering at Akola, Patancheru and pooled data. Minimum 50% was recorded by SPV-1201, AKms-14B, SPV-946, SRT-26B at Akola, Patancheru and in pooled data.

Among crosses ICSB-101 B \times IS-6375, SPV-104 \times IS-6335 and IS-2284 \times IS-6335 recorded maximum 75% glume covering at Akola, Patancheru and pooled data. AKms-14B \times GJ-35-15-15 and AKms-14B \times SPV-946 recorded maximum glume covering at Patancheru.

In F_2 progenies (Table 9) maximum 66.50, 75.00 and 70.75 % glume covering was exhibited in SPV-946 \times IS-6335 at Akola, Patancheru and pooled data, respectively. However, minimum 37.50% was recorded in SPV-1201 \times SPV-946 and AKms- 14B \times IS-946 at Akola. Whereas, SPV-1201 \times SPV-104 recorded minimum 37.50 and 41.25% glume covering at Patancheru and in pooled data, respectively.

In F_1 diallel set (Table 13) none of the crosses recorded significant positive heterosis. However, 23 crosses recorded positive heterosis, the highest (25%) exhibited by AKms- 14 B \times GJ 35-15-15 and AKms- 14 B \times SPV -946. None of crosses exhibited significant positive heterobeltiosis. However, above crosses exhibited highest 25% heterobeltiosis in positive direction.

In F_2 diallel set (Table 15) 18 progenies exhibited positive heterosis, but it was not significant. The highest positive heterosis was exhibited in progenies SPV-104 \times IS 2284 (20.47%) followed by GJ-35-15-15 \times IS 2284 (16.71%). Whereas, 16 progenies exhibited positive heterobeltiosis, the range was 0.00 to 13.78%. The highest positive heterobeltiosis was exhibited in SPV 104 \times IS 2284 (13.78%) followed by GJ-35-15-15 \times IS-2284 (10.22 %).

4.2.9 Mesocarp thickness

The parental lines ranged was from 45.45 to 97.52 μm at Akola 59.91 to 96.57 μm at Patancheru and 54.02 μm to 97.05 μm in pooled data. Lowest mesocarp thickness was recorded in parental line SPV-946 (45.45 μm) at Akola and pooled data and IS-9471 (59.91 μm) at Patancheru. Among crosses the mean values ranged from 32.03 μm , 31.32 μm and 31.68 μm (SPV-104 \times IS-6335) to 96.79 μm , 99.56 μm and 98.13 μm (AKms-14B \times SRT-26B) at Akola, Patancheru and in pooled data, respectively (Table 6). Mesocarp thickness of susceptible, elite (moderately resistant), resistant parental lines and red and white cross are shown in Plate 4.

The ranged for desirable heterosis was from -3.03 to -61.14% (Table 13). Out of 45 crosses in F_1 generation 29 exhibited significant negative heterosis, the highest (-61.14%) recorded by SPV-104 \times IS-6335 followed by SPV-104 \times IS-2284 (-57.19%) and ICSB-101 B \times IS -6365 (-56.61%).

Heterobeltiosis in desired direction (negative and significant) were exhibited by 32 crosses. The ranged of heterobeltiosis was from 13.74 to -64.39%. The highest heterobeltiosis (-64.39%) was indicated by SPV-104 \times IS-6335 followed by ICSB-101 B \times IS- 6335 (-61.74%). Both crosses were significantly superior to other crosses for heterosis in negative direction.

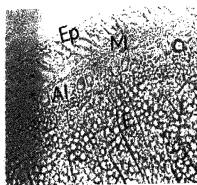
4.2.10 Threshed grain mold rating

At Akola, the range of variation for parental lines was form 2.00 to 4.00, whereas it was 1.50 to 5.00 and 1.75 to 4.50 at Patancheru and in pooled data, respectively. (Table 6). The parental lines IS-9471, IS- 6335 and IS-2284 exhibited lowest (2.0) TGM at Akola, whereas, IS-9471 exhibited (1.5) TGM at Patancheru and (1.75)

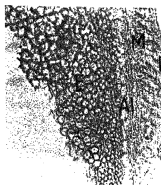
Plate : 4 : Mesocarp thickness (Transverse section, 66.6 X)

I Parental Lines		
A	Akms 14-B	White (S)
B	SPV-946	White (MR)
C	IS 9471	Red (R)
II F₁ Cross (red)		
D	SPV-104	White (Parental line)
E	IS-6335	Red (Parental line)
F	SPV-104 x IS-6335	Red (Cross)
III F₁ (Cross (white))		
G	SPV-1201	White (Parental line)
H	ICSB-101B	White (Parental line)
I	SPV-1201 x ICSB-101B	White (Cross)

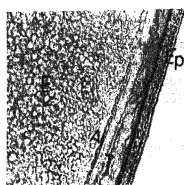
(Ep = Epicarp, M=Mesocarp, Al-Aleurone layer,
E-Endosperm and T-testa)



A

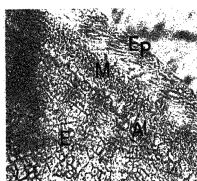


B

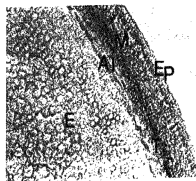


C

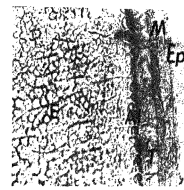
I Parental lines



D

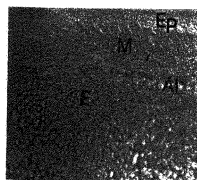


E

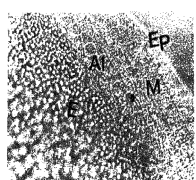


F

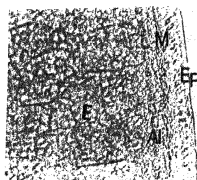
II F₁ cross (red)



G



H



I

III F₁ cross (white)

TGMR in pooled data. Among crosses the range was from (2.0 to 3.50), cross SPV-946 \times IS-6335 exhibited the lowest and AKms-14B \times SRT-26B exhibited the highest TGMR at Akola, at Patancheru TGMR varied between (1.50 and 5.00) cross GJ 35-15-15 \times IS-9471, IS-2284 \times IS-6335 and IS-6353 \times IS-9471 exhibited lowest (1.50) TGMR and SPV-946 \times SPV-104 and SRT 26 B \times GJ 35-15-15 exhibited highest (5.0) TGMR. However, in pooled data IS-6335 \times IS-9471 and IS 2284 \times IS-6335 recorded lowest (1.75) TGMR.

In F_2 diallel (Table 9) lowest TGMR was exhibited by progeny IS-2284 \times IS-9471 (1.70, 2.0 and 1.85) at Akola, Patancheru and pooled data respectively. However highest TGMR was exhibited by AKms-14B \times SPV-104 (3.05) and (3.28) at Akola and pooled data, respectively. Whereas at Patancheru progeny AKms-14B \times SRT-26B (4.00) exhibited highest TGMR.

In F_1 diallel set (Table 13) the range for desirable heterosis was from 2.86 to -46.15% Out of 45 crosses 22 crosses exhibited significant negative heterosis, the highest (-46.15%) recorded by AKms-14 B \times IS-6335 followed by (44.00 %) by AKms-14 B \times IS-9471.

Heterobeltiosis in negative direction (negative and significant) were exhibited by 25 crosses. The range of heterobeltiosis was from 5.56 to 61.11%. The highest heterobeltiosis was exhibited (-61.11 %) in AKms-14B \times IS-6335 and AKms-14B \times IS-9471 followed by AKms-14 B \times IS-2284 (-55.56%). First two crosses were significantly superior to other crosses for heterosis in negative direction.

In F_2 diallel set (Table 15) 44 progenies exhibited negative heterosis, of which 25 exhibited significant negative heterosis, the highest negative heterosis exhibited by AKms-14B \times IS-6335 (-30.00%) followed by AKms-14 B \times GJ 35-15-15 (-29.70%) and AKms-14 B \times IS-9471 (-29.60 %). Whereas, all 45 progenies exhibited negative heterobeltiosis of which 41 exhibited significant negative heterobeltiosis. The range was from - 51.11% to - 2.50%. The highest negative heterobeltiosis was exhibited in AKms-14B \times IS-9471 (-51.11%) followed by AKms-14 B \times IS-6335 (-49.44%) and AKms-14B \times IS-2284 (-46.67%).

4.2.11 Seed germination

It is seen from the (Table 7), that for parental lines the range of variation for seed germination was from 29.00 to 86.00% at Akola, whereas, at Patancheru it ranged between 2.50 and 61.00%. However, when pooled over locations parental lines ranged between 15.75 to 71.75%. Among crosses the range of variation for germination was from 32.50 to 98.00%, and 14.00 to 88.50% at Akola and Patancheru, respectively. Whereas, pooled data the crosses ranged between 25.25 to 90.00%. The highest seed germination at Akola was exhibited by parental line IS 6335 (86.00%), whereas, at Patancheru and pooled data parental line IS-2284 exhibited highest germination 61.00 and 71.75%, respectively.

Among crosses SPV-1201 \times IS-2284 exhibited highest 98.00% seed germination at Akola, ICSB-101B \times IS-6335 exhibit highest 88.50% germination at Patancheru, whereas, cross ICSB-101B \times IS-2284 exhibited highest germination (90.00%) in pooled data.

In F_2 progenies (Table 9) the mean values for germination ranged from 53.00 to 87.90% at Akola, 9.75 to 83.50% at Patancheru and 38.00 to 84.70% in pooled data. At Akola progeny IS-2284 \times IS-6335 exhibited highest germination (87.90%) followed by IS-6335 \times IS-9471 (86.40%) and SPV-946 \times IS-6335 (86.15%), while progeny IS-2284 \times IS-9471 exhibited highest seed germination (83.50 and 84.70%) at Patancheru and pooled data, respectively.

The range of heterosis in positive direction was from 4.14 to 76.76% (Table 14). All crosses exhibit positive heterosis, however 37 crosses exhibit significant. The highest heterosis (76.76%) was indicated in cross AKms-14B \times IS-6335, followed by cross AKms-14 B \times IS-9471 (71.44%).

Regarding heterobeltiosis the range was 1.20 to 40.56% in positive direction, whereas, 17 crosses out of 45 exhibits significant positive heterobeltiosis. The highest (40.56%) was indicated by SPV-1201 \times SRT-26B followed by ICSB-101B \times SPV-946 (33.89%) and ICSB-101B \times IS-9471 (33.45 %).

In F_2 diallel set (Table 16) all the progenies exhibited positive heterosis, of which 23 progenies exhibited significant positive heterosis. The highest positive heterosis was exhibited by progeny ICSB-101B \times AKms-14B (78.73%) followed by AKms-14B \times GJ-35-15-15 (62.84%) and AKms-14B \times IS-6335 (57.66%). However, 36 progenies exhibited positive heterobeltiosis, of which only four progenies exhibited significant positive heterobeltiosis. The highest positive heterobeltiosis was exhibited by, ICSB-101B \times AKms-14B (38.93%) followed by SPV-1201 \times SRT-26B (37.20%).

4.2.12 Infection by *F. moniliforme*

It is seen from the (Table 7) that, the range of variation for parental lines in F_1 diallel set, was from 5.33 to 38.67% at Akola, whereas, at Patancheru the same ranged between 34.00 to 71.34%. However, in crosses the range was from 5.99 to 35.99 and 21.34 to 70.67% at Akola and Patancheru, respectively. The lowest fungal load of *F. moniliforme* at Akola, Patancheru and in pooled data was exhibited by IS-9471, (5.33, 34.00 and 19.67%), respectively. Among crosses lowest fungal load of *F. moniliforme* at Akola was exhibited by cross SPV-104 \times IS-6335 (5.99%), whereas, at Patancheru and pooled data, cross ICSB-101B \times IS-6335 exhibited lowest infection by *F. moniliforme* (21.34 and 15.66%), respectively.

In F_2 diallel progenies (Table 10) mean values ranged from 18.50 (ICSB-101B \times IS-6335) to 52.52% (SPV-104 \times IS-9471) at Akola, where as it was 25.34 (IS-6335 \times IS-9471) to 67.50% (AKms-14B \times SPV-104) at Patancheru. However, in pooled data infection by *F. moniliforme* ranged between 26.88 to 54.25%, lowest 26.88% *F. moniliforme* load was recorded in progeny (SPV- 946 \times IS-6335).

In F_1 diallel set 40 crosses exhibited negative heterosis of which only five crosses exhibited significant. The range was 53.47% to 0.02% in negative direction. However, the highest negative heterosis was in cross ICSB-101B \times IS-6335 (-53.47%) followed by AKms-14B \times IS-6335 (-50.81 %) and SPV-104 \times IS-6335 (-48.39%) (Table 14).

However, 43 progenies exhibited negative heterobeltiosis of which 13 progenies exhibited significant. The range was -62.41 to -0.83 % in negative direction. The highest negative heterobeltiosis was observed in AKms-14B \times IS-6335 (-62.41%) followed by SPV-104 \times IS-6335 (-61.22%) and ICSB-101B \times IS- 6335 (-60.51%).

In F_2 diallel set (Table 16) only three progenies exhibited negative heterosis, but it was not significant. The highest negative heterosis was exhibited in progenies SRT-26B \times SPV-946 (-12.50%) followed by ICSB-101B \times AKms-14 B (-7.59%) and SRT -26B \times SPV-104 (-1.93%). Three progenies recorded significant positive heterosis.

However, 13 progenies exhibited negative heterobeltiosis, but it was not significant. The range was -17.43 to 0.04%. The highest negative heterobeltiosis exhibited in SRT-26B \times SPV-946 (-17.43%) followed by AKms-14 B \times IS-9471 (-16.13%) and SPV-946 \times IS-6335 (-16.00 %), but it was not significant. Only one progeny IS-2284 \times IS-6335 exhibited (104.41%) significant positive heterobeltiosis.

4.2.13 Infection by *F. pallidoroseum*

The range of variation for parental lines for *F. pallidoroseum* was exhibited from 0.71 to 2.67% at Akola, 7.34 to 19.34% at Patancheru and 4.02 to 10.51% in pooled data. Lowest (0.71%) *F. pallidoroseum* load was exhibited by parental lines SPV-1201, ICSB-101B, AKms-14B and GJ-35-15-15 at Akola, whereas, AKms-14B exhibited lowest 7.34 and 4.02% *F. pallidoroseum* load at Patancheru and in pooled data, respectively. In crosses lowest (0.71, 5.34 and 3.02%) *F. pallidoroseum* load was exhibited in cross ICSB-101 B \times IS 2284 at Akola, Patancheru and pooled data, respectively (Table 7).

In F_2 progeny lowest (0.77%) fungal load of *F. pallidoroseum* was exhibited in GJ-35-15-15 \times SPV-946 at Akola, however, progeny AKms-14B \times GJ-35-15-15 exhibited lowest (1.50 and 1.53%) fungal load at Patancheru and pooled data, respectively (Table 10).

Out of 45 crosses, 13 crosses exhibited negative heterosis out of which only one cross ICSB-101 B \times IS-2284 (-64.85%) exhibited significant heterosis in F_1 diallel set (Table 14).

The range of heterobeltiosis was from -71.25 % to -0.04%, 22 crosses exhibited negative heterobeltiosis, out of which only one cross ICSB-101 B \times IS-2284 (-71.25%) exhibited significant heterobeltiosis. The same cross was found significantly superior to other crosses for heterosis in negative direction.

In F_2 diallel 37 progenies exhibited negative heterosis of which only one progeny GJ 35-15-15 \times SPV-946 (-74.43%) exhibited significant negative heterosis. The range was -74.43 to -1.71% in negative direction. Thirty eight progenies exhibited heterobeltiosis in negative direction and out of them, only one progeny GJ-35-15-15 \times SPV-946 (-79.88%) was significant. The same progeny was found significantly superior to other crosses for heterosis in negative direction (Table16).

4.2.14 Infection by *C. lunata*

At Akola location, the range of variation for parental lines was from 24 (IS-9471) to 60% (GJ-35-15-15), whereas, at Patancheru and in pooled data the range of variation for parental lines was from 15.33 and 20.67 (IS-6335) to 37.33 and 46.67% (SRT 26B), respectively. Among crosses the range was from 18.00 (SPV-1201 \times IS-6335 and SPV-1201 \times IS-9471) to 68% (ICSB-101B \times SRT-26B) at Akola, whereas, at Patancheru 8.0 (ICSB-101B \times SPV-104) to 37.33% (SPV-1201 \times SPV-104) and in pooled data the range was from 20.33 (ICSB-101B \times IS-6335) to 46.33% (SPV-1201 \times SPV-104) (Table 7).

In F_2 progenies lowest (19.26%) fungal load of *C. lunata* was exhibited in IS-6335 \times IS-9471 at Akola, whereas, IS 2284 \times IS 9471 exhibited lowest values (18.50 and 19.71%) at Patancheru and in pooled data, respectively (Table 10).

In F_1 diallel set (Table 14), 36 crosses exhibited negative heterosis, but it was not significant. The range was -36.13 to -0.55% , highest negative heterosis was exhibited in ICSB-101 B \times IS-6335 (-36.13%) followed by SRT-26B \times IS-9471 (-34.29%) and SPV-1201 \times IS-9471 (-33.01%). The range of heterobeltiosis was from -52.72 to -2.87% , 41 crosses exhibited negative heterobeltiosis, out of which 11 crosses were exhibited significant. The highest negative heterobeltiosis was exhibited by cross ICSB-101 B \times IS-6335 (-52.72%) followed by SRT-26B \times IS-9471 (-50.71%) and SPV-1201 \times IS-9471 (-49.26%).

In F_2 diallel set (Table 16), 34 progenies exhibited negative heterosis. The range was from -38.72 (IS-2284 \times IS-9471) to 1.17% (SRT-26B \times IS-2284), none of them was significant. However, 41 progenies exhibited negative heterobeltiosis but it was not significant. The range was from -45.25 to 3.94% . Highest negative heterobeltiosis was exhibited in progeny IS-2284 \times IS-9471 (-45.25%) followed by IS-2284 \times IS-6335 (-41.39%).

4.2.15 Infection by other fungi

At Akola location the range of variation for parental lines was from 9.34 (AKms-14B) to 24.67% (IS-2284), whereas, at Patancheru the range of variations was from 0.71 (SPV-104) to 6.00% (ICSB-101B), however in pooled data the range of variation was from 5.18 (AKms-14B) to 15.00% (IS-2284). Among crosses lowest (12.00 , 0.71 and 9.00%) fungal load of other fungi was exhibited in AKms-14B \times SPV-104, ICSB-101B \times SPV-104, and AKms-14B \times SPV-104 at Akola, Patancheru and pooled data, respectively (Table 7).

In F_2 progenies lowest (27.00 , 20.52 and 8.45%) fungal load of other fungi was exhibited in ICSB-101 B \times SRT-26B, SPV-1201 \times IS-2284 and AKms-14B \times SRT-26B at Akola, Patancheru and pooled data, respectively (Table 10).

In F_1 diallel four crosses exhibited negative heterosis but it was not significant. The highest negative heterosis exhibited by GJ-35-15-15 \times IS-2284 (-15.66%) followed by IS-6335 \times IS-9471 (-13.94%) and ICSB 101B \times SPV-104 (-5.14%). However, 11 crosses exhibited negative heterobeltiosis, the range was from

0.02 to -22.22%. The highest negative heterobeltiosis exhibited in GJ-35-15-15 \times IS 2284 (-22.22%) followed by SRT-26 B \times IS 9471 (-19.52). None of them exhibited significant negative heterobeltiosis (Table 14).

In F_2 diallel set 14 progenies exhibited negative heterosis, but it was not significant. The highest negative heterosis was exhibited in progeny GJ-35-15-15 \times SPV-104 (-29.64%) followed by SPV-946 \times SPV-104 (-21.40 %) and AKms-14B \times GJ-35-15-15 (-18.18 %), whereas, 20 progenies exhibited negative heterobeltiosis. The range was -40.00 to -4.15%. The highest negative heterobeltiosis exhibited in progeny AKms-14B \times GJ-35-15-15 (-40.00%) followed by GJ-35-15-15 \times SPV-104 (-39.80%). None of the progeny exhibited significant negative heterobeltiosis (Table 16).

Biochemical characters

4.2.16 Proteins

The data relative to mean performance of parental lines and crosses for biochemical parameters is presented in (Table 11).

Data revealed, that the range of variation in parental lines for proteins in sorghum grains varied from 8.93 to 12.94%, however, for crosses the range was 5.23 to 10.56%. The highest proteins was exhibited by parental lines ICSB-101B (12.94 %) followed by SRT-26 B (12.41%) and SPV-1201 (12.35%). Among crosses the highest proteins was exhibited by cross SPV-104 \times IS-6335 (10.56%) followed by SPV-1201 \times IS-9471 (10.11%).

Among 45 crosses none of cross exhibited the heterosis in positive direction, a negative heterosis exhibited in all 45 crosses (Table 17) The highest negative heterosis was exhibited by cross AKms-14B \times SRT 26 B (-27.43%), followed by AKms-14 B \times SPV-946 (-27.27%).

Significant heterobeltiosis was reported in all 45 crosses in negative direction, and none of crosses exhibited positive heterobeltiosis. The highest heterobeltiosis in negative direction was exhibited by cross SRT-26B \times IS-9471 (-30.36%) followed by SRT-26 B \times IS-2284 (-29.15 %).

4.2.17 Soluble sugars

The mean performance of parental lines and crosses for soluble sugars content is presented in (Table 11).

Data revealed that parental lines mean values for this character ranged from 1.2 (SPV-104) to 2.3% (AKms-14B). Among crosses SPV-1201 \times IS-6335 and SRT- 26B \times IS-2284 (1.8%) exhibited highest and SPV-946 \times SPV-104 and SPV-104 \times IS-6335. (0.8%) recorded lowest value.

Among 45 crosses (Table 17), only four exhibited the heterosis in positive direction, the highest being 2.53% (GJ-35-15-15 \times IS-9471) followed by 2.29% (SPV-1201 \times IS-6335). Negative heterosis exhibited in 41 crosses. The highest significant negative heterosis was exhibited by cross AKms-14B \times SRT-26B (-28.67%).

None of crosses exhibited significant and positive heterobeltiosis. Only one cross SPV-1201 \times IS-6335 (2.29%) exhibited positive heterobeltiosis. Significant heterobeltiosis was reported in 34 crosses in negative direction.

4.2.18 Tannins

The mean performance of parental lines and crosses for tannins content is presented in (Table-11).

Parental lines range for tannins content was varied from 0.02 (ICSB-101B) to 5.84 CE % (IS-2284). However, crosses range was varied from 0.01 (ICSB-101B \times GJ-35-15-15) to 3.41 CE % (AKms 14B \times IS 2284).

Among 45 crosses 20 exhibited positive heterosis (Table 17), of which 16 were significant. Cross (SRT-26B \times IS-9471) recorded highest 38.64 CE % followed by (AKms-14B \times IS-9471) 36.84 CE %.

Only one cross (SRT-26B \times SPV-104) exhibited positive heterobeltiosis (0.07%). Twenty seven crosses exhibited significant negative heterobeltiosis.

4.2.19 Flavan-4-ols

The data relative to mean performance on flavan-4-ols both extract (methanol + H+/methanol) content is presented in (Table 11 and Fig. 3).

The range for parental lines for this characters varied from 0.00 to 18.00 A_{550g}^{-1} , however, for crosses the range was 0.00 to 12.00 A_{550g}^{-1} . The highest flavan-4-ols content was exhibited by parental lines IS-9471 (18.0) followed by IS-2284 (7.9) and IS-6335 (7.6 A_{550g}^{-1}). Among crosses the highest flavan-4-ols content was exhibited by IS-6335 \times IS-9471 (12.00) followed by IS-2284 \times IS-9471 (11.90 A_{550g}^{-1}).

Among 45 crosses 25 exhibited positive and significant heterosis, the range was 52.10 to 5.67% in positive direction (Table 17). Cross SRT-26B \times IS-6335 exhibited the highest positive heterosis (52.10 %) followed by SPV-946 \times IS-6335 (52.07%). However, only one crosses recorded negative significant heterosis, IS-2284 \times IS-6335 (-14.01%).

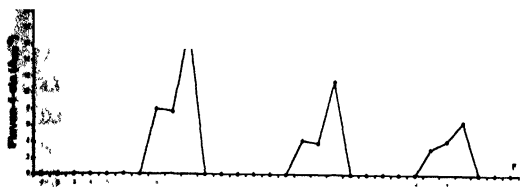
Out of 45 crosses 21 crosses exhibited positive heterobeltiosis of which four were significant. The highest positive heterobeltiosis was observed in GJ-35-15-15 \times SPV-946 (48.37 %) followed by SRT-26B \times SPV-946 (41.30%)

Mean performance at Akola, 1995 and 1996

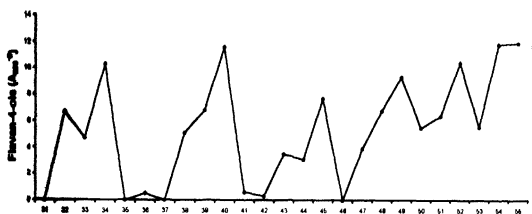
4.3.1 100-grain weight

Data presented in (Table 18) revealed that, parental line SPV-1201 exhibited highest 100-grain weight (3.04, 2.80, and 2.92 g) in both season and in pooled data. However, lowest (1.61 g) value was recorded in parental line IS-2284 in Akola 95 season, whereas, parental line IS-6335 exhibited lowest (1.69 and 1.66 g) grain weight in Akola 96 season and in pooled data, respectively.

Among crosses maximum 100-grain weight was recorded in SPV-946 \times SPV-104 (3.17 g) in Akola 1995. whereas, cross SPV-1201 \times SPV-104 recorded maximum (3.40 and 3.28 g) grain weight in Akola 1996 and in pooled data, respectively. Cross IS-2294 \times IS-6335 recorded minimum (1.62, 1.76 and 1.69 g) grain weight in Akola 1995, Akola 1996 and in pooled data, respectively.



Parents (1-10) F₁ Crosses (11-20)



F₁ Crosses (31-55)

F – Flavan-4-ols

Figures 3. Flavan-4-ols of parents and F₁ crosses at Akola 1995

1 BPV-1201	11 SPV 1201 x ICSB 101B	26 ICSB 101B x JS-6335	41 GJ 35 15 15 x SPV 946
2 ICSB-101B	12 SPV 1201 x AKms 14B	27 ICSB 101B x JS 9471	42 GJ 35 15 15 x SPV 104
3 AKms-14B	13 SPV 1201 x SRT 26B	28 AKms 14B x SRT 26B	43 GJ 35 15 15 x JS 2284
4 SRT-26B	14 SPV 1201 x GJ 35 15 15	29 AKms 14B x GJ 35 15 15	44 GJ 35 15 15 x JS-6335
5 GJ-35-15-15	15 SPV 1201 x SPV 946	30 AKms 14B x SPV 946	45 GJ 35 15 15 x JS 2471
6 BPV-946	16 SPV 1201 x SPV 104	31 AKms 14B x SPV 104	46 SPV 946 x SPV 104
7 BPV-104	17 SPV 1201 x JS 2284	32 AKms 14B x JS 2284	47 SPV 946 x JS 2284
8 JS-2284	18 SPV 1201 x JS-6335	33 AKms 14B x JS-6335	48 SPV 946 x JS-6335
9 JS-6335	19 SPV 1201 x JS 9471	34 AKms 14B x JS 9471	49 SPV 946 x JS 9471
10 JS-9471	20 ICSB 101B x AKms 14B	35 SRT 26B x GJ 35 15 15	50 SPV 104 x JS 2284
	21 ICSB 101B x SRT 26B	36 SRT 26B x SPV 946	51 SPV 104 x JS-6335
	22 ICSB 101B x GJ 35 15 15	37 SRT 26B x SPV 104	52 SPV 104 x JS 9471
	23 ICSB 101B x SPV 946	38 SRT 26B x JS 2284	53 JS 2284 x JS-6335
	24 ICSB 101B x SPV 104	39 SRT 26B x JS-6335	54 JS 2284 x JS 9471
	25 ICSB 101B x JS 2284	40 SRT 26B x JS 9471	55 JS-6335 x JS 9471

4.3.2 Grain hardness

Data presented in (Table 18) revealed that parental line SPV-1201 exhibited highest (9.09, 8.67 and 8.88 kg) grain hardness in Akola 1995, Akola 1996 and in pooled data, respectively. Cross GJ-35-15-15 \times SPV-946 exhibited highest (8.47, 8.70, 8.59 kg) grain hardness in Akola 1995, Akola 1996 and in pooled data, respectively.

4.3.3 Endosperm texture

It is observed from (Table 18), that parental line SPV-946 recorded (24.10 and 22.00%) corneous endosperm texture in Akola 1995 and in pooled data, respectively, however, parental line GJ-35-15-15 exhibited (18.00 %) corneous endosperm texture in Akola 1996. Whereas, floury endosperm texture (66.95 and 61.40%) recorded in parental line IS-6335 in Akola 1995 and in pooled data, respectively, and (56.95%) in IS-2284 in Akola 1996. Cross SPV-1201 \times GJ-35-15-15 exhibited (27.25, 21.15 and 24.00%) corneous endosperm texture in Akola 1995, Akola 1996 and in pooled data, respectively, however, cross SPV-104 \times IS-9471 recorded (73.85 and 71.20%) floury endosperm texture in Akola 1995 and in pooled data, respectively and in SPV-104 \times IS-6335 (74.00 %) in Akola 1996.

4.3.4 Electrical conductivity

It is observed from the (Table 18), that lowest electrical conductivity (97.00, 105.00 and 101.00 dSm^{-1}) was exhibited in parental line IS-9471 in Akola 1995, Akola 1996 and in pooled data, respectively. However, highest electrical conductivity was exhibited by parental line SPV-104 (350.00 and 305.00 dSm^{-1}) in Akola 1995 and in pooled data, respectively and SRT-26B (300.50 dSm^{-1}) in Akola 1996.

4.3.5 Days to 50% flowering

It is seen from (Table 18), that parental line IS-6335 exhibited minimum (60.00, 66.00 and 63.00 days) flowering and maximum (80.50, 80.00 and 80.25 days) flowering was observed in SPV-104 at Akola 1995, Akola 1996 and in pooled data, respectively. Cross SPV-1201 \times SRT-26 indicated maximum value (89.00 and 82.50 days) in Akola 1995 and in pooled data respectively. However, cross SPV-946 \times IS-9471 recorded

maximum flowering 79.00 days in Akola 1996 season and minimum flowering was recorded in ICSB-101B \times IS-2284, ICSB-101 B \times IS-6335, and AKms-14B \times IS-6335 (57 days) in Akola 1995. However, in Akola 1996 and in pooled data cross SRF-26B \times IS-6335 exhibited minimum (59.50 and 58.7 days) flowering, respectively.

4.3.6 Plant height

Data presented in (Table 19) revealed that parental line AKms-14B recorded minimum plant height (142.50, 136.00 and 139.25 cm) in both season and in pooled data, respectively. However, maximum (265.00 cm) was recorded in parental line IS-2284 in Akola 1995 and (251.00 and 252.25 cm) in parental line IS-9471 in Akola 1996 and in pooled data, respectively. Cross ICSB-101B \times SPV-104 recorded minimum (150.50 cm) plant height in Akola 1995, however, cross ICSB-101 B \times AKms-14B recorded minimum plant height (183.00 and 171.50 cm) in Akola 1996 and in pooled data, respectively.

4.3.7 Cob length

It is observed from (Table 19), that parental line ICSB-101B indicated maximum (26.60, 26.20 and 26.40 cm) and cross GJ-35-15-15 \times IS-6335 recorded maximum (30.90, 30.60 and 30.75 cm) cob length in both seasons and in pooled data, respectively.

4.3.8 TGM/R

Data presented in (Table 19) revealed that parental line IS-9471 recorded lowest (1.50, 2.00 and 1.75) and parent AKms-14B recorded highest (4.00, 4.00 and 4.00) TGM/R in both seasons and in pooled data, respectively. Among crosses 25.23 and 22 cross recorded minimum (2.00) TGM/R at both locations and in pooled data, respectively.

4.3.9 Seed germination

In F_1 diallel set (Table 19) the range of variation for parental line was exhibited from 51.00 to 97.00, 29.00 to 86.00 and 40.00 to 90.00 % in Akola 1995, Akola 1996 and in pooled data, respectively. Parental line GJ-35-15-15 exhibited highest (97.00 %) seed germination in Akola 1995, Akola 1996 and in pooled data, parental line IS-6335 recorded highest 86.00 and 90.00% germination, respectively. Parental line AKms-

98

14B recorded lowest 51.00, 29.00 and 40.00% seed germination in both seasons and in pooled data, respectively

Among crosses, the highest germination was recorded in GJ-35-15-15 \times IS-6335 and SPV-946 \times IS-6335 (93.00%) in Akola 1995 season. In Akola 1996 and in pooled data, cross SPV-1201 \times IS-2284 recorded highest seed germination 98.00 and 94.50%, respectively

4.3.10 Infection by *F. moniliforme*

It is seen from (Table 20), that the range of variation in F_1 diallel set was from 0.21 (IS-2284 and GJ-35-15-15) to 34.45% (SRT-26B) in Akola 1995, whereas, in Akola 1996 the range was 5.33 (IS-9471) to 46.00% (AKms-14B). However, in pooled data the range was 4.51 (IS-9471) to 30.23% (SRT-26B).

Among crosses minimum (3.34, 6.00 and 4.67%) *F. moniliforme* load was exhibited in IS-6335 \times IS-9471 at both locations and in pooled data, respectively. However, maximum (73.50, and 51.08%) load was recorded in cross AKms-14B \times SRT-26B in Akola 1995 and in pooled data respectively. Whereas cross SPV-1201 \times SPV-946 indicated maximum (35.99%) load in Akola 96 season.

4.3.11 Infection by *F. pallidoroseum*

It is observed from (Table 20) that, all the parental lines in Akola 1995 recorded minimum (0.71%) load, however, five parents viz. SPV-1201, IC SB-101B, AKms-14B, GJ-35-15-15 and SPV-104 recorded minimum load in Akola 1996, and in pooled data.

Among crosses, all the crosses except SPV-1201 \times IS-2284 recorded minimum (0.71%) *F. pallidoroseum* load in Akola 1995, 12 crosses each in Akola 1996 and in pooled data.

4.3.12 Infection by *C. lunata*

It is observed from (Table 20), that in Akola 1995 the range of variation for parental lines was from 1.46 (SPV-1201) to 45.55% (SPV-104), whereas in Akola 1996 it was from 24.00 (IS-9471) to 60.00% in (GJ-35-15-15), however in pooled data the range was 15.89 (IS-9471) to 50.66% (SPV-1201).

Among crosses minimum (12.34%) *C. lunata* load was exhibited in cross SPV-946 × IS-9471 in Akola 1995 season, however, cross SPV-1201 × IS-9471 exhibited minimum (18.00 and 19.00%) load in Akola 1996 and in pooled data. Whereas, cross ICSB-101B × SRT-26B indicated maximum (58.90, 68.00 and 63.45%) load in Akola 1995, Akola 1996 and in pooled data, respectively.

4.3.13 Infection by other fungi

Data presented in (Table 20) revealed that in Akola 1995 the range of variation for parental lines was from 16.67 (SPV-1201) to 88.89% in (GJ-35-15-15), in Akola 1996, 9.33 (AKms-14B) to 26.00% (SPV-946) and in pooled data 18.55 (SRI-26B) to 55.11% in (GJ-35-15-15). Among crosses the range of variation was from 8.88 (AKms-14B × SRT-26B) to 46.67% (GJ-35-15-15 × SPV-946) in Akola 1995, however, it was from 12.00 (AKms-14B × SPV-104) to 37.34% (AKms-14B × IS-6335) in Akola 1996 and 14.11 (AKms-14B × SRI-26B) to 36.66% (ICSB-101B × IS-2284) in pooled data.

4.4 Combining ability analysis

Combining ability analysis was carried out for 15 characters of F_1 and 11 characters of F_2 diallel progenies and results are presented in (Tables 21 and 22) respectively.

The variance due to treatments was further partitioned using appropriate expectations of the observed mean squares, into components of variation attributable to general combining ability (gca) variance and specific combining ability (sca) variance. The characteristic results of the above aspects are presented here with

4.4.1 100-grain weight

In F_1 diallel set, significant variance due to gca and sca indicated role of additive as well as non-additive gene action. The ratio of additive to non-additive variance was more (1.936) than unity indicating predominance of additive gene action (Table 21).

4.4.2 Grain hardness

In F_1 diallel set, variance due to gca and sca were significant indicating that additive as well as non-additive gene actions were important in expression of this character.

The ratio of additive to non-additive variance was more (1.700) than unity indicating **predominance** of additive gene action (Table 21)

Variance due to gca and sca were significant and thus additive as well as non-additive gene action for this character was important in F_2 progenies (Table 22). Lower ratio of $\delta^2 \text{ gca} / \delta^2 \text{ sca}$ than unity (0.378) proved predominance of non-additive gene action

4.4.4 Electrical conductivity

In F_1 diallel set variance components due to gca and sca were significant indicating presence of additive as well as non additive gene action in the inheritance of this character. Lower ratio of $\delta^2 \text{ gca} / \delta^2 \text{ sca}$ (0.728) revealed that the character is predominantly controlled by non-additive gene action (Table 21)

In F_2 diallel progenies the variance of gca and sca were significant proving the importance of additive and non-additive gene action for the control of this character. But higher ratio of $\delta^2 \text{ gca} / \delta^2 \text{ sca}$ (1.716) indicated that the additive gene action is more predominant than non-additive gene action (Table 22)

4.4.5 Days to 50% flowering

Variance due to gca and sca were significant and thus additive as well as non-additive gene action for this character was important in F_1 diallel set (Table 21). Lower ratio of $\delta^2 \text{ gca} / \delta^2 \text{ sca}$ than unity (0.974) proved predominance of non-additive gene action

4.4.6 Plant height

In F_1 diallel set, variance due to gca and sca were significant indicating that additive and non-additive gene action were important in expression of this character. However, lower ratio of $\delta^2 \text{ gca} / \delta^2 \text{ sca}$ (0.443) indicated that non-additive gene action was important than additive (Table 21)

In F_2 diallel progenies, the variance of gca and sca were significant proving the importance of additive as well as non-additive gene action for the control of this character. But higher ratio of $\delta^2 \text{ gca} / \delta^2 \text{ sca}$ (1.227) indicated that the additive gene action is more predominant than non additive gene action (Table 22)

4.4.7 Cob length

In F_1 diallel analysis, significant variance component for gca and sca indicated additive and non-additive gene action. Higher ratio of $\delta^2_{gca}/\delta^2_{sca}$ (1.256) than unity proved that the character is primarily governed by additive gene action (Table 21).

In F_2 diallel set the variance components due to gca and sca were significant. This indicated the existence of additive and non-additive gene action. Higher ratio of $\delta^2_{gca}/\delta^2_{sca}$ (1.967) than unity showed the importance of additive genetic variance than non-additive genetic variance (Table 22).

4.4.8 Glume covering

In F_1 diallel set variance due to gca was significant, whereas sca was non-significant. It is therefore concluded that the character is predominantly controlled by additive gene action only. This is further proved by higher ratio (1.293) of $\delta^2_{gca}/\delta^2_{sca}$ (Table 21).

Variance due to gca and sca were significant in F_1 diallel set (Table 22). It is therefore, proved that the character is governed by additive as well as non-additive gene action. In inheritance of this character, lower ratio of $\delta^2_{gca}/\delta^2_{sca}$ (0.480) revealed that the character is predominantly controlled by non-additive gene action.

4.4.9 Mesocarp thickness

In F_1 diallel set, significant variance due to gca and sca indicated role of additive as well as non-additive gene action. Lower ratio of $\delta^2_{gca}/\delta^2_{sca}$ than unity (0.376) proved predominance of non-additive gene action (Table 21).

4.4.10 Threshed grain mold rating

In F_1 diallel set, significant variance due to gca and sca indicated role of additive and non-additive gene action. Higher ratio of $\delta^2_{gca}/\delta^2_{sca}$ (1.961) than unity proved that the character is primarily governed by additive gene action (Table 21).

In F_2 diallel progenies the variance of gca and sca were significant proving the importance of additive as well as non-additive gene action for the control of this character. But higher ratio of δ^2gca/δ^2sca (1.187) indicated that the additive gene action is more predominant than non-additive gene action (Table 22).

4.4.11 Seed germination

In F_1 diallel set, variance due to gca and sca were significant indicating presence of additive as well as non-additive gene action in inheritance of this character. Lower ratio of δ^2gca/δ^2sca (0.646) revealed that the character is predominantly controlled by non-additive gene action (Table 21).

In F_2 diallel progenies variance components due to gca and sca were significant. This proved the presence of additive and non-additive gene action for control of germination and there was a lot of useful variability for the character also. Similarly from the ratio of δ^2gca/δ^2sca (0.772) it was clear that additive as well as non-additive gene action was equally operating (Table 22).

4.4.12 Infection by *F. moniliforme*

In F_1 diallel set, variance due to gca and sca were observed significant indicating that additive as well as non-additive gene actions were important in expression of this character. The ratio of additive to non-additive variance was more (1.573) than unity indicating predominance of additive gene action (Table 21).

In F_2 diallel set variance due to gca was significant, whereas sca was non-significant. It is therefore concluded that the character is predominantly controlled by additive gene action only. However, lower ratio (0.725) of δ^2gca/δ^2sca revealed that the character is predominantly controlled by non-additive gene action also (Table 22).

4.4.13 Infection by *F. pallidoroseum*

In F_1 diallel set variance due to gca was non-significant and sca was significant. It is therefore, appears that the character is controlled by non-additive gene action only. This is further proved by lower ratio (0.043) of δ^2gca/δ^2sca (Table 21).

In F_2 diallel set variance due to gca and sca was non-significant. However, the ratio of $\delta^2_{\text{gca}}/\delta^2_{\text{sca}}$ was observed to be (0.045) which is indicative of non additive gene action (Table 22)

4.4.14 Infection by *C. lunata*

The variance due to gca was significant and sca was non-significant in I_1 diallel set (Table 21). It is therefore concluded that the character is predominantly controlled by additive gene action only. This is further proved by higher ratio (4.068) of $\delta^2_{\text{gca}}/\delta^2_{\text{sca}}$

In F_2 diallel set variance due to gca and sca was significant and non significant, respectively (Table 22). It is therefore concluded that the character is predominantly controlled by additive gene action only. This is further proved by very high ratio (49.780) of $\delta^2_{\text{gca}}/\delta^2_{\text{sca}}$

4.4.15 Infection by other fungi

In F_1 diallel set variance due to gca was non significant and sca was significant. The ratio of $\delta^2_{\text{gca}}/\delta^2_{\text{sca}}$ was also observed to be less than unity (0.082) which indicated that this character is predominantly governed by non additive gene action (Table 21)

Variance due to gca was significant however sca was non significant in F_2 diallel set (Table 22). The ratio of $\delta^2_{\text{gca}}/\delta^2_{\text{sca}}$ was higher (1.845) which indicated the predominant of additive gene action in the inheritance of this trait. It is therefore concluded that the character is predominantly controlled by additive gene action only.

Biochemical characters

4.4.16 Proteins

In F_1 diallel set significant variance due to gca and sca indicated role of additive as well as non additive gene action in the inheritance of this character. The ratio of additive to non-additive variance was (0.101) lower than unity revealed that the character is predominantly controlled by non additive gene action (Table 23)

4.4.17 Soluble sugars

Variance due to gca and sca were significant and thus additive as well as non-additive gene action for this trait was important in F_1 diallel set (Table 23). However, ratio of $\delta^2_{gca}/\delta^2_{sca}$ than unity (0.313) proved predominance of non-additive gene action.

4.4.18 Tannins

In F_1 diallel set, variance due to gca and sca were to be significant indicating that, additive as well as non-additive gene actions were important in expression of this character. The ratio of additive to non-additive variance was more (3.192) than unity indicating predominance of additive gene action (Table 23).

4.4.19 Flavan-4-ols

In F_1 diallel set variance components due to gca and sca were significant indicating presence of additive and non-additive gene action in inheritance of this character. But higher ratio of $\delta^2_{gca}/\delta^2_{sca}$ (4.821) indicated that the additive gene action is more predominant than non-additive gene action (Table 23).

4.5 Estimates of general combining ability effects

The general combining ability effects were calculated for all the 15 characters in I_1 and 11 characters in I_2 for pooled over locations. The character wise results are presented below.

4.5.1 100 grain weight

The data relative to gca effects for 100-grain weight are presented in (Table 24).

Among 10 parental lines in I_1 diallel set only one parental line (J-35 15-15 (0.132) exhibited significant gca in positive direction. Remaining parental lines, exhibited either non-significant gca or significant gca in negative direction. Thus, they may not be of any use in breeding program as far as 100-grain weight is concerned.

4.5.2 Grain hardness

Hard grain is a desirable character for grain mold resistance. In I_1 diallel set, five parental lines recorded significant positive gca effects. Parental line IS-9471 exhibited higher (0.843) significant gca estimates, followed by AKms-14B (0.784), SR I-26B

(0.677), IS-6335 (0.464) and GJ-35-15-15 (0.429). These parental lines can be said as good general combiners (Table 24).

Rest of parents have exhibited non significant gca or significant gca in negative direction and hence, may not be useful in breeding program for this trait.

In F_2 (Table 25) progenies all the parental lines showed significant positive gca effects. SRT-26B recorded highest gca (1.568), followed by IS 9471 (1.388), SPV-1201 (1.301) and IS-6335 (1.295).

4.5.3 Endosperm texture

Comeous endosperm texture is desirable character for grain mold resistance. Therefore, negative and significant gca effects are desirable. It is revealed from the data (Table 24) that, all the 10 parental lines exhibited significant negative gca effects. Higher significant negative gca effects was exhibited by parental lines AKms 14B (-17.316) followed by SPV-104 (-8.400), SRT 26B (-7.858) and GJ-35-15-15 (-6.212). Thus these parental lines can be identified as best general combiners for endosperm texture.

4.5.4 Electrical conductivity

Low electrical conductivity is desirable character for grain mold resistance. Therefore, negative and significant gca effects are desirable. It is revealed from the data (Table 24), that none of the parental lines exhibited significant negative gca effects in F_1 crosses. However, minimum significant positive effect was observed in GJ 35 15 15 (44.042) and SPV-946 (65.792).

However, in F_2 diallel (Table 25) seven parents exhibited significant negative gca effects. The highest significant negative gca effects exhibited by GJ 35 15-15 (-121.617) followed by SPV-1201 (-101.451), SPV 946 (-78.284) and IS 2284 (-59.117) all are significantly superior to first parental line SPV 104 (-21.992).

4.5.5 Days to 50% flowering

In sorghum breeding program, earliness is one of the objective for selection of genotypes and negative gca effects of higher magnitude are always favoured. In F_1 diallel set parental line SPV-1201 (-8.375) appeared to be the best general combiner for earliness exhibiting highly significant and negative gca effects (Table 24).

4.5.6 Plant height

Dwarf to mid dwarf plant type is a desirable character. Therefore negative gca effects are appreciated. The gca estimates for plant height for F_1 diallel set is presented in (Table 24). It is revealed from the data that all the parental lines exhibited significant negative gca effects. The highest significant negative gca effects was exhibited by SPV-1201 (-68.667) followed by IC SB-101B (-66.042) IS 2284 (-59.167) and AKms-14B (-57.167). Thus these parental lines can be identified as the best general combiners for introducing dwarfness. In F_2 diallel set four parental lines SRT-26 B (41.734), AKms-14 B (-34.722) IC SB-101 B (15.434) and IS 2284 (-14.005) exhibited significant negative gca effects (Table 25).

4.5.7 Cob length

In F_1 diallel set general combining ability effects the range for the character cob length was 0.446 to 2.774 in all the parental lines. None of the parental line exhibited significant positive gca effects (Table 24).

In F_2 diallel set (Table 25) only one parental line SPV-1201 (0.072) exhibited positive gca effects while all others recorded significant negative gca effects except SRT-26 B (-0.349).

4.5.8 Glume covering

Maximum glume covering is a desirable characters for grain mold resistance therefore positive gca effects of higher magnitude are desirable. In F_1 diallel (Table 24) four parental lines exhibited significant positive gca effects the highest being recorded by IC SB-101 B (10.511) followed by IS-2284 (8.428) IS 9471 (7.386) and SPV 1201 (3.220). Thus these parental lines can be identified as best general combiners for glume covering. In F_2 diallel set (Table 25) five parental lines recorded significant gca effects in positive direction viz IS-6335 (11.701) SPV-1201 (5.576) IC SB-101 B (4.451) AKms-14 B (2.742) and SRT 26 B (1.742).

4.5.9 Mesocarp thickness

Thin mesocarp thickness is desirable character for grain mold resistance. Therefore, negative gca effects are appreciated. The gca estimates for mesocarp thickness for F_1 diallel set is presented in (Table 24). It is revealed from the data that, four parental lines exhibited significant negative gca effects SRT-26 B (-6.767), AKms-14B (-3.499), SPV-946 (-3.475) and SPV-1201 (-2.605). Thus these parental lines can be identified as the best general combiners for introducing thin mesocarp.

4.5.10 Threshed grain mold rating

Low thresh grain mold rating is desirable character, negative gca effect of higher magnitude are desirable. In F_1 diallel (Table 24) none of the parental line exhibited significant negative gca effects. However, parental line SPV-946 exhibited negative gca effects (-0.038). In F_2 diallel set (Table 25) none of parental line exhibited significant negative gca effects.

4.5.11 Seed germination

In F_1 diallel set (Table 24) all the parental lines exhibited significant negative gca effects. The range was from (-3.858) in SPV-946 to (-17.589) in IC5B-101 B. The maximum significant gca effects was exhibited parental lines by IC5B-101B (-17.589) followed by IS-6335 (-15.505). In F_2 diallel set (Table 25) all the parental lines recorded significant negative gca effects. The range varies from (-5.933) in IS-2284 to (-16.271) in AKms-14 B.

4.5.12 Infection by *F. moniliforme*

In F_1 diallel set only two parental lines GJ-35-15-15 (-0.934) and SPV-946 (-0.708) exhibited negative gca effects and can be considered as good general combiners for this traits (Table 24).

In F_2 diallel set (Table 25) seven parental lines viz. IS-9471 (-15.954), IS-6335 (-14.268), IS-2284 (-13.818), SPV-104 (-10.343), GJ-35-15-15 (-7.828), SPV-1201 (-3.558) and SPV-946 (-2.989) exhibited negative significant gca effects. Thus these parents can be identified as good general combiners for grain mold resistance against the infection *F. moniliforme*. Parental lines IS-9471 (-15.954), IS-

6335 (-14.268) and **IS-2284 (-13.818)** exhibited highest significant *gca* effects proving its significance over other parental lines and hence can be considered as best general combiners for this traits.

4.5.13 Infection by *F. pallidoroseum*

In **F₁ diallel set**, five parental lines **AKms- 14 B (-3.555)**, **SPV-104 (-2.195)**, **IS-2284 (-1.164)**, **IS-9471 (-1.002)** and **SPV-946 (-0.985)** exhibited significant negative *gca* estimates and hence can be identified as good general combiners (Table 24). In **F₂ diallel set** (Table 25) only two parental lines **IS-2284 (-0.493)** and **IS-6335 (-0.329)** were the good general combiners giving highest negative *gca* effects.

4.5.14 Infection by *C. lunata*

In **F₁ diallel set**, only one parental line **SPV-104 (-2.941)** exhibited significant negative *sca* effects, and hence can be considered as best general combiner for this trait (Table 24). In **F₂ diallel set** two parental lines **SRT -26 B(-1.061)** and **IS-6335 (-0.505)** exhibited negative *gca* effects, hence can be considered good general combiners for this trait (Table 25).

4.5.15 Infection by other fungi

In **F₁ diallel set**, nine parental lines exhibited significant negative *gca* effects. First three higher magnitude parents were **SPV-1201 (-6.727)**, **AKms-14B (-6.336)** and **SRT-26B (-4.896)**. Thus these can be used for best combines for this trait (Table 24). In **F₂ diallel set**, among 10 parental lines five **ICSB-101B (-5.706)**, **SRT-26B (-3.696)**, **AKms- 14 B (-3.168)**, **SPV-1201 (-2.463)** and **IS-6335 (-1.619)** exhibited negative and significant *gca* effects and thus can be identified as a best general combiners for this trait (Table 25).

Biochemical characters

The general combining effects estimates were computed for biochemical characters viz. proteins, soluble sugars, tannins and flavan-4-ols for Akola 1995 location and are presented in (Table 26).

4.5.16 Protein

It is observed from (Table 26), that all the parental lines exhibited highly significant *gca* in desirable direction however, parental lines **AKms-14B** exhibited the highest

(0.801) significant gca in desirable direction followed by SRT-26 B (0.787), ICSB-101B (0.637), GJ-35-15-15 (0.619), and SPV-946 (0.616) and thus they can be identified as good general combiners for proteins content.

4.5.17 Soluble sugars

All parental lines except two exhibited highly significant gca in desirable direction however, highest (0.280) significant gca in desirable direction was exhibited by parental lines AKms-14B, followed by SRT-26B (0.197), SPV-104 (0.167), SPV-946 (0.112) and ICSB 101 B (0.109) indicating that they are best general combiners for this character (Table 26).

4.5.18 Tannins

Out of 10 parents only three parental lines IS-2284 (0.815), IS-6335 (0.790), and IS-9471 (0.590) exhibited significant gca in desirable direction, indicating thereby, that they are best general combiners. Remaining parents exhibited significant or non significant gca effects in negative direction. Hence they can be said as poor general combiners (Table 26).

4.5.19 Flavan-4-ols

Out of 10 parental lines, only three parental line viz. IS-9471 (1.851), IS-6335 (0.578) and IS-2284 (0.469), exhibited significant gca effects in desirable direction and hence considered as good combiners. However, all others seven parental lines exhibited significant gca effects in negative direction and hence can be said as poor general combiners (Table 26).

4.6 Estimates of specific combining ability effects

The specific combining effects were calculated for 11 characters in F_1 and seven characters in F_2 for pooled over locations. The character wise results are presented below.

4.6.1 100-grain weight

The relative estimate due to sca effects for 100 grain weight is presented in (Table 27). In F_1 diallel set four crosses exhibited significant positive sca effects for 100-grain weight. The highest significant positive sca effects was exhibited by SPV-1201x SRT-26 B (0.336) followed by AKms-14B x IS-6335 (0.296), ICSB-101B x SPV-104

(0.227) and SPV-1201x SPV-104(0.207) and hence these crosses can be identified as a **best specific combinations** for 100-grain weight. Negative and significant sca estimates were exhibited by three crosses SPV-1201x SPV-946 (-0.326), ICSB-101 B x AKms-14 B (-0.227) and ICSB-101B x SRT-26B (-0.224) which proved poorer. Rest crosses recorded non significant sca effects.

4.6.2 Grain hardness

Specific combining ability effects with positive and significant values were exhibited by two crosses (Table 27) out of 45 crosses in F₁ diallel set. The highest positive sca effects was observed in SPV-1201 x GJ-35-15-15 (0.948) followed by AKms-14B x IS-2284 (0.760) hence, these two crosses appeared to be **best specific combinations for grain hardness**. On the contrary three crosses exhibited negative and significant sca effects and hence these are the poorer for grain hardness. In F₂ diallel set (Table 29) only one progeny SPV-946 x IS-2284 (1.098) exhibited significant and positive sca effects, hence can be observed as best specific combination for grain hardness.

4.6.3 Endosperm texture

Corneous endosperm texture is considered as a desirable character for grain mold resistance. Significant and negative sca effects are therefore considered for assessing crosses. In F₁ diallel set three crosses exhibited significant negative sca effect for endosperm texture. The highest significant negative sca effect was exhibited by cross IS-6335 x IS-9471 (-7.604) followed by IS-2284 x IS-6335 (-7.354) and SPV-1201 x GJ-35-15-15 (-6.283). These three crosses appeared to be best specific combinations for this trait (Table 27).

4.6.4 Electrical conductivity

Lower electrical conductivity grain leachates is considered as a desirable character for grain mold resistance. Significant and negative sca effects are therefore considered for assessing crosses. In F₁ diallel set 11 crosses exhibited significant and negative sca effects (Table 27). The cross SPV-104 x IS-9471 (-120.063) was found most promising and significantly superior to remaining 10 crosses. Some of these ten showing better combinations were SPV-104 x IS-6335 (-76.813), AKms-14B x IS-6335 (-68.00) AKms-14B x IS-9471 (-68.00), SRT 26-B x SPV-104 (-66.917) and

SRT-26B \times IS-6335 (61.146). Three crosses yielded positive and significant sca effects. Rest of crosses did not show any significance in their sca effects for this trait.

In F_2 diallel set (Table 29) seven crosses exhibited significant and negative sca effects. These crosses were AKms- 14 B \times IS-6335 (-76.284), followed by SRT-26 \times IS-9471 (-66.722), SRT-26B \times GJ-35-15-15 (-50.451), SPV-1201 \times IS-6335 (-49.909), SPV-104 \times IS-9471 (-46.576), GJ-35-15-15 \times IS-2284 (-45.867) and SPV-946 \times IS-9471 (-38.347). As against this nine crosses exhibited significant positive sca effects and hence appears poor combinations for this trait.

4.6.5 Days to 50% flowering

Since selection of genotype towards early maturity is desirable in sorghum, negative and significant sca effects are therefore considered for assessing crosses. In F_1 diallel set 11 crosses exhibited significant and negative sca effects (Table 27). The highest significant negative sca effects for first five crosses were exhibited by SPV-104 \times IS-9471 (-7.813) followed by SPV-104 \times IS-6335 (-4.750), SPV-946 \times IS-2284 (-4.646), SPV-946 \times IS-9471 (-4.646) and SRT-26-B \times IS-6335 (-4.542) hence, these crosses may be good for earliness. Seven crosses exhibited significant and positive sca effects, while remaining crosses exhibited non-significant sca effect.

4.6.6 Plant height

Among 45 crosses, negative and significant sca effects were found in 10 crosses (Table 27) in F_1 diallel set. The highest negative and significant sca effects was exhibited in SPV-1201 \times SPV - 946 (-54.208) followed by SRT-26B \times GJ-35-15-15 (-34.000) and ICSB-101 B \times AKms-14B (-30.359) exhibited significantly superior negative sca effects over seven crosses. Considering higher magnitude of negative sca effects these crosses proved to be best specific combinations.

Twenty two crosses exhibited significant and positive sca effects, the highest recorded by SRT-26B \times IS-2284 (49.688) and lowest by SPV-104 \times IS-6335 (14.333). Rest parents have exhibited non-significant sca effects. In F_2 diallel set (Table 29) only two progenies exhibited significant and negative sca effects SRT 26 \times GJ-35-15-15 (-31.463) and SPV-946 \times IS-2284 (-21.617) which proved best specific

combinations. However, eight progenies recorded significant and positive sca effects, **which appears** poor combinations for this trait. The highest significant positive sca effects was recorded in SPV-946 \times IS-6335 (50.985) and lowest by SPV-1201 \times GJ-35-15-15(20.222).

4.6.7 Cob length

In F_1 diallel set the range for sca effects was from -0.037 (GJ-35-15-15 \times SPV-104) to 2.455 (SPV-104 \times IS-6335). Four crosses exhibited significant and positive sca effects SPV-104 \times IS-6335 (2.455), AKms-14B \times GJ-35-15-15 (2.316), SPV-1201 \times IS-9471 (2.280) and ICSB 101 B \times SRT -26B (2.103) which appeared best specific combinations. Three crosses exhibited significant and negative sca effects GJ-35-15-15 \times SPV-946 (-2.972), IS-2284 \times IS-9471 (-2.166) and SPV-1201 \times ICSB-101B (-2.161), categorised as undesirable for sca effects combinations. Rest of parents have exhibited non significant positive or negative sca effects (Table 27).

In F_2 diallel set (Table 29) three progenies exhibited significant and positive sca effects GJ-35-15-15 \times IS-2284 (3.722), SPV-946 \times IS-6335 (2.645) and ICSB-101B \times AKms-14B, (1.872) all these crosses may be considered as better specific combinations for this trait. Only one progeny exhibited significant and negative sca combination SRT-26B \times IS-2284 (-2.628).

4.6.8 Glume covering

Maximum glume covering is considered as desirable character. Significant and positive sca effects. The highest significant positive sca effects was exhibited ICSB-101B \times IS-6335 (11.553) followed by AKms- 14 B \times SPV-946 (11.032) and AKms-14B \times GJ-35-15-15 (10.511) thus these crosses appeared best combinations for glume covering trait. Four crosses exhibited significant negative sca effects. While rest did not exhibited any significant sca effects.(Table 27).

Among 45 progenies in F_2 diallel set three exhibited significant sca effects in positive direction. The highest was exhibited by SPV-946 \times IS-6335 (11.617) followed by SPV-104 \times IS-2284 (10.180) and GJ-35-15-15 \times IS-2284 (8.388). Four progenies recorded significant negative sca effects. (Table 29)

4.6.9 Mesocarp thickness

Thin mesocarp thickness is desirable character for grain mold resistance. Therefore, **negative gca** effects are appreciated. The sca effects for mesocarp thickness for F_1 diallel set is presented in (Table 28). It is revealed from the data that, 22 crosses exhibited significant and negative sca effects. Few of these higher magnitude crosses were SPV-104 \times IS-2284 (-26.604), ICSB-101B \times IS-6335 (-22.195), ICSB-101B \times IS-2284 (-22.031) and SPV-104 \times IS-6335 (-21.485). Thus these crosses appeared **best combinations** for mesocarp thickness. Twelve crosses exhibited significant and **positive sca** effects, cross SPV-1201 \times AKms-14B exhibited highest (19.318) while, cross GJ-35-15-15 \times SPV-104 exhibited lowest (6.334) value. Thus these crosses are **of no use** for this trait. Rest did not exhibited any significant sca effects.

4.6.10 Threshed grain mold rating

Low TGMR is a desirable character, negative sca effect of higher magnitude are **desirable**. In F_1 diallel set Among 45 crosses, 11 crosses exhibited significant and **negative sca** effects which range from -0.913 in (AKms- 14B \times IS-6335) to -0.319 (SPV-1201 \times GJ-35-15-15). Crosses, AKms-14B \times IS-6335 (-0.913), AKms- 14 \times IS-9471 (-0.871) and AKms-14B \times IS-2284 (-0.767) of higher magnitude appeared **best specific combinations** for this trait. Eight crosses exhibited significant and **positive sca effects** which range from 0.733 in (IS-2284 \times IS-9471) to 0.420 in (GJ-35-15-15 \times SPV-946). Hence these crosses can be grouped under poor specific combinations for this trait (Table 28).

In F_2 diallel set, only four progenies exhibited significant and negative **sca effects** the highest exhibited by SPV-1201 \times ICSB-101B (-0.388) followed by **AKms-14B \times IS-9471** (-0.381), ICSB-101 B \times SPV-104 (-0.367) and AKms- 14 B \times IS-6335 (-0.346) hence proved to be best specific combinations for this trait (Table 29). Rest all others have exhibited negative and positive non-significant sca effects, hence these progenies appear undesirable.

4.6.11 Seed germination

Positive and significant sca effects for germination percentage was exhibited in 15 crosses in F_1 diallel set (Table 28). The highest being 15.185 (AKms-14B \times IS-6335), followed by AKms-14B \times IS-9471 (12.769), IS-2284 \times IS-6335 (10.839), ICSB-101B \times IS-9471 (10.526) and SPV-104 \times IS-9471 (10.111) ranked 1st to 5th and exhibited the highest positive and significant sca estimates, hence these crosses can therefore be identified as best specific combinations for improving germination. Negative and significant sca effects were exhibited by four crosses which appear to be poor combinations for this trait.

In F_2 diallel set eight progenies exhibited positive and significant sca effects (Table 29). Highest significant sca effects were exhibited by AKms-14B \times IS-6335 (9.286) followed by AKms-14B \times GJ-35-15-15 (8.175), SPV-1201 \times SRT-26B (8.170), SPV-946 \times IS-6335 (8.017), ICSB-101B \times AKms-14B (7.184), SPV-946 \times SPV-104 (7.053), AKms-14B \times IS-9471 (6.604) and SPV-1201 \times SPV-104 (6.478). Hence these crosses proved best specific combinations for improving germination. All others have exhibited non significant positive and negative sca effects.

4.6.12 Infection by *F. moniliforme*

Minimum fungal load is a desirable character for grain mold resistance. Significant and negative sca effects are therefore, considered for assessing crosses.

In F_1 diallel set, four crosses ICSB-101B \times IS-6335 (-10.017), SPV-104 \times IS-6335 (-9.795), AKms-14B \times IS-6335 (-9.259) and SPV-104 \times IS-2284 (-8.392) exhibited significant negative sca effects hence considered as best specific combinations for introducing resistance against *F. moniliforme*. Rest crosses exhibited non-significant and negative sca effects or positive sca effects, hence cannot be considered for this trait (Table 28).

In F_2 diallel set sixteen progenies exhibited negative sca effects. Considering higher magnitude of negative sca effects, first three good sca

combinations for this trait were SRT-26B \times SPV-104 (-6.582), SRT-26B \times SPV-946 (-6.425) and SPV-1201 \times IS-2284 (-5.028) (Table 29).

4.6.13 Infection by *F. pallidroseum*

Negative and significant sca effects are considered for evaluating the crosses in respect of fungal load of *F. pallidroseum*. In F_1 diallel set, three crosses ICSB-101B \times IS-2284, SPV-946 \times SPV-104 and GJ-35-15-15 \times SPV-946 exhibited negative and significant sca effects, hence these can be considered, good for specific combinations for introducing resistance against *F. pallidroseum* (Table 28). In F_1 diallel set only one progeny (ICSB-101 B \times IS-6335) exhibited negative and significant sca effects (-2.567) and hence this can be considered the best specific combinations (Table 29).

4.6.14 Infection by *C. lunata*

Thirty crosses exhibited negative sca effect in F_1 diallel set (Table 28) considering higher magnitude of negative sca effects, first three good sca crosses considers for this trait were, ICSB-101B \times SPV-104, SPV-946 \times IS-2284) and SPV-1201 \times IS-9471 (Table 29.) In F_2 diallel set, two progenies SRT-26B \times GJ-35-15-15 and AKms-14B \times SPV-104 exhibited significant negative sca effects, which can be considered as good sca combinations for introducing resistance against *C. lunata* (Table 29)

4.6.15 Infection by other fungi

Twenty crosses exhibited negative sca effects in F_1 diallel set (Table 28) considering higher magnitude of negative sca effects, first three good combinations for this trait were GJ-35-15-15 \times IS-2284, ICSB-101B \times SPV-104 and IS-6335 \times IS-9471. Three crosses SPV-1201 \times SRT-26B, AKms- 14 B \times IS-6335 and SPV-1201 \times SPV-946 exhibited significant positive sca effects, hence were poor combinations. In F_2 diallel set 21 progenies exhibited negative sca effects. Considering higher magnitude of negative sca effects, first three good combinations for this trait were AKms-14B \times SRT-26 B, AKms-14B \times GJ-35-15-15 and AKms-14B \times SPV-946. One progeny AKms-14B \times IS-9471 exhibited significant positive sca effects (Table 29).

Biochemical characters

The specific combining ability effect were calculated for 4 biochemical characters viz. **Proteins, soluble sugars, tannins and flavan-4-ols** for Akola 1995 location and are presented in (Table 30).

4.6.16 Proteins

Data (Table 30), revealed that the highest positive sca effect was observed in SPV-1201 \times IS-9471 whereas, lowest was observed in SPV-1201 \times GJ-35-15-15. Among 45 crosses eight crosses exhibited positive and significant sca effect. SPV-1201 \times IS-9471 exhibited highest (0.278) sca effect followed by IS-6335 \times IS-9471(0.240), SPV-1201 \times IS-2284) (0.181), AKms-14B \times IS-6335 (0.175), SPV-104 \times IS-6335 (0.175), SPV-104 \times IS-2284 (0.119), IS-2284 \times IS-9471 (0.106) and SPV-946 \times SPV-104 (0.070). Therefore these crosses can be identified as a best specific combines can be identified as best specific combinations for protein content. On the contrary 31 crosses exhibited negative and significant sca effects, hence these can be treated as poor specific combinations for protein content.

4.6.17 Soluble sugars

Twenty crosses exhibited significant sca effects. Out of which only three exhibited significant positive sca effects GJ-35-15-15 \times SPV-104 (0.095), SPV-1201 \times IS-6335 (0.089) and AKms- 14B \times IS-9471 (0.080), hence these crosses can be identified as good specific combinations for soluble sugars (Table 30). On the contrary 11 crosses exhibited negative and significant sca effects, the highest being observed in AKms-14B \times SRT-26B followed by AKms-14B \times GJ-35-15-15 and SPV-104 \times IS-6335, hence these crosses can be treated as poor specific combinations for soluble sugars content.

4.6.18 Tannins

Out of 45 crosses, 20 crosses exhibited significant and positive sca effects and 20 crosses exhibited significant sca effect n negative direction. Crosses IS-2284 \times IS-6335 exhibited highest positive and significant sca effects (0.763) followed by AKms-14B \times IS-2284 (0.344), SRT-26 \times IS-9471 (0.302) SRT-26 \times IS-6335 (0.299), AKms-

14B \times IS-9471 (0.283), SPV-1201 \times IS-9471 (0.244), SPV-104 \times IS-6335 (0.234) and SPV-104 \times IS-9471 (0.233). Therefore these crosses can be identified as a best specific combinations for tannin content (Table 30). On the other hand out of 20 crosses which exhibited negative and significant sca effects, the highest being observed in IS-2284 \times IS-9471 (-0.408) followed by IS-6335 \times IS-9471 (-0.291), AKms-14B \times SRT-26B (-0.146), and SPV-1201 \times ICSB-101B (-0.125) can be treated as poor specific combinations for tannins content.

4.6.19 Flavan-4-ols

Significant sca effect in positive direction were exhibited by 17 crosses (Table 30). These crosses exhibited the range of 0.072 to 0.635 sca effects, the highest being exhibited by AKms-14B \times IS-2284 (0.635), followed by SRT-26B \times IS-6335 (0.596), SPV-946 \times IS-6335 (0.586), SPV-104 \times IS-6335 (0.530), SPV-104 \times IS-9471 (0.426), AKms-14B \times IS-9471 (0.405) and SPV-104 \times IS-2284 (0.391), hence these crosses can be identified as best specific combinations for flavan-4-ols content.

However, 16 crosses exhibited significant negative sca effects. These crosses exhibits the range of -0.075 to -0.749 sca effects, the highest being indicated by IS-2284 \times IS-6335 and lowest by ICSB 101B \times SPV-104. Hence these can be treated as poor specific combinations for flavan-4-ols content.

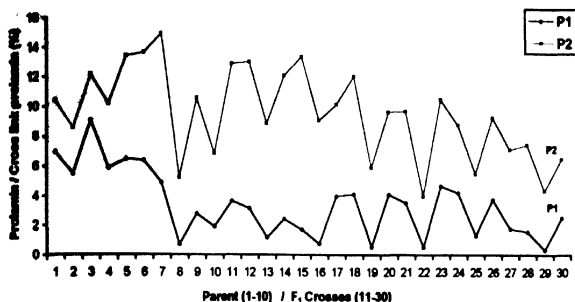
4.7 Protein fractions

4.7.1 Albumin and Globulin

Protein fractionation was carried out for all the parental lines and F_1 crosses (20) of Akola 1995 season. Results are presented in (Table 31). It is observed from the data that, the range of variation for albumin and globulin for parental lines was from 1.81 (IS-2284) to 20.63% in SRT-26B. For crosses, the range was from 7.90 (IS-2284 \times IS-9471) to 16.74% (SPV-946 \times SPV-104).

4.7.2 Prolamin

It is observed from (Table 31 and Fig. 4), that maximum (9.09 %) prolamin was recorded in parental lines AKms-14B followed by SPV-1201 (6.96 %) and GJ-35-15-15 (6.47 %), however, minimum prolamin was observed in IS-2284 (0.70 %).



P1 = Prolamin P2 = Cross link prolamin

Figure 4. Prolamin and Cross link prolamin of parents and F₁ crosses at Akola 1995

1 SPV-1201	11 SPV-1201 x SPV-946	21 SR I-2613 x SPV-104
2 ICSB-101B	12 SPV-1201 x SPV-104	22 SR I-2613 x IS-9471
3 AKms-14B	13 SPV-1201 x IS-9471	23 GJ-35-15-15 x SPV-946
4 SRT-26B	14 ICSB-101B x SPV-946	24 GJ-35-15-15 x SPV-104
5 GJ-35-15-15	15 ICSB-101B x SPV-104	25 GJ-35-15-15 x IS-9471
6 SPV-946	16 ICSB-101B x IS-9471	26 SPV-946 x SPV-104
7 SPV-104	17 AKms-14B x SPV-946	27 SPV-946 x IS-9471
8 IS-2284	18 AKms-14B x SPV-104	28 SPV-104 x IS-9471
9 IS-6335	19 AKms-14B x IS-9471	29 IS-2284 x IS-9471
10 IS-9471	20 SR I-2613 x SPV-946	30 IS-6335 x IS-9471

Among crosses, cross (GJ-35-15-15 \times SPV-946) exhibited maximum prolamin (4.72 %), however, minimum prolamin was recorded in (IS-2284 \times IS-9471) 0.38 %.

4.7.3 Cross-link prolamin

It is observed (Table-31, Fig. 4), that parental lines SPV-104 exhibited maximum (14.92 %) cross-link prolamin followed by SPV-946 (13.62 %) and GJ-35-15-15 (13.39 %), minimum was observed in IS-2284 (5.24 %). Among crosses, maximum (13.09 %) cross-link prolamin was recorded in SPV-1201 \times SPV-104 followed by SPV-1201 \times SPV-946 (12.89 %), minimum (3.99 %) was recorded in SRT-26B \times IS-9471.

4.7.4 Glutelin-like

It is observed (Table 31) that parental lines IS-6335 exhibited maximum (12.69%) glutelin-like, followed by IS-2284 (12.39%), minimum value (2.46%) was recorded in AKms-14B. Among crosses, maximum (10.88%) glutelin-like was indicated in SPV-1201 \times IS-9471 and minimum (0.96 %) in IS-6335 \times IS-9471.

4.7.5 Glutelin

It is observed from the data (Table 31), that, the range of variation for parental lines was from 27.90 (ICSB-101 B) to 68.27% (IS-2284). Among crosses, the range of variation was from 26.62 (SRT-26B \times IS-104) to 51.96% (SPV-946 \times IS-9471).

4.7.6 Residues

It is indicated (Table 31) that the range of variation for parental lines was from 7.26 (IS-2284) to 11.46% (AKms-14B). Among crosses, the range of variation was from 2.70 (SPV-1201 \times SPV-104) to 11.46% (SRT-26B \times IS-9471).

4.8 Glume color, grain color and testa

It is observed from data (Table 32) that three parental lines IS-2284, IS-6335 and IS-9471 with colored grain had a testa layer or sub-coat, whereas, seven parental lines with white grain did not show black reaction with bleach test indicating absence of testa. Among F_1 crosses, 24 (crosses with colored parental lines) out of 45 crosses observed colored grain and exhibited black reaction with bleach test indicating presence of testa. Remaining 21 crosses with white grain did not show black reaction

with bleach test thus confirm the absence of testa layer. The above observations were confirmed by section cutting, where colored grain observed to have testa layer or sub-coat and white grain did not have the testa layer. Observations on glume cover were recorded just to provide additional information to the breeder.

4.9 Fungal load on germinated and ungerminated seeds (pre-treated and untreated)

Effect of seed pre-treatment (0.1% HgCl_2) on germination and seed mycoflora of both germinated and ungerminated seed of parental lines, F_1 crosses and F_2 progenies from Akola and Patancheru centres during 1996 were studied. Comparison of two seasons at Akola centre of parental lines and F_1 crosses during 1995 and 1996 were also studied. The details of observations have been narrated under following subheads.

- Fungal load on germinated seeds, Akola and Patancheru, 1996
- Fungal load on ungerminated seeds, Akola and Patancheru, 1996
- Fungal load on germinated seeds, Akola, 1995 and 1996
- Fungal load on ungerminated seeds, Akola, 1995 and 1996

4.9.1 Fungal load on germinated seed, Akola and Patancheru, 1996

4.9.1.1 Seed germination

Higher germination of the seed is the desired trait for the breeders as well as farmers. Grain mold is an important factor in reduction in germination in sorghum. Many molding fungi remain confined to the surface and hence effect of pre treatment of seed was studied. Method of pre-treatment of seed is described in materials and methods.

Germination percent recorded on standard blotter tests method on treated and untreated seed. In F_1 diallel set (Table 33), the range of variation for parental lines was exhibited from 59.57 to 93.67, 5.00 to 80.67 and 33.00 to 87.00% at Akola, Patancheru and in pooled data, respectively. Parental line IS-9471 exhibited highest 93.67 % germination followed by IS-6335 (93.33%) at Akola location, at Patancheru parental line IS-6335 recorded highest (80.67%) germination followed by IS-2284 (74.33%). In pooled data, parental IS-6335 recorded highest germination (87.00%) followed by IS-9471 (83.83%).

Among crosses the highest seed germination was recorded in AKms 14B × IS-6335, IS-6335 × IS-9471 (95.33%) followed by SPV-1201 × IS-6335 (94.67%) at Akola location. At Patancheru cross ICSB-101B × IS-6335 recorded highest seed germination (92.00%) followed by SRT-26B × IS-6335 (91.00%) and ICSB-101B × IS-9471 (89.67%). However, in pooled data ICSB 101B × IS-6335 recorded highest seed germination (91.83%) followed by AKms-14B × IS-6335 (91.67%) and ICSB-101B × IS-9471 (91.34%). Over all, there was significant increase in germination 2.28, 5.51 and 3.64% at Akola, Patancheru and in pooled data, respectively, in treated seed with mercuric chloride over untreated control, and there was 27.39 % increased in germination at Akola over Patancheru.

In F₂ diallel (Table 45) at Akola, progeny IS-2284 × IS-6335 recorded highest germination (92.13%) followed by ICSB-101 B × IS-9471 (91.90 %) and IS-6335 × IS-9471 (91.79%). At Patancheru, progeny GJ-35-15-15 × IS-2284 recorded highest seed germination (91.50%) and SPV-946 × IS-6335 (89.50%). However, in pooled data IS-2284 × IS-9471 recorded highest seed germination (90.53%) followed by SPV-946 × IS-6335 (90.45 %) and GJ-35-15-15 × IS-2284 (89.35%). Over all, there was significant increase in germination 4.45, 6.56 and 5.35 % at Akola, Patancheru and in pooled data, respectively, in treated seed with mercuric chloride over untreated control and there was 24.69 % reduction in germination at Patancheru over Akola.

4.9.1.2 Infection by *F. moniliforme*

In grain mold of sorghum low fungal load is considered as a desirable characters. It is seen from the data (Table-34), that the range of variation for parental lines was exhibited from 3.33 in (IS-9471) to 23.67 % in (SPV-104) at Akola, 2.67 in (AKms-14B) to 22.33% (IS-6335) at Patancheru and 10.7 (IS-9471) to 17.17% (IS-2284) in pooled data. Among crosses the lowest fungal load of *F. moniliforme* was noticed in SPV-104 × IS-6335 (3.33%) followed by IS-6335 × IS-9471 (4.01%) and IS-2284 × IS-6335 (4.34%) at Akola, whereas, at Patancheru AKms 14 B × SPV-104 (9.33)

exhibited lowest *F. moniliforme* load followed by AKms 14 B × SPV-946 (11.33 %) in pooled data IS-6335 × IS-9471 (9.84%) recorded lowest *F. moniliforme* load.

Over all, there was significant reduction in *F. moniliforme* load 14.75, 44.75 and 35.06 % at Akola, Patancheru and in pooled data, respectively in treated seed with mercuric chloride over untreated control and there was 86.28% increased *F. moniliforme* load at Patancheru over Akola location.

In F_2 diallel (Table 46) progeny SRT-26B × SPV-946 exhibited (10.93) *F. moniliforme* load followed by ICSB-101 B × IS-6335 (11.43%) at Akola, whereas, at Patancheru IS-6335 × IS-9471 recorded lowest (14.50%) *F. moniliforme* count followed by ICSB-101 B × SPV-104 (15.50%), in pooled data AKms 14 B × SRT-26 B recorded lowest (16.2%) *F. moniliforme* load followed by SRT-26 B × SPV-104 (16.70%).

Over all, there was significant reduction in *F. moniliforme* load 27.10, 17.98 and 22.98% at Akola, Patancheru and in pooled data, respectively in treated seed over untreated control and in general there was 28.32 % increase in load at Patancheru over Akola.

4.9.1.3 Infection by *F. pallidoroseum*

In F_1 diallel it is observed from data (Table 35), that the range of variation for parental lines was from 0.33% to 3.33% at Akola, 0.0 to 9.00% at Patancheru. However in pooled data the range of variation was 1.33 to 5.50%. Parental line SPV-1201 exhibited lowest *F. pallidoroseum* load at Akola (0.33%) and AKms 14B (1.33%) in pooled data. For crosses the lowest *F. pallidoroseum* load was recorded in cross ICSB 101B × GJ-35-15-15 and SRT-26B × GJ-35-15-15 (0.00%) at Akola and AKms-14 B × SPV-946 (2.67%) at Patancheru. Cross AKms 14 B × SPV-946 (1.50%) also exhibited lowest *F. pallidoroseum* load in pooled data.

Over all, there was significant reduction in *F. pallidoroseum* load 3.64, 36.28 and 21.75% at Akola, Patancheru and in pooled data, respectively in treated seed with mercuric chloride over untreated control and there was 165.27 % increase *F. pallidoroseum* load at Patancheru over Akola.

In F_2 diallel (Table 47) progeny GJ-35-15-15 \times SPV-946 exhibited lowest *F. pallidoroseum* load (0.34%) followed by SPV-1201 \times SPV-946 and AKms 14 B \times SPV-946 (0.60%) at Akola whereas, at Patancheru AKms 14 B \times SRT 26B exhibited lowest *F. pallidoroseum* load (0.50%) followed by ICsB 101 B \times SPV-104 (0.75%) and AKms 14 B \times GJ-35-15-15 (0.92%). Progeny GJ-35-15-15 \times SPV-946 exhibited lowest *F. pallidoroseum* load (0.80%) followed by SPV-1201 \times AKms-14B (0.99%) and AKms 14 B \times GJ-35-15-15 (1.06%) in pooled data.

Over all, at Akola, Patancheru and pooled data exhibited 4.80, 35.48 and 21.30% reduction in fungal load of *F. pallidoroseum*, respectively in treated seed with mercuric chloride over untreated control and in general, there was 36.60% increase in *F. pallidoroseum* load at Patancheru over Akola.

4.9.1.4 Infection by *C. lunata*

In F_1 diallel set, it is seen from data (Table 36), that the range of variation for *C. lunata* in parental lines was exhibited from 18.00 to 45.35% at Akola, whereas, it was 1.33 to 18.00 and 12.00 to 31.17% at Patancheru and in pooled data, respectively. Parental line IS-6335 exhibited lowest *C. lunata* load (18.00%) at Akola, however parental line SPV-104 recorded lowest (1.33 and 12.00%) load at Patancheru and in pooled data. Among crosses the lowest *C. lunata* load was exhibited by SPV-1201 \times IS-6335 (12.67%) and SPV-1201 \times IS-9471 (12.68%) at Akola. Crosses AKms-14B \times GJ-35-15-15 and AKms-14B \times SPV-104 exhibited lowest (8.67%) followed by ICsB-101B \times SPV-104 (9.33%) at Patancheru. Whereas, SPV-1201 \times IS-9471 exhibited lowest (14.01%) *C. lunata* load followed by SPV-1201 \times IS-6335 (14.67%) in pooled data.

Over all, there was 20.69 and 7.44% reduction in *C. lunata* load at Akola and in pooled data, respectively in treated seed with mercuric chloride over untreated control. However, at Patancheru there was 21.73 increased in *C. lunata* load and there was 108.95% increase in *C. lunata* load at Akola over Patancheru.

In F_2 diallel set (Table 48) the range was from 15.17 to 47.03%, progeny IS-6335 \times IS-9471 exhibited lowest (15.17%) *C. lunata* load followed by IS-2284 \times IS-9471 (17.71%) at Akola. At Patancheru the range varied from 4.00 to 29.76%, lowest (4.00%) *C. lunata* load was exhibited in AKms 14 B \times SRT 26B. However, progeny IS-6335 \times IS-9471 exhibited lowest *C. lunata* load (15.50 %) in pooled data.

Over all, there was reduction in *C. lunata* load 19.36, 2.44 and 13.81 % at Akola, Patancheru and in pooled data, respectively in treated seed over untreated control and in general there was 46.43 % increase in *C. lunata* load at Akola over Patancheru.

4.9.1.5 Infection by other fungi

In F_1 diallel set it is observed from (Table 37) that the range of variation for parental lines was from 6.33 to 19.33% at Akola, 0.00 to 5.33% at Patancheru and 3.17 to 11.00% in pooled data. Parental line AKms - 14 B recorded lowest other fungi load (6.33, 0.00 and 3.17%) at both locations and in pooled data, respectively. Among crosses lowest other fungi load was exhibited by AKms- 14B \times SPV-104 (0.33 and 5.83%) at Akola and in pooled data. However, cross KCSB-101B \times SPV-104 exhibited lowest (0.00%) followed by AKms 14 B \times SRT-16B (0.33%) at Patancheru.

Over all, there was significant reduction in other fungi load 39.16, 22.89 and 36.09 % at Akola, Patancheru and in pooled data, respectively in treated seed with mercuric chloride over untreated control, and there was 283.44% increase in other fungi load at Akola over Patancheru. In F_2 diallel (Table 49), progeny IS-6335 \times IS- 9471 exhibited lowest other fungi load (9.17%) followed by SPV-104 \times IS-9471 (9.73 %) and IS- 2284 \times IS- 6335 (9.94%) at Akola. Whereas, progeny AKms - 14 B \times SRT -26B exhibited lowest other fungi load (0.50%) and (6.85%) at Patancheru and in pooled data, respectively. In treated seed with mercuric chloride over untreated control, over all, there was 6.37, 4.91 and 14.44% reduction in other fungi load at Akola, Patancheru and in pooled data, respectively and there was 46.50% reduction in other fungi load at Patancheru over Akola.

4.9.1.6 Score (Germinated seed)

It is observed from data (Table 38), that parental lines IS- 9471 and IS- 6335 recorded minimum (1.00 score) at both locations and in pooled data, however, parental line AKms -14 B exhibited maximum (4.25 and 3.96 score) at Akola and in pooled data and parental lines SPV -1201 and ICSB -101 B (4.00 score) at Patancheru. Thirteen crosses recorded minimum (1.00 score) and seven recorded more than (3.00 score) at Akola, at Patancheru nine crosses recorded minimum (1.00 score) and 19 crosses recorded more than (3.00) score at Patancheru, however in pooled data only four crosses recorded minimum (1.00 score) and more than (3.0 score) recorded in 14 crosses.

Over all, there was reduction in score 13.27 , 0.83 and 6.83% at Akola, Patancheru and in pooled data, respectively, in treated seed over untreated control, and there was 12.74 % increase in score at Patancheru over Akola.

In F₂ diallel set (Table 50) 27 progenies recorded more than (3.00 score) maximum (3.70 score) recorded by AKms -14 B × SPV- 104, SRT -26 B × SPV- 104 and SPV- 946 × SPV -104 and progeny IS- 2284 × IS- 9471 exhibited minimum (2.10 score) at Akola, at Patancheru, however 36 progenies exhibited more than (3.00 score), maximum (3.94 score) and minimum (1.75 score) recorded by, AKms- 14 B × SPV 104 and IS -6335 × IS-9471 at Patancheru, respectively.

Over all, there was reduction in score 17.71, 11.14 and 14.41 % in treated over untreated control at Akola, Patancheru and in pooled data, respectively. However there was 3.63 % increase in score at Patancheru over Akola.

4.9.2 Fungal load on ungerminated seed.

4.9.2.1 Ungerminated Seed

It is observed from data (Table 39) in F₁ diallel for ungerminated seed, the range of variation for parental lines was from 6.33 to 40.33% at Akola, 19.33 to 95.33% at Patancheru and 13.00 to 67.00% in pooled data. Parental line IS-9471 exhibited minimum (6.33%) and SPV-104 exhibited maximum (40.33%) ungerminated seed at Akola. At Patancheru, parental line IS-6335 exhibited minimum (19.33%) and

AKms-14B maximum (95.33 %) ungerminated seed. In pooled data parental line **IS-6335** recorded minimum (13.00%) and **SPV-104** and **AKms-14 B** recorded (67%) ungerminated seed.

For crosses the range for ungerminated seed was from (4.67%) in **AKms-14B** × **IS-6335** and **IS-6335** × **IS-9471** to (40.00%) in **AKms-14B** × **SPV-104** at Akola and (8.00%) in **ICSB 101B** × **IS-9471** to (72.67%) in **Akms-14B** × **SRT-26 B** at Patancheru.

Over all, there was 12.77, 9.06 and 10.09% decrease in ungerminated seed in treated over untreated control at Akola, Patancheru and in pooled data, respectively, and 165.24% increase in ungerminated seed at Patancheru over Akola.

4.9.2.2 Infection by *F. moniliforme*

In **F₁** diallel set, load of *F. moniliforme* on ungerminated seed for parental lines ranged from 2.33 to 18.67%, 11.67 to 54.00 and 7.17 to 35.33% at Akola, Patancheru and in pooled data. (Table 40). Parental line (**IS-9471**) recorded lowest 2.33 and 7.17% fungal load at Akola and in pooled data, however, (**IS-6335**) recorded lowest 11.67% at Patancheru. Highest fungal load was indicated by (**AKms-14B**) 18.67 % at Akola, whereas, parental line (**SPV-104**) recorded highest fungal load 54.00 and 35.34 % at Patancheru and in pooled data, respectively. Among crosses lowest fungal load was recorded by **AKms-14B** × **IS-6335** and **IS-6335** × **IS-9471** (1.67 %) at Akola, **ICSB-101 B** × **IS-9471** (5.00 %) at Patancheru and **ICSB 101B** × **IS-6335** (4.17%) in pooled data.

Over all, there was reduction in *F. moniliforme* load 13.77, 25.66 and 28.05% at Akola, Patancheru and in pooled data, respectively in treated seed with mercuric chloride over untreated control, and 241.04% more *F. moniliforme* fungal load was recorded at Patancheru over Akola. In **F₂** diallel (Table 52) progeny **ICSB-101B** × **IS-6335** recorded lowest load (2.73%) at Akola, **GJ-35-15-15** × **IS-2284** (5.50%) at Patancheru and **SPV-946** × **IS-6335** (5.31%) in pooled data. In treated seed with mercuric chloride over untreated control, over all, there was 18.36, 4.45 and 6.67% reduction in *F. moniliforme* load at Akola, Patancheru and in pooled data,

respectively. In general 165.56% more *F. moniliforme* load on ungerminated seed was recorded at Patancheru over Akola.

4.9.2.3 Infection by *F. pallidoroseum*

In F_1 diallel (Table 41) six parental lines out of 10 parental lines recorded lowest (0.00%) fungal load IS-6335, IS-2284, SPV-946, GJ-35-15-15, SRT-26B and ICSB-101B at Akola. At Patancheru and in pooled data parental lines IS-6335 recorded lowest (2.00 and 1.00%) and SPV-104 recorded highest (10.33 and 7.17%) *F. pallidoroseum* load respectively. Among crosses 21 recorded lowest (0.00%) load, at Akola, however only one cross SRT-26B \times IS-6335 exhibited lowest fungal load (0.00 and 0.17%) at Patancheru and in pooled data, respectively. Over all, there was 58.33% increase, 33.09 and 27.68% significant decrease in fungal load at Akola, Patancheru and in pooled data, respectively in treated seed with mercuric chloride over untreated control, and there was 918.89 % increase in fungal load at Patancheru over Akola.

In F_2 diallel (Table 53) 22 progenies exhibited lowest fungal load (0.00%) at Akola, however, at Patancheru and in pooled data four and three progenies recorded lowest (0.00%) *F. pallidoroseum* load, respectively. Over all, there was 87.50 and 26.52% increase in fungal load at Akola and in pooled data, respectively and 88.15% decrease in fungal load at Patancheru in treated seed with mercuric chloride over untreated control and 882.60% more fungal load was exhibited at Patancheru over Akola.

4.9.2.4 *C. lunata*

In F_1 diallel (Table 42) at Akola parental lines IS-9471 and IS-6335 exhibited lowest fungal load (3.33%) and highest in AKms-14B and SPV-104 (16.00%). At Patancheru and in pooled data parental line IS-6335 exhibited lowest (5.00 and 4.17%) and parental lines AKms-14B exhibited highest (43.00 and 29.50 %) respectively. Among crosses the lowest *C. lunata* load was exhibited by crosses SPV-1201 \times IS-6335 (1.67 %) at Akola, whereas, at Patancheru and in pooled data cross ICSB-101 B \times IS-6335 exhibited lowest fungal load (2.33 and 2.67%), respectively.

Over all, there was 8.27 % reduction in fungal load in treated seed over untreated control at Akola. However, at Patancheru and in pooled data exhibited 43.44 and 21.09% increase in fungal load in treated over untreated control, respectively, Patancheru exhibited 66.91% increase in fungal load over Akola. In F_2 diallel (Table 54) progeny IS-6335 \times IS-9471 exhibited lowest (1.68%) *C. lunata* load at Akola, however, IS-2284 \times IS-9471 recorded lowest *C. lunata* load (1.75 and 1.76%) at Patancheru and in pooled data. All the treatment exhibited 17.59, 19.93 and 15.58% reduction in fungal load at Akola, Patancheru and in pooled data, respectively in treated over untreated control and exhibited 27.92% reduction fungal load at Patancheru over Akola.

4.9.2.5 Other fungi

In F_1 diallel (Table 43) parental lines IS-2284 and IS-6335 exhibited lowest other fungi load i.e. 0.00, 0.67 and 0.33% at Akola, Patancheru and in pooled data, respectively. Among crosses AKms-14B \times IS-6335 indicated lowest (0.00%) fungal load at both locations and in pooled data.

Over all, the treatments exhibited 60.00 and 22.30% reduction in fungal load at Akola and in pooled data, respectively, however, at Patancheru location exhibited 26.89% increase in fungal load in treated over untreated control. In general Patancheru exhibited 20.53% increase in fungal load over Akola.

In F_2 diallel (Table 55) progeny IC5B-101B \times IS-9471 exhibited lowest fungal load (0.00%) at Akola, GJ-35-15-15 \times IS-2284 and SRT 26B \times IS-2284 (0.00%) at Patancheru and SPV-946 \times IS-6335 and IS-2284 \times IS-9471 (0.15%) in pooled data, over all, the treatments exhibited 19.93, 21.05 and 20.62% reduction in other fungi load in treated over untreated control. Patancheru location exhibited 26.78 % more other fungi load over Akola.

4.9.2.6 Score (Ungerminated seed)

In F_1 diallel set. (Table 44) parental lines IS-9471 exhibited minimum 1.17, 2.50 and 1.84 score and AKms-14B exhibited maximum 4.75, 5.00 and 4.88 score at Akola,

Patancheru and in pooled data, respectively. Among crosses the range of variation was from 0.92 to 4.58 at Akola, 1.42 to 5.00 at Patancheru and 1.38 to 4.71 in pooled data

Over all, there was 3.97 and 0.63 % reduction in score at Akola and in pooled data, respectively and 3.38 % increase in score at Patancheru in treated over untreated control and there was 23.32% increase score at Patancheru over Akola.

In F_2 diallel, (Table 56) all the progenies at Akola and Patancheru except one each IS-2284 \times IS-9471 (2.91) and IS-6335 \times IS-9471 (2.92), respectively recorded more than 3.00 score. However, all the progenies in pooled data recorded more than 3.00 score. Over all there was, 13.22, 6.96 and 9.72% significant reduction in score at Akola, Patancheru and in pooled data, respectively, in treated seed over untreated control and there was 11.31% increase in score at Patancheru over Akola.

4.9.3 Fungal load on germinated seed, Akola 1995 and 1996

4.9.3.1 Seed germination

Seed germination (%) recorded on standard blotter test method on treated and untreated seed were used for study. In F_1 diallel set (Table 57) parental line IS-6335 and IS-9471 exhibited highest germination (97.23%) followed by SPV-1201 (96.67 %) and lowest in parental line AKms-14B (53.34%) in Akola 1995 and Akola 1996 parental line IS-9471 recorded highest (93.67%) and lowest in parent SPV-104 (59.67%) however, in pooled data parental lines IS-9471 recorded highest (95.45%) and lowest in AKms-14B (57.33%) seed germination.

Among crosses the highest seed germination (%) were recorded in crosses SPV-946 \times IS-2284, SPV-946 \times IS-6335 and GJ-35-15-15 \times IS-6335 (97.78 %) and lowest in GJ-35-15-15 \times SPV-104 (50.00%) in Akola 1995. In Akola 1996, crosses AKms-14B \times IS-6335 and IS-6335 \times IS-9471 recorded highest (95.33%) seed germination and lowest in AKms-14B \times SPV-104 (59.67%), however, in pooled data highest % seed germination was recorded in cross SPV-946 \times IS-6335 (96.06%) followed by IS-6335 \times IS-9471 (95.45%) and lowest in GJ-35-15-15 \times SPV-104 and SPV-946 \times SPV-104 (67.33%).

Over all, there was significant increase in seed germination 7.07, 2.33 and 4.63% at Akola 1995, Akola 1996 and in pooled data, respectively, in treated seed with mercuric chloride over untreated control and there was 3.52% increase in % germination at Akola 1996 over Akola 1995.

4.9.3.2 Infection by *F. moniliforme*

In sorghum grain mold low fungal load is considered as desirable characters. In F_1 diallel set, (Table 58) parental lines GJ-35-15-15 and IS-6335 recorded lowest (1.11%) *F. moniliforme* load and highest in ICSB-101B (23.33%) in Akola 1995. In Akola 1996 the lowest load was recorded in IS-9471 (3.33%) and highest in SPV-104 (23.67%), however, in pooled data parental lines IS-9471 recorded lowest (2.50%) and ICSB-101B recorded highest (20.34%) *F. moniliforme* load.

Among crosses lowest load was exhibited by GJ-35-15-15 \times IS-6335 (1.11%) followed by ICSB-101 B \times IS-9471 (1.67%) in Akola 1995. In Akola 1996 crosses SPV-104 \times IS-9471 (3.33%) followed by IS-6335 \times IS-9471 (4.01%), however, crosses ICSB-101B \times IS-9471 exhibited lowest (3.34%) *F. moniliforme* load followed by SPV-104 \times IS-9471 (3.61 %) in pooled data.

Over all, there was significant reduction in *F. moniliforme* load 26.51, 78.49 and 54.87% in Akola 1995, Akola 1996 and in pooled data, respectively in treated seed over untreated control. However, there was 19.80 % decrease in fungal load in Akola 1996 over Akola 1995.

4.9.3.3 Infection by *F. pallidoroseum*

There was no occurrence of *F. pallidoroseum* at Akola during 1995 in all the entries except one tested (Table 59), hence no conclusion could be drawn.

4.9.3.4 *C. lunata*

It is observed from (Table 60) that the range of variation for parental lines was from 4.44% (SPV-1201) to 34.45% (SPV-104) in Akola 1995. In Akola 1996 the same was observed to be 18.00% (IS-6335) to 45.35% (SPV-1201), whereas, in pooled data range was 14.73% (IS-9471) to 35.33% (ICSB-101B). Among crosses in Akola 1995 the range was 10.00% in (SPV-946 \times IS-6335) and (SPV-104 \times IS-9471) to 38.34 %

in (ICSB-101 B \times GJ-35-15-15), however in Akola 1996 and in pooled data the range was 12.67% to (SPV-1201 \times IS-6335) to 53.33% (ICSB-101B \times GJ-35-15-15). Over all, there was 13.34, 22.36 and 17.82% reduction in fungal load in Akola 1996 in treated over untreated control and 46.85% increase in *C. lunata* load Akola 1996 over Akola 1995.

4.9.3.5 Infection by other fungi

In F_1 diallel (Table 61) in Akola 1995 season the range for parental line was from 15.00% (SPV-1201) to 64.65% (GJ-35-15-15) in Akola 1996 the range was from 6.33% (AKms-14B) to 19.33% (SPV-1201), however, in pooled data the range was 15.95% (AKms-14B) to 40.89% in (GJ-35-15-15).

Among crosses in Akola 1995 cross AKms-14B \times SR1-26B recorded lowest (10.00%) other fungi fungal load and highest in SPV-104 \times IS-2284 (32.78%) in Akola 1996 the lowest fungal load recorded in AKms-14B \times SPV-104 (10.33%) and highest in SPV-1201 \times SR1-26 B (27.33%), however, in pooled data cross ICSB-101B \times SR1-26B recorded lowest (12.28%) fungal load and highest in SPV-104 \times IS-2284 (25.89%).

4.9.3.6 Score (GS)

In F_1 diallel (Table 62) it is observed from data that the range of variation for parental lines was from 1.50 (SPV-1201) to 3.83 (ICSB-101B) in Akola 1995 from 1.00 (IS-9471 and IS-6335) to 4.25 (AKms-14B) in Akola 1996 and from 1.50 (IS-9471 and IS-6335) to 3.71 (AKms-14B) in pooled data. Among crosses the range was from 1.25 (SPV-946 \times IS-6335) to 4.24 (ICSB-101B \times SPV-104) in Akola 1995 from 1.00 (ICSB-101 B \times IS-6335) to 3.75 (AKms-14B \times SPV-104) in Akola 1996 and 1.25 (IS-6335 \times IS-9471) to 3.63 (AKms-14B \times SPV-104) in pooled data. Over all there was 14.26, 13.27 and 13.75% reduction in score in Akola 1995 Akola 1996 and in pooled data, respectively in treated over untreated control and there was 11.84% increase in score over Akola 1996.

4.9.4 Fungal load on ungerminated seed, Akola, 1995 and 1996

4.9.4.1 Ungerminated seed

In parental line minimum ungerminated seed were recorded in IS-9471 (2.22, 6.33, **4.28%**) in Akola 1995, Akola 1996 season and in pooled data, respectively. However **maximum** ungerminated seed were recorded in AKms-14B (45.00%) in Akola 1995 **SPV-104** (40.33%) in Akola 1996 and AKms-14B (41.83%) in pooled data (Table 63).

Among crosses minimum ungerminated seed was recorded in SPV-1201 × GJ-35-15-15 (2.78%) and maximum in IC5B 101 × SR1-26B (51.11%) in Akola 1995. In Akola 1996 minimum ungerminated was recorded in AKms-14B IS-6335 and IS-6335 × IS-9471 (4.67%) and maximum in SPV-1201 × SPV-946 (30.00%), however, in pooled data cross IS-6335 × IS-9471 recorded minimum (4.28%) and GJ-35-15-15 × SPV-104 recorded maximum (32.67%) ungerminated seed.

Over all there was 28.47, 12.83 and 21.71% reduction in ungerminated seed in Akola 1995, 1996 and in pooled data, respectively in treated seed over untreated control and 16.74% reduction in ungerminated seed in Akola 1996 over Akola 1995.

4.9.4.2 Infection by *F. moniliforme*

Four parental lines GJ-35-15-15, IS-2284, IS-6335 and IS-9471 recorded lowest (1.11%) and highest in SR1-26B (14.45%) in Akola 1995 (Table 64) however parental lines IS-9471 recorded lowest 2.33 and 1.72% and highest in AKms-14B 15.67 and **12.67%** *F. moniliforme* load at Akola 1996 and in pooled data, respectively.

Among crosses four crosses recorded lowest *F. moniliforme* load (1.11%) in Akola 1995, cross IS-6335 × IS-9471 also recorded lowest load 1.67 and **1.39 %** in Akola 1996 season and in pooled data, respectively. Highest load was recorded in GJ-35-15-15 × SPV-104 (30.56%) and (17.95%) in Akola 1995 and in pooled data, respectively and SPV-1201 × SPV-946 (17.00%) in Akola 1996.

Over all, there was significant decrease *F. moniliforme* load 19.76, 13.64 and 16.95% in Akola 1995, Akola 1996 and in pooled data, respectively in treated seed over untreated control and 14.88% increase in fungal load in Akola 1995 over Akola 1996.

4.9.4.3 Infection by *F. pallidoroseum*

The fungal load of *F. pallidoroseum* in Akola 1995 and Akola 1996 season were recorded nil in most of entries (Table 65), hence conclusion could not be drawn. There was no occurrence of *F. pallidoroseum* at Akola during 1995. In all the entries and in majority entries tested during 1996 season, hence no conclusion could be drawn (Table 65).

4.9.4.4 Infection by *C. lunata*

Table 66, revealed that parental line IS-9471 recorded lowest 0.00, 3.33 and 1.67% *C. lunata* load in Akola 1995, Akola 1996 and in pooled data, respectively. SPV-946 recorded highest 16.67% in Akola 1995 and AKms-14B exhibited highest 16.00 and 13.00% load in Akola 1996 and in pooled data, respectively. Among crosses lowest (0.00 %) load were recorded in GJ-35-IS-15 × IS-6335 and SPV-946 × IS-6335 in Akola 1995 cross SPV-1201 × IS-6335 (1.67%) in Akola 1996 and cross SPV-946 × IS-6335 (1.17%) in pooled data. Over all, there was 27.55, 8.27 and 17.99% reduction in fungal load in Akola 1995, Akola 1996 and in pooled data, respectively in treated over untreated control and 4.26% reduction in *C. lunata* load in Akola 1996 over Akola 1995.

4.9.4.5 Infection by other fungi

Data presented in Table 67 revealed that, the range for parental lines was from 0.00 (SPV-1201) to 26.11% (AKms-14B) in Akola 1995 season however, in Akola 1996 was from 0.00% in (IS-9471 and IS-2284) to 3.67% in (SPV-104) and in pooled data 0.28 % (IS-9471) to 14.72% (AKms-14B).

Among crosses SPV-946 × IS-2284 recorded lowest (0.00%) other fungi load at both locations and in pooled data, however, crosses SPV-946 × SPV-104 recorded highest (7.78%) fungal load in Akola 1995, cross AKms-14B × SRT-26B

and SPV-1201 \times SPV-104 (3.00 %) on Akola 1996 and cross SPV-1201 \times AKms-14B (5.28%) in pooled data, respectively.

Over all, there was 49.52, 60.00 and 52.41% significant reduction of other fungi load in Akola 1995, Akola 1996 and in pooled data, respectively in treated over untreated control and 64.55% reduction in fungal load in Akola 1996 over Akola 1995.

4.9.4.6 Score

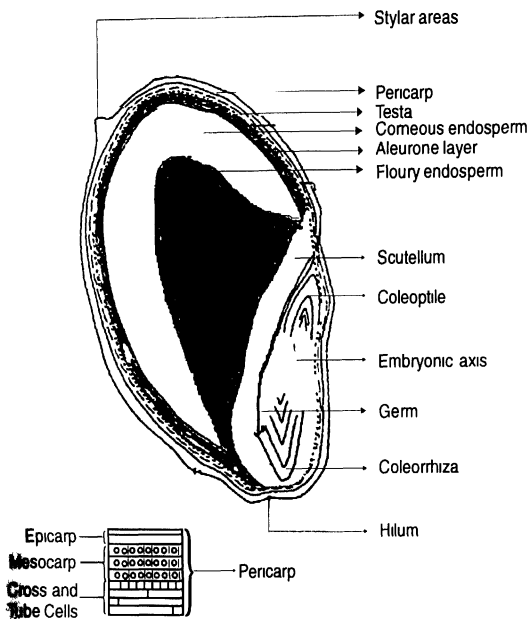
It is observed from data (Table 68), that the score range in parental lines was from 0.83 (SPV-1201) to 4.08 (ICSB-101B) in Akola 1995, from 1.17 (IS-9471) to 4.75 (AKms-14B) in Akola 1996 and from 1.04 (IS-9471) to 4.21 (ICSB-101B) in pooled data. Among crosses the range was from 0.58 (SPV-946 \times IS-6335) to 4.75 (AKms-14B \times SRT-26B) in Akola 1995 from 0.92 (SPV-1201 \times IS-2284) to 4.58 (AKms-14B \times SPV-104) in Akola 1996 and from 1.13 (SPV-946 \times IS-6335) to 4.54 (AKms-14B \times SPV-104) in pooled data. AKms-14B \times SPV-104 exhibited maximum score in both locations and in pooled data.

4.10. Histopathology of infection of molding fungi (Microtomy)

Histopathology of infection and colonization of major molding fungi viz. *F. moniliforme* and *F. pallidoroseum* and *C. lunata* at different stages of grain development i.e., at anthesis, 10, 20, 30 and 40 DAF were investigated.

Histologically sorghum seed consist of outer covering pericarp (seed coat), the storage tissue endosperm and embryo (germ). Pericarp is sub-divided into outer epicarp, two to three cells layers, middle mesocarp contains starch granules, usually thickest opposite embryo; innermost layer of pericarp is endocarp consist of cross and tube cells (Plate 5).

Just beneath the pericarp, some sorghum kernels have highly pigmented layer called testa or sub-coat, testa usually thickest at the crown of the kernel and thinnest over embryo. The endosperm of sorghum seed consist of aleurone layer and peripheral (earlier so called stele layer), cornecous and floury portions. Aleurone cell layer located beneath the pericarp or testa if it is present, block-like



Median longitudinal section of sorghum grain showing its parts

rectangular cells. The peripheral endosperm is beneath the aleurone layer, consisting of first two to six endosperm cell. The corneous endosperm (hard/ horny) located beneath the peripheral endosperm made up of starch and proteins. The floury endosperm area has loose packed endosperm cells. The starch granules are sphericals and they are held together by protein matrix.

Embryo or germ lies at the base of the kernel and consist of scutellum, plumule and radicle, plumule, radicle and a part between is known as embryonic axis. Coleoptile and coleorrhiza are the fused parts around the plumule and radicle toward embryonic leaf and root cap respectively.

4.10.1. Infection by *F. moniliforme*

Small size shriveled grains and in some cases no grain formed in the earhead, infection in all parts when the inoculations were made at anthesis (Plate 6A). When the earhead were inoculated at 10 DAF infection observed in all parts except in some parts of endosperm (Plate 6B). The grain from the earhead inoculated 20 DAF colonization observed in endosperm scutellum, embryonic region and at hilar areas (Plate 6C). Earhead inoculated at 30 DAF fungus colonizing in embryonic region involving coleoptile, coleorrhiza and embryonic axis (Plate D). When the earhead were inoculated at 40 DAF fungus colonizing in pericarp, aleurone and peripherals region fungus mycelium and spores were observed on pericarp (Plate 6E).

2. Infection by *F. pallidoroseum*

Shriveled grain and infection in all parts when the inoculations were made at anthesis (Plate 7A). Infection in all parts except some parts of endosperm, when the inoculations were made at 10 DAF (Plate 7B). At 20 DAF infection observed in hilar areas, embryonic region and pericarp (Plate 7C). Pericarp and embryonic region colonized when inoculations were made at 30 DAF (Plate 7D). At 40 DAF colonization were observed in endosperm, pericarp and aleurone layer (Plate 7E).

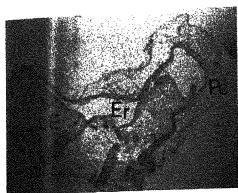
4.10.3. Infection by *C. lunata*

Shriveled grain formed and infection observed in embryonic region when the inoculations were made at anthesis (Plate 8A). At 10 DAF infection noticed in embryonic region, this infection takes place through hilar region (Plate 8B). Infection

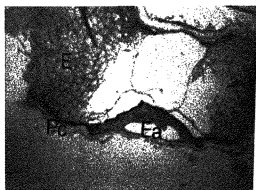
Plate : 6 Infection by *F. moniliforme*

- A Anthesis - (LS) Shriveled grain
Infection in all parts (13.2 x)
- B 10 DAA - (LS) Infection in all parts (13.2 x)
Except endosperm
- C 20 DAA - (LS) Hilar, pericarp, embryonic
Region (33.3 x)
- D 30 DAA - (LS) Embryonic infection (66.6 x)
- E 40 DAA - Aleurone layer and pericarp (66.6 x)

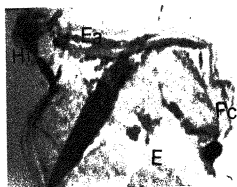
(Pe-Pericarp, E-Endosperm, Al-Aleurone layer,
Er-Embryonic region, Ea-Embryonic axis, Hi-Hilar
F-Fungus mycelium, F's-Fungus mycelium and spores)



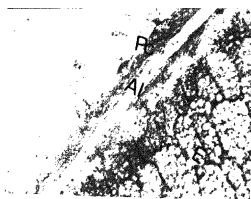
A



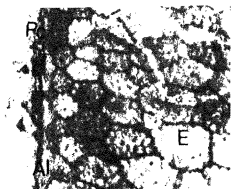
B



C



D



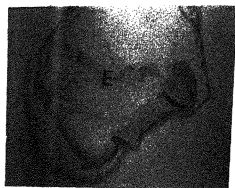
E

Plate 6. Infection by *F. moniliforme*

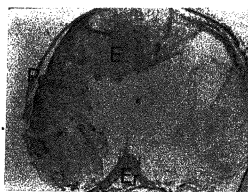
Plate : 7 Infection by *F. pallidoroseum*

A- Anthesis	(L.S) Shriveled grain infection in all parts (13.2 x)
B-10 DAA	(LS) Infection in all parts (33.3 x)
C-20DAA	(L.S) Hilum, pericarp, endosperm and embryonic region (33.3 x)
D-30 DAA	(LS) Pericarp and embryo (33.3 x)
E-40DAA	(L.S) Infection in endosperm and pericarp (66.6 x)

(Pc- Pericarp, E-Endosperm, Er-Embryonic region,
Ea-Embryonic axis, Al-Aleurone layer, F-fungus mycelium)



A



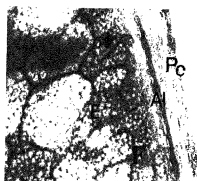
B



C



D



E

Plate 7. Infection by *F. pallidorozeum*

was observed in embryonic axis, scutellum near hilar voids, when inoculations were made at 20 DAF (Plate 8C). At 30 DAF, infection were observed in pericarp with spores and mycelium, aleurone layer and endosperm intercellular mycelium was observed (Plate 8D). Infection observed in pericarp and aleurone layer only with spores and mycelium on pericarp when inoculation were made at 40 DAF (Plate 8E).

4.10.4. Infection by different fungi (Fluorescence microscopy)

Shriveled grain was formed when the inoculation were made at anthesis and infection was observed in all region except endosperm (Plate 9A). Infection was observed in hilar areas when the inoculations were made at 10 DAF (Plate 9B). Infection observed in embryonic axis with coleoptile and coleorrhiza infection at 20 DAF (Plate 9C). Infection in pericarp, aleurone layer and endosperm with mycelium. Infection A to D by *F. moniliforme* (Plate 9D). Embryonic infection was observed when the inoculations were made at 20 DAF (Plate 9E). Pericarp infection when inoculated at 40 DAF by *C. lunata* (Plate 9F). E and F infection by *C. lunata*.

4.11 Correlation studies

Correlation describes the inter relationship between the variables. In any biological entity the variables are generally associated with each other. With a view to find out the association between grain molding contributing physical and biochemical characters. Simple correlation coefficient have been worked out on following aspects.

- Correlation between fungal load on germinated and ungerminated F_1 seed (Untreated), Akola and Patancheru. 1996
- Correlation between fungal load on germinated and ungerminated F_1 seed (Treated), Akola and Patancheru. 1996
- Correlation between physical characters, Akola and Patancheru. 1996
- Correlation between biochemical characters
- Correlation between protein fractions and physical characters

Plate : 8 by *C. lunata*

- | | | | |
|---|----------|---|--|
| A | Anthesis | - | (LS) Shriveled grain, embryonic region (13.2 x) |
| b | 10 DAA | - | (LS) Hilar areas and embryo (33.3 x) |
| C | 20DAA | - | (TS) Embryonic axis, scutellum and near hilar
viod (33.3 x) |
| D | 30DAA | - | (TS) Pericarp, aleurone layer, endosperm (66.6 x) |
| E | 40 DAA | - | (TS) Pericarp (33.3 x) |

(Pc-Pericarp, Al-Aleurone layer, E-Endosperm, Er-Embryonic region,
Ea-Embryonic axis, Hv-Hilar void, F-Fungus mycelium, Fs-Fungus
mycelium and spores)

Plate : 8 by *C. lunata*

- | | | |
|---|----------|---|
| A | Anthesis | (LS) Shriveled grain, embryonic region (13.2 x) |
| B | 10 DAA | (LS) Hilar areas and embryo (33.3 x) |
| C | 20DAA | (TS) Embryonic axis, scutellum and near hilar void (33.3 x) |
| D | 30DAA | (TS) Pericarp, aleurone layer, endosperm (66.6 x) |
| E | 40 DAA | (TS) Pericarp (33.3 x) |

(Pc-Pericarp, Al-Aleurone layer, E-Endosperm, Er-Embryonic region, Ea-Embryonic axis, Hv-Hilar void, F-Fungus mycelium, Fs-Fungus mycelium and spores)



A



B



C



D



E

Plate 8. Infection by *C. lunata*

Plate : 9 Infection by different fungi (fluorescence microphoto)

- A (L.S) At anthesis all region except endosperm *F. moniliforme* (20 x)
- B (L.S) 10 DAA, Hilar, *F. moniliforme* (100 x)
- C (L.S) Embryonic region *F. moniliforme* (100 x)
- D (TS) Endosperm (intercellular) *F. moniliforme* (100 x)
- E (L.S) Embryonic region *C. lunata* (100 x)
- F (TS) Pericarp, *C. lunata* (100 x)

(Pc-Pericarp, E-Endosperm, Ea-Embryonic axis, Hi-Hilar,
Cp-Coleoptile, Ch-Coleorrhiza, F-Fungus mycelium,
Fs-Fungus mycelium and spores).

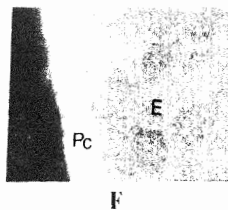
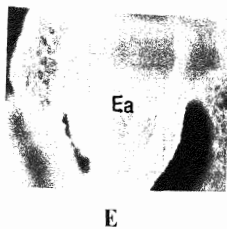
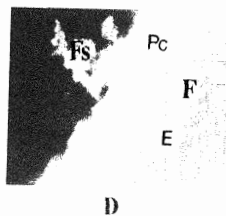
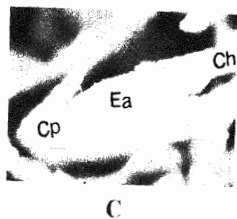
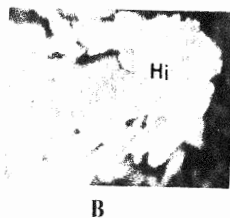
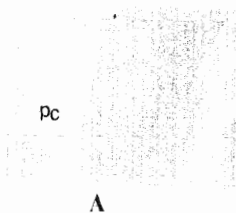


Plate 9. Infection by *F. moniliforme* and *C. lunata*
 Fluorescence microphoto)

4.11.1 Correlation between fungal load on germinated and ungerminated F_1 seed (Untreated), Akola and Patancheru, 1996

Fungal load viz. *F. moniliforme*, *F. pallidoroseum*, *C. lunata* other fungi and their score on germinated (GS) and ungerminated (UGS) (untreated) F_1 seed were recorded at both locations and correlation between them with TGMR as a dependent variable were worked out and results are presented in Table 69.

4.11.1.1 Germinated seed

Out of 12 characters studied, germinated seed had a significant negative correlation with seven characters at both the locations viz. ungerminated seed, score (GS), *F. moniliforme* (UGS), *C. lunata* (UGS), other fungi (UGS), score (UGS), and TGMR. however, *F. moniliforme* (GS) at Akola and *F. pallidoroseum* (UGS) at Patancheru location. Whereas, significant positive correlation exhibited in other fungi (GS) at Akola and *F. moniliforme* (GS), *F. pallidoroseum* (GS), *C. lunata* (GS) and other fungi (GS), at Patancheru.

4.11.1.2 Ungerminated seed

Ungerminated seed had a significant positive association with six characters score (GS), *F. moniliforme* (UGS), *C. lunata* (UGS), other fungi (UGS), score (UGS), and TGMR at both the location, and *F. pallidoroseum* (UGS) at Patancheru only. However, significant negative association was exhibited in *F. moniliforme* (GS) and other fungi (GS) at both location and *F. pallidoroseum* (GS), and *C. lunata* (GS) at Patancheru location only.

4.11.1.3 Infection by *F. moniliforme* (GS)

F. moniliforme (GS) exhibited significant positive correlation with score (GS) (0.76), *F. moniliforme* (UGS) (0.72), score (UGS) (0.71) *C. lunata* (UGS) (0.67), TGMR (0.65), other fungi (UGS) (0.38) and *C. lunata* (GS) (0.28) at Akola location. whereas, with *F. pallidoroseum* (GS) (0.78), *C. lunata* (GS) (0.48), Patancheru. However, significant negative association was exhibited with other fungi (GS) (-0.49) at Akola and *C. lunata* (UGS) (-0.64), *F. moniliforme* (UGS) (-0.52), TGMR (-0.38), *F. pallidoroseum* (UGS) (-0.31), score (UGS) (-0.30) and other fungi at Patancheru.

4.11.1.4 Infection by *F. pallidroseum* (GS)

F. pallidroseum (GS) had a significant positive association with *F. pallidroseum* (UGS) (0.55) at Akola and *C. lunata* (GS) (0.56), other fungi (UG) (0.28) at Patancheru location. However, *F. pallidroseum* (GS) exhibited significant a negative association with *C. lunata* (GS), score (GS), score (UGS), and TGMR at Akola location and score (GS), *F. moniliforme* (UGS), *F. pallidroseum* (UGS), *C. lunata* (UGS), other fungi (UGS), score UGS, and TGMR at Patancheru location.

4.11.1.5 Infection by *C. lunata* (GS)

C. lunata (GS) exhibited significant positive correlation with score (UGS), score (GS), and TGMR at Akola and other fungi (GS) at Patancheru, significant negative correlation was exhibited with all characters except other fungi (GS) at Patancheru.

4.11.1.6 Infection by other fungi (GS)

Other fungi (GS) exhibited significant and negative association with all characters at both locations except *F. pallidroseum* (UGS) at Akola and score (GS) at Patancheru.

4.11.1.7 Score (GS)

Score (GS) also exhibited significant positive correlation at both locations for all the characters except *F. pallidroseum* (UGS) at Akola, where, significant negative correlation was noticed.

4.11.1.8 Infection by *F. moniliforme* (UGS)

F. moniliforme (UGS) had a significant positive association with all the characters at both the locations, except *F. pallidroseum* (UGS) at Akola.

4.11.1.9 Infection by *F. pallidroseum* (UGS)

F. pallidroseum (UGS) had a significant positive association with all the characters at Patancheru, however TGMR (-0.28) exhibited significant negative association at Akola.

4.11.1.10 Infection by *C. lunata* (UGS)

C. lunata (UGS) exhibited significant positive correlation with all the characters at both the locations.

4.11.1.11 Infection by other fungi (UGS)

Other fungi (UGS) also exhibited significant positive association with all the characters at both the locations.

4.11.1.12 Score (UGS)

Score (UGS) also exhibited highly significant positive correlation with TGM_R at both the locations.

4.11. 2 Correlation between fungal load on germinated and

ungerminated F₁ seed (Treated). Akola and Patancheru, 1996

Fungal load viz. *F. moniliforme*, *F. pallidoroseum* , *C. lunata* and other fungi and their score on germinated and ungerminated F₁ (Treated) seed were recorded at both locations and correlation between them were worked out and results are presented in Table 70.

4.11.2.1 Germinated Seed

Out of 12 characters studied, germinated seed had a significant negative association with eight characters UGS, score (GS), *F. moniliforme* (UGS), *F. pallidoroseum* (UGS), *C. lunata* (UGS), other fungi (UGS), score (UGS) and TGM_R (UGS) at both locations. However, significant negative association was exhibited in *F. moniliforme* (-0.72), *C. lunata* (GS) (-0.40) and *F. pallidoroseum* (GS) (-0.36) at Akola location and significant positive association with *C. lunata* (GS) (0.60), *F. pallidoroseum* (GS) (0.46), *F. moniliforme* (GS) (0.45) and other fungi (GS) (0.43) at Patancheru.

4.11.2.2 Ungerminated seed

Out of 11 characters, ungerminated seed had a significant positive association with seven characters score (GS), *F. moniliforme* (UGS), *F. pallidoroseum* (UGS), *C. lunata* (UGS), other fungi (UGS) score (UGS) and TGM_R at both locations. However, significant positive association exhibited with *F. moniliforme* (GS) (0.72), *C. lunata* (GS) (0.40), *F. pallidoroseum* (GS) (0.36) and other fungi (GS) (0.26) at Akola and significant negative association with *C. lunata* (GS) (-0.60), *F. pallidoroseum* (GS) (-0.47), *F. moniliforme* (GS) (-0.45) and other fungi (GS) (-0.43) at Patancheru.

4.11.2.3 Infection by *F. moniliforme* (GS)

F. moniliforme (GS) exhibited significant positive correlation for all characters except other fungi (GS) at Akola location, but only for *F. pallidoroseum* (GS) (0.60), *C. lunata* (GS) (0.59) and other fungi (GS) (0.46) at Patancheru. Significant negative correlation with *C. lunata* (UGS) (-0.52), *F. pallidoroseum* (UGS) (-0.44), other fungi (UGS) (-0.39) and *F. moniliforme* (UGS) (-0.32) was exhibited at Patancheru.

4.11.2.4 Infection by *F. pallidoroseum* (GS)

F. pallidoroseum had a significant positive association with *F. moniliforme* (UGS) (0.41), other fungi (UGS) (0.36) and *F. pallidoroseum* (UGS) (0.30) at Akola, and with *C. lunata* (GS) (0.48) and other fungi (GS) (0.36) at Patancheru. Five characters viz. *C. lunata* (UGS) (-0.53), other fungi (UGS) (-0.45), *F. pallidoroseum* (UGS) (-0.41), *F. moniliforme* (UGS) (-0.34) and TGM (UGS) (-0.28) exhibited significant negative correlation at Patancheru. The remaining characters exhibited non-significant positive and negative association at both locations.

4.11.2.5 *C. lunata* (GS)

All the characters except *F. pallidoroseum* (UGS) and other fungi (UGS) and only one character i.e. other fungi (GS) have shown significant positive correlation with *C. lunata* (GS) at Akola and Patancheru locations respectively. Seven out of eight characters exhibited negative correlation of which five exhibited significant negative correlation with *C. lunata* (GS).

4.11.2.6 Infection by other fungi (GS)

Other fungi (GS) exhibited significant positive with score (UGS) (0.60), TGM (UGS) (0.59), score (GS) (0.56), and *C. lunata* (UGS) (0.31) at Akola location. However, at Patancheru all characters exhibited negative association of which four viz. *F. pallidoroseum* (UGS) (-0.44), *C. lunata* (UGS) (-0.43), other fungi (UGS) (-0.38) and *F. moniliforme* (UGS) (-0.36) were significant.

4.11.2.7 Score (GS)

Score (GS) exhibited highly significant positive correlation with all the traits at both the locations.

4.11.2.8 Infection by *F. moniliforme* (UGS)

F. moniliforme (UGS) also exhibited significant positive correlation at both the locations for all the traits.

4.11.2.9 Infection by *F. pallidoroseum* (UGS)

Four characters viz. *C. lunata* (UGS), other fungi (UGS), score (UGS) and TGMR exhibited significant positive correlation with *F. pallidoroseum* (UGS) at both the locations.

4.11.2.10 Infection by *C. lunata* (UGS)

C. lunata exhibited highly significant positive correlation with TGMR (0.76, 0.74), score (UGS) (0.70, 0.68) and other fungi (UGS) (0.69, 0.84) at Akola and Patancheru, respectively.

4.11.2.11 Infection by other fungi (UGS)

Other fungi (UGS) exhibited significant positive correlation with TGMR (0.68, 0.66) and score (UGS) (0.54, 0.57) at Akola and Patancheru, respectively.

4.11.2.12 Score (UGS)

Score (UGS) exhibited highly significant positive correlation with TGMR (0.85, 0.85) at both locations.

4.11.3 Correlation between physical characters, Akola and Patancheru, 1996

Twelve characters at both the locations were correlated using germination percent as a dependent variable Table 71.

4.11.3.1 100-grain weight

The study of results revealed that, 100 grain weight had significant and positive correlation with TGMR (0.40) and *F. moniliforme* (0.28) and significant negative correlation with glume covering (-0.26) and *F. pallidoroseum* (-0.26) at Akola

location. however. at Patancheru location none of the characters is having significant positive and negative association.

4.11.3.2 Grain hardness

Grain hardness had a significant positive association with DTF (0.42) and significant negative association with endosperm texture (-0.61) and *F. pallidoroseum* (-0.32) at Akola, whereas, at Patancheru, significant positive association was observed with other fungi (0.50), DTF (0.39), germination (0.30), whereas, significant and negative association was observed with endosperm texture (-0.33) and *F. moniliforme* (-0.26).

4.11.3.3 Endosperm texture

Endosperm texture had a significant and positive correlation with *F. pallidoroseum* (0.41) and significant negative correlation with DTF (-0.54), *C. lunata* (-0.54), electrical conductivity (-0.51), germination (-0.41), mesocarp thickness (-0.39), TGMR (-0.39) and *F. moniliforme* (-0.35) at Akola location, whereas, at Patancheru it had significant and positive correlation with germination (0.52) and significant negative correlation with TGMR (-0.87), *F. moniliforme* (-0.59), mesocarp thickness (-0.55), electrical conductivity (-0.54) and DTF (-0.51).

4.11.3.4 Electrical conductivity

Out of 9 characters studied, 5 characters viz. *F. moniliforme* (0.67, 0.72), TGMR (0.64, 0.82), *C. lunata* (Akola, 0.61), mesocarp thickness (0.56, 0.59) and Days to 50% flowering (0.50, 0.40), exhibited significant positive correlation at Akola and Patancheru respectively. However, germination (-0.66, -0.82), other fungi (-0.44, -0.31) and *F. pallidoroseum* (Akola, -0.33) indicated significant negative correlation at Akola and Patancheru, respectively.

4.11.3.5 Days to 50% flowering

Days to 50% flowering at Akola had significant positive association with mesocarp thickness (0.41), *F. moniliforme* (0.37), *C. lunata* (0.37) and TGMR (0.29) whereas, significant negative correlation were established with germination (-0.38), glume covering (-0.33) and *F. pallidoroseum* (-0.32). However, at Patancheru significant positive correlation was exhibited with *C. lunata* (0.46), mesocarp thickness (0.39).

TGMR (0.34), *F. moniliforme* (0.31) and *F. pallidoroseum* (0.26) and negative association with germination (-0.20) and glume covering (-0.05).

4.11.3.6 Glume covering

Glume covering had no significant correlation (positive or negative) with most of the variables at both the locations. However positive correlation was noticed with germination at Akola (0.06) and Patancheru (0.11) whereas, negative correlation was exhibited with mesocarp thickness, *F. moniliforme*, *C. lunata* and TGMR at both locations.

4.11.3.7 Mesocarp thickness

Mesocarp thickness at Akola location has a significant positive association with *C. lunata* (0.68), TGMR (0.65) and *F. moniliforme* (0.51), however, significant negative correlation, was exhibited with germination (-0.57) and *F. pallidoroseum* (-0.38). At Patancheru TGMR (0.64) and *F. moniliforme* (0.58) exhibited significant positive association however, significant negative correlation exhibited with germination only (-0.53).

4.11.3.8 Infection by *F. moniliforme*

F. moniliforme had a significant positive correlation with TGMR (0.68) and *C. lunata* (0.45) and significant negative correlation with germination (-0.82) and other fungi (-0.52) at Akola, however at Patancheru significant positive correlation was exhibited with TGMR (0.77) and *F. pallidoroseum* (0.49) and significant negative in germination (-0.76) and other fungi (-0.50).

4.11.3.9 Infection by *F. pallidoroseum*

F. pallidoroseum at Akola had exhibited significant negative correlation with TGMR (-0.41) and *C. lunata* (-0.40) and significant positive correlation with germination (0.28), however, at Patancheru germination (-0.19), other fungi (-0.18) and *C. lunata* (-0.17) exhibited negative correlation and TGMR (0.19) indicated positive correlation

4.11.3.10 Infection by *C. lunata*

C. lunata exhibited significant positive correlation with TGMR (0.73) and significant negative correlation with germination (-0.59) at Akola. Whereas, at Patancheru other

fungi (0.23) and TGM_R (0.19) exhibited positive but non significant and germination (-0.23) indicated non significant negative correlation.

4.11.3.11 Infection by other fungi

Other fungi had significant positive association with germination (0.56, 0.41) at Akola and Patancheru respectively. other fungi had a significant negative correlation with TGM_R (-0.31) at Akola and non significant negative association (-0.24) at Patancheru.

4.11.3.12 TGM_R

TGM_R exhibited highly significant negative correlation with germination (-0.78) and (-0.83) at both the locations.

4.11.4 Correlation between biochemical characters

For biochemical characters viz. proteins, soluble sugars, tannins and flavan-4-ols were correlated with grain hardness and TGM_R as dependent variable and results are presented in Table 72.

4.11.4.1 Protein

Protein exhibited significant positive correlation grain hardness (0.29), total sugars (0.27), non significant negative correlation with flavan-4-ols (-0.15). Correlation with the remaining characters could not be established.

4.11.4.2 Soluble sugars

Total sugars exhibited significant positive correlation with tannins (0.35) and grain hardness (0.26), significant negative association with TGM_R (-0.26) and non-significant positive with flavan-4-ols (0.24).

4.11.4.3 Tannins

Tannins exhibited highly significant positive correlation (0.73) with flavan-4-ols, however grain hardness (-0.37) and TGM_R (-0.75) exhibited significant negative correlation.

4.11.4.4 Flavan-4-ols

Flavan-4-ols was found to be negatively associated with grain hardness (-0.59) and TGM_R (-0.78).

4.11.4.5 Grain hardness

Grain hardness had a significant positive correlation with TGM_R.

4.11.5 Correlation between protein fractions and physical characters

Correlation between six protein fractions viz. Albumin and globulin, prolamin, cross-link prolamin, glutelin like, glutelin, and residues with physical characters viz. mesocarp thickness, endosperm texture, germination and grain hardness were correlated and the results are presented in Table 73.

4.11.5.1 Albumin and globulin.

Out of nine characters only two characters viz. prolamin (0.55) and cross link prolamin (0.36) exhibited significant positive correlation. whereas, three characters viz. glutelin-like (-0.73), glutelin (-0.73) and endosperm texture (-0.45) indicated significant negative correlation other characters recorded positive and negative non-significant association with albumin and globulin.

4.11.5.2 Prolamin

Prolamin had a highly significant correlation with cross-link prolamin (0.64) and grain hardness (0.63), however significant negative association were exhibited in endosperm texture (-0.73), glutelin like (-0.69), germination (-0.69) and glutelin (-0.66). Other two characters viz. mesocarp thickness (0.35) and residues (0.34) recorded positive non significant association.

4.11.5.3 Cross-link prolamin

Cross-link prolamin exhibited significant positive correlation with grain hardness (0.63) and mesocarp thickness (0.42). However, four characters viz. germination (-0.60), glutelin (-0.59), glutelin-like (-0.58) and endosperm texture (-0.57) exhibited significant negative association. Residues (0.11) exhibited positive but non-significant association.

4.11.5.4 Glutelin-like

Glutelin-like had a significant positive association with glutelin (0.72), endosperm texture (0.62) and germination (0.57), whereas, with grain hardness had a significant negative correlation (-0.47). Mesocarp thickness (-0.20) and residues (-0.18) also reported negative association but non-significant.

4.11.5.5 Glutelin

Glutelin exhibited significant and positive correlation with germination (0.56) and endosperm texture (0.52), whereas, negative significant correlation with grain hardness (-0.42). Mesocarp thickness (-0.34) and residues (-0.07) also reported statistically non significant negative association.

4.11.5.6 Residues

All the characters associated with residues exhibited non-significant positive and negative association. Highest negative association was recorded with germination (-0.32).

4.11.5.7 Mesocarp thickness

Mesocarp thickness exhibited positive and negative significant association with grain hardness (0.41) and germination (-0.48) respectively. Endosperm texture recorded negative association (-0.33) but non-significant.

4.11.5.8 Seed germination

Seed germination exhibited significant negative (-0.38) association with grain hardness.

CHAPTER – V

DISCUSSION

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CHAPTER V

DISCUSSION

Parental diversity

Genetic divergence of parental lines is an important pre-requisite for obtaining desirable segregants in the progenies. Three red resistant, five white elite and two white susceptible parental lines of sorghum selected for the study provided sufficient variability for various quantitative traits. The analysis of variance also exhibited significant variation for all the characters except infection by *F. pallidoroseum* and other fungi under study which is indicative of their genetic diversity.

Combining ability effects

Combining ability is necessary in selection of appropriate parents in hybridization. Since, it gives an idea whether a particular parents combines well in a cross and also denotes specific performance of cross combination against the expectations from the general combining ability of parents.

The results (Table 74) revealed that none of the parental lines was a good general combiner for all the traits under study. However, IS-9471 had desirable gene for nine (F_1) and three (F_2) characters. This line could transmit higher grain hardness, corneous endosperm texture, lower plant height, more glume covering, low fungal infection by *F. pallidoroseum*, more proteins, more soluble sugars, more tannins and more flavan-4-ols in F_1 crosses and whereas, more grain hardness, lower plant height and low fungal infection by *F. moniliforme* in F_2 progenies. Similarly parental lines SPV-1201, IS-6335, GJ-35-15-15 and IS-2284 transmitted favourable genes for various traits to F_1 crosses and F_2 progenies notable among these are the increased grain hardness, corneous endosperm texture, tannins, flavan-4-ols and reduced infection by *F. moniliforme* and *F. pallidoroseum*. In general transmission of traits in F_1 crosses was more pronounced than those in F_2 progenies.

The results indicated that the first five parents IS-9471, SPV-1201, IS-6335, GJ-35-15-15 and IS-2284 can be categorised as good general combiners for various physical and biochemical characters. Since higher general combining effects correspond with additive and for additive to additive interaction (Griffings, 1956b) and represent the fixable genetic

components of variation, these parental lines appear to be worthy of exploitation in a recombination breeding program.

Specific combining ability effects is the indicative of heterosis and also the dominance and epistatic gene action. Denis and Girard (1977) reported loss in viability to be so important part of the grain mold syndrome, that they recommend a germination test as a part of standard evaluation for identification of grain mold resistance.

First cross AKms-14B x IS-6335 exhibited significant sca effects for germination per cent in both F_1 and F_2 diallel progenies. The same cross exhibited significant desirable sca effects for other component characters viz., 100-grain weight, electrical conductivity, days to 50% flowering, mesocarp thickness, TGM, *F. moniliforme*, protein, tannins, and flavan-4-ols in F_1 diallel, and electrical conductivity and TGM in F_2 diallel (Table 75). Significant sca effects on germination both in F_1 and F_2 diallel progenies of other crosses were also noted. In addition, significant sca effects were recorded for other important grain mold resistance components, such as TGM, grain hardness, endosperm texture, infection by *F. moniliforme*, and *F. pallidoroseum*.

Among the 10 specific combinations (Table 75) first, second and ninth were most desirable, since these had significant positive sca effects for germination in both F_1 and F_2 progenies. Of the parental lines involved in above three crosses, IS-6335 IS-9471, and SPV-1201 were identified as good general combiners. Therefore one can expect good segregants in further generation from these crosses. It can therefore, be suggested that biparental mating among the segregants of these crosses may be attempted to evolve line with better grain mold tolerance.

Gene action governing inheritance of characters

The gea and sca variances for the traits studied indicate the gene action associated with them. Knowledge gained on the relative magnitude of various types of gene actions is useful in deciding the most appropriate breeding procedures.

In the present investigation the variance due to gea were larger than that of sca for the characters, like 100-grain weight, endosperm texture, tannins and flavan-4-ols in F_1 progenies, and for cob length, TGM, and infection by *F. moniliforme* and *C. lunata* both in F_1 and F_2 progenies. Therefore, additive gene action was more predominant for these characters. Ghorade (1995) reported the role of additive gene effects for 100-grain weight,

abholkar and Baghel (1982) reported similar results for tannin content. Karale *et al.*, (1984) reported similar results for cob length. No literature is available on inheritance of TGMR. abholkar and Baghel (1980). Narayana and Prasad (1983), Kataria *et al.* (1990) reported similar results for infection by *Fusarium* and *Curvularia*.

For grain hardness and glume covering both gca and sca estimates were significant. Their ratio of more than 1 demonstrated preponderance of additive type gene action in F_1 diallel. In F_2 diallel, though both estimates were significant, the ratio of gca/δ^2sca works out to be less than 1 indicating predominance of more non-additive gene action. The role of additive gene action for these traits have been reported by Ghorade (1995) and the role of additive gene action for grain hardness was reported by Rana *et al.* (1978).

In case of plant height lower ratio of δ^2gca/δ^2sca indicated non-additive gene action in F_1 diallel. These results are similar to those of Nimabalkar and Bapat (1987) and Jeger *et al.* (1988). The higher ratio of δ^2gca/δ^2sca indicated predominance of additive gene action in controlling plant height in F_2 progenies. These findings were in confirmation with the observation of Patil and Thombre (1986) and Shekar *et al.* (1987).

The results obtained for electrical conductivity and other fungi fungal load were similar to those of plant height, which is supported by Ghorade (1995) for F_2 diallel, but contradictory to F_1 diallel.

For days to 50% flowering, mesocarp thickness, proteins and soluble sugars, in diallel, though both the estimates were significant, the ratio of δ^2gca/δ^2sca works out to be less than 1 indicating non-additive gene action. These results are in agreement with Kanakaiah (1982) and Huger *et al.* (1988). For germination and infection by *F. pallidoroseum* though the estimates were significant, the ratio of δ^2gca/δ^2sca comes out to be less than unity in F_1 F_2 diallel progenies, indicating predominance of non-additive gene action. Contradictory results were reported by Ghorade (1995) for germination and for *F. pallidoroseum* no literature is available.

In the present investigation, it could be concluded that gca variance were dominant for most of the characters studied, i.e., 100-grain weight, grain hardness, perm texture, cob length, glume covering, TGMR, *F. moniliforme*, *C. lunata*, tannins, flavan-4-ols. However, sca variance was predominant for electrical conductivity, days to flowering, mesocarp thickness, germination, *F. pallidoroseum*, proteins and soluble

sugars. Studies on gene action governing inheritance of endosperm texture, flavan-4-ols and TGMR have been made for the first time, and thus these are new information.

Heterosis

In the present investigation, of the several agronomic, physical and bio-chemical parameters studied, desirable significant heterosis and heterobeltiosis were observed for 100-grain weight, electrical conductivity, DTF, plant height, cob-length, mesocarp thickness, TGMR, seed germination, infection by *F. moniliforme*, *F. pallidoroseum*, *C. lunata*, tannins and flavan-4-ols (Tables 12-17). These parameters can significantly contribute towards reducing or preventing infection and colonization by mold fungi. Similar results were reported by Kanaka (1982) for 100-grain weight; Ghorade (1995) for electrical conductivity and seed germination, Quinby (1963), Goyal and Joshi (1976) and Saradaman (1981) for cob-length; Naryana and Prasad (1983) and Ghorade (1995) for fungal load.

Biochemical characters

For proteins and soluble sugars none of the crosses exhibited significant desirable heterosis or heterobeltiosis, however, for tannins significant heterosis exhibited from 2.38 to 38.64%, whereas for flavan-4-ols the range for heterosis was from 5.67 to 52.10% and for heterobeltiosis 26.45 to 48.37%. From the data of heterosis and heterobeltiosis for seed germination, it is observed that crosses with high magnitude of heterosis or heterobeltiosis have not necessarily showed better *per se* performance or vice-versa. Hence the selection of superior crosses should necessarily be based not only on magnitude of heterosis but also on actual performance of crosses for germination so that appropriate selection can be made without errors.

On the basis of their *per se* performance for germination and the extent of heterosis and heterobeltiosis, it is concluded that out of 10 crosses, first four crosses i.e. ICSB-101B x IS-2284, ICSB-101 B x IS-6335, ICSB-101B x IS-9471 and SPV-1201 x IS-2284 could be stated as a desirable ones for mold resistance breeding. These crosses need to be exploited in further generations for isolating mold resistance lines

Physical characters

hundred-grain weight of kernels was less in all the entries tested at Patancheru during 1996 under controlled conditions. Over all 86.25% increase in *F. moniliforme* load was recorded at Patancheru over Akola in the same year. Bhatnagar (1971), Gray *et al.* (1971) and Mathur *et*

(1975) reported that *F. moniliforme* and *C. lunata* interfere with carbohydrate translocation developing kernels causing reduction in size and weight of seed. Similar results were also reported by Glueck and Rooney (1976), Castor (1977), Castor and Frederiksen (1980), Singh and Agrawal (1989) and Somani (1992). Seventy per cent loss in grain weight was reported by Ray *et al.* (1971) in central Kentucky USA. Sundaram *et al.* (1972) reported losses up to 50% due to head molds in hybrid sorghum in experimental plots at Coimbatore. Glueck and Rooney (1976) reported test weight loss from 62.2 lb/bu to 47.3 lb/bu under severe weathering conditions at College Station, Texas. Singh and Agrawal (1989) reported that sorghum seed infected with *C. lunata*, *F. moniliforme* and *Phoma sorghina* reduced 100-grain weight by 69, 35 and 40%, respectively. Significant specific combining ability for 100-grain weight was detected in two F_1 crosses (AKms-14B \times IS-6335 and SPV-1201 \times SRT-26B), as well as significant heterosis for grain weight in two crosses (AKms-14B \times IS-6335 and SRT-26B \times IS-6335) out of 10 superior F_1 crosses. Gene action in F_1 diallel was additive for 100-grain weight.

Rana *et al.* (1984), Bandyopadhyay (1988), Stenhouse *et al.* (1990) Reddy *et al.* (1991), Jambunathan *et al.* (1992) and Kumari *et al.* (1992) reported that in white-grained sorghums hardness contribute to resistance. Hardness was measured by force required to break the grain. From the present studies it is observed that, parent SPV-1201 exhibited maximum grain hardness followed by GJ-35-15-15 and ICSB-101 B at both the locations. In all areas, cross SPV-1201 \times GJ-35-15-15 exhibited highest grain hardness i.e. 7.85 and 7.43 kg/cm² and (Av. 7.64 kg/cm²) at Akola and Patancheru, respectively. Colored parental lines and crosses having more floury endosperm were brittle and required less breaking strength. Overall less grain hardness was recorded at Patancheru than at Akola during 1996, indicating that controlled conditions cause more deterioration of grain than natural infection. Jambunathan *et al.* (1992) reported higher grain hardness in grain grown in post rainy season than in grain grown in rainy season. Mukuru (1992), Jambunathan *et al.* (1992) and Somani (1992) also reported that mold resistance in white grain advanced selections was associated with grain thinness, whereas, resistance in red-grain types was associated with flavan-4-ols and grain thinness. The present results are in confirmation of these results. Grain hardness is governed by additive and non-additive gene action in F_1 and F_2 diallel, respectively.

In the present investigation among 48 crosses eight crosses exhibited corneous endosperm six crosses exhibited floury endosperm and 31 crosses indicated intermediate (Table 5). Clark *et al.* (1973) and Ellis (1972) noticed that corneous endosperm characteristic is not necessary for resistance to weathering however if all other things were equal a line with more corneous grain would resist deterioration more than would a floury endosperm line because of more dense structure and organization. Glueck *et al.* (1977) and Glueck and Rooney (1980) noticed that grain with more corneous endosperm were more likely to resist deterioration than floury endosperm lines. Similar results were reported by Garud (1992) and Somani (1992). Significant specific combining ability was noticed in F_1 diallel in only one cross (IS-2284 IS-6335) out of 10 and this character is governed by additive gene action in F_1 diallel.

Seventeen of the 45 crosses exhibited electrical conductivity less than 150 dSm⁻¹ and six between 150 and 200 dSm⁻¹ and 22 more than 200 dSm⁻¹. Electrical conductivity on higher sides was recorded at Patancheru than at Akola this might be due to more electrolytes leakage because of more deterioration under controlled conditions at Patancheru. Measuring of seed leachates and correlation with germination could become an efficient techniques for studying the effects of grain mold severity on seed viability (Forbes 1986). Glueck and Rooney (1980) reported that cultivars with thick mesocarp and floury endosperm texture increased water absorption and richer leachates. Somani (1992) noticed that the electrical conductivity of seed leachates was more in susceptible cultivars. Crosses ICSB-101B IS-2284 ICSB-101B IS-6335 and ICSB-101B IS-9471 selected on the basis of germination indicated lower electrical conductivity. Electrical conductivity is governed by non-additive and additive gene action in F_1 and F_2 diallel respectively.

Glumes appear to be plants first defense against fungal invasion and colonization (Waniska *et al.* 1992). But it was also reported that even on completely covered but not compact glumes, mold development was noticed (Somani 1992). It was also reported that except *F. moniliforme* infection of other fungi takes place from uncovered portion of kernels by glumes. Murty (1975) noticed *C. nivul* infection on a portion of seed not covered by glume and concluded that open head with seed completely enclosed in long papery glumes are relatively resistant to field deterioration. Gangadharan *et al.* (1978) noticed similar observation and concluded that fully enclosed grain by glume and loose heads are the factors

associated with resistance. Glueck and Rooney (1980) and Somani (1992) reported similar results. Mansuetus *et al.* (1988) showed that disease incidence was negatively associated with glume cover ($r = -0.56$) and glume length ($r = -0.56$). Pigmentation in the pericarp and glume may impart some degree of resistance (Waniska *et al.*, 1992). Sorghum contains polyphenols (tannins) in colored pericarp and tannins are deterrents to mold Harris and Burns (1973). In the present study, 11 F_1 crosses showed more than 50% glume covering at Akola, and 21 F_1 crosses at Patancheru. More glume covering was noticed in three crosses of the 10 crosses (Table 77) which exhibited resistant reaction to mold. This character is governed by additive and non-additive gene action in F_1 and F_2 diallel respectively.

With regards to mesocarp thickness, 15 out of 45 crosses at both locations exhibited less than 50 μm thickness (thin). Thin mesocarp is considered as resistant trait for grain mold. The mesocarp is thin when the gene is dominant (Z-) and thick when the gene is recessive (zz). Ten out of 10 crosses based on germination and grain mold resistance recorded less than 50 μm mesocarp thickness (Table 77). Glueck and Rooney (1980) reported similar results, and stated further that thin mesocarp sorghum withstand weathering better than those with thick mesocarp, thick mesocarp contains starch and protein which support more fungal colonies than thin mesocarp. Castor (1981) provided additional evidence that infection takes place by relatively few species of fungi during anthesis. Also a number of different tissues appear to be involved in resistance to colonization. Since the infection takes place at such an early stage, the presence or absence of testa has little effect on initial colonization. His work substantiate the fact that mesocarp provide an ideal environment for early colonization. Miller (1981) reported similar results. This trait is governed by non additive gene action in F_1 diallel.

Visual appraisal has been the most common means of quantifying grain mold to date. Visual appraisal involves a complex of factors and can estimate severity, incidence and damage, depending upon the method of assessment. Visual assessment, obviously the quickest and easiest method of disease assessment, is used for screening large number of samples (Bandyopadhyay and Mughogho, 1988 a). According to Frederiksen *et al.* (1982) comparing threshed grain is the most accurate method of visual assessment of grain mold. Hence TGMR is an important character for evaluation of crosses, less TGMR is considered as favourable for grain mold resistance. As TGMR grade exhibited significant positive association with different types of fungal load indicated that low TGMR grade is favourable. Forbes (1986) reported

positive associations with visual assessment as percentage of kernels moldy ($r = 0.98$) with mony forming with of *F. moniliforme*. Additive gene action was exhibited for this trait in F_1 and F_2 diallel.

Poor seed germination is positively correlated with infection level by mold fungi. There was 86.26% more load of *F. moniliforme* and 165.27% more load of *F. illidorozeum* at Patancheru over Akola in F_1 , that may have resulted in lower germination. However, 108.95% more *C. lunata* load was recorded at Akola over Patancheru. Germination considered desirable character for mold resistance. Denis and Girard (1977) and Castor and Frederiksen (1980) recommended germination test as a part of standard evaluation for identification of grain mold resistance. Many fungi were reported responsible for loss in germination Tarr (1962) reported *Aspergillus*, *Fusarium* and *Rhizoctonia* spp. were responsible for poor emergence of sorghum seedlings, since these fungi destroy starchy endosperm of seed and thus deprive young seedling of its food. Tripathi (1974) reported, reduced germination by (42%) due to *Colletotrichum graminicola* followed by *C. lunata* (0%), *F. moniliforme* (37%) and *Phoma insidiosa* (26%). Mathur *et al.* (1975) reported maximum *F. moniliforme* and *F. semitectum* from 70 samples collected from U.P., Gujarat and Rajasthan (northern India). *F. moniliforme* affected both germination and seedling growth and was mainly observed in embryos. Castor (1975) reported maximum *Fusarium* and *Curvularia* as a principal pathogens and caused discoloration and reduction in viability of sorghum seed.

Dayan and Dalmacio (1982) reported *F. moniliforme* and *C. lunata* as most predominant fungi invading embryo and endosperm thereby reduced seed viability in milippines. Wu (1983) isolated 16 genera of which *F. moniliforme* was most prominent together with *C. lunata* in Taiwan. Granja and Zambolim (1984) reported low germination in sorghum due to *F. moniliforme*. Deshmukh (1989) noticed that *C. lunata*, *F. moniliforme* and *Serophilum halodes* caused considerable reduction in germination from Vidarbha. Since these fungi destroy starchy endosperm of seed and thus deprive young seedling of its food. In the present investigation better heterotic crosses for germination showing heterosis and sterobeltiosis for other components were: ICSB-101B \times IS-2284, ICSB-101B \times IS-6335, ISB-101B \times IS-9471, SPV-1201 \times IS-2284, AKms-14B \times IS-6335, GJ-35-15-15 \times IS-6335, RT-26B \times IS 9471, IS-2284 \times IS-9471, SRT-26B \times IS-6335 and SPV-104 \times IS-6335, which

hibited 81.75 to 90.00% germination. The significant heterosis exhibited for other components characters were electrical conductivity, mesocarp thickness, TGM, *F. coniforme*, *C. lunata*, *F. pallidroseum*, tannins and flavan-4-ols contributed towards mold resistance in above crosses (Table 77). Germination is governed by non-additive gene action in F_1 and F_2 diallel.

Based on *F. pallidroseum* infestation at two locations it appears that *F. pallidroseum* is comparatively of minor importance in grain mold at Akola, whereas, it is an important mold fungus next to *C. lunata* at Patancheru. This variation possibility might be due to variations in the ecological conditions. Only one cross ICSB-101B \times IS-2284 out of 10 selected on basis of germination exhibited significant heterosis for this trait. This character is governed by non-additive gene action in both F_1 and F_2 diallel.

C. lunata is another major fungus associated with grain mold. It is number one fungus in some parts, causing reduction in weight, germination and viability (Bhatnagar, 1971; Mathur *et al.*, 1967, Tripathi, 1974; and Khare *et al.*, 1976). *C. lunata* has been found important also in Taiwan (Wu and Cheng, 1990), Thailand (Boon-Long, 1992) and also from different parts of India (Bhale and Khare, 1982; Deshmukh, 1989; Somani, 1992). At Akola 39 out of 45 crosses recorded more than 31% *Curvularia* load. However, at Patancheru 35 out of 45 crosses recorded 16-30% *C. lunata* load and over all, there was 108.95% increase at Akola over Patancheru. Significant heterosis was recorded in cross SRT-26 \times IS-6335 and sterobeltiosis in five crosses (ICSB-101B \times IS-6335, ICSB-101B \times IS-9471, GJ-35-15-15 \times IS-6335, SRT-26B \times IS-9471 and SRT-26B \times IS-6335) of the 10 crosses selected on basis of germination in F_1 diallel. This character is governed by additive gene action in F_1 and F_2 diallel.

The protein content of the parental lines ranged in between 8.93 and 12.94% and those of 45, F_1 crosses, it ranged between 5.23 and 10.56%. The cultivars presently under wide cultivation are medium hard with moderate protein content. The results obtained are in concurrence with those of Desai *et al.*, (1992) and Somani (1992). In the present studies, a significant specific combining ability (sca) was noticed in three F_1 crosses viz., Akms-14B \times IS-6335, SPV-104 \times IS-2284 and SPV-104 \times IS-6335. It is also revealed that this specific constituent is governed by non additive gene inheritance in F_1 diallel.

In this investigation soluble sugars content varied from 1.20 to 2.30% in parents and 0.80 to 1.80% in F_1 crosses. Only one cross SPV-104 x IS-6335 exhibited significant heterotic effects. It is also concluded that this trait is controlled by non additive gene. Glueck *et al.*, (1997) reported that in deteriorated grains, soluble carbohydrates are greatly decreased as they are used to provide energy for the growth and development of fungi. On the contrary Somani (1992) opined that due to enzymolysis soluble sugars increased in old grains.

In the present investigation the TGMR was 1.75 to 2.50 in colored grain that suggest the tannin is deterrent to molding fungi and these observations confirm the findings of earlier workers (Ellis, 1972; Murty, 1975; Glueck and Rooney, 1980; Rooney and Miller, 1981; Hahn *et al.*, 1984; Bandyopadhyay, 1988; Mansuetus, 1990). Nine of the 10 crosses selected for higher germination (Table 77) exhibited significant heterosis for tannin or flavan-4-ol or both. These characters is governed by additive gene inheritance in F_1 diallel.

Hagerman and Butler (1981) noticed tannin associated protein consist of three or components, two of which are high molecular weight prolamins and one of these was rich in proline. Guiragossian *et al.*, (1978) noticed that mutation in P-721 sorghum increased quality of kafirin (prolamin) with and increased albumin and globulin. It suggest the prolamins are inversely correlate with albumin and globulin. Similar trend was noticed in present studies also Subramanian *et al.*, (1990). Similar observations have been noticed for all of crosses in present investigation. However, the percentage of this fraction was on quite lower side i.e. 65.16% in IS-9471 whereas, it was bit less in other colored grain parents.

Prolamin content was maximum in AKms-14B followed by SPV-1201 and GJ-15-15 whereas, crosslink prolamin was maximum in SPV-104 followed by SPV-946 and GJ-15-15. All these parental lines have white grain. It is interesting to note that prolamin and crosslink prolamin were on quite lower side in all colored grain parental lines and hence resistance to grain mold in colored grain cultivars can be attributed to tannins and flavan-4-ols rather than grain hardness. Similar trend was noticed in F_1 crosses where colored grained parental line was used and gene for colored testa was dominant. These findings are in accordance with those of (Glueck and Rooney 1980; Bandyopadhyay 1986; Jambunathan 1986; Murty 1992). In white grained sorghum the prolamin and cross-like prolamin were on higher

in SPV-946, SPV-104 and AKms-14B all these parental lines have more breaking strength by indicating that the endosperm is vitreous and prolamin and cross link prolamin attributes for vitreousness. Clark *et al.*, 1973; Ellis 1972, Somani 1992) noticed five to six dense layer beneath the aleurone layer was present in tolerant to resistant cultivars. Sani and Chandrashekar to had similar observations and reported that, prolamin and cross-prolamin are more in the dense layer which contributed to resist deterioration by grain mold. The present results are with tolerant cultivars are in concurrence with the results of other workers. However, the mold in SPV-104 and AKms-14B was more even though the prolamin and cross prolamin content are high, this might be due to higher mesocarp thickness hence the infestation was more but restricted mere to pericarp only.

In F_1 crosses prolamin and cross link prolamin content was increased where, in one of parent from white grain had more prolamin and cross link prolamin suggesting that gene for this particular traits is dominant.

Effect of pre-treatment

Overall, there was significant increase in germination in pre-treated over untreated control in parental lines, F_1 crosses and F_2 progenies at Akola and Patancheru grown during 1996 and parental lines and F_1 crosses of two seasons at Akola grown during 1995,1996. There was increase in germination at Akola over Patancheru during 1996. There was significant reduction in *F. moniliforme* load at both locations in pre-treated seeds over untreated control, however overall, there were 86.26 and 28.32% increase in *F. moniliforme* infection at Patancheru over Akola in F_1 and F_2 diallel, respectively. These results are in concurrence with results of other workers (Bhagwat and Datar, 1974; Mathur *et al.*, 1975; Castor and Frederiksen, 1981; Jayasekaran, 1983; Granja and Zambolim, 1984; Wu, 1983). The variation between two locations at Akola was due to infection of seed due to *F. moniliforme* at the time of anthesis due to use of continuous rains (Somani 1992). For *F. pallidoroseum* there was a significant increase in load in pre-treated over control whereas, there was an increase of 165.27 and 6% at Patancheru over Akola in F_1 and F_2 diallel on germinated seed. However, two seasons at Akola 1996 recorded more load over Akola 1995 season.

Significant reduction in ungerminated seed was noticed in pre-treated seed over untreated control at both locations and both seasons at Akola. There was significant reduction in fungal load of *F. moniliforme* and *F. pallidoroseum* in F_1 and F_2 progenies, however for *C.*

lunata there was increase in load was recorded in F_1 crosses at Patancheru. This might be due to more superficial infestation of *C. lunata* at Akola under natural condition, whereas due to deep seated infestation the pre-treatment did not revealed any effect on *C. lunata* at Patancheru.

Histopathology

From the infection and colonization studies it was seen that the Fusarial penetration (both species) usually take place from hilar end and no pericarpal infection was observed. Curvularial infection however, is usually through both kernel ends i.e. stylar and hilar as well directly through the pericarp wall. These findings are similar with those of Castor (1977) and Castor and Frederiksen (1980). Inoculations at anthesis has shown development of Fusaria in developing kernels in all the parts of spikelets and denser growth around the ovary base and progress acropetally between aleurone layer and pericarp subsequently endosperm, embryonic tissue. For *C. lunata* infection was observed on ovary wall, pericarp, endosperm and embryonic tissue. Similar observations were recorded by Forbes (1986) and Bandyopadhyay (1986).

Correlation studies

The knowledge of association of different grain mold contributing components is of significant importance in grain mold resistance breeding programme. This study provides reliable information on nature, extent and effectivity of selection. The simple correlation studies between fungal load of untreated and pre-treated, germinated and ungerminated seeds and contribution of physical and biochemical characters if any, towards resistance have been studied in the present investigation since many researchers had putforth different physical and biochemical characters contributing towards grain mold resistance. The available literature however, revealed that no appropriate correlation between various factors for mechanism and genetics of grain mold resistant had been established.

Germinated seed (untreated) has shown negative association with *F. moniliforme*, *C. lunata*, score (GS) and TGMR at Akola location, indicating that with the low fungal load of above fungi there was increased germination and reduced TGMR. Whereas, at Patancheru, fungal load (GS) had significant positive association thereby indicating that, fungal load has no positive effect on germination. However, ungerminated seed had positive correlation with fungal load of all fungi and score, indicating that on ungerminated seed the

ungal load of *F. moniliforme* and *C. lunata* was more that ultimately might have reduced the germination and increased the TGMR. Wu and Cheng (1990) reported similar results for *C. lunata*, *F. moniliforme* and some other fungi.

Positive significant correlation was establish between *F. moniliforme* (GS) and *C. lunata* (GS) with TGMR at Akola. Forbes *et al.* (1989) too established similar correlation regards to *F. moniliforme* only. However, at Patancheru negative significant association of th the fungi with TGMR was noticed. This might be due to continuous congenial condition infection by moldy fungi. Significant association between molding fungi and TGMR was ticed in ungerminated seed, it proved that with increasing fungal load there is decrease in germination and increase in TGMR. Similar observation have been made by Bhatnagar (1971), Tripathi (1974), Castor (1977), Rao and Williams (1977), Castor and Frederiksen (1980), Vidyasekaran (1982) and Forbes (1986).

Correlation studies of fungal load on germinated and ungerminated (treated) seed revealed that, germinated seed had significant negative correlation with *F. moniliforme* (GS), *F. pallidoroseum* (GS), *C. lunata* (GS), other fungi (GS), and TGMR at Akola that, indicates that with the minimum load of these fungi there is improvement in seed germination. However, significant positive association exhibited with germinated (treated) seed and *F. moniliforme*, *F. pallidoroseum*, *C. lunata* and other fungi at Patancheru revealing that the presence of molding fungi did not effect seed germination. It indicates that, there is no effect pre-treatment of seed in impairing germination since controlled conditions at Patancheru ulted in total loss of viability. All molding fungi on germinated seed increased score and reased TGMR at both the locations. Fungal load on ungerminated seed had a significant ociation with all fungi, score and TGMR at both the locations indicating that, presence of re fungal load reduces germination, increase score and TGMR. Forbes *et al.*, (1989) also rorted similar results but for *F. moniliforme* and TGMR.

Correlation between physical characters

ndred grain weight has shown positive correlation with days to 50% flowering at Akola. nilar results were also reported by Patil *et al.* (1980) and Patil and Thombre (1985). However, no correlation studies between grain hardness and mold contributing characters viz. lossperm texture, electrical conductivity, days to 50% flowering, glume covering, mesocarp

less, fungal load of different fungi, disease severity and seed germination has been studied so far. The present studies tried to correlate these factors.

As anticipated the endosperm texture has negative significant correlation with hardness, because the grain hardness is usually exhibited by vitreous endosperm. Similar variation was also reported by Maxson *et al.* (1971). As regards to correlation with electrical conductivity, it is observed that there was positive correlation at one location, where ring was done under natural conditions, even when there was good mold development, however, a negative correlation was established at another location i.e., Patancheru which may be due to totally controlled conditions i.e., due to providing more favourable condition for infection, establishment and deterioration by molding fungi at Patancheru.

Grain hardness exhibited significant positive correlation at both locations with maturity indicating that late maturing crosses escape mold attack and exhibited more grain hardness. Grain hardness has shown significant positive correlation with germination at Patancheru, thereby confirm the finding of Ghorade (1995). As regard to endosperm texture, increasing corneousness of endosperm there is decrease in electrical conductivity, maturity period, mesocarp thickness, *F. moniliforme*, *C. lunata*, TGMR and germination. Joshi *et al.* (1977), Glueck and Rooney (1980), Garud (1992), Kumari *et al.* (1992) and Joshi (1992) recorded similar observation for TGMR.

As regard to electrical conductivity, with the increased maturity period, mesocarp thickness, *F. moniliforme*, *C. lunata* load and TGMR there was increase in electrical conductivity. Whereas, fully covered grain, other fungi and germination exhibited negative correlation with electrical conductivity. Glueck and Rooney (1980) reported similar variation for mesocarp thickness. Forbes (1989) reported similar observations for *F. moniliforme*, TGMR and germination while Ghorade (1995) reported similar trend for germination, *F. moniliforme* and *C. lunata*. As regard to glume covering, increase in glume covering decrease fungal load of all fungi and TGMR and increase in germination was observed. Similar observations were made by Murty (1975) Gangadharan *et al.* (1978), Glueck and Rooney (1980), Mansuetus (1988) and Somani (1992) for TGMR; and Narayana and Joshi (1980) for TGMR. With the increase mesocarp thickness there is increase in fungal load of *F. moniliforme*, *C. lunata* and TGMR and decrease germination. Glueck and Rooney (1980), Castor (1981) and Miller (1981) also reported similar observations.

Increase in fungal load of *F. moniliforme* increases TGM and seed germination at both locations. These results are in accordance with those of Rao and Williams (1977), Denis and Girard (1977), Vidyasekaran (1983), Forbes (1986), Deshmukh (1989) for *C. lunata* and germination. With the increase in fungal load of *C. lunata*, there is increase in TGM and decrease in germination. Similar observations were reported by Tripathi (1974) and Williams (1977).

Correlation between biochemical characters

Results exhibited positive significant correlation with flavan-4-ols indicating that, with the increase in tannins there is corresponding rise in flavan-4-ols. However, tannins and flavan-4-ols had negative significant correlation with grain hardness and TGM indicating that grain hardness is not an important contributing character for grain mold resistance if tannins and flavan-4-ols are present. However, tannins and flavan-4-ols are highly correlated with mold resistance (Mukuru, 1992).

In the present investigation grain hardness and soluble sugars are significantly positively correlated with protein, it indicates that increase in grain hardness is correlated with increase in protein and it might be due to some protein fractions contributing towards hardness.

Tannin is commonly present in the pigmented pericarp and testa layer in sorghum. Condensed tannins are formed by polymerisation of molecular units, of flavanoids as flavan-3-ols (catechin) and flavan-3,4-diols (leucoanthocyanidins). Flavan-3-ols give rise to anthocyanidins Swain and Bate-Smith (1962), Harris (1969), Weinges *et al.* (1969) and Peterson and Butler (1983), Ellis (1972), Harris and Burns (1973), Murty (1975) and Jonathan *et al.* (1986) reported that sorghum seed tannin content is strongly and negatively correlated with seed molding indices.

The present study confirms the finding of above researchers that flavan-4-ols were detected only in coloured grained crosses, and mold infestation was also less and there by results exhibited positive significant correlation with flavan-4-ols and mold resistance. Mukuru (1992) also reported similar finding.

Correlation between protein fractions and physical characters

Results indicated from the present study that albumin and globulin has shown significant positive correlation with prolamin and cross-link prolamin, however, highly negative significant correlation was established with glutelin, glutelin-like, endosperm texture and germination.

Kumari *et al.* (1992) noticed intense deposition of protein in hard grain than in soft grain. Indira *et al.* (1991) and Kumari and Chandrashekar (1992) reported that prolamin is contributing to hardness of grain and corneousness (vitreous) endosperm contains higher protein and such grains are less deteriorated due to grain mold fungi. Glueck and Rooney (1982), Mukuru (1982), Bandyopadhyay (1988) and Somani (1992) noticed that non-pigmented compact cell layer beneath the aleurone layer was present in resistant cultivars. Kumari *et al.* (1992) noticed more prolamin in these 5-6 compact layers. Similarly Abdelrahman and El-Sayed (1984) reported role of cross-link prolamin in grain mold resistance.

It is observed from the present study that glutelin-like and glutelin have a significant positive correlation with endosperm texture and germination and negative correlation with grain hardness. It is seen that with more floury endosperm, there is an increase in grain hardness and increase in germination.

CHAPTER – VI

SUMMARY

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CHAPTER VI

SUMMARY

Grain mold is a major production constraint of early maturing, high yielding sorghum cultivars, in Vidarbha region of Maharashtra. There are number of fungal species involved in the grain mold complex, but the important pathogenic fungi are: *Fusarium moniliforme*, *F. pallidoroseum*, and *C. lunata*.

The present investigation was carried out to determine the mechanism of host resistance, including physical, agronomic and biochemical parameters, and understand genetics and heritability of various parameters imparting resistance to grain mold in selected sorghum lines.

For incorporation of mold resistance into elite material of good grain quality, following aspects were investigated after performing 10 x 10 diallel, excluding reciprocals, viz., amount of heterosis and heterobeltiosis, the general and specific combining ability for selection of potential parents and crosses, and to ascertain the inheritance of some important quantitative (physical, biochemical) characters associated with grain mold resistance.

The experimental material was selected from germplasm collection of Sorghum Research Unit, Dr. PDKV, Akola and ICRISAT, Patancheru, India, comprised of 10 parental lines, SPV-1201, ICSB-101B, SPV-946, GJ-35-15-15, SRI-26B, (elite), SPV-04, AKms-14B (susceptible), IS-2284, IS-6335 and IS-9471 (resistant), having wide range of variability. During the *Kharif* 1994-95 and 1995-96 these parents were crossed to make half diallel (excluding reciprocals) to obtain 45 crosses. An experiment was conducted during *Kharif* 1995 at Akola (10 parents + 45 crosses) to study the mold reaction under natural condition besides agronomic traits, plant characters and to obtain F_1 seed. Another experiment was carried out during *Kharif* 1996 at both locations, Akola under natural conditions and at Patancheru under controlled condition using 10 parents, 45 F_1 crosses and F_2 progenies in a randomized block design.

Data were recorded on plant height, days to 50% flowering, cob length, thresh in mold rating (TGMR), 100-grain weight, glume color, glume covering, grain hardness, endosperm texture, pericarp, mesocarp thickness, presence of testa layer and electrical conductivity of grain leachates. Biochemical characters, such as proteins, proteins fractions

(albumin and globulin, prolamin, cross-link prolamin, glutelin-like, glutelin and residues) soluble sugars, tannins and flavan-4-ols were also studied.

Data were also recorded on grain infection and colonization by *F. moniliforme*, *F. pallidoroseum*, *C. lunata* and other fungi, on pre-treated and untreated germinated and ungerminated seed of parents and F_1 grown at Akola, Kharif 1995 and parents, F_1 and F_2 grown both at Akola and Patancheru, Kharif 1996. Seed germination was recorded using Ragdoll's (rolled paper towel) method and standard blotter plate method of IATA (1976). Observations were recorded on five plants from F_1 and 15 plants from F_2 progenies in each replication. Arithmetic averages of scores were used for further computation. Data were analysed as per Griffings (1956 b) method 2 model 1 as further attended by Singh (1973 a, 1973 b) and standard method suggested by Panse and Sukhatme (1954). The salient features of the results pertaining to mean performance, heterosis, combining ability, physical, agronomic, biochemical traits effect of pre-treatment on fungal load, histopathology of infection and correlation are summarized below :

Significant gca in positive direction for 100-grain weight indicated that parent SPV-1201 and sca for crosses, SPV-1201 \times SRT-26B and AKms-14B \times IS-6335 were giving high value for this character along with superior performance for many other characters. Similarly for germination sca effects for crosses AKms-14B \times IS-6335, AKms-3 \times IS-9471 and SPV-1201 \times SRT-26B were found to possess high germination percentage and were resistant to grain mold with many other grain mold resistance contributing factors in F_2 diallel.

Heterosis and heterobeltiosis were studied for all the characters. The highest estimates of heterobeltiosis were up to 18.75% for 100-grain weight, 5.49% for grain length, -19.63% for endosperm texture, -80.89% for electrical conductivity, -22.04% for days to 50% flowering, -19.04% for plant height, 19.54% for cob length, 25.00% for glume length, -64.39% for mesocarp thickness, -61.11% for TGM, -62.41% for *F. moniliforme*, 25% for *F. pallidoroseum*, -52.72% for *C. lunata*, -22.22% for other fungi, 2.29% for soluble sugars, 0.07% for tannins and 48.37% for flavan-4-ols. Crosses having significant positive heterobeltiosis for germination were recorded in 17 and first four were SPV-1201 \times SRT-26B, SPV-1201 \times ICSB-101B, ICSB-101B \times SPV-946 and ICSB-101B \times IS-9471. Altogether, 37 crosses exhibited significant positive heterosis for germination, first four higher

itude crosses were AKms-14B \times IS-6335, AKms-14B \times IS-9471, ICSB-101B \times AKms-14B and ICSB-101B \times IS-9471.

The mean squares due to genotypes were significant which indicated the presence of substantial degree of diversity for all the characters studied. Variance due to gca were significant for almost all the characters. This indicates the importance of additive as well as non-additive type of gene action in the expression of these characters. But the magnitude of variance due to gca revealed that additive gene action was predominant for characters like cob length, TGMR, *F. moniliforme* and *C. lunata* in F_1 crosses and F_2 diallel progenies. Additive gene action was also predominant for the characters like 100-grain weight, grain hardness, endosperm texture, glume covering, tannins and flavan-4-ols in F_1 crosses. For germination and *F. pallidoseum* the variance due to gca was lower than indicating non additive gene inheritance in F_1 and F_2 progenies; electrical conductivity, DTF, plant height, mesocarp thickness other fungi, proteins and soluble sugars indicated non additive gene action in F_1 crosses and grain hardness, glume covering in F_2 progenies.

Germination test is a standard evaluation method of grain mold resistance. Based on this, the crosses having significantly high specific combining ability effects for germination percentage are considered useful for the purpose of resistance to grain mold. Additive and significant sca effect for germination percentage was obtained in 15 crosses of F_1 diallel set. The sca effects of AKms-14B \times IS-6335 and AKms-14B \times IS-9471 crosses were higher magnitude exhibiting significant superiority over other crosses. The next better group of crosses included five crosses ICSB-101B \times IS-9471, IS-2284 \times IS-6335, SPV-104 \times IS-9471, SPV-104 \times IS-2284, IS-101B \times IS-2284, SPV-104 \times IS-6335 and SPV-1201 \times T-26B. Only eight progenies exhibited significant positive sca effects for germination percentage of F_2 diallel of which first four higher magnitude progenies were AKms-14B \times IS-6335, SPV-1201 \times SRT-26B, AKms-14B \times GJ-35-15-15 and SPV-946 \times IS-6335.

On the basis of average over two locations the highest 100-grain weight, in treatments was recorded in SPV-104 (2.72 g) and (2.66 g), however, cross SPV-1201 \times SPV-104 (2.94 g) exhibited highest grain weight in F_1 diallel. Hundred grain weight was less in all the entries tested at Patancheru during 1996, under controlled condition.

Maximum grain hardness was exhibited in parents SPV-1201 (7.35 kg/cm²) and cross SPV-1201 × GJ-35-15-15 (7.64 kg/cm²). Colored parents and crosses exhibited less breaking strength as compared to white. Over all, less breaking strength was recorded at Patancheru than at Akola. In general, parental line GJ-35-15-15 (24.15%) and F₁ cross SPV-1201 × GJ-35-15-15 (25.13%) recorded corneous endosperm texture. Corneous endosperm contribute toward resistance in white grain type.

In general, more electrical conductivity was recorded at Patancheru over Akola, because grain deterioration was more under controlled condition. Five crosses viz., ICSB-101B × IS-6335, AKms-14B × IS-6335, GJ-35-15-15 × IS-6335, SRT-26B × IS-6335 and SPV-104 × IS-6335 exhibited earliness. More glume covering was recorded in six of 10 crosses viz., AKms-14B × IS-6335, GJ-35-15-15 × IS-6335, SRT-26B × IS-6335, ICSB-101 × IS-2284, ICSB-101B × IS-6335 and SPV-104 × IS-6335, which indicated resistant reaction to mold.

Mesocarp thickness is an important character for grain mold resistance. Thin mesocarp less than (50 µm) was observed in all crosses selected on basis of germination, these are ICSB-101B × IS-2284, ICSB-101B × IS-6335, ICSB-101B × IS-9471, AKms-14B × IS-6335, GJ-35-15-15 × IS-6335 and SPV-104 × IS-6335.

Thresh grain mold rating is an important character for evaluating of cross. Nine out of 10 crosses selected on basis of germination exhibited 2.00 or less than 2.00 TMR grade indicating resistance to grain mold.

Thirty and nine crosses exhibited more than 75% germination at Akola and Patancheru, respectively. On pooled basis 24 crosses recorded germination more than 75% which had only two white crosses viz. SPV-1201 × SRT-26B (77.25%) and SPV-1201 × ICSB-101B (75.50%) exhibited tolerant reaction to mold on germination basis. In general, 39% decrease germination and 86.26 % increase *F. moniliforme* load at Patancheru was recorded.

F. moniliforme is a major grain mold fungus which interfere with carbohydrate translocation causing reduction in kernel size, seed viability and germination. Patancheru location 86.26% and 28.32 % more *F. moniliforme* load was observed over Akola in F₁ and F₂ diallel. Five out of 10 crosses viz., ICSB-101B × IS-6335, ICSB-101B ×

171, AKms-14B × IS-6335, SRT-26B × IS-9471, and SPV-104 × IS-6335 exhibited less *oniliforme* load over better parents and are resistant to grain mold. Over all there was 27 and 36.60% increase in *F. pallidoroseum* load in F₁ and F₂ progenies at Patancheru Akola. It is comparatively of minor importance in grain molding at Akola, whereas, it is important molding fungus next to *C. lunata* at Patancheru.

C. lunata was observed a major molding fungus at Akola recording 108.95 and 14% more load in F₁ and F₂ progenies over Patancheru. Five out of 10 cross viz., ICSB-B × IS-6335, ICSB-101B × IS-9471 GJ-35-15-15 × IS-6335, SRT-26B × IS-9471 and SRT-26B × IS-6335 estimated less *C. lunata* load over better parents and are resistant to grain mold. *C. lunata* was observed a major mold fungus at Akola 108.95 and 46.34% more load over Patancheru. Positive heterosis of highest magnitude was exhibited in cross ICSB-101B × IS-6335, however, significant heterobeltiosis was exhibited in five out of 10 crosses viz., ICSB-101B × IS-6335, ICSB-101B × IS-9471 GJ-35-15-15 × IS-6335, SRT-26B × IS-9471 and SRT-26B × IS-6335. In general at Akola 283.44 and 46.50 % more other fungi load was observed in F₁ and F₂ progenies, respectively over Patancheru.

As regards to biochemical characters, protein fractions albumin and globulin, prolamin, cross-link prolamin, glutelin-like, glutelin and residual protein, soluble sugars, tannins and flavan-4-ols was investigated from Akola 1995 seed samples. Tannins ranged between 0.02 and 5.84 CE% in parents and F₁ crosses and flavan-4-ols between 0.00 and 0.15 A_{550g}⁻¹. Tannins and flavan-4-ols were positively correlated with colored pericarp and endosperm layer. In colored grain TGMR scores of 1.75 to 2.50 suggest that tannin is deterrent to grain mold fungi. Nine out of the 10 crosses selected on the basis of germination percent exhibited significant heterosis and heterobeltiosis for tannin or flavan-4-ols or both.

In white grained sorghum the prolamin and cross-link prolamin were higher in SPV-104 and AKms 14B and all these parental lines had greater seed hardness there by indicating that the endosperm is vitreous and prolamin and cross link prolamin contribute to seed hardness. It is interesting to note that prolamin and cross link prolamin were quite lower in the colored grain parents and crosses, hence resistant to grain mold in colored grain parents and crosses can be attributed to tannins and flavan-4-ols rather than grain hardness.

In present investigation effect of (0.1% Hg Cl₂) on germination and mycoflora of both germinated and ungerminated seed was studied. Over all there was significant

increase in germination in pre-treated over untreated control was observed at both locations. However, there was 27.39% in F_1 crosses and 24.69% in F_2 progenies increased in germination at Akola over Patancheru 1996. For *F. moniliforme* there was significant reduction in load at both locations in pre-treated seeds over untreated control. In general, there was 86.26 and 28.32% increase load of *F. moniliforme* was observed at Patancheru over Akola in F_1 and F_2 diallel, respectively. There was 165.27 and 36.36% increase load of *F. pallidoroseum* was recorded at Patancheru over Akola in F_1 and F_2 respectively. At Akola, higher load was recorded during 1996 than in 1995. For *C. lunata* on germinated seed 0.69% reduction in load was recorded at Akola, however, there was 21.73% increase in load at Patancheru in pre-treated over untreated control. Over all, there was 108.95% and 46.43% increase in *C. lunata* load at Akola over Patancheru in F_1 and F_2 diallel respectively. For other fungi more load was recorded at Akola than Patancheru. In F_1 crosses reduction in fungal load was recorded on ungerminated pre-treatment seed particularly of *F. moniliforme*, however significant increase in load of *C. lunata* was recorded at Patancheru.

Germinated seed had negative correlation with fungal load of all fungi and TMR at Akola, whereas at Patancheru significant positive correlation was noticed in pre-treated and untreated seed. Hundred-grain weight had significant positive correlation with TMR and *F. moniliforme* and significant negative with glume covering and *F. pallidoroseum* at Akola. Grain hardness recorded significant positive association with DTF, significant negative with endosperm texture and *F. pallidoroseum* at Akola. Grain hardness showed negative association with endosperm texture (-0.61) and (-0.33) at both locations. Endosperm texture had significant negative correlation with DTF (-0.52), *C. lunata* (-0.54), critical conductivity (-0.51) germination (-0.41), mesocarp thickness (-0.39), TGM (-0.39) and *F. moniliforme* (-0.35) at Akola, similar trend was exhibited at Patancheru. Electrical conductivity had positive correlation with *F. moniliforme*, TGM, *C. lunata* (Akola), mesocarp thickness and days to 50% flowering at both locations, however significant positive with *F. pallidoroseum* at Akola.

Days to 50% flowering had significant positive correlation with mesocarp thickness, *F. moniliforme*, *C. lunata* and TGM, *F. moniliforme* exhibited significant positive association with germination and positive association with TGM at both locations. *C. lunata* also exhibited positive correlation with TGM and negative with germination at

cola. TGMR has shown highly significant negative correlation with germination (-0.78) and (-0.83) at Akola and Patancheru, respectively.

Protein content had positive correlation with grain hardness (0.29). Total gars exhibited significant correlation with tannins (0.35); tannins and flavan-4-ols had highly significant negative correlation with grain hardness and TGMR. Prolamin and cross-k prolamin exhibited positive correlation with grain hardness (0.63, 0.55), respectively, never negative with germination, glutelin, glutelin-like and endosperm texture. Glutelin-; and glutelin exhibited significant positive association with germination, endosperm texture, whereas, negative with grain hardness, and mesocarp thickness.

Some physical and biochemical characteristics contribute to resistance to grain molds. In white grained sorghum greater grain hardness, low electrical conductivity, thin mesocarp, more glume covering are some of the desired traits besides more of prolamin and amin-like protein fraction and for some of the characters respective gene (s) are dominant and have additive action. However, in red grains, condensed tannins, flavan-4-ols and presence of testa layer are some of the biochemical characters contributing towards resistance to mold. In Indian sorghums are white grained sorghum, some additive desirable gene can be incorporated in white background source. Histopathological studies, however, confirmed the earlier results of various workers. The study has clearly established the genetics, inheritance of physical and biochemical traits for resistance.

Implication of these findings for genetic improvement of mold resistance in sorghum is considered and the following suggestions are offered on breeding methodology for better utilization of the experimental material.

Parental lines IS-9471, SPV-1201, IS-6335, GJ-35-15-15 and IS-2284 with high gca effect for most of the characters contributing towards resistance to grain mold may be utilized in a hybrid breeding program.

The crosses: Akms 14B \times IS 6335, ICSB-101 B \times IS-9471, ICSB-101B \times IS-2284, SPV-104 \times IS-6335, GJ-35-15-15 \times IS-6335 and SPV-1201 \times SRT-26B exhibited high gca effects, desirable heterosis for most of the characters and additive gene action for important traits (agronomic, physical and biochemical) related to grain mold resistance. Therefore it is suggested that these crosses and their progenies may be utilized to generate better tolerance to grain mold infection.

Important traits imparting resistance to grain mold fungi are: low electrical conductivity, low mesocarp, low TGMR, grain hardness, more glume covering, more tannins, more

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I - 1

Table: 1 Weekly weather data for year 1995 and 1996 recorded at Agro-meteorological laboratory, Akola

1995							1996						
Week	Date	Rainfall in (mm)	Rainy days	Temperature (°C)	Relative humidity (%)		Week	Date	Rainfall in (mm)	Rainy days	Temperature (°C)	Relative humidity (%)	
				Max.	Min.						Max.	Min.	Morning Evening
23	4-10 June	0.00	NIL	44.30	28.40	63.00	23	4-10 June	0.00	NIL	42.00	26.40	£6.00 31.00
24	11-17	13.20	3	41.30	27.00	71.00	24	11-17	0.00	NIL	41.50	25.40	£8.00 34.00
25	18-24	10.00	2	38.90	26.30	73.00	25	18-24	24.40	1	37.90	24.10	77.00 41.00
26	25-1 July	31.60	3	35.00	23.70	86.00	26	25-1 July	0.00	NIL	33.20	25.50	77.00 41.00
27	2-8	4.40	1	38.20	25.50	80.00	27	2-8	9.40	2	37.00	25.40	79.00 41.00
28	9-15	201.60	5	33.40	22.90	92.00	28	9-15	4.80	NIL	35.60	24.70	£0.00 48.00
29	16-22	63.20	4	29.60	23.00	90.00	29	16-22	32.60	2	34.70	24.40	£4.00 58.00
30	23-29	54.0	4	30.60	22.80	91.00	30	23-29	68.40	6	27.80	22.60	£1.00 85.00
31	30-5 Aug.	8.00	2	31.40	23.40	90.00	31	30-5 Aug.	25.20	3	30.00	23.10	£1.00 75.00
32	6-12	3.60	1	31.80	23.50	83.00	32	6-12	11.50	1	28.60	22.70	£1.00 79.00
33	13-19	0.00	NIL	33.30	23.00	78.00	33	13-19	5.60	1	32.20	23.60	£6.00 57.00
34	20-26	1.00	NIL	33.90	23.70	79.00	34	20-26	74.90	1	31.20	22.90	£8.00 71.00
35	27-2 Sep.	78.30	1	33.40	24.50	85.00	35	27-2 Sep.	51.80	3	29.40	22.60	£2.00 74.00
36	3-9	45.60	1	30.30	21.90	89.00	36	3-9	133.90	2	31.00	22.80	£2.00 70.00
37	10-16	7.50	2	33.60	22.40	91.00	37	10-16	55.10	3	31.80	23.10	£2.00 69.00
38	17-23	8.70	1	32.40	22.00	85.00	38	17-23	1.80	NIL	30.90	21.80	£8.00 66.00
39	24-30	9.80	1	35.30	21.60	83.00	39	24-30	20.40	2	34.10	22.90	£4.00 54.00
40	1-7 Oct.	0.00	NIL	36.20	20.80	81.00	40	1-7 Oct.	51.40	4	31.10	22.30	£8.00 65.00
41	8-14	0.00	NIL	34.30	20.80	82.00	41	8-14	0.00	NIL	32.80	17.60	£0.00 34.00
42	15-21	2.00	NIL	32.90	19.80	90.00	42	15-21	0.00	NIL	32.60	17.40	£7.00 37.00
43	22-28	0.00	NIL	34.00	13.60	76.00	43	22-28	97.40	5	29.80	20.90	£3.00 79.00
44	29-4 Nov.	0.00	NIL	33.50	11.60	81.00	44	29-4 Nov.	8.50	1	31.00	14.90	£9.00 33.00
45	5-12	0.00	NIL	32.30	11.20	80.00	45	5-12	0.00	NIL	31.70	16.80	£7.00 39.00

Sources and degrees of freedom

Sr. No.	Characters	Sources and degrees of freedom									
		Environ- ments (1)	Treat- ments (54)	Parents (9)	Hybrids (44)	Parents vs hybrids (1)	Treat - ments x Env. (54)	Parents x Env. (9)	Hybrids x Env. (44)	Parents vs. Hybrids x Env. (1)	Error (108)
1	100 grain weight	170.26**	7.25**	12.03**	6.15**	12.45**	1.91**	2.36*	1.78**	3.32	0.05
2	Grain hardness	302.43**	4.77**	4.38**	4.80**	7.04**	1.59**	2.16*	1.37	6.18*	0.60
3	Endosperm texture	11.71**	13.26**	18.67**	11.04**	62.38**	1.80**	0.55	2.09**	0.56	37.10
4	Ele. conductivity	419.70**	14.69**	27.16**	8.48**	175.73**	1.60**	11.84**	4.44**	54.24**	2111.85
5	DTF	122.67**	25.55**	18.42**	26.84**	33.16**	5.78**	3.09**	6.11**	15.33**	5.18
6	Plant height	0.26	39.60**	45.36**	28.08**	494.63**	2.55**	4.56**	2.11**	3.88	206.43
7	Cob length	17.43**	4.49**	4.24**	3.89**	33.02**	0.87	0.99	0.87	0.31	5.32
8	Glume covering	19.99**	4.11**	4.86**	4.01**	1.70	1.29	0.95	1.32	2.70	88.80
9	Mesocarp thickness	0.67	54.57**	28.44**	55.34**	255.80**	1.64*	2.38*	1.49*	1.69	25.46
10	TGMR	192.86**	27.42**	31.12**	26.04**	54.92**	8.27**	7.48**	8.26**	15.64**	0.12
11	Germination	726.47**	25.22**	24.70**	19.40**	285.70**	4.66**	4.40**	4.27**	24.43**	22.87
12	<i>F. moniliforme</i>	637.64**	4.73**	5.60**	4.10**	24.35**	1.65**	0.99	1.82**	0.08	76.78
13	<i>F. pallidoroeseum</i>	603.45**	1.84**	1.17	1.95**	3.21	1.71**	1.10	1.87**	0.05	10.17
14	<i>C. lunata</i>	333.63**	3.49**	5.00**	3.04**	9.70**	2.70**	1.47	3.02**	0.01	68.76
15	Other fungi	426.93**	1.81**	0.99	1.64*	16.55**	1.43**	0.77	1.59**	0.32	36.48

Figures in parenthesis indicate degrees of freedom

**** Significant at 5%**

Significant at 1%

II - 2

Table 3 Analysis of variance (Pooled F₂) for physical, agronomic and pathological traits, Akola and Patancheru, 1996

Sr. No.	Characters	Sources and degrees of freedom									Error
		Environ- ments (Locations)	Treat- ments	Parents	F ₂ crosses	Parents vs F ₂ crosses	Treat- ments × Env.	Parents × Env.	F ₂ crosses × Env.	Parents vs. F ₂ crosses × Env.	
		(1)	(54)	(9)	(44)	(1)	(54)	(9)	(44)	(1)	(108)
1	Grain hardness	341.10**	5.42**	4.90**	3.53**	93.71**	3.48**	2.65**	3.55**	7.55**	0.50
2	Ele. conductivity	269.45**	33.45**	39.75**	31.97**	42.06**	9.76**	17.56**	5.01**	148.62**	1443.08
3	Plant height	308.94**	6.80**	18.16**	4.56**	3.24	3.15**	1.59	2.68**	37.70**	458.94
4	Cob length	0.30	5.44**	6.06**	4.86**	25.39**	1.55*	1.47	1.48*	5.72**	3.77
5	Glume covering	0.11	3.37**	6.09**	2.79**	4.81*	0.87	0.85	0.90	0.02	41.04
6	TGMR	140.22**	10.17**	28.31**	4.23**	108.05**	2.36**	5.00**	1.61*	11.23**	11.82
7	Germination	215.10**	7.63**	12.40**	4.44**	104.86**	2.57**	1.86	2.08**	30.52**	46.24
8	<i>F. moniliforme</i>	130.25**	1.81**	2.87**	1.13	22.02**	1.01	1.26	0.95	1.15	121.77
9	<i>F. pallidoroseum</i>	101.49**	1.16	1.01	0.97	11.18**	1.41*	1.49	0.50	40.89**	314.27
10	<i>C. lunata</i>	116.39**	2.59**	3.59**	2.35**	4.28*	1.47*	1.80	1.40	1.92	94.42
11	Other fungi	70.46**	1.15	1.16	1.08	4.02*	1.52*	0.80	1.10	26.66**	29.77

Figures in parenthesis indicate degrees of freedom

* Significant at 5%

** Significant at 1%

II - 3

Table. 4 Analysis of variance for biochemical characters of parental lines and F₁ crosses at Akola, 1995.
Akola 1995

Sr.No.	Characters	Sources and degrees of freedom					
		Replications	Treatments	Parents	Hybrids	Parents vs. Hybrids	Error
		(1)	(54)	(9)	(44)	(1)	(54)
1	Protein	0.302	211.761**	65.035**	104.552**	6249.488**	0.00103
2	Soluble sugars	0.507	8.629**	9.018**	6.565**	95.940**	0.00324
3	Tannins	3.206	2061.853**	3809.883**	1747.343**	168.055**	0.00028
4	Flavan-4-ols	3.558	846.121**	1267.962**	769.066**	439.937**	0.0028

Figures in parenthesis indicate degrees of freedom.

* Significant at 5%

** Significant at 1%

Table 5 Mean values of parents and F₁ crosses for different characters in 10 × 10 diallel of sorghum Akola and Patancheru, 1996

Sr. No	Parents/Crosses	100 grain weight (g)				Grain hardness kg/cm ²				Endosperm texture (%)				Elec. conductivity (millimhos)				Days* to 50% flowering			
		L1	L2	Pooled	L1	L2	Pooled	L1	L2	Pooled	L1	L2	Pooled	L1	L2	Pooled	L1	L2	Pooled		
1	SPV-1201	2.81	2.51	2.66	8.67	6.03	7.35	30.80	27.90	29.35	159.50	397.50	372.00	278.50	73.00	307.50	73.00	78.00	73.00		
2	ICSB-101B	2.51	1.61	2.06	6.94	5.58	6.26	36.15	29.70	27.93	242.50	372.00	372.00	307.25	72.00	307.25	72.00	73.00	73.00		
3	AKma-14B	2.57	1.36	1.97	6.95	3.15	5.05	32.80	34.00	33.40	188.50	371.00	371.00	379.75	68.50	379.75	68.50	70.00	69.25		
4	SRT-26B	2.06	1.47	1.76	6.56	4.44	5.50	32.10	40.55	36.33	300.50	445.00	445.00	372.75	69.50	372.75	69.50	73.00	71.50		
5	GI-35-15-15	2.24	1.58	1.91	8.11	4.52	6.31	22.40	25.90	24.15	218.20	342.50	342.50	280.25	77.50	280.25	77.50	79.00	78.25		
6	SPV-946	2.19	1.73	1.96	7.25	3.75	5.50	35.05	27.65	31.35	151.50	446.50	446.50	299.00	77.50	299.00	77.50	78.75	77.75		
7	SPV-104	2.90	2.54	2.72	6.08	2.78	4.43	46.80	52.25	49.53	260.00	846.50	846.50	553.25	80.00	553.25	80.00	76.50	74.25		
8	IS-2284	1.76	1.45	1.60	5.91	4.48	5.20	36.95	56.25	55.60	108.00	226.50	226.50	167.25	70.00	167.25	70.00	76.50	73.25		
9	IS-6335	1.69	1.59	1.64	6.68	5.52	6.10	55.85	55.95	55.90	122.00	214.00	214.00	168.00	66.00	168.00	66.00	59.00	62.50		
10	IS-9471	1.91	1.73	1.82	6.10	5.32	5.72	55.30	59.15	57.23	105.00	231.50	231.50	168.25	71.50	168.25	71.50	70.50	71.00		
11	SPV-1201 x ICSB - 101B	2.62	2.38	2.50	7.14	7.05	7.09	32.10	29.25	30.67	138.00	292.00	292.00	215.00	74.50	215.00	74.50	83.50	78.00		
12	SPV-1201 x AKma - 14 B	2.61	1.85	2.23	5.57	5.24	5.40	38.75	36.55	37.65	150.00	399.50	399.50	274.75	75.00	274.75	75.00	83.50	79.25		
13	SPV-1201 x SRT - 26B	3.25	2.08	2.66	7.23	5.25	6.24	43.25	40.80	42.03	155.00	287.50	287.50	221.25	76.00	221.25	76.00	83.50	88.75		
14	SPV-1201 x GI-35-15-15	2.53	1.80	2.17	7.85	7.43	7.64	21.15	29.10	25.13	135.00	275.00	275.00	203.00	74.50	203.00	74.50	83.50	88.00		
15	SPV-1201 x SPV - 946	1.99	2.06	2.02	6.97	6.88	6.92	33.95	35.80	34.88	237.50	278.00	278.00	257.75	76.00	257.75	76.00	80.50	78.25		
16	SPV-1201 x SPV - 104	3.41	2.47	2.94	6.46	6.22	6.34	45.05	48.35	46.70	144.00	408.00	408.00	276.00	74.50	276.00	74.50	83.50	88.00		
17	SPV-1201 x IS - 2284	2.26	1.77	2.02	7.30	4.55	5.93	48.00	57.75	52.88	118.00	201.50	201.50	259.75	71.00	259.75	71.00	82.00	76.50		
18	SPV-1201 x IS-6335	2.29	1.72	2.00	7.60	5.05	6.32	45.35	50.15	47.25	102.50	209.50	209.50	156.00	71.50	156.00	71.50	82.00	78.50		
19	SPV-1201 x IS - 9471	2.71	2.09	2.40	6.08	5.23	5.65	51.05	56.20	53.63	112.50	217.00	217.00	164.75	74.00	164.75	74.00	80.50	77.25		
20	ICSB - 101B x AKma 14 B	1.92	1.88	1.90	6.86	3.95	5.40	45.00	42.95	45.47	156.00	358.50	358.50	257.25	68.00	257.25	68.00	71.00	68.50		
21	ICSB - 101B x SRT - 26 B	1.95	1.79	1.87	5.27	3.80	4.53	41.30	35.15	38.22	167.00	284.50	284.50	225.75	68.50	225.75	68.50	70.00	69.25		
22	ICSB - 101B x GI-35-15-15	1.95	1.77	1.87	6.64	5.12	5.88	31.10	25.55	28.33	164.00	257.50	257.50	210.75	72.00	210.75	72.00	73.00	73.00		
23	ICSB - 101B x SPV - 946	2.43	1.82	2.13	7.32	5.60	6.46	35.80	38.35	37.07	124.50	339.50	339.50	232.00	70.50	232.00	70.50	80.50	73.50		
24	ICSB - 101B x SPV - 104	2.88	2.58	2.72	6.64	3.90	5.28	39.75	41.95	40.85	150.50	400.00	400.00	275.25	71.00	275.25	71.00	73.00	72.00		
25	ICSB - 101B x IS - 2284	2.08	2.20	2.14	6.44	4.61	5.53	46.10	55.15	50.63	106.50	167.00	167.00	136.75	69.50	136.75	69.50	71.00	78.25		
26	ICSB - 101B x IS - 6335	2.08	2.04	2.06	6.70	5.94	6.32	43.55	50.65	47.10	99.00	122.00	122.00	110.50	62.00	110.50	62.00	60.00	61.00		
27	ICSB - 101B x IS - 9471	2.17	2.17	2.17	5.88	5.07	5.48	34.90	53.15	54.03	94.00	154.00	154.00	126.50	66.00	126.50	66.00	70.50	68.25		
28	AKma - 14 B x SRT - 26B	2.11	1.94	2.03	3.96	2.85	3.40	54.00	52.20	53.10	230.00	343.00	343.00	286.50	66.50	286.50	66.50	70.50	68.50		
29	AKma - 14 B x GI-35-15-15	1.79	1.77	1.70	4.64	2.82	3.73	38.25	44.70	41.47	137.50	342.50	342.50	250.00	68.00	250.00	68.00	70.00	68.25		
30	AKma - 14 B x SPV - 946	2.49	2.06	2.28	6.32	3.63	4.97	55.35	50.20	52.78	150.50	347.50	347.50	249.00	66.00	249.00	66.00	70.50	68.50		
31	AKma - 14 B x SPV - 104	2.88	2.22	2.55	4.89	3.45	4.17	56.30	57.20	56.75	177.00	485.00	485.00	331.00	68.00	331.00	68.00	71.50	68.75		
32	AKma - 14 B x IS - 2284	2.38	1.76	2.07	6.19	4.77	5.48	55.75	66.60	61.17	114.00	172.50	172.50	143.25	65.00	143.25	65.00	71.50	68.50		
33	AKma - 14 B x IS - 6335	2.42	2.05	2.23	6.72	4.22	5.47	64.55	63.25	63.90	98.50	122.50	122.50	110.50	60.50	110.50	60.50	59.00	58.75		
34	AKma - 14 B x IS - 9471	2.25	2.32	2.29	5.35	4.04	4.70	38.80	62.15	60.48	100.50	134.50	134.50	117.50	65.00	117.50	65.00	60.00	62.50		

Table 5 Condt.

2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17
SRT - 26B x GJ-35-15-15	2.16	1.59	1.88	6.95	4.52	5.74	41.30	33.25	37.28	149.50	407.00	278.25	70.00	73.00	71.50
SRT - 26B x SPV - 946	2.38	2.02	2.20	5.84	4.53	5.18	46.35	41.90	44.13	174.00	267.50	220.75	72.00	75.00	73.50
SRT - 26B x SPV - 104	2.72	2.13	2.42	6.45	3.99	5.22	52.55	50.65	51.60	166.50	329.00	247.50	66.50	71.50	69.50
SRT - 26B x IS - 2284	2.27	1.81	2.04	5.36	4.92	5.14	46.75	57.70	52.22	119.00	140.00	129.50	60.50	71.50	66.00
SRT - 26B x IS - 6335	2.31	1.87	2.09	6.09	4.18	5.13	59.65	60.15	59.90	123.00	84.50	103.75	59.50	60.00	59.75
SRT - 26B x IS - 9471	2.15	2.15	2.15	5.63	3.53	4.58	34.50	64.00	49.25	201.00	113.00	157.00	70.50	71.00	70.75
GJ-35-15-15 x SPV - 946	2.50	1.72	2.11	7.68	3.77	5.73	36.15	30.45	33.30	184.50	318.50	251.50	76.00	78.50	77.25
GJ-35-15-15 x SPV 104	2.26	2.12	2.19	7.28	3.83	5.55	48.35	41.10	44.72	186.00	387.00	286.50	72.50	74.00	73.25
GJ-35-15-15 x IS - 2284	1.92	1.56	1.74	7.03	4.27	5.65	46.30	59.10	52.70	129.00	168.00	128.50	65.50	77.00	71.25
GJ-35-15-15 x IS - 6335	2.03	1.51	1.77	5.77	4.06	4.91	46.85	61.10	53.97	106.50	120.00	113.25	63.00	67.00	65.00
GJ-35-15-15 x IS - 9471	2.02	1.84	1.93	5.91	4.24	5.07	42.90	61.60	52.25	157.00	172.00	164.50	72.00	71.50	71.25
SPV - 946 x SPV - 104	2.79	2.13	2.46	6.41	3.85	5.13	43.40	44.30	43.85	138.50	475.00	306.75	72.00	78.50	75.25
SPV - 946 x IS - 2284	2.48	1.74	2.11	6.22	4.52	5.37	35.50	55.70	45.60	108.00	115.50	111.75	66.00	71.50	68.75
SPV - 946 x IS - 6335	1.99	1.72	1.85	6.99	4.72	5.86	36.05	53.80	44.92	115.50	126.50	121.00	67.50	93.50	75.50
SPV - 946 x IS 9471	2.04	2.07	2.06	6.42	4.35	5.39	45.75	56.75	51.25	122.50	82.00	102.25	74.50	59.00	66.75
SPV - 104 x IS 2284	2.32	2.03	2.17	4.45	3.58	4.01	72.25	56.35	64.30	118.50	252.50	185.50	68.50	74.00	71.25
SPV - 104 x IS 6335	2.34	1.99	2.16	5.33	3.44	4.38	74.00	64.25	69.13	114.50	169.50	142.00	63.00	59.00	61.00
SPV - 104 x IS 9471	2.71	2.59	2.65	4.69	3.58	4.13	68.55	63.20	65.88	107.50	104.00	105.75	64.00	59.00	61.50
IS 2284 x IS - 6335	1.76	1.63	1.69	6.83	4.36	5.59	53.20	55.85	54.53	108.00	188.50	148.25	62.00	70.50	66.25
IS 2284 x IS - 9471	1.94	1.76	1.86	5.63	4.75	5.19	53.65	62.05	57.85	114.00	187.00	150.50	70.00	74.00	72.00
IS 6335 x IS 9471	2.19	1.75	1.97	5.32	4.37	4.85	53.40	56.65	55.03	90.00	159.00	124.50	62.00	59.00	60.50
Parental range	1.69	1.36	1.60	5.91	2.78	4.43	22.40	25.90	24.15	105.00	214.00	167.25	66.00	59.00	62.50
	to	to	to	to	To	to	to	to	to	to	to	to	to	to	to
	2.90	2.54	2.72	8.67	6.03	7.35	56.95	59.15	57.23	300.50	846.50	553.25	80.00	79.00	78.25
Hybrid range	1.76	1.51	1.68	3.96	2.82	3.40	21.15	25.55	25.13	90.00	82.00	102.25	59.50	59.00	59.75
	to	to	to	to	To	to	to	to	to	to	to	to	to	to	to
	3.41	2.58	2.94	7.85	7.43	7.64	74.00	66.60	69.13	237.50	485.00	331.00	76.00	85.50	80.75
SE (m)	0.155	0.090	0.249	0.479	0.549	1.051	2.628	3.402	3.730	17.909	42.758	85.400	0.852	0.639	3.452
CD at 5%	0.442	0.257	0.710	1.367	1.567	3.000	7.502	9.711	10.650	51.110	122.028	243.721	2.433	1.825	9.854

L₁ = AkolaL₂ = Patancheru

Table 6 Condi..... 2

	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17
35	SRT - 26B x GI-35-15-15	206.50	197.50	202.00	26.70	24.70	23.70	50.00	50.00	62.66	64.15	63.40	3.00	5.00	4.00
36	SRT - 26B x SPV - 946	253.00	254.00	253.50	23.90	27.50	23.70	50.00	50.00	85.91	93.56	89.73	2.75	4.00	3.38
37	SRT - 26B x SPV - 104	213.50	213.50	213.50	22.10	24.80	23.45	50.00	50.00	87.56	86.00	86.78	3.00	5.00	4.00
38	SRT - 26B x IS - 2284	304.50	307.00	304.25	24.10	27.30	25.70	50.00	50.00	63.61	62.16	62.89	2.50	2.00	2.25
39	SRT - 26B x IS - 6335	274.50	248.50	261.50	23.40	26.55	24.98	62.50	62.50	59.67	62.79	61.23	2.50	2.00	2.25
40	SRT - 26B x IS - 9471	259.50	282.50	271.00	22.80	30.60	26.70	50.00	50.00	57.32	49.58	53.45	2.50	1.50	2.00
41	GI-35-15-15 x SPV - 946	275.00	244.50	260.50	23.20	22.20	22.70	50.00	62.50	56.25	75.72	61.34	68.53	2.50	3.75
42	GI-35-15-15 x SPV 104	313.50	255.50	279.50	23.00	24.90	23.95	50.00	50.00	67.44	80.61	74.03	2.25	5.00	3.63
43	GI-35-15-15 x IS - 2284	309.50	320.50	300.00	27.60	26.70	27.15	62.50	62.50	45.97	53.74	49.85	2.00	2.00	2.00
44	GI-35-15-15 x IS - 6335	329.00	303.50	316.25	29.70	26.50	28.10	62.50	68.75	36.55	35.90	36.06	2.00	2.00	2.00
45	GI-35-15-15 x IS - 9471	291.50	308.00	299.75	25.70	31.05	28.38	50.00	50.00	48.55	33.89	41.22	2.50	1.50	2.00
46	SPV - 946 x SPV - 104	311.50	288.50	300.00	25.10	25.00	25.05	50.00	62.50	49.41	68.85	59.13	2.00	5.00	3.75
47	SPV - 946 x IS - 2284	279.50	297.50	288.50	21.40	28.20	24.80	50.00	37.50	43.75	56.92	53.81	2.50	2.00	2.25
48	SPV - 946 x IS - 6335	297.50	322.00	309.25	26.00	27.10	26.55	62.50	50.00	56.25	43.94	38.51	41.22	2.00	2.00
49	SPV - 946 x IS 9471	328.50	288.00	308.25	26.10	29.20	27.65	62.50	50.00	33.78	35.56	34.67	2.00	2.00	2.00
50	SPV - 104 x IS 2284	288.50	302.00	295.25	22.70	22.60	22.65	37.50	75.00	56.25	37.09	39.47	38.28	2.00	2.50
51	SPV - 104 x IS 6335	297.50	269.50	293.50	26.50	27.60	27.05	75.00	75.00	32.03	31.32	31.68	2.00	2.00	2.00
52	SPV - 104 x IS 9471	283.50	256.00	269.75	25.30	28.40	26.85	37.50	50.00	43.75	47.26	46.95	2.00	2.00	2.00
53	IS 2284 x IS - 6335	280.50	302.50	291.50	25.60	26.30	25.95	75.00	75.00	61.75	62.57	62.16	2.00	1.50	1.75
54	IS 2284 x IS - 9471	270.00	301.00	285.50	23.30	25.30	24.30	37.50	37.50	66.72	64.73	65.73	2.00	2.00	2.00
55	IS 6335 x IS 9471	287.50	274.00	280.75	29.40	28.55	28.98	50.00	50.00	32.20	33.82	33.01	2.00	1.50	1.75
Parental range															
	136.00	140.00	138.00	21.20	17.00	19.10	50.00	50.00	50.00	45.45	59.91	54.02	2.00	1.50	1.75
	to	to	to	to	to	to	to	to	to	to	to	to	to	to	to
	283.00	285.50	273.75	26.20	28.90	27.55	75.00	75.00	75.00	97.52	96.57	97.05	4.00	5.00	4.50
Hybrid range															
	183.00	183.00	183.00	20.40	21.00	20.95	37.50	37.50	37.50	32.20	33.82	33.01	2.00	1.50	1.75
	to	to	to	to	to	to	to	to	to	to	to	to	to	to	to
	343.00	340.00	341.50	29.70	32.10	30.80	75.00	75.00	75.00	96.79	99.56	98.13	3.50	5.00	4.25
SE (m)															
	3.708	6.047	11.12	1.634	1.434	1.169	6.306	6.077	5.899	2.926	6.773	2.040	0.207	0.252	0.565
CD at 5%															
	10.582	17.257	37.713	4.664	4.093	3.338	17.998	17.343	16.835	8.352	19.330	5.822	0.593	0.719	1.614

Table : 7 Mean values of parents and F_1 crosses for different characters in 10×10 diallel of sorghum Akola and Patancheru, 1996

Sr. No	Parents/Crosses	Germination (%)			<i>P. mouliformis</i> (%)						<i>P. pellidivocans</i> (%)						<i>C. lunata</i> (%)						Other fungi (%)		
		L1	L2	Pooled	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17				
1	SPV-1201	67.50 (35.25)	30.00 (33.07)	48.75 (44.15)	20.67	54.67	37.67	0.71	12.00	6.36	56.00	34.67	45.33	23.33	1.02	12.18									
2	ICSB-101B	47.00 (43.25)	26.50 (30.98)	36.75 (37.11)	26.67	52.67	39.67	0.71	12.00	6.69	58.00	28.00	43.00	15.34	6.00	10.67									
3	AKma-14B	29.00 (32.53)	1.50 (8.64)	15.75 (20.38)	46.00	58.67	52.33	0.71	7.34	4.02	44.00	34.00	9.00	9.34	1.02	5.18									
4	SRT-26B	81.50 (64.62)	12.00 (20.12)	46.75 (42.37)	26.01	51.34	38.67	2.36	10.00	6.18	56.00	37.33	46.67	16.00	1.69	8.85									
5	GI-35-15-15	72.50 (51.40)	29.50 (32.89)	51.00 (42.15)	18.66	52.67	35.66	0.71	13.34	7.02	60.00	30.00	45.00	21.34	4.00	12.67									
6	SPV-946	75.50 (50.49)	18.00 (25.49)	46.00 (42.69)	18.00	62.67	40.34	1.02	14.67	7.84	57.34	26.00	41.67	26.00	1.02	13.51									
7	SPV-104	50.00 (45.03)	61.00 (13.98)	28.00 (29.49)	38.67	71.34	55.00	0.71	10.00	5.36	44.67	18.67	31.67	16.00	0.71	8.36									
8	IS-2284	82.50 (65.03)	53.00 (55.03)	71.75 (58.19)	20.00	48.00	33.99	1.69	19.34	10.51	46.67	24.00	35.33	24.67	5.33	15.00									
9	IS-6335	86.00 (68.04)	48.00 (46.73)	69.50 (57.39)	7.99	47.33	27.66	2.67	10.67	6.67	26.00	15.33	20.67	20.00	5.34	12.67									
10	IS-9471	79.00 (63.41)	48.50 (44.14)	63.75 (53.38)	5.33	34.00	27.66	1.69	12.00	5.51	24.00	22.67	23.34	24.00	7.34	13.67									
11	SPV-1201 x ICSB - 101B	85.00 (67.26)	66.00 (54.34)	75.50 (60.00)	18.01	53.33	35.67	1.69	12.00	6.84	56.67	29.33	43.00	24.00	5.34	14.67									
12	SPV-1201 x AKma - 14 B	62.50 (52.23)	33.00 (33.65)	47.25 (42.95)	13.34	55.00	34.34	0.71	16.67	8.69	61.33	21.34	41.33	25.33	8.00	16.67									
13	SPV-1201 x SRT - 26B	69.00 (67.63)	69.00 (56.49)	77.25 (62.06)	15.34	36.00	25.67	0.71	10.00	5.36	50.67	30.00	41.33	34.00	24.00	29.00									
14	SPV-1201 x GI-35-15-15	64.50 (53.47)	58.00 (49.90)	61.50 (51.68)	29.33	55.99	42.00	38.99	0.71	10.67	5.69	50.00	26.00	38.00	20.00	20.00									
15	SPV-1201 x SPV - 946	77.00 (64.05)	32.00 (34.36)	55.00 (48.24)	14.67	37.34	22.83	0.71	7.34	4.02	55.33	37.33	46.33	32.67	9.34	21.00									
16	SPV-1201 x ISV - 104	98.00 (82.14)	75.50 (60.43)	86.75 (71.28)	8.36	37.00	31.33	2.35	16.67	9.51	28.00	32.00	40.33	32.67	9.34	21.00									
17	SPV-1201 x IS - 2284	97.00 (80.17)	66.00 (54.60)	81.00 (67.98)	7.34	40.00	23.67	0.71	16.00	8.35	18.00	31.34	24.67	18.67	8.66	15.00									
18	SPV-1201 x IS - 9471	94.00 (76.72)	77.50 (55.23)	80.75 (65.98)	8.66	6.00	17.33	2.00	12.66	7.30	18.00	28.00	23.00	17.33	10.00	14.34									
19	ICSB - 101B x AKma 14 B	66.50 (54.70)	42.90 (38.65)	54.50 (47.69)	19.34	56.00	37.67	1.69	14.66	8.59	59.33	16.66	36.00	19.34	6.00	12.67									
20	ICSB - 101B x SRT - 26 B	67.50 (55.25)	39.00 (38.63)	53.25 (46.93)	11.99	64.67	39.33	0.71	18.67	9.69	68.00	14.00	41.00	21.34	1.33	11.33									
21	ICSB - 101B x GI-35-15-15	72.50 (58.42)	58.00 (49.61)	65.25 (54.02)	9.34	53.33	32.33	0.71	16.00	8.36	67.33	22.67	45.00	23.34	6.00	14.67									
22	ICSB - 101B x SPV - 946	77.50 (61.75)	63.00 (52.51)	70.25 (57.16)	14.67	38.00	36.33	3.02	16.66	9.89	49.33	22.00	35.67	34.00	2.00	18.00									
23	ICSB - 101B x SPV - 104	38.50 (49.89)	33.00 (37.44)	47.75 (43.67)	34.01	70.67	52.34	0.71	21.34	11.92	49.33	8.00	28.67	17.34	0.71	9.02									
24	ICSB - 101B x IS - 2284	95.00 (77.16)	85.00 (67.50)	90.00 (72.32)	14.66	32.00	23.33	0.71	5.34	3.02	38.00	20.66	29.33	33.33	8.67	21.00									
25	ICSB - 101B x IS - 6335	91.00 (72.56)	88.00 (70.54)	89.75 (71.35)	9.99	21.34	15.66	1.02	6.67	3.85	26.66	14.00	20.33	34.00	7.33	20.66									
26	ICSB - 101B x IS - 9471	96.50 (79.24)	86.00 (63.78)	88.25 (71.31)	10.68	22.66	16.67	1.69	8.00	4.84	27.34	21.34	24.33	26.67	8.00	17.33									
27	AKma-14 B x SRT - 26B	62.50 (52.23)	26.00 (30.62)	44.25 (41.43)	28.67	54.67	41.67	1.69	11.34	6.51	50.67	24.66	37.67	19.34	5.33	12.33									
28	AKma-14 B x GI-35-15-15	69.50 (54.71)	34.00 (35.34)	41.50 (40.03)	17.32	33.33	38.33	4.01	15.33	7.66	59.33	30.66	45.00	23.34	5.33	14.33									
29	AKma-14 B x SPV - 946	64.00 (54.93)	23.00 (29.01)	44.00 (41.97)	17.34	33.33	33.33	0.71	14.67	7.69	52.67	26.67	39.67	19.33	5.33	12.33									
30	AKma-14 B x SPV - 104	82.50 (66.65)	55.00 (25.04)	25.25 (29.85)	15.34	34.67	45.00	4.67	18.67	11.67	48.00	20.00	34.00	12.00	6.00	9.00									
31	AKma-14 B x IS - 2284	89.00 (74.67)	88.00 (72.87)	72.00 (59.26)	35.33	56.67	26.00	6.0	13.67	9.67	47.33	31.34	29.33	37.34	10.00	18.67									
32	AKma-14 B x IS - 6335	94.00 (76.26)	77.00 (61.63)	85.50 (68.91)	14.01	25.34	19.67	3.33	8.67	6.0	30.67	20.00	25.33	37.34	6.00	21.67									
33	AKma-14 B x IS - 9471	83.00 (65.69)	77.00 (61.48)	80.00 (63.38)	16.67	37.34	27.00	2.35	10.66	6.51	39.33	22.66	31.00	22.00	8.00	15.00									
34	AKma-14 B x IS - 9471																								

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17
35	SRT - 26B x GI-35-15-15	88.50(70.80)	37.00(37.46)	67.25(53.82)	12.68	67.33	40.01	0.71	16.66	8.69	55.33	12.00	33.67	34.00	3.69	18.84	
36	SRT - 26B x SPV - 946	71.00(57.44)	44.50(41.83)	57.75(49.64)	26.01	62.67	44.33	0.71	16.66	8.69	55.33	12.00	33.67	34.00	3.69	18.84	
37	SRT - 26B x SPV - 104	58.50(49.93)	44.00(41.56)	51.25(45.74)	30.67	52.00	41.33	1.69	15.34	8.51	53.33	26.67	40.00	15.34	4.00	9.67	
38	SRT - 26B x IS - 2284	98.50(75.24)	64.00(53.22)	78.75(64.23)	11.32	40.67	26.00	1.69	10.00	5.84	46.00	22.67	34.33	34.00	5.34	19.67	
39	SRT - 26B x IS - 6335	92.00(73.57)	73.50(59.14)	82.75(66.35)	10.01	34.00	22.00	3.34	7.33	5.33	36.67	15.33	26.00	25.34	6.00	15.67	
40	SRT - 26B x IS - 9471	89.00(64.87)	79.50(63.46)	84.25(64.17)	9.34	31.33	20.34	3.02	9.33	6.17	31.34	14.66	23.00	17.34	4.66	11.00	
41	GI-35-15-15 x SPV - 946	52.00(56.10)	28.50(32.25)	40.25(44.17)	20.00	60.00	40.00	1.02	10.00	5.51	56.00	25.33	40.67	24.66	5.34	15.00	
42	GI-35-15-15 x SPV 104	69.00(60.25)	24.00(29.31)	46.50(44.78)	22.66	59.33	41.00	2.67	15.33	9.00	50.00	20.00	35.00	18.66	6.66	12.66	
43	GI-35-15-15 x IS - 2284	86.50(67.21)	73.00(58.79)	79.75(63.00)	9.34	40.00	24.67	4.67	10.66	7.67	38.67	16.67	27.67	18.67	4.66	11.67	
44	GI-35-15-15 x IS - 6335	90.00(74.32)	79.50(63.12)	84.75(68.72)	13.99	44.67	29.33	1.33	17.34	9.33	32.00	19.34	25.67	22.00	8.00	15.00	
45	GI-35-15-15 x IS - 9471	89.00(61.13)	65.00(53.74)	77.00(57.44)	7.99	46.67	27.33	1.33	17.34	9.33	32.00	19.34	25.67	22.66	7.34	15.00	
46	SPV - 946 x SPV - 104	81.50(68.04)	14.00(21.94)	47.75(44.98)	11.33	50.00	32.67	4.67	17.34	11.00	30.67	22.00	26.33	23.34	6.00	14.67	
47	SPV - 946 x IS - 2284	87.50(64.22)	68.50(55.85)	78.00(60.10)	15.34	50.00	32.67	4.67	17.34	11.00	30.67	22.00	26.33	23.34	6.00	14.67	
48	SPV - 946 x IS - 6335	96.00(77.21)	60.00(50.85)	78.00(64.03)	11.99	52.00	32.00	2.35	21.34	11.85	29.34	20.67	25.00	26.00	4.00	15.00	
49	SPV - 946 x IS 9471	85.00(72.35)	31.00(33.70)	58.00(53.03)	11.99	42.67	27.33	3.34	16.66	10.00	30.66	18.67	24.67	23.34	7.34	15.34	
50	SPV - 104 x IS 2284	90.50(72.10)	73.00(59.09)	81.75(65.60)	15.99	35.33	25.66	7.33	13.34	10.33	34.67	26.67	30.67	27.34	7.34	17.33	
51	SPV - 104 x IS 6335	89.50(71.09)	74.00(59.49)	81.75(65.60)	5.99	36.67	21.33	1.69	11.34	6.51	36.00	22.67	29.33	29.37	7.34	18.33	
52	SPV - 104 x IS 9471	91.50(73.41)	65.00(53.93)	78.25(63.67)	7.99	44.00	26.00	2.35	10.66	6.51	38.67	16.67	27.67	22.66	5.34	14.00	
53	IS 2284 x IS - 6335	85.50(67.65)	67.50(55.25)	76.50(61.45)	6.01	36.67	21.34	3.02	9.34	6.18	31.33	20.67	26.00	24.66	4.66	14.66	
54	IS 2284 x IS - 9471	90.50(72.20)	77.50(61.75)	84.00(66.97)	12.01	34.00	23.00	8.00	12.00	10.00	30.00	26.67	28.33	26.00	6.67	16.34	
55	IS 6335 x IS 9471	89.50(70.56)	63.50(52.85)	76.25(61.75)	6.00	30.00	18.00	2.35	6.66	4.51	23.33	22.00	22.67	17.33	5.34	11.33	
Parental range		29.00 to 86.00	2.50 to 61.00	15.75 to 71.75	5.33 to 38.67	34.00 to 71.34	19.67 to 55.00	0.71 to 2.67	7.34 to 19.34	4.02 to 10.51	24.00 to 60.00	15.33 to 37.33	20.67 to 46.67	9.34 to 24.67	0.71 to 6.00	5.18 to 15.00	
Hybrid range		32.50 to 98.00	14.00 to 88.50	25.25 to 90.00	5.99 to 35.99	21.34 to 70.67	15.66 to 52.34	0.71 to 8.00	5.34 to 21.34	3.02 to 11.85	18.00 to 68.00	8.00 to 37.33	20.33 to 46.33	12.00 to 37.34	0.71 to 24.00	9.00 to 21.67	
SE (m)		3.606	4.750	14.911	0.961	2.520	1.512	0.17	0.738	0.564	2.561	1.985	2.215	1.213	0.368	1.592	
CD at 5%		10.293	13.556	42.554	1.903	4.992	2.993	0.33	1.501	1.117	5.073	3.932	4.387	2.406	0.729	3.153	

Figures in parenthesis indicated arcsin values

L2= Panancheru

L1= Akola

P = Pooled

Table 8 Mean values of F_2 progenies for different characters in 10×10 diallel of sorghum, Akola and Patancheru, 1996

Sr No	Crosses	Grain hardness Kg/cm ²			Electrical conductivity (millimhos)				Plant height (cm)				Cob length (cm)			
		L1	L2	Pooled	L1	L2	Pooled	8	L1	L2	Pooled	11	L1	L2	Pooled	14
1	SPV-1201 x ICSB - 101B	5.81	5.52	5.67	374.50	386.50	380.50	380.50	174.00	186.00	180.00	24.65	24.80	24.73		
2	SPV-1201 x AKms - 14 B	6.85	3.57	5.21	374.50	413.00	393.75	393.75	189.15	248.75	218.95	21.75	22.60	22.17		
3	SPV-1201 x SRT - 26B	6.67	4.16	5.41	404.50	454.00	429.25	429.25	173.75	235.00	204.38	22.50	23.60	23.05		
4	SPV-1201 x GI-35-15-15	6.97	4.15	5.56	494.50	685.00	589.50	589.50	209.40	275.00	242.20	24.25	23.50	23.88		
5	SPV-1201 x SPV - 946	5.27	3.75	4.51	310.50	404.00	357.25	357.25	200.75	252.00	226.38	23.15	22.35	22.75		
6	SPV-1201 x SPV - 104	5.00	3.25	4.12	451.50	488.00	469.75	469.75	182.00	196.65	189.32	18.95	23.65	21.30		
7	SPV-1201 x IS - 2284	5.63	3.66	4.64	355.50	342.00	348.75	348.75	229.15	233.75	231.15	23.25	24.00	23.63		
8	SPV-1201 x IS-6335	7.17	3.41	5.29	228.00	222.00	225.00	225.00	226.40	230.00	228.20	26.20	26.15	26.17		
9	SPV-1201 x IS - 9471	7.31	2.43	4.87	233.50	187.00	210.25	210.25	230.15	226.25	228.20	23.10	25.25	24.17		
10	ICSB - 101B x AKms 14 B	5.66	3.83	4.74	214.00	488.00	351.00	351.00	157.90	184.15	171.02	30.05	27.75	28.90		
11	ICSB - 101B x SRT - 26 B	5.81	3.23	4.52	381.50	398.00	389.75	389.75	171.50	241.25	206.38	29.00	26.15	27.58		
12	ICSB - 101B x GI-35-15-15	5.90	3.96	4.93	366.50	410.00	388.25	388.25	181.50	233.15	207.07	29.60	26.25	27.92		
13	ICSB - 101B x SPV - 946	5.79	4.37	5.08	276.50	479.00	377.75	377.75	174.00	258.75	216.38	29.30	25.90	27.60		
14	ICSB - 101B x SPV - 104	5.72	3.52	4.62	382.00	490.50	436.25	436.25	170.50	196.25	183.38	29.50	24.00	25.25		
15	ICSB - 101B x IS - 2284	6.41	4.49	5.45	196.00	390.50	293.25	293.25	218.75	280.00	249.38	28.20	26.15	27.17		
16	ICSB - 101B x IS - 6335	6.03	4.92	5.48	240.00	297.00	268.50	268.50	183.50	255.65	219.57	28.40	26.65	27.52		
17	ICSB - 101B x IS - 9471	4.90	4.22	4.56	255.50	186.00	220.75	220.75	195.75	267.50	231.63	27.45	27.25	27.35		
18	AKms-14 B x SRT - 26B	7.23	2.13	4.68	311.00	438.50	374.75	374.75	174.50	240.00	207.25	26.80	25.25	26.02		
19	AKms-14 B x GI-35-15-15	6.35	2.62	4.48	334.00	389.50	361.75	361.75	166.50	217.50	192.00	28.70	24.25	26.48		
20	AKms-14 B x SPV - 946	6.93	2.71	4.82	340.50	481.00	410.75	410.75	189.00	233.50	213.25	25.90	23.25	24.58		
21	AKms-14 B x SPV - 104	4.62	2.89	3.75	442.50	533.50	488.00	488.00	162.75	210.00	186.86	24.80	21.50	23.15		
22	AKms-14 B x IS - 2284	5.12	3.01	4.07	255.50	370.00	287.75	287.75	172.00	283.75	227.88	25.20	23.50	24.35		
23	AKms-14 B x IS - 6335	5.88	3.80	4.84	172.50	205.50	189.00	189.00	196.60	261.25	228.93	26.50	25.25	25.88		
24	AKms-14 B x IS - 9471	5.72	3.25	4.48	181.00	277.50	229.25	229.25	172.25	277.50	224.88	27.40	25.50	26.45		

Condt.

Table : 9 Mean values of F_2 progenies for different characters in 10×10 diallel of sorghum, Akola and Patancheru, 1996

Sr. No	Crosses	Glume covering (%)			TGMRR			Germination (%)		
		L1	L2	Pooled	L1	L2	Pooled	L1	L2	Pooled
1	SPV-1201 x ICSCB - 101B	47.50	50.00	48.75	2.05	3.15	2.60	80.40(63.77)	66.45(54.61)	73.43(59.19)
2	SPV-1201 x AKms - 14 B	50.00	50.00	50.00	2.55	3.85	3.20	70.50(57.41)	43.00(40.77)	56.75(49.09)
3	SPV-1201 x SRT - 26B	45.00	50.00	47.50	2.40	3.15	2.78	81.75(64.94)	64.10(56.23)	79.93(60.58)
4	SPV-1201 x GJ-35-15-15	42.50	50.00	46.25	2.50	3.45	2.97	80.00(63.92)	59.15(50.51)	69.38(57.23)
5	SPV-1201 x SPV - 946	37.50	50.00	43.75	2.40	3.55	2.97	74.75(59.91)	56.15(48.55)	65.45(54.23)
6	SPV-1201 x SPV - 104	45.00	37.50	41.25	2.60	3.40	3.00	74.35(59.84)	50.50(45.28)	62.43(52.56)
7	SPV-1201 x IS - 2284	50.00	54.00	52.00	2.20	2.45	2.33	73.85(59.26)	76.15(60.77)	75.00(60.01)
8	SPV-1201 x IS-6335	50.00	47.00	48.75	2.20	2.20	2.20	75.95(60.73)	76.65(61.15)	76.30(60.94)
9	SPV-1201 x IS - 9471	50.00	44.00	47.00	2.10	2.30	2.20	76.50(61.00)	48.50(44.14)	62.50(52.57)
10	ICSB - 101B x AKms 14 B	50.00	50.00	50.00	2.70	3.25	2.97	71.40(57.89)	50.40(45.24)	60.90(51.56)
11	ICSB - 101B x SRT - 26 B	50.00	50.00	50.00	2.25	3.40	2.83	79.60(63.20)	45.40(43.56)	62.50(53.38)
12	ICSB - 101B x GJ-35-15-15	50.00	50.00	50.00	2.35	3.05	2.70	79.50(63.14)	67.75(55.41)	73.63(59.28)
13	ICSB - 101B x SPV - 946	50.00	50.00	50.00	2.45	2.95	2.70	70.50(57.28)	51.00(45.61)	60.75(51.45)
14	ICSB - 101B x SPV - 104	50.00	50.00	50.00	2.25	3.55	2.90	76.95(61.31)	21.50(26.48)	49.23(43.90)
15	ICSB - 101B x IS - 2284	59.00	41.50	50.25	2.35	2.55	2.45	81.50(64.44)	65.25(53.90)	73.38(59.19)
16	ICSB - 101B x IS - 6335	54.00	54.00	54.00	2.40	2.70	2.55	81.40(64.46)	75.50(60.36)	78.45(62.41)
17	ICSB - 101B x IS - 9471	58.00	50.00	54.00	2.20	3.05	2.63	82.50(65.39)	69.50(56.48)	76.00(60.94)
18	AKms-14 B x SRT - 26B	50.00	50.00	50.00	2.50	4.00	3.25	71.20(57.70)	9.75(17.65)	40.48(37.67)
19	AKms-14 B x GJ-35-15-15	50.00	50.00	50.00	2.70	3.10	2.90	69.20(56.71)	60.00(51.16)	64.90(53.93)
20	AKms-14 B x SPV - 946	50.00	58.50	54.25	2.55	3.55	3.05	65.80(54.26)	24.50(29.57)	45.15(41.91)
21	AKms-14 B x SPV - 104	50.00	50.00	50.00	3.05	3.50	3.28	53.00(46.53)	23.00(28.39)	38.00(37.56)
22	AKms-14 B x IS - 2284	47.50	50.00	48.75	2.15	2.65	2.40	80.85(64.05)	52.00(46.17)	66.43(55.11)
23	AKms-14 B x IS - 6335	41.00	50.00	45.50	2.15	2.40	2.28	84.15(66.55)	68.75(56.38)	76.45(61.46)
24	AKms-14 B x IS - 9471	37.50	50.00	43.75	1.75	2.65	2.20	81.10(64.34)	51.25(45.78)	66.18(55.06)

Contd.

Table 9 Contd.

	1	2	3	4	5	6	7	8	9	10	11
25	SRT - 26B x GJ-35-15-15	50.00	50.00	50.00	50.00	2.05	3.40	2.72	75.00(65.05)	51.50(45.86)	63.25(55.46)
26	SRT - 26B x SPV - 946	50.00	50.00	50.00	50.00	2.60	2.80	2.70	77.65(54.16)	64.55(53.52)	71.10(53.84)
27	SRT - 26B x SPV - 104	54.00	50.00	50.00	52.00	2.90	3.25	3.08	63.50(59.56)	23.75(29.11)	43.63(44.34)
28	SRT - 26B x IS - 2284	47.50	50.00	50.00	48.75	2.80	2.65	2.72	79.15(46.29)	74.25(58.15)	76.70(52.22)
29	SRT - 26B x IS - 6335	50.00	50.00	50.00	50.00	2.20	2.30	2.25	73.40(64.25)	80.25(66.64)	76.83(63.94)
30	SRT - 26B x IS - 9471	50.00	50.00	50.00	50.00	2.65	2.50	2.58	76.70(60.07)	60.55(51.15)	68.62(55.61)
31	GJ-35-15-15 x SPV - 946	52.50	50.00	50.00	51.25	2.45	3.10	2.78	69.80(57.55)	58.85(50.19)	64.33(53.87)
32	GJ-35-15-15 x SPV 104	50.00	50.00	50.00	50.00	2.80	3.65	3.72	66.00(54.80)	25.50(30.31)	45.75(42.55)
33	GJ-35-15-15 x IS - 2284	61.50	62.50	62.00	62.00	2.10	2.35	2.17	81.75(64.76)	81.50(64.57)	81.62(64.67)
34	GJ-35-15-15 x IS - 6335	56.00	50.00	50.00	52.00	2.15	2.40	2.28	80.45(63.88)	69.00(56.19)	74.73(60.03)
35	GJ-35-15-15 x IS - 9471	50.00	47.50	48.75	48.75	1.90	2.50	2.20	67.30(55.88)	61.50(51.68)	64.40(53.76)
36	SPV - 946 x SPV - 104	56.50	50.00	53.25	53.25	2.50	3.25	2.88	69.20(57.13)	49.00(44.40)	59.10(50.77)
37	SPV - 946 x IS - 2284	60.00	60.50	60.25	60.25	2.30	2.15	2.22	74.80(60.40)	67.10(55.08)	70.95(57.74)
38	SPV - 946 x IS - 6335	66.50	75.00	70.75	70.75	2.40	2.25	2.33	86.15(68.21)	83.15(65.79)	84.56(67.00)
39	SPV - 946 x IS 9471	60.00	50.00	55.10	55.10	2.00	2.20	2.10	78.60(62.93)	59.00(50.32)	68.80(56.63)
40	SPV - 104 x IS 2284	59.00	69.00	64.00	64.00	2.40	3.25	2.83	68.35(55.93)	62.75(52.80)	65.50(54.36)
41	SPV - 104 x IS 6335	56.50	54.00	55.25	55.25	2.75	2.15	2.45	65.35(54.36)	54.00(47.53)	59.68(50.95)
42	SPV - 104 x IS 9471	50.00	44.00	47.00	47.00	2.30	2.65	2.47	62.55(52.58)	53.75(47.15)	58.15(49.87)
43	IS 2284 x IS - 6335	54.00	50.00	52.00	52.00	1.75	2.00	1.88	87.90(69.74)	63.50(53.81)	75.70(61.77)
44	IS 2284 x IS - 9471	50.00	50.00	50.00	50.00	1.70	2.00	1.85	85.90(68.10)	83.50(66.03)	84.70(67.07)
45	IS 6335 x IS 9471	54.00	66.50	60.25	60.25	1.90	2.00	1.95	86.40(68.45)	77.15(63.23)	81.75(65.84)

F₂ progenies range

38.00

to

53.00

to

9.75

to

83.50

84.70

SE (m)

38.00

to

53.00

to

9.75

to

83.50

84.70

CD at 5%

13.424

to

8.586

to

24.505

to

38.312

p = Pooled L1 = Akola L2 Palancheru

Figures in parenthesis indicate arcsin values.

Table 10 Mean infection % of F_2 progenies for pathological characters in 10x10 dialled of sorghum, Akola and Patancheru, 1996

St.	No.	Crosses	<i>F. moniliforme</i>				<i>F. pallidoseum</i>				<i>C. lunata</i>				Other fungi			
			<i>F. moniliforme</i>		<i>F. pallidoseum</i>		<i>F. pallidoseum</i>		<i>F. pallidoseum</i>		<i>C. lunata</i>		<i>C. lunata</i>		Other fungi		Other fungi	
			L1	L2	Pooled	L1	L2	Pooled	L1	L2	Pooled	L1	L2	Pooled	L1	L2	Pooled	L2
1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19
1		SPV-1201 x ICSB - 101B	32.22	37.86	35.64	1.50	3.34	2.42	46.52	34.06	40.29	17.40	14.80	16.10				
2		SPV-1201 x AKms - 14 B	31.82	44.82	38.32	1.76	4.34	3.05	45.36	27.66	36.51	17.50	10.34	13.92				
3		SPV-1201 x SRT - 26B	32.60	46.16	39.38	2.15	6.16	4.16	47.10	38.66	39.88	20.80	13.48	17.14				
4		SPV-1201 x GI-35-15-15	21.40	48.50	34.95	1.56	7.50	4.53	61.02	32.26	46.64	16.60	8.26	12.43				
5		SPV-1201 x SPV - 946	22.72	50.20	36.46	1.35	7.10	4.23	47.20	31.60	39.40	18.56	11.10	14.83				
6		SPV-1201 x SPV - 104	28.20	58.34	43.27	2.50	2.35	2.43	48.26	28.00	38.13	18.46	9.66	14.06				
7		SPV-1201 x IS - 2284	28.32	31.00	29.66	1.56	3.22	2.39	46.66	36.52	41.59	17.46	20.52	18.99				
8		SPV-1201 x IS-6335	30.56	39.50	35.03	3.36	7.00	5.18	36.59	38.50	37.53	17.02	11.52	14.27				
9		SPV-1201 x IS - 9471	31.70	41.50	31.60	1.56	6.50	4.03	42.00	41.00	41.50	16.70	12.50	14.60				
10		ICSB - 101B x AKms 14 B	21.58	46.50	34.04	1.60	4.68	3.14	64.26	39.00	51.33	12.60	11.18	11.89				
11		ICSB - 101B x SRT - 26 B	19.00	58.84	38.92	3.40	4.84	4.12	61.80	27.68	44.74	27.00	8.16	17.58				
12		ICSB - 101B x GI-35-15-15	29.40	38.26	33.83	1.60	3.26	2.43	51.20	44.76	47.98	21.40	14.26	17.83				
13		ICSB - 101B x SPV - 946	29.80	50.26	40.03	2.76	5.85	4.30	54.20	31.00	42.60	13.20	14.76	13.98				
14		ICSB - 101B x SPV - 104	31.30	57.50	44.40	4.20	11.00	7.60	45.60	29.00	37.30	18.70	6.50	12.60				
15		ICSB - 101B x IS - 2284	35.10	50.86	42.98	3.25	3.70	3.53	41.82	31.76	36.79	19.50	13.10	16.30				
16		ICSB - 101B x IS - 6335	18.50	45.76	32.13	1.66	2.37	2.02	59.30	23.26	41.28	15.76	16.02	15.89				
17		ICSB - 101B x IS - 9471	27.36	42.00	34.68	4.26	5.82	5.04	45.82	25.50	35.51	14.40	14.32	14.36				
18		AKms-14 B x SRT - 26B	39.40	51.00	40.20	2.40	6.00	4.20	56.60	38.00	47.30	11.40	5.50	8.45				
19		AKms-14 B x GI-35-15-15	26.00	56.34	41.17	1.56	1.50	1.53	61.20	35.00	48.35	12.00	6.00	9.00				
20		AKms-14 B x SPV - 946	23.40	50.00	36.70	1.35	7.50	4.43	61.80	43.50	52.65	13.80	7.00	10.40				
21		AKms-14 B x SPV - 104	41.06	67.50	54.25	2.15	5.50	3.83	45.00	20.00	32.50	12.82	7.00	9.91				
22		AKms-14 B x IS - 2284	29.06	53.50	41.28	1.56	6.50	4.03	49.72	24.50	32.11	21.30	10.50	15.90				
23		AKms-14 B x IS - 6335	27.82	41.00	34.41	1.35	7.50	4.43	42.34	26.00	34.17	26.76	7.50	16.83				
24		AKms-14 B x IS - 9471	29.30	38.00	32.15	1.35	3.00	2.18	39.30	34.50	36.90	18.40	19.50	18.95				

Contd.

III - 12

Table 10 Cond.....

1	2	3	4	5	6	7	8	9	10	11	12	13	14
25	SRT - 26B x GJ-35-15-15	33.40	50.50	41.95	3.60	4.35	3.98	42.60	31.00	36.80	16.00	15.50	15.75
26	SRT - 26B x SPV - 946	18.70	40.76	29.76	1.10	6.30	3.70	63.30	38.66	50.98	18.10	6.06	17.08
27	SRT - 26B x SPV - 104	24.00	52.50	38.25	1.75	3.50	2.63	58.60	37.50	48.05	16.60	8.00	12.30
28	SRT - 26B x IS - 2284	25.62	37.50	31.56	0.96	3.73	4.73	47.36	35.00	41.18	19.76	9.00	14.38
29	SRT - 26B x IS - 6335	22.06	50.02	36.04	2.56	4.26	3.41	59.34	27.76	43.50	15.74	14.26	15.0
30	SRT - 26B x IS - 9471	34.00	47.82	40.91	3.15	4.34	3.75	43.20	35.82	39.51	17.80	7.68	12.74
31	GJ-35-15-15 x SPV - 946	25.70	45.60	35.65	0.77	2.52	1.64	48.50	34.50	41.50	19.08	14.16	16.62
32	GJ-35-15-15 x SPV 104	30.86	67.00	48.93	2.86	5.00	3.93	66.36	24.50	45.33	15.06	3.00	9.03
33	GJ-35-15-15 x IS - 2284	33.06	45.00	39.03	4.40	8.00	6.23	47.92	28.00	37.96	9.34	16.00	12.67
34	GJ-35-15-15 x IS - 6335	30.26	48.16	39.21	2.20	5.00	3.60	37.06	31.50	34.28	14.06	11.84	12.95
35	GJ-35-15-15 x IS - 9471	38.92	38.26	38.59	3.58	5.26	4.42	37.90	30.26	34.08	14.26	11.26	12.76
36	SPV - 946 x SPV - 104	27.00	60.50	43.75	2.20	6.50	4.35	54.60	19.00	36.80	13.20	7.50	10.35
37	SPV - 946 x IS - 2284	38.00	47.26	42.63	4.60	4.50	4.55	38.80	36.66	37.73	23.20	12.34	17.77
38	SPV - 946 x IS - 6335	20.42	33.34	26.88	3.42	3.00	3.21	42.42	22.16	32.29	18.02	21.68	19.85
39	SPV - 946 x IS 9471	36.26	42.00	39.13	3.6	5.60	4.68	30.26	33.40	31.83	11.06	13.40	12.23
40	SPV - 104 x IS 2284	33.82	53.50	43.66	2.56	6.16	4.36	47.02	29.16	38.09	10.06	11.50	10.78
41	SPV - 104 x IS 6335	44.06	47.50	45.18	4.96	8.00	6.48	34.86	33.50	34.18	12.76	11.50	12.13
42	SPV - 104 x IS 9471	52.52	53.00	52.76	5.92	7.50	6.71	28.04	26.50	27.27	10.72	12.50	11.61
43	IS 2284 x IS - 6335	33.66	59.00	46.33	6.04	8.00	7.02	22.20	20.00	21.10	12.12	12.35	12.74
44	IS 2284 x IS - 9471	34.42	48.50	41.46	3.74	4.50	4.12	20.92	18.50	19.71	12.00	14.50	13.25
45	IS 6335 x IS 9471	38.92	25.34	32.13	5.84	5.34	5.59	19.26	25.32	22.29	10.60	12.66	11.63

F₂ progenies range

SE (m)

CD at 5%

P = Pooled

L1 = Akola

L2 = Patancheru

18.50	25.34	26.28	0.77	1.50	1.53	16.26	18.50	19.71	9.34	3.00	8.45
52.52	67.50	54.25	6.04	11.00	7.60	66.36	44.76	52.65	27.00	20.52	19.85
1.788	2.097	2.249	0.88	1.154	1.606	2.670	1.253	2.095	0.648	0.384	0.549
3.554	1.541	4.455	1.75	2.861	3.181	5.298	2.482	4.150	1.283	0.760	1.087

III - 13
Table : 11 Mean value of parents and F₁ crosses for biochemical characters in 10x10 diallel of sorghum, Akola, 1995

Sr.No	Parents/Crosses	Protein (%)				Tannins (CE %)				Flavan-4-ols A ₃₀			
		1	2	3	4	5	6	7	8	9	10	11	12
1	SPV-1201	12.35(3.59)	1.70(1.49)	0.06(0.75)	0.00(0.71)	0.06(0.75)	0.00(0.71)	0.06(0.75)	0.00(0.71)	0.06(0.75)	0.00(0.71)	0.06(0.75)	0.00(0.71)
2	ICSB-101B	12.94(3.66)	1.50(1.41)	0.02(0.72)	0.00(0.71)	0.02(0.72)	0.00(0.71)	0.02(0.72)	0.00(0.71)	0.02(0.72)	0.00(0.71)	0.02(0.72)	0.00(0.71)
3	AKMS-14B	11.78(3.51)	2.30(1.67)	0.06(0.75)	0.00(0.71)	0.06(0.75)	0.00(0.71)	0.06(0.75)	0.00(0.71)	0.06(0.75)	0.00(0.71)	0.06(0.75)	0.00(0.71)
4	SRT-26B	12.41(3.59)	2.20(1.64)	0.05(0.74)	0.00(0.71)	0.05(0.74)	0.00(0.71)	0.05(0.74)	0.00(0.71)	0.05(0.74)	0.00(0.71)	0.05(0.74)	0.00(0.71)
5	CI-35-15-15	10.51(3.32)	1.40(1.38)	0.06(0.75)	0.00(0.71)	0.06(0.75)	0.00(0.71)	0.06(0.75)	0.00(0.71)	0.06(0.75)	0.00(0.71)	0.06(0.75)	0.00(0.71)
6	SPV-946	11.01(3.39)	1.40(1.38)	0.06(0.75)	0.00(0.71)	0.06(0.75)	0.00(0.71)	0.06(0.75)	0.00(0.71)	0.06(0.75)	0.00(0.71)	0.06(0.75)	0.00(0.71)
7	SPV-104	11.81(3.51)	1.20(1.30)	0.06(0.75)	0.00(0.71)	0.06(0.75)	0.00(0.71)	0.06(0.75)	0.00(0.71)	0.06(0.75)	0.00(0.71)	0.06(0.75)	0.00(0.71)
8	IS-2284	9.95(3.23)	2.00(1.58)	5.84(2.52)	4.93(2.33)	5.84(2.52)	4.93(2.33)	5.84(2.52)	4.93(2.33)	5.84(2.52)	4.93(2.33)	5.84(2.52)	4.93(2.33)
9	IS-6335	11.11(3.41)	1.60(1.45)	2.53(1.74)	0.03(0.73)	2.53(1.74)	0.03(0.73)	2.53(1.74)	0.03(0.73)	2.53(1.74)	0.03(0.73)	2.53(1.74)	0.03(0.73)
10	IS-9471	8.93(3.01)	1.30(1.34)	0.01(0.71)	0.00(0.71)	0.01(0.71)	0.00(0.71)	0.01(0.71)	0.00(0.71)	0.01(0.71)	0.00(0.71)	0.01(0.71)	0.00(0.71)
11	SPV-1201 x ICSB - 101B	9.11(3.10)	1.20(1.30)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)
12	SPV-1201 x AKMS - 14 B	6.35(2.62)	1.40(1.38)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)
13	SPV-1201 x SRT - 26B	6.62(2.67)	1.60(1.45)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)
14	SPV-1201 x CI-35-15-15	8.27(2.86)	1.40(1.38)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)
15	SPV-1201 x SPV - 946	7.68(2.88)	1.40(1.38)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)
16	SPV-1201 x SPV - 104	8.90(3.07)	1.20(1.30)	0.04(0.73)	0.00(0.71)	0.04(0.73)	0.00(0.71)	0.04(0.73)	0.00(0.71)	0.04(0.73)	0.00(0.71)	0.04(0.73)	0.00(0.71)
17	SPV-1201 x IS - 2284	9.87(3.22)	1.60(1.45)	2.25(1.66)	4.20(2.17)	2.25(1.66)	4.20(2.17)	2.25(1.66)	4.20(2.17)	2.25(1.66)	4.20(2.17)	2.25(1.66)	4.20(2.17)
18	SPV-1201 x IS-6335	8.08(2.93)	1.80(1.52)	2.70(1.79)	11.40(3.45)	2.70(1.79)	11.40(3.45)	2.70(1.79)	11.40(3.45)	2.70(1.79)	11.40(3.45)	2.70(1.79)	11.40(3.45)
19	SPV-1201 x IS - 9471	10.11(3.26)	1.70(1.48)	2.16(1.63)	0.00(0.71)	2.16(1.63)	0.00(0.71)	2.16(1.63)	0.00(0.71)	2.16(1.63)	0.00(0.71)	2.16(1.63)	0.00(0.71)
20	ICSB - 101B x AKMS 14 B	7.01(2.74)	1.00(1.22)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)
21	ICSB - 101B x SRT - 26 B	8.01(2.92)	1.10(1.26)	0.02(0.72)	0.00(0.71)	0.02(0.72)	0.00(0.71)	0.02(0.72)	0.00(0.71)	0.02(0.72)	0.00(0.71)	0.02(0.72)	0.00(0.71)
22	ICSB - 101B x CI-35-15-15	7.32(2.80)	1.30(1.34)	0.01(0.71)	0.00(0.71)	0.01(0.71)	0.00(0.71)	0.01(0.71)	0.00(0.71)	0.01(0.71)	0.00(0.71)	0.01(0.71)	0.00(0.71)
23	ICSB - 101B x SPV - 946	6.61(2.67)	1.20(1.30)	0.04(0.73)	0.00(0.71)	0.04(0.73)	0.00(0.71)	0.04(0.73)	0.00(0.71)	0.04(0.73)	0.00(0.71)	0.04(0.73)	0.00(0.71)
24	ICSB - 101B x SPV - 104	7.61(2.85)	1.00(1.22)	0.01(0.71)	0.00(0.71)	0.01(0.71)	0.00(0.71)	0.01(0.71)	0.00(0.71)	0.01(0.71)	0.00(0.71)	0.01(0.71)	0.00(0.71)
25	ICSB - 101B x IS - 2284	7.19(2.77)	1.50(1.41)	2.25(1.66)	4.20(2.17)	2.25(1.66)	4.20(2.17)	2.25(1.66)	4.20(2.17)	2.25(1.66)	4.20(2.17)	2.25(1.66)	4.20(2.17)
26	ICSB - 101B x IS - 6335	7.75(2.87)	1.40(1.38)	1.78(1.51)	6.50(2.65)	1.78(1.51)	6.50(2.65)	1.78(1.51)	6.50(2.65)	1.78(1.51)	6.50(2.65)	1.78(1.51)	6.50(2.65)
27	ICSB - 101B x IS - 9471	6.80(2.70)	1.30(1.34)	1.41(1.38)	0.00(0.71)	1.41(1.38)	0.00(0.71)	1.41(1.38)	0.00(0.71)	1.41(1.38)	0.00(0.71)	1.41(1.38)	0.00(0.71)
28	AKMS - 14 B x SRT - 26B	6.13(2.8)	0.90(1.18)	0.02(0.72)	0.00(0.71)	0.02(0.72)	0.00(0.71)	0.02(0.72)	0.00(0.71)	0.02(0.72)	0.00(0.71)	0.02(0.72)	0.00(0.71)
29	AKMS - 14 B x CI-35-15-15	6.10(2.57)	1.00(1.22)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)	0.03(0.73)	0.00(0.71)
30	AKMS - 14 B x SPV - 946	5.78(2.51)	1.40(1.38)	0.01(0.71)	0.00(0.71)	0.01(0.71)	0.00(0.71)	0.01(0.71)	0.00(0.71)	0.01(0.71)	0.00(0.71)	0.01(0.71)	0.00(0.71)
31	AKMS - 14 B x SPV - 104	6.53(2.65)	1.00(1.22)	0.01(0.71)	0.00(0.71)	0.01(0.71)	0.00(0.71)	0.01(0.71)	0.00(0.71)	0.01(0.71)	0.00(0.71)	0.01(0.71)	0.00(0.71)
32	AKMS - 14 B x IS - 2284	6.24(2.60)	1.70(1.48)	3.41(1.98)	4.70(2.68)	3.41(1.98)	4.70(2.68)	3.41(1.98)	4.70(2.68)	3.41(1.98)	4.70(2.68)	3.41(1.98)	4.70(2.68)
33	AKMS - 14 B x IS - 6335	8.88(3.06)	1.30(1.34)	2.83(1.82)	10.30(3.29)	2.83(1.82)	10.30(3.29)	2.83(1.82)	10.30(3.29)	2.83(1.82)	10.30(3.29)	2.83(1.82)	10.30(3.29)
34	AKMS - 14 B x IS - 9471	5.56(2.46)	1.70(1.48)	2.40(1.70)	Concl.	2.40(1.70)	Concl.	2.40(1.70)	Concl.	2.40(1.70)	Concl.	2.40(1.70)	Concl.

III - 14

1	2	3			4		5		6	
		35	36	37	38	39	40	41	42	43
35	SRT - 26B x GI-35-15-15	6.25(2.60)	6.58(2.66)	8.49(3.00)	5.98(2.55)	7.10(2.76)	5.76(2.50)	6.99(2.74)	7.78(2.88)	6.23(2.59)
36	SRT - 26B x SPV - 946	6.58(2.66)	8.49(3.00)	5.98(2.55)	7.10(2.76)	5.76(2.50)	6.99(2.74)	7.78(2.88)	6.23(2.59)	6.26(2.60)
37	SRT - 26B x SPV - 104	8.49(3.00)	5.98(2.55)	7.10(2.76)	5.76(2.50)	6.99(2.74)	7.78(2.88)	6.23(2.59)	6.26(2.60)	5.23(2.39)
38	SRT - 26B x IS - 2284	5.98(2.55)	7.10(2.76)	5.76(2.50)	6.99(2.74)	7.78(2.88)	6.23(2.59)	6.26(2.60)	5.23(2.39)	8.96(3.08)
39	SRT - 26B x IS - 6335	7.10(2.76)	5.76(2.50)	6.99(2.74)	7.78(2.88)	6.23(2.59)	6.26(2.60)	5.23(2.39)	8.96(3.08)	6.68(2.68)
40	SRT - 26B x IS - 9471	5.76(2.50)	6.99(2.74)	7.78(2.88)	6.23(2.59)	6.26(2.60)	5.23(2.39)	8.96(3.08)	6.68(2.68)	6.72(2.69)
41	GI-35-15-15 x SPV - 946	6.99(2.74)	7.78(2.88)	6.23(2.59)	6.26(2.60)	5.23(2.39)	8.96(3.08)	6.68(2.68)	6.72(2.69)	6.12(2.57)
42	GI-35-15-15 x SPV 104	7.78(2.88)	6.23(2.59)	6.26(2.60)	5.23(2.39)	8.96(3.08)	6.68(2.68)	6.72(2.69)	6.12(2.57)	9.54(3.17)
43	GI-35-15-15 x IS - 2284	6.23(2.59)	6.26(2.60)	5.23(2.39)	8.96(3.08)	6.68(2.68)	6.72(2.69)	6.12(2.57)	9.54(3.17)	10.56(3.32)
44	GI-35-15-15 x IS - 6335	6.26(2.60)	5.23(2.39)	8.96(3.08)	6.68(2.68)	6.72(2.69)	6.12(2.57)	9.54(3.17)	10.56(3.32)	8.22(2.95)
45	GI-35-15-15 x IS - 9471	5.23(2.39)	8.96(3.08)	6.68(2.68)	6.72(2.69)	6.12(2.57)	9.54(3.17)	10.56(3.32)	8.22(2.95)	7.42(2.81)
46	SPV - 946 x SPV - 104	8.96(3.08)	6.68(2.68)	6.72(2.69)	6.12(2.57)	9.54(3.17)	10.56(3.32)	8.22(2.95)	7.42(2.81)	7.97(2.91)
47	SPV - 946 x IS - 2284	6.68(2.68)	6.72(2.69)	6.12(2.57)	9.54(3.17)	10.56(3.32)	8.22(2.95)	7.42(2.81)	7.97(2.91)	9.42(3.15)
48	SPV - 946 x IS - 6335	6.72(2.69)	6.12(2.57)	9.54(3.17)	10.56(3.32)	8.22(2.95)	7.42(2.81)	7.97(2.91)	9.42(3.15)	1.20
49	SPV - 946 x IS 9471	6.12(2.57)	9.54(3.17)	10.56(3.32)	8.22(2.95)	7.42(2.81)	7.97(2.91)	9.42(3.15)	1.20	to
50	SPV - 104 x IS 2284	9.54(3.17)	10.56(3.32)	8.22(2.95)	7.42(2.81)	7.97(2.91)	9.42(3.15)	1.20	to	to
51	SPV - 104 x IS 6335	10.56(3.32)	8.22(2.95)	7.42(2.81)	7.97(2.91)	9.42(3.15)	1.20	to	to	to
52	SPV - 104 x IS 9471	8.22(2.95)	7.42(2.81)	7.97(2.91)	9.42(3.15)	1.20	to	to	to	to
53	IS 2284 x IS - 6335	7.42(2.81)	7.97(2.91)	9.42(3.15)	1.20	to	to	to	to	to
54	IS 2284 x IS - 9471	7.97(2.91)	9.42(3.15)	1.20	to	to	to	to	to	to
55	IS 6335 x IS 9471	9.42(3.15)	1.20	to	to	to	to	to	to	to

Parental range

8.93
to
12.941.20
to
2.300.02
to
5.840.00
to
18.00

Hybrid range

5.23
to
10.560.80
to
1.800.01
to
3.410.00
to
12.00

SE (m)

0.132

0.107

0.111

0.208

CD at 5%

0.377

0.305

0.389

0.595

Figures in parenthesis indicate square root values.

Table : 12 Heterosis (H₁) and Heterobeltiosis (H₂) in % for F₁ crosses in 10 × 10 diallel of sorghum (Pooled), Akola and Patancheru, 1996

Sr. No.	Crosses	100 grain weight		Grain hardness		Endosperm texture		Electrical conductivity		DTF	
		H1	H2	H1	H2	H1	H2	H1	H2	H1	H2
1	2										
1	SPV-1201 x ICSB - 101B	6.26	-5.74	4.22	-3.50	7.11	4.51	-26.59	-30.02*	6.76*	4.64
2	SPV-1201 x AKms - 14 B	-3.46	-16.01	-12.86	-26.50*	20.00	12.72	-16.52	-27.65*	9.50*	4.97
3	SPV-1201 x SRT - 26B	20.61*	0.28	-2.88	-15.10	27.98	15.69	-32.05**	-40.64**	9.86**	6.95*
4	SPV-1201 x GJ-35-15-15	-4.96	-18.36*	11.84	3.95	-6.07	-14.40	-26.62	-26.85	4.07	2.24
5	SPV-1201 x SPV - 946	-12.30	-23.82**	7.76	-5.82	14.91	11.24	-10.74	-13.80	2.12	0.64
6	SPV-1201 x SPV - 104	9.30	8.00	7.64	-13.74	18.42	-5.70	-33.63**	-50.11**	4.07	2.24
7	SPV-1201 x IS - 2284	-5.11	-23.92**	-5.54	-19.39	23.04	-6.58	-28.32	42.64*	2.86	1.32
8	SPV-1201 x IS-6335	-6.64	-24.48**	-5.97	-13.98	12.02	-14.58	-30.12	-43.99**	11.23**	1.66
9	SPV-1201 x IS - 9471	7.32	-9.60	-13.47	-23.10*	23.88	-6.29	-26.25	-40.84*	5.46*	2.32
10	ICSB - 101B x AKms 14 B	-5.35	-7.42	-4.51	-13.74	48.31**	36.15*	-25.11*	-32.26**	-1.94	-4.14
11	ICSB - 101B x SRT - 26 B	-1.97	-9.00	-22.87*	-27.56*	18.99	5.23	-33.60**	-39.44**	-3.82	-4.48
12	ICSB - 101B x GJ-35-15-15	-5.81	-9.25	-6.46	-6.85	8.79	1.43	-28.26*	-31.41*	-3.15	-6.71*
13	ICSB - 101B x SPV - 946	5.92	3.41	9.93	3.23	25.09	18.26	-23.46	-24.49	-0.50	-2.89
14	ICSB - 101B x SPV - 104	14.14	0.18	-1.31	-15.73	5.49	-17.52	-36.03**	-50.25**	-4.48	-7.99**
15	ICSB - 101B x IS - 2284	17.02	4.14	-3.54	-11.74	19.79	-10.56	-42.36*	-55.49**	-3.60	-4.10
16	ICSB - 101B x IS - 6335	11.50	0.24	2.29	0.96	12.38	-15.74	-53.50**	-64.04**	-9.63**	-15.86**
17	ICSB - 101B x IS - 9471	11.94	5.47	-8.52	-12.50	26.89*	-5.59	-46.79**	-58.83**	-4.88	-5.86
18	AKms - 14 B x SRT - 26B	8.72	3.05	-35.50**	-38.14**	52.31**	46.18**	-23.85*	-24.56*	-2.66	-4.20
19	AKms - 14 B x GJ-35-15-15	-7.88	-9.29	-34.35**	-40.91**	44.14*	24.18	-24.24	-34.17**	-6.10*	-11.50**
20	AKms - 14 B x SPV - 946	16.25	16.03	-5.66	-9.50	58.01**	58.01**	-26.63*	-34.43**	-7.14**	-12.22**
21	AKms - 14 B x SPV - 104	8.75	-6.34	-12.03	-17.43	36.87**	14.59	-29.05**	-40.17**	-5.42*	-10.86**
22	AKms - 14 B x IS - 2284	16.05	5.34	6.98	5.49	35.94**	8.08	-47.65**	-70.28**	-6.48*	-13.72**
23	AKms - 14 B x IS - 6335	23.72*	13.49	-1.82	-10.25	43.11**	14.31	-59.65**	-70.90**	-9.30**	-13.72**
24	AKms - 14 B x IS - 9471	20.95*	16.41	-12.77	-17.85	33.46**	5.68	-57.12**	-69.06**	-10.87**	-11.97**

Contd. . . .

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Table 12 Contd. ...

1	2	3	4	5	6	7	8	9	10	11	12
25	SRT - 26B x GJ-35-15-15	2.32	-1.57	-2.90	-9.15	23.27	2.62	-14.78	-25.35*	-4.51	-8.63**
26	SRT - 26B x SPV - 946	18.36	12.39	-5.71	-5.73	30.40	21.47	-34.28**	-40.78**	-1.51	-5.47
27	SRT - 26B x SPV - 104	8.15	-10.94	5.09	-5.14	20.21	4.19	-46.49**	-55.22**	-7.85**	-11.82**
28	SRT - 26B x IS - 2284	21.34	15.91	-3.83	-6.50	12.40	-7.73	-52.04**	-65.26*	-8.81**	-9.90**
29	SRT - 26B x IS - 6335	22.94*	18.75	-11.49	-15.83	29.90*	7.16	61.63**	-72.17**	-10.82**	-16.43**
30	SRT - 26B x IS - 9471	20.20	18.29	-18.32	-19.86	5.29	-13.94	-41.96**	-57.88**	-0.70	-1.05
31	GJ-35-15-15 x SPV - 946	9.26	-7.79	-3.01	-9.27	20.00	6.22	-13.16	-15.89	-0.96	-1.28
32	GJ-35-15-15 x SPV 104	-5.30	-19.49*	3.42	-12.00	21.41	-9.69	-31.25**	-48.22**	-6.39*	-6.39*
33	GJ-35-15-15 x IS - 2284	-0.93	-8.79	-1.80	-10.50	30.53*	-6.89	-33.63	-47.01**	-5.94*	-8.95**
34	GJ-35-15-15 x IS - 6335	0.00	-6.96	-20.79	-22.14	34.85**	-3.44	-49.47**	-59.59**	-7.64**	-16.93**
35	GJ-35-15-15 x IS - 9471	3.69	1.31	-15.61	-19.60	28.42*	-8.69	-26.64	-41.30*	-3.85	-8.31**
36	SPV - 946 x SPV - 104	5.29	-9.47	3.40	-6.64	8.44	-11.46	-28.01**	-44.55**	-3.53	-3.83
37	SPV - 946 x IS - 2284	18.26	7.54	0.40	-2.36	3.70	-19.43	-52.06**	-62.63**	-8.94**	-11.58**
38	SPV - 946 x IS - 6335	2.99	-5.36	0.99	-3.98	2.98	-19.63	-48.18**	-59.53**	7.66**	-2.29
39	SPV - 946 x IS 9471	8.87	4.98	-3.95	-5.77	15.72	-10.44	-56.23**	-65.80**	-10.25**	-14.15**
40	SPV - 104 x IS 2284	0.64	-20.04*	-16.57	-22.71	21.18*	13.60	-48.51**	-66.47**	-5.94*	-8.95**
41	SPV - 104 x IS 6335	-0.80	-20.50*	-16.74	-28.13*	31.14**	23.66*	-60.62**	-74.33**	-13.32**	-22.04**
42	SPV - 104 x IS 9471	16.69	-2.67	-18.48	-27.65*	23.42*	15.12	-70.69**	-80.89**	-17.59**	-21.41**
43	IS 2284 x IS -6335	4.55	3.35	-0.95	-8.28	-3.07	-3.67	-11.56	-11.76	-2.39	-9.56**
44	IS 2284 x IS - 9471	8.48	2.06	-4.81	-9.14	1.65	1.09	10.28	-10.55	-0.17	-1.71
45	IS 6335 x IS 9471	14.10	8.53	-17.87	-20.54	-2.72	-3.84	-25.95	-26.00	-9.36**	-14.79**
SE (m)		0.193	0.223	0.671	0.755	5.274	6.091	39.798	45.955	1.972	2.277
CD at 5%		0.382	0.441	1.330	1.536	10.456	12.073	78.887	91.090	3.908	4.513

* Significant at 5%

** Significant at 1%

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Table : 13 Heterosis (H₁) and Heterobeltiosis (H₂) in % for F₁ crosses in 10 x10 diallel of sorghum (Pooled) Akola and Patancheru, 1996

Sr. No.	Crosses	Plant height		Cob length		Glume covering		Mesocarp thickness			TGM	
		H1	H2	H1	H2	H1	H2	H1	H2	H1	H1	H2
1	2	3	4	5	6	7	8	9	10	11	11	12
1	SPV-1201 x ICSB - 101B	54.40**	31.63**	-3.34	-13.43	-20.00	-33.33**	-28.25**	-38.02**	-20.69*	-23.33*	-23.33*
2	SPV-1201 x AKms - 14 B	56.81**	24.49**	-2.42	-6.33	-12.50	-12.50	32.60**	30.20**	-12.50	-12.50	-22.22**
3	SPV-1201 x SRT - 26B	33.71**	14.70*	-4.23	-9.07	-12.50	-12.50	17.42**	8.85	-13.79	-13.79	-16.67
4	SPV-1201 x GJ-35-15-15	22.66**	19.04**	12.94	8.19	0.00	0.00	21.58**	20.50**	-17.86*	-17.86*	-17.86*
5	SPV-1201 x SPV - 946	-3.93	-10.78*	9.05	5.11	-12.50	-12.50	5.25	-7.11	-3.70	-3.70	-7.14
6	SPV-1201 x SPV - 104	33.49**	23.11**	2.44	-3.90	5.88	0.00	2.68	-7.91	-3.23	-3.23	-11.76
7	SPV-1201 x IS - 2284	26.44**	22.94**	7.83	6.74	-11.11	-20.00	-23.06**	-31.31**	-27.27*	-27.27*	-42.86**
8	SPV-1201 x IS-6335	36.51**	32.86**	14.93	7.37	10.00	-8.33	-40.10**	-41.51**	-27.27*	-27.27*	-42.86**
9	SPV-1201 x IS - 9471	34.98**	25.90**	16.22	6.27	0.00	0.00	-40.67**	-43.88**	-23.81*	-23.81*	-42.86**
10	ICSB - 101B x AKms 14 B	20.59*	10.57	16.10*	7.99	-20.00	-33.33**	-3.08	-17.32**	3.03	3.03	-5.56
11	ICSB - 101B x SRT - 26 B	32.53**	31.55**	13.71	6.90	10.00	-8.33	-3.03	-10.21	6.67	6.67	6.67
12	ICSB - 101B x GJ-35-15-15	23.37**	2.61	11.59	3.99	-10.00	-25.00*	9.49	-5.94	-10.34	-10.34	-13.33
13	ICSB - 101B x SPV - 946	33.86**	7.40	5.58	-2.18	-10.00	-25.00*	8.52	-15.54**	-17.86*	-17.86*	-23.33**
14	ICSB - 101B x SPV - 104	15.88*	6.31	8.25	-8.35	-23.81	-33.33**	-9.99*	-13.74**	-12.50	-12.50	-17.65*
15	ICSB - 101B x IS - 2284	48.79**	23.94**	10.24	-0.36	-18.18	-25.00*	-47.56**	-49.50**	-30.43**	-30.43**	-46.67**
16	ICSB - 101B x IS - 6335	46.67**	22.28**	11.87	6.90	0.00	0.00	-56.61**	-61.74**	-30.43**	-30.43**	-46.67**
17	ICSB - 101B x IS - 9471	37.15**	10.41	14.39	11.80	-20.00	-33.33**	-37.69**	-48.63**	-27.27*	-27.27*	-46.67**
18	AKms - 14 B x SRT - 26B	32.68**	20.83*	6.57	5.36	0.00	0.00	29.88**	18.76**	3.03	3.03	-5.56
19	AKms - 14 B x GJ-35-15-15	13.29*	-12.02*	19.79*	19.54*	25.00	25.00	-5.20	-6.00	0.00	0.00	-11.11
20	AKms - 14 B x SPV - 946	36.49**	2.65	13.77	13.29	25.00	25.00	4.11	-6.91	9.68	9.68	-5.56
21	AKms - 14 B x SPV - 104	18.51*	0.50	9.58	-1.05	-5.88	-11.11	-23.45**	-32.25**	-2.86	-2.86	-5.56
22	AKms - 14 B x IS - 2284	43.21**	11.37	9.03	5.70	-22.22	-30.00*	-27.12**	-35.78**	-38.46**	-38.46**	-55.56**
23	AKms - 14 B x IS - 6335	41.71**	10.28	8.20	5.18	0.00	-16.67	-35.05**	-37.49**	-46.15**	-46.15**	-61.11**
24	AKms - 14 B x IS - 9471	43.19**	8.02	13.60	7.98	-12.50	-12.50	-19.95**	-23.20**	-44.00**	-44.00**	-61.11**

Contd.

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Table 13 Contd.....

1	2	3	4	5	6	7	8	9	10	11	12
25	SRT - 26B x GI-35-15-15	-3.23	-19.04**	6.97	5.98	0.00	0.00	-16.77**	-23.30**	10.34	6.67
26	SRT - 26B x SPV - 946	14.77*	-7.40	7.64	5.98	0.00	0.00	31.30**	8.55	-3.57	-10.00
27	SRT - 26B x SPV - 104	16.59*	7.69	8.19	-3.30	-5.88	-11.11	1.12	-2.45	0.00	-5.88
28	SRT - 26B x IS - 2284	46.10**	22.43**	10.54	5.98	-11.11	-20.00	-27.11**	-30.03**	-21.74*	-40.00**
29	SRT - 26B x IS - 6335	25.72**	5.44	1.22	-0.50	0.00	-16.67	-21.87**	-25.93**	-21.74*	-40.00**
30	SRT - 26B x IS - 9471	23.39**	-0.09	5.64	1.52	0.00	0.00	-26.60**	-35.35**	-27.27*	-46.67**
31	GI-35-15-15 x SPV - 946	-0.43	-4.84	-4.02	-4.62	12.50	12.50	10.79	-1.68	11.11	7.14
32	GI-35-15-15 x SPV 104	24.85**	12.02*	1.66	0.63	-5.88	-11.11	-6.68	-16.78**	-6.45	-14.71
33	GI-35-15-15 x IS - 2284	28.51**	28.26**	17.92*	14.08	11.11	0.00	-37.52**	-44.53**	-27.27*	-42.86**
34	GI-35-15-15 x IS - 6335	27.14**	26.75**	14.93	11.95	10.00	-8.33	-49.84**	-51.32**	-27.27*	-42.86**
35	GI-35-15-15 x IS - 9471	15.12**	10.51*	13.27	7.89	0.00	0.00	-37.86**	-40.86**	-23.81*	-42.86**
36	SPV - 946 x SPV - 104	27.12**	9.59	17.61	6.63	17.65	11.11	-17.29**	-33.54**	0.00	-11.76
37	SPV - 946 x IS - 2284	10.48*	5.39	8.42	5.53	-22.22	-30.00*	-25.21**	-40.93**	-14.29	-30.77**
38	SPV - 946 x IS - 6335	18.74**	13.15*	9.26	5.78	10.00	-8.33	-35.63**	-44.35**	-23.81*	-38.46**
39	SPV - 946 x IS 9471	13.12**	12.60*	11.04	5.13	12.50	12.50	-40.72**	-44.93**	-20.00	-38.46**
40	SPV - 104 x IS 2284	32.18**	18.81**	9.55	1.80	-5.26	-10.00	-57.19**	-57.41**	-20.00	-41.18**
41	SPV - 104 x IS 6335	27.06**	14.31*	22.40*	7.77	14.29	0.00	-61.14**	-64.39**	-36.00	-52.94**
42	SPV - 104 x IS 9471	14.91**	-0.55	18.28*	2.09	-17.65	-22.22	-38.19**	-47.22**	-33.33**	-52.94**
43	IS 2284 x IS - 6335	17.42**	17.30**	9.61	3.39	9.09	0.00	-24.18**	-30.84**	-12.50	-12.50
44	IS 2284 x IS - 9471	9.86*	5.25	0.10	-7.60	-33.33*	-40.00**	-13.99*	-26.87**	6.67	0.00
45	IS 6335 x IS 9471	8.14	3.50	12.74	10.17	-20.00	-33.33**	-51.82**	-55.43**	-6.67	-12.50
SE (m)		12.443	14.368	1.998	2.307	8.616	9.424	4.370	5.046	0.303	0.350
CD at 5%		24.663	28.479	3.961	4.574	16.177	18.679	8.662	10.002	0.600	0.693

* Significant at 5%

** Significant at 1%

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Table : 14 Heterosis (H₁) and Heterobeltiosis (H₂) in % for F₁ crosses in 10 × 10 diallel of sorghum (Pooled) Akola and Patancheru, 1996

Sr. No.	Crosses	Germination		F. moniliforme		F. pallidroseum		C. lunata		Other fungi	
		H1	H2	H1	H2	H1	H2	H1	H2	H1	H2
1	SPV-1201 x ICSB-101B	49.62**	37.70**	5	6	7	8	9	10	11	12
2	SPV-1201 x AKms - 14 B	35.77**	-0.46	-23.69	-34.39*	67.48	36.74	-2.64	-5.15	28.42	20.47
3	SPV-1201 x SRT - 26B	43.45**	40.56**	-32.75	-33.62	-14.54	-15.74	-1.97	-8.82	92.08	36.88
4	SPV-1201 x GJ-35-15-15	19.77*	17.05	-5.45	-7.96	2.34	-2.53	-12.32	-13.57	175.93**	138.15**
5	SPV-1201 x SPV - 946	16.37	14.44	-0.02	-3.32	-19.85	-27.45	-14.39	-14.71	61.01	57.88
6	SPV-1201 x SPV - 104	31.00**	9.25	-32.37	-43.03**	-31.30	-36.70	-12.64	-16.17	45.34	38.16
7	SPV-1201 x IS - 2284	39.30**	22.50**	-36.24	-39.34	0.93	-19.02	20.32	2.21	104.60*	72.51
8	SPV-1201 x IS-6335	32.73**	17.43*	-27.53	-37.16	28.32	25.31	-25.62	-33.82	10.38	-00.02
9	SPV-1201 x IS - 9471	35.01**	23.13*	-39.55	-53.99*	23.60	15.38	-25.25	-45.59*	15.41	13.16
10	ICSB - 101B x AKms 14 B	65.32**	28.51*	-18.11	-28.02	52.71	22.28	-33.01	-49.26**	5.76	-0.02
11	ICSB - 101B x SRT - 26 B	18.13	10.80	0.41	-0.84	50.64	44.90	-7.32	-11.63	59.89	18.75
12	ICSB - 101B x GJ-35-15-15	36.30**	28.17*	-14.15	-18.48	21.88	18.97	-8.55	-12.14	16.16	6.23
13	ICSB - 101B x SPV - 946	43.25**	33.89**	-9.17	-9.92	35.48	25.50	2.27	0.00	25.71	15.89
14	ICSB - 101B x SPV - 104	31.11*	17.65	10.57	-4.85	83.06	64.82	-15.75	-17.05	48.90	33.23
15	ICSB - 101B x IS - 2284	51.76**	24.28**	-36.66	-41.19	-64.85*	-71.25*	-23.21	-33.33	-5.14	-15.42
16	ICSB - 101B x IS - 6335	51.44**	24.69**	-53.47*	-60.51**	-42.42	-42.56	-25.11	-31.78	63.63	40.00
17	ICSB - 101B x IS - 9471	57.68**	33.45**	-43.81	-57.98**	-20.56	-27.55	-36.13	-52.72**	77.12	63.13
18	AKms - 14 B x SRT - 26B	31.60*	2.23	-8.43	-20.38	27.70	5.42	-26.63	-43.41*	42.47	26.83
19	AKms - 14 B x GJ-35-15-15	27.61*	-5.03	-12.88	-26.76	75.01	37.63	-12.06	-19.29	75.90	39.43
20	AKms - 14 B x SPV - 946	32.64*	-1.70	-28.05	-36.30*	29.62	-1.94	-5.55	-11.85	38.19	-2.66
21	AKms - 14 B x SPV - 104	19.20	1.20	-16.16	-18.19	148.84*	117.88	11.56	7.99	53.39	6.09
22	AKms - 14 B x IS - 2284	50.44**	1.83	-39.76*	-50.31**	33.02	-8.04	-3.77	-12.82	33.01	7.72
23	AKms - 14 B x IS - 6335	76.76**	20.08*	-50.81**	-62.41**	12.25	-10.01	5.84	0.86	85.03	24.45
24	AKms - 14 B x IS - 9471	71.44**	18.65*	-24.99	-48.40**	36.59	18.15	-15.08	-35.04	142.84*	71.05
								-0.55	-20.52	99.19	9.75

Contd. ...

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Table 14 Contd.

	1	2	3	4	5	6	7	8	9	10	11	12
25	SRT - 26B x GI-35-15-15	27.36**	27.02*	7.63	3.45	31.63	23.71	-26.55	-27.86	75.20	48.77	
26	SRT - 26B x SPV - 946	16.70	16.26	12.23	9.92	23.93	10.77	-17.73	-22.14	0.00	-17.26	
27	SRT - 26B x SPV - 104	27.29*	7.95	-11.75	-24.85	47.63	37.80	2.14	-14.28	12.41	9.30	
28	SRT - 26B x IS - 2284	27.74**	10.38	-28.46	-32.78	-29.96	-44.40	-16.25	-26.42	64.96	31.12	
29	SRT - 26B x IS - 6335	33.03**	15.63	-33.66	-43.11	-16.67	-20.02	-22.78	-44.29*	45.66	23.63	
30	SRT - 26B x IS - 9471	33.73**	19.74*	-30.28	-47.42*	5.67	-00.04	-34.29	-50.71**	-2.28	-19.52	
31	GI-35-15-15 x SPV - 946	4.14	3.47	5.27	-0.83	-25.87	-29.74	-6.16	-9.63	14.60	11.03	
32	GI-35-15-15 x SPV 104	25.03*	6.26	-9.56	-25.46	45.38	28.12	-8.69	-22.22	20.49	-0.02	
33	GI-35-15-15 x IS - 2284	25.58**	8.26	-29.17	-30.82	-12.55	-27.06	-31.11	-38.51*	-15.66	-22.22	
34	GI-35-15-15 x IS - 6335	38.09**	19.76*	-7.36	-17.76	26.63	23.42	-21.82	-42.96*	18.41	18.41	
35	GI-35-15-15 x IS - 9471	20.00*	7.18	-1.22	-23.37	48.93	32.89	-21.95	-40.74*	13.92	9.75	
36	SPV - 946 x SPV - 104	24.63*	5.36	-16.09	-27.28	-36.73	-46.76	9.09	-4.00	57.12	27.15	
37	SPV - 946 x IS - 2284	19.14*	3.27	-12.10	-19.00	19.89	4.66	-31.60	-36.80	2.89	-2.22	
38	SPV - 946 x IS - 6335	27.96**	11.58	-5.88	-20.67	63.27	51.04	-19.78	-40.00*	14.60	11.03	
39	SPV - 946 x IS 9471	10.13	-1.06	-8.90	-32.24	49.78	27.51	-24.10	-40.80*	12.85	12.20	
40	SPV - 104 x IS 2284	49.62**	12.72	-42.33*	-53.34**	30.23	-1.71	-8.45	-13.20	48.45	15.57	
41	SPV - 104 x IS 6335	50.32**	13.79	-48.39**	-61.22**	8.34	-2.32	12.11	-7.37	74.43	44.74	
42	SPV - 104 x IS 9471	53.27**	18.82*	-30.36	-52.73**	19.83	18.15	0.61	-12.62	27.14	2.43	
43	IS 2284 x IS - 6335	6.33	5.59	-30.79	-37.24	-28.08	-41.24	-7.14	-26.41	6.01	-2.23	
44	IS 2284 x IS - 9471	19.83**	15.09	-14.27	-32.34	24.62	-4.88	-3.40	-19.80	13.96	8.90	
45	IS 6335 x IS 9471	11.29	7.61	-23.92	-34.92	-25.93	-32.36	3.02	-2.87	-13.94	-17.08	

* Significant at 5%

** Significant at 1%

Table 15 Heterosis (H_1) and Heterobeltiosis (H_2) in % F_2 progenies 10×10 diallel (Pooled), Akola and Patancheri, 1996

Crosses	Grain hardness		Ele. Conductivity		Plant height		Cob length		Glume covering		TGM	
	H1	H2	H1	H2	H1	H2	H1	H2	H1	H2	H1	H2
	3	4	5	6	7	8	9	10	11	12	13	14
SPV-1201 x ICSB-101B	16.69	-22.89*	29.92**	23.84	-10.06	-23.32*	0.20	-10.25	-8.24	-13.33	-28.28**	-30.67**
SPV-1201 x AKms-14 B	-16.01	-29.15**	19.64	3.69	17.48	-6.73	-2.53	-6.43	0.00	0.00	-20.00*	-28.80**
SPV-1201 x SRT-26B	-15.76	-26.37**	13.82**	15.16	1.49	-12.94	0.11	-4.95	-5.00	-5.00	-23.45**	-26.00**
SPV-1201 x GI-35-15-15	-15.31	-24.35*	104.95**	98.65**	2.52	1.87	6.70	-4.03	-7.50	-7.50	-17.93*	-20.67*
SPV-1201 x SPV-946	-31.43**	-38.67**	19.68	12.17	-7.60	-11.31	1.22	-1.73	-12.50	-12.50	-15.00	-15.00
SPV-1201 x SPV-104	-30.01**	-43.91**	12.95	-15.09*	-12.55	-19.35*	4.16	2.29	-17.50	-17.50	-22.58**	-29.41**
SPV-1201 x IS-2284	-25.95**	-36.80**	56.48**	25.22	-4.34	-6.98	7.26	6.18	-2.12	-7.56	-15.45	-33.57**
SPV-1201 x IS-6335	-21.29*	-27.99**	0.78	-19.21	-5.46	-7.98	11.62	4.28	-22.00*	-35.00**	-20.00	-37.14**
SPV-1201 x IS-9471	-25.45**	-33.74**	-5.88	-24.51	-9.80	-15.87*	0.52	-8.08	-6.00	-6.00	-16.19	-37.14**
ICSB-101B x AKms 14 B	-16.10	-24.18*	2.18	-7.57	12.70	3.34	12.78	4.90	-5.88	-11.11	-27.88**	-33.89**
ICSB-101B x SRT-26 B	-23.10*	-27.74*	14.63	4.56	23.76*	22.84	6.47	0.09	-5.88	-11.11	-21.67**	-24.67**
ICSB-101B x GI-35-15-15	-18.07	-21.18	28.56*	26.36*	2.70	-12.90	10.59	1.36	-5.88	-11.11	-28.00**	-28.00**
ICSB-101B x SPV-946	-15.70	-18.78	20.74	18.60	0.85	-15.23	8.88	0.18	-5.88	-11.11	-23.52**	-28.00**
ICSB-101B x SPV-104	-13.52	-26.14*	1.39	-21.15**	0.82	-7.50	8.25	-8.35	-5.88	-11.11	-27.50**	-31.76**
ICSB-101B x IS-2284	-4.80	-12.87	23.60	-4.56	20.47*	0.35	9.14	-1.36	-10.67	-10.67	-14.78	-34.67**
ICSB-101B x IS-6335	-11.31	-12.43	12.99	-12.61	6.20	-11.46	4.56	-0.09	-17.71*	-28.00**	-11.30	-32.00**
ICSB-101B x IS-9471	-23.81*	-27.10*	-7.15	-28.15*	6.07	-14.61	1.58	-0.73	1.65	-4.00	-1.55	-30.00**
AKms-14 B x SRT-26B	-11.28	-14.91	-0.40	-1.32	35.46**	23.36	8.55	7.32	0.00	0.00	-21.21**	-27.76**
AKms-14 B x GI-35-15-15	-17.27	-22.49	0.95	-4.74	2.20	-19.24*	13.50	11.71	0.00	0.00	-29.70**	-35.56**
AKms-14 B x SPV-946	-11.13	-16.86	17.65	8.16	8.46	-16.45	4.91	3.69	8.50	8.50	-23.75**	-32.22**
AKms-14 B x SPV-104	-20.83	-25.69	4.61	-11.79	10.86	-5.99	8.18	-2.32	0.00	0.00	-25.14**	-27.22**
AKms-14 B x IS-2284	-20.60	-21.70	5.21	-24.23*	17.92	-8.30	5.95	2.74	-8.24	-13.33	-26.15**	-46.67**
AKms-14 B x IS-6335	-13.16	-20.62	-30.99*	-50.23**	18.61	-7.69	6.05	3.09	-27.20**	-39.33**	-30.00**	-49.44**
AKms-14 B x IS-9471	-16.72	-21.57	-16.33	-39.63**	9.90	-17.10*	5.80	0.57	-12.50	-12.50	-29.60**	-51.11**

Contd.....

Table 15 Contd.

	1	2	3	4	5	6	7	8	9	10	11	12	13	14
5	SRT - 26B x GI-35-15	-28.55**	-30.28*	6.87	-4.02	-8.13	-21.61*	7.52	4.64	0.00	0.00	0.00	-27.33**	-27.33**
6	SRT - 26B x SPV - 946	-29.76**	-31.57*	29.48**	20.05	7.60	-10.79	-0.84	-3.09	0.00	0.00	0.00	-25.52**	-28.00**
7	SRT - 26B x SPV - 104	-33.43**	-39.91**	4.86	-12.25	22.02*	12.71	5.07	-6.08	4.00	4.00	4.00	-23.12**	-27.65**
8	SRT - 26B x IS - 2284	-33.57**	-35.41**	35.74**	-1.68	25.15**	4.88	-7.20	-11.03	-8.24	-13.33	-13.33	-5.22	-27.33**
9	SRT - 26B x IS - 6335	-46.62**	-43.87**	8.00	-21.66*	24.40**	4.33	8.61	6.77	-20.00*	-33.33**	-33.33**	-21.74	-40.00**
10	SRT - 26B x IS - 9471	-36.47**	-37.66**	-33.09**	-51.44**	17.34*	-5.00	8.80	4.56	0.00	0.00	0.00	-6.36	-31.33**
11	GI-35-15-15 x SPV - 946	-17.73	-17.85	69.04**	63.27**	-8.80	-11.93	13.12	12.63	2.50	2.50	2.50	-23.45**	-26.00**
12	GI-35-15-15 x SPV 104	-15.33	-25.22*	32.00**	1.40	-4.32	-12.27	13.20	3.70	0.00	0.00	0.00	-19.38*	-24.12**
13	GI-35-15-15 x IS - 2284	-10.52	-15.05	19.18	-6.82	6.25	3.95	27.54**	25.60**	16.71	10.22	10.22	-24.35*	-42.00**
14	GI-35-15-15 x IS - 6335	-30.25**	-32.06**	24.26	-2.70	-6.23	-8.17	12.07	7.27	-16.80	-30.67**	-30.67**	-20.87	-39.33**
15	GI-35-15-15 x IS - 9471	-34.36**	-34.73*	9.46	-14.24	-4.42	-10.32	14.92*	7.60	-2.50	-2.50	-2.50	-20.00	-41.33**
16	SPV - 946 x SPV - 104	-12.64	-22.94	13.56	-10.53	0.99	-10.28	0.59	-8.21	6.50	6.50	6.50	-25.81**	-32.35**
17	SPV - 946 x IS - 2284	7.94	2.33	40.50**	7.14	7.67	-8.89	5.95	3.89	13.41	7.11	7.11	-19.09	-36.43**
18	SPV - 946 x IS - 6335	-14.38	-16.48	9.66	-16.25	4.79	3.30	17.62*	13.05	13.20	5.67	5.67	-15.45	-33.57**
19	SPV - 946 x IS 9471	-12.31	-12.94	-14.23	-34.46**	-8.47	-11.17	4.85	-1.43	10.00	10.00	10.00	-2.00	-40.00**
20	SPV - 104 x IS 2284	-28.16*	-33.45*	4.58	-31.90**	9.90	-1.21	17.26	-0.34	20.47	13.78	13.78	-9.60	-33.53**
21	SPV - 104 x IS 6335	-21.49	-32.23**	17.64	-23.32**	-4.76	-14.31	11.54	-1.79	-11.60	-26.33**	-26.33**	-21.60*	-42.35**
22	SPV - 104 x IS 9471	-18.83	-27.95*	-19.68*	-47.63**	-9.30	-21.50**	8.59*	-6.27	-6.00	-6.00	-6.00	-17.50	-41.76**
23	IS 2284 x IS - 6335	-29.25**	-34.48**	46.01*	-45.68*	-4.70	-4.80	3.48	-2.39	-20.76*	-30.67**	-30.67**	-6.25	-6.25
24	IS 2284 x IS - 9471	-31.67**	-34.78**	1.04	0.74	-5.24	-9.22	11.84	3.23	-5.88	-11.11	-11.11	-1.33	-7.50
25	IS 6335 x IS 9471	-30.88**	-33.05**	-1.56	-1.63	-9.27	-13.16	7.00	4.56	-3.60	-19.67*	-19.67*	4.00	-2.50
SE		0.613	0.708	32.899	37.988	18.553	21.423	1.684	1.944	5.548	6.406	6.406	0.372	0.372
CD at 5%		1.216	1.404	65.210	75.299	36.775	42.464	3.337	3.854	10.997	12.699	12.699	0.638	0.737

* Significant at 5%

** Significant at 1%

Table : 16 Heterosis (H₁) and Heterobeltiosis (H₂) in % F₂ progenies 10 x 10 dialled (Pooled), Akola and Patancheru, 1996

Sr. No.	Crosses	Germination		<i>F. moniliforme</i>		<i>F. pallidorozeum</i>		<i>C. lunata</i>		Other fungi	
		H1	H2	H1	H2	H1	H2	H1	H2	H1	H2
1	2	3	4	5	6	7	8	9	10	11	12
1	SPV-1201 x ICSB - 101B	45.67**	34.06*	8.37	-0.83	-62.90	-65.54	-16.64	-18.88	47.42	20.70
2	SPV-1201 x AKms - 14 B	5.66**	11.18	13.26	-0.04	-41.22	-49.36	-24.98	-27.46	36.94	4.43
3	SPV-1201 x SRT - 26B	40.03**	37.20*	20.54	9.37	-35.38	-39.26	-15.45	-15.74	46.94	28.58
4	SPV-1201 x GJ-35-15-15	57.41**	25.31	17.15	15.23	-15.45	-24.82	-4.17	-7.34	-12.25	-17.13
5	SPV-1201 x SPV - 946	29.78**	22.81	18.89	13.94	-40.46	-48.30	-14.66	-16.17	2.28	-5.35
6	SPV-1201 x SPV - 104	42.74**	19.04	21.31	3.02	-58.55	-59.69	-13.99	-18.87	17.18	5.48
7	SPV-1201 x IS - 2284	17.27	3.13	14.08	1.11	-51.61	-60.36	0.22	-11.51	34.05	26.58
8	SPV-1201 x IS-6335	20.03	6.20	60.46	19.41	0.02	-13.99	3.77	-20.15	7.04	7.03
9	SPV-1201 x IS - 9471	7.57	-1.90	40.44	7.72	-30.11	-33.08	10.18	-11.70	13.78	9.53
10	ICSB - 101B x AKms 14 B	78.73**	38.93*	-7.59	-11.20	-44.85	-55.32	3.26	2.58	53.30	39.68
11	ICSB - 101B x SRT - 26 B	34.31*	25.98	9.11	8.10	-40.58	-41.33	-7.75	-9.92	89.93	75.80
12	ICSB - 101B x GJ-35-15-15	43.24**	29.84*	3.04	-4.26	-58.50	-65.40	-4.04	-4.68	51.66	18.87
13	ICSB - 101B x SPV - 946	34.45*	30.53	18.90	13.29	-43.36	-47.36	-10.32	-14.23	15.63	-10.77
14	ICSB - 101B x SPV - 104	31.81	18.27	14.83	5.71	-19.57	8.22	-18.32	-24.90	31.39	18.12
15	ICSB - 101B x IS - 2284	24.22	1.72	48.21	21.64	-35.08	-49.77	-14.11	-25.92	38.63	8.65
16	ICSB - 101B x IS - 6335	32.08*	8.76	29.39	-9.07	-64.47	-71.27	10.08	-16.88	45.48	19.18
17	ICSB - 101B x IS - 9471	34.36**	13.71	36.00	-1.85	19.57	-28.23	-8.95	-28.50	37.78	16.44
18	AKms - 14 B x SRT - 26B	19.68	-11.09	8.15	4.87	-25.00	-38.64	-3.14	-6.02	-0.59	-15.50
19	AKms - 14 B x GJ-35-15-15	62.84**	18.13	19.92	7.40	-66.22	-67.41	-3.94	-3.94	-18.18	-40.00
20	AKms - 14 B x SPV - 946	39.71*	6.34	4.36	-4.27	-29.34	-45.86	10.07	4.60	-8.24	-33.62
21	AKms - 14 B x SPV - 104	50.03*	27.37	35.13	29.24	-23.79	-32.73	-29.35	-35.43	12.18	-7.10
22	AKms - 14 B x IS - 2284	39.91**	-5.30	35.34	7.68	-1.71	-7.46	-25.61	-36.20	44.53	5.98
23	AKms - 14 B x IS - 6335	57.66**	7.11	30.69	-10.24	1.90	1.66	-9.68	-32.11	65.55	26.23
24	AKms - 14 B x IS - 9471	48.48**	2.76	19.07	-16.13	-55.85	-60.48	-6.18	-26.69	96.04	53.66

Contd.

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Table 16 Contd.

	1	2	3	4	5	6	7	8	9	10	11	12
25	SRT - 26B x GJ-35-15-15	26.00	21.47	26.48	16.51	-31.02	-41.89	-24.64	-26.89	26.00	5.00	
26	SRT - 26B x SPV - 946	31.66*	27.07	-12.57	-17.43	-50.74	-54.75	10.03	7.71	33.09	9.02	
27	SRT - 26B x SPV - 104	23.40	4.64	-1.93	-8.93	-58.08	-61.61	7.98	1.52	19.03	15.30	
28	SRT - 26B x IS - 2284	3.85	-10.27	7.58	-12.35	-30.26	-45.54	-1.17	-13.00	15.03	-4.15	
29	SRT - 26B x IS - 6335	28.19*	11.43	43.21	0.10	-39.04	-50.22	19.87	-7.99	28.58	12.51	
30	SRT - 26B x IS - 9471	15.90	3.77	58.35	13.62	-39.34	-45.25	4.43	-16.33	14.09	3.30	
31	GJ-35-15-15 x SPV - 946	26.64	17.99	14.39	11.41	-74.43*	-79.88*	-13.24	-17.55	8.39	6.08	
32	GJ-35-15-15 x SPV 104	13.24	-6.80	35.30	16.50	-24.26	-30.93	-1.24	-9.74	-29.64	-39.80	
33	GJ-35-15-15 x IS - 2284	24.54*	11.12	47.30	28.68	46.03	32.91	-12.06	-24.54	-15.54	-15.55	
34	GJ-35-15-15 x IS - 6335	16.53	4.62	75.60	29.28	-20.20	-23.20	-9.39	-31.90	-8.59	-13.67	
35	GJ-35-15-15 x IS - 9471	8.35	0.33	67.80	27.23	-13.31	-19.78	-13.36	-32.29	-6.63	-14.93	
36	SPV - 946 x SPV - 104	47.35**	28.80	18.24	4.17	-37.26	-46.81	-15.40	-10.83	-21.40	-33.94	
37	SPV - 946 x IS - 2284	18.30	-0.79	55.97	33.22	-24.31	-44.36	-7.22	-16.78	15.88	13.42	
38	SPV - 946 x IS - 6335	38.44**	16.76	16.04	-16.00	-48.69	-60.75	-8.61	-28.77	36.90	26.70	
39	SPV - 946 x IS 9471	21.78	5.67	64.19	22.28	-31.62	-42.77	-13.58	-29.79	-12.64	-21.94	
40	SPV - 104 x IS 2284	24.00	-6.58	35.03	3.95	-23.57	-8.55	-23.57	-1.91	-8.58	-16.01	
41	SPV - 104 x IS 6335	17.29	-11.21	62.55	9.00	29.28	13.88	2.03	-17.96	1.08	-9.02	
42	SPV - 104 x IS 9471	20.04	-6.95	82.99*	25.62	19.82	17.93	-22.08	-34.55	0.96	-5.86	
43	IS 2284 x IS - 6335	6.89	6.15	150.48**	104.41**	71.64	61.94	-31.19	-41.39	-10.09	-15.10	
44	IS 2284 x IS - 9471	20.00	15.26	116.33*	82.93	-11.92	-25.23	-38.72	-45.25	-3.05	-11.68	
45	IS 6335 x IS 9471	18.66	14.74	114.25	105.11	13.56	1.45	-16.93	-21.33	-9.37	-12.77	
SE		5.899	6.800	9.56	11.03	2.40	2.77	8.42	9.72	4.72	5.46	
CD at 5%		11.673	13.478	18.94	21.87	4.76	5.50	16.68	19.26	9.37	10.81	

* Significant at 5%

** Significant at 1%

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Table 17 Heterosis (H_1) and heterobeltiosis (H_2) in % for 10×10 diallel of sorghum biochemical character Akola, 1995

Sr. No.	Crosses	Protein		Soluble sugars		Tannins		Flavan-4-ols	
		H1	H2	H1	H2	H1	H2	H1	H2
1	SPV-1201 x ICSB - 101B	-14.49**	-15.44**	-7.34*	-9.51*	-0.88	-2.67	0.00	0.00
2	SPV-1201 x AKms - 14 B	-26.15**	-26.99**	-17.40**	-22.09**	-4.58*	-4.61*	0.00	0.00
3	SPV-1201 x SRT - 26B	-25.65**	-25.73**	-11.82**	-16.10**	-2.25	-2.67	0.00	0.00
4	SPV-1201 x GJ-35-15-15	-14.20**	-17.39**	1.31	-2.33	-2.67	-2.67	0.00	0.00
5	SPV-1201 x SPV - 946	-18.01**	-20.21**	-3.67	-7.08	-2.71	-2.74	0.00	0.00
6	SPV-1201 x SPV - 104	-13.57**	-14.49**	-6.44	-12.10**	-1.80	-1.80	0.00	0.00
7	SPV-1201 x IS - 2284	-5.50**	-10.18**	-5.50	-8.42*	1.53	-34.15**	20.28**	-25.19**
8	SPV-1201 x IS-6335	-16.22**	-18.30**	2.29	2.29	16.23**	-23.24**	18.04**	-26.32**
9	SPV-1201 x IS - 9471	-2.13**	-9.15**	1.06	-0.10	31.08**	-6.29**	37.72**	-19.82**
10	ICSB - 101B x AKms 14 B	-23.56**	-25.25**	-20.82**	-26.96**	-0.92	-2.74	0.00	0.00
11	ICSB - 101B x SRT - 26 B	-19.62**	-20.42**	-17.38**	-23.14**	-1.40	-2.76	0.00	0.00
12	ICSB - 101B x GJ-35-15-15	-19.92**	-23.72**	-3.87	-5.13	-2.79	-4.55*	0.00	0.00
13	ICSB - 101B x SPV - 946	-24.45**	-27.26**	-6.75	-7.92	-0.03	-1.87	0.00	0.00
14	ICSB - 101B x SPV - 104	-20.62**	-22.33**	-9.90**	-13.41**	-2.79	-4.55*	0.00	0.00
15	ICSB - 101B x IS - 2284	-19.57**	-24.36**	-5.54	-10.53**	2.38*	-34.15	6.69*	-33.64**
16	ICSB - 101B x IS - 6335	-18.78**	-21.64**	-4.85	-7.08	-1.03	-35.21**	22.04**	-23.82**
17	ICSB - 101B x IS - 9471	-19.80**	-26.31**	-6.31	-7.45	12.29**	-20.60**	5.67**	-38.48**
18	AKms - 14 B x SRT - 26B	-27.43**	-28.36**	-28.67**	-29.32**	-3.22	-3.67	0.00	0.00
19	AKms - 14 B x GJ-35-15-15	-24.68**	-26.68**	-19.72**	-26.84**	-3.64	-3.67	30.62**	30.62**
20	AKms - 14 B x SPV - 946	-27.27**	-28.42**	-9.67**	-17.63**	-2.74	-2.74	0.00	0.00
21	AKms - 14 B x SPV - 104	-24.38**	-24.43**	-17.76**	-26.84**	-4.58*	-4.61	0.00	0.00
22	AKms - 14 B x IS - 2284	-22.89**	-25.91**	-8.84**	-11.36**	21.05**	-21.49**	48.88**	-7.40**
23	AKms - 14 B x IS - 6335	-11.38**	-12.60**	-14.96**	-19.78**	18.51**	-21.71**	28.34**	-19.89**
24	AKms - 14 B x IS - 9471	-25.19**	-29.81**	-5.0	-11.36**	36.84**	-2.15*	31.23**	-23.60**

Contd.

Table 17 Contd.

1	2	3	4	5	6	7	8	9	10
25	SRT - 26B x GJ-35-15-15	-24.82**	-27.69**	-11.10**	-18.30**	-0.37	-0.80	0.00	0.00
26	SRT - 26B x SPV - 946	-23.81**	-25.94**	-8.86**	-16.19**	-2.28	-2.74	41.30**	41.30**
27	SRT - 26B x SPV - 104	-15.57**	-16.56**	-16.90**	-25.48**	0.50	0.07	0.00	0.00
28	SRT - 26B x IS - 2284	-25.38**	-29.15**	-5.86	-7.64**	11.12**	-28.08**	31.26**	-18.36**
29	SRT - 26B x IS - 6335	-21.23**	-23.27**	-7.31*	-11.81**	24.02**	-18.26**	52.10**	-5.06**
30	SRT - 26B x IS - 9471	-24.91**	-30.36**	-4.06	-9.71**	38.64**	-1.15	33.03**	-22.55**
31	GJ-35-15-15 x SPV - 946	-18.43**	-19.32**	-14.14**	-14.79**	-0.84	-0.87	48.37**	48.37**
32	GJ-35-15-15 x SPV 104	-15.70**	-17.98**	-0.04	-2.69	-3.61	-3.61	26.45**	26.45**
33	GJ-35-15-15 x IS - 2284	-20.75**	-21.81**	-2.00	-8.32*	1.71	-34.03**	10.96**	-30.99**
34	GJ-35-15-15 x IS - 6335	-22.68**	-23.70**	-1.14	-4.69	-1.90*	-35.21**	6.81*	-33.33**
35	GJ-35-15-15 x IS - 9471	-25.07**	-27.86**	2.53	-0.03	10.19**	-21.23**	14.38**	-33.41**
36	SPV - 946 x SPV - 104	-10.87**	-12.34**	-15.5**	-17.34**	-3.64	-3.67	0.00	0.00
37	SPV - 946 x IS - 2284	-19.07**	-21.02**	-11.88**	-17.53**	0.23	-34.99**	28.88**	-19.84**
38	SPV - 946 x IS - 6335	-20.97**	-21.14**	-6.19	-9.51*	11.17**	-26.56**	52.07**	-5.08**
39	SPV - 946 x IS 9471	-20.38**	-24.16**	-10.54**	-12.73**	11.61**	-20.20**	25.64**	-26.85**
40	SPV - 104 x IS 2284	-5.95**	-9.69**	-12.31**	-19.99	2.63**	-33.44**	35.87**	-15.49**
41	SPV - 104 x IS 6335	-3.88**	-5.26**	18.18**	-23.13**	17.65**	-22.29**	47.85**	-7.71**
42	SPV - 104 x IS 9471	-10.24**	-15.83**	-14.08**	-18.39**	30.36**	-6.81**	32.17**	-22.88**
43	IS 2284 x IS - 6335	-15.19**	-17.40**	-12.39**	-15.09**	-33.63**	-36.10**	-14.01**	-14.79**
44	IS 2284 x IS - 9471	-7.61**	-9.88**	-9.03**	-12.81**	-16.54**	-29.43**	-2.18	-18.14**
45	IS 6335 x IS 9471	-2.77**	-7.57**	-8.53*	-9.58*	-6.23**	-19.85**	-1.08	-17.80**

SE (m)

CD at 5%

* Significant at 5%

** Significant at 1%

0.028

0.032

0.049

0.057

0.014

0.017

0.046

0.053

0.106

H1 Heterosis over mean of parents

H2 Heterosis over mean of better parents

III - 27

ble 18 Mean values of parents and F₁ crosses for different characters in 10 x 10 dialled of sorghum, Akola 1995 and Patancheru, 1996

Parents/ Crosses	100 grain weight (g)		Grain hardness kg/cm ²		Endosperm texture (%)		Ele. conductivity (mulinhoise)		DTF	
	L1	L2	Pooled	L1	L2	Pooled	L1	L2	L1	L2
SPV-1201	3.04	2.81	2.92	9.10	8.67	8.88	27.75	30.80	29.28	194.50
ICSB-101B	2.51	2.51	2.51	7.74	6.94	7.34	30.30	26.15	28.23	287.50
AKms-14B	2.03	2.57	2.30	8.00	6.95	7.48	29.70	32.80	31.25	335.00
SRT-26B	2.54	2.06	2.30	7.44	6.56	7.00	33.70	32.10	32.90	275.00
GI-35-15-15	2.01	2.14	2.07	8.44	7.04	7.74	27.60	18.00	22.80	135.00
SPV-946	2.27	2.03	2.15	8.77	7.85	8.31	24.10	19.90	22.00	125.00
SPV-104	2.60	2.90	2.75	6.60	6.08	6.34	51.60	46.90	49.20	350.00
IS-2284	1.61	1.76	1.69	6.82	5.91	6.37	58.55	56.95	57.75	185.50
IS-6335	1.64	1.69	1.67	6.97	6.68	6.83	66.95	55.85	61.40	198.00
IS-9471	2.06	1.91	1.99	5.29	6.11	5.70	61.40	55.30	58.35	97.00
SPV-1201 x ICSB-101B	2.50	2.62	2.56	8.18	7.14	7.66	30.05	32.10	31.08	257.50
SPV-1201 x AKms-14 B	2.40	2.61	2.51	6.90	5.57	6.23	40.00	38.75	39.38	295.00
SPV-1201 x SRT - 26B	2.43	3.25	2.84	6.90	7.23	7.06	33.30	34.25	38.28	262.50
SPV-1201 x GI-35-15-15	2.13	2.54	2.33	7.70	7.85	7.78	27.25	21.15	24.20	235.00
SPV-1201 x SPV - 946	2.50	1.99	2.25	6.95	6.97	6.96	35.15	33.95	34.55	247.50
SPV-1201 x SPV - 104	3.16	3.41	3.29	5.95	6.46	6.21	48.10	45.05	46.58	397.50
SPV-1201 x IS-2284	2.36	2.27	2.31	6.61	7.30	6.95	59.50	48.00	53.75	184.00
SPV-1201 x IS-6335	2.33	2.29	2.31	7.28	7.60	7.44	57.45	45.35	51.40	209.00
SPV-1201 x IS - 9471	2.59	2.71	2.65	6.54	6.08	6.31	60.10	51.05	55.58	230.00
ICSB - 101B x AKms 14 B	2.18	1.93	2.05	6.75	6.85	6.80	35.75	45.00	40.38	293.00
ICSB - 101B x SRT - 26 B	2.54	1.95	2.25	6.45	5.27	5.86	29.25	41.30	35.28	412.50
ICSB - 101B x GI-35-15-15	2.20	1.96	2.08	7.60	6.64	7.12	25.40	31.10	28.25	277.00
ICSB - 101B x SPV - 946	2.25	2.44	2.34	7.23	7.32	7.28	29.45	35.80	32.63	280.00
ICSB - 101B x SPV - 104	3.11	2.88	2.99	6.69	6.65	6.67	40.00	39.75	39.88	303.50
ICSB - 101B x IS - 2284	2.08	2.08	2.08	7.27	6.44	6.86	52.25	46.10	49.18	129.00
ICSB - 101B x IS - 6335	2.10	2.08	2.09	7.73	6.70	7.21	49.75	43.55	46.65	112.50
ICSB - 101B x IS - 9471	2.25	2.17	2.21	6.13	5.89	6.01	57.55	54.90	56.23	99.00
AKms - 14 B x SRT - 26B	2.14	2.11	2.13	6.40	3.96	5.18	51.70	54.00	52.85	313.50
AKms - 14 B x GI-35-15-15	1.88	1.80	1.84	5.57	4.64	5.10	49.10	38.25	43.68	320.00
AKms - 14 B x SPV - 946	2.55	2.50	2.52	6.98	6.32	6.65	48.15	55.35	51.75	167.50
AKms - 14 B x SPV - 104	3.10	2.88	2.99	5.59	4.89	5.24	54.95	56.30	55.63	178.00
AKms - 14 B x IS - 2284	2.12	2.38	2.25	6.09	6.19	6.18	68.10	55.75	61.93	180.00
AKms - 14 B x IS - 6335	2.30	2.42	2.36	6.91	6.73	6.82	64.25	64.55	64.40	175.00
AKms - 14 B x IS - 9471	1.96	2.25	2.11	6.16	5.35	5.76	66.55	58.80	62.68	121.50

Contd.....

Table 19. Mean values of parents and F_1 crosses for different characters in 10×10 diallel of sorghum, Akola, 1995 and 1996

No.	Parents / Crosses	Plant height (cm)					Cob length (cm)					TGM/R					Germination (%)					
		L1	L2	Pooled	5	6	L1	L2	Pooled	7	8	L1	L2	Pooled	9	10	L1	L2	Pooled	11	12	L1
1	SPV-1201	225.00	226.00	225.50	23.60	21.10	22.35	2.50	3.00	2.75	95.50	67.50	(78.26)	55.24	81.50	(66.75)						
2	ICSB-101B	155.00	164.50	159.75	26.60	26.20	26.40	3.00	3.00	3.00	74.00	47.00	(43.25)	60.50	(51.30)							
3	AKma-14B	142.50	136.00	139.25	24.60	24.10	24.35	4.00	4.00	4.00	51.00	(45.58)	29.00	(32.53)	40.00	(39.05)						
4	SRT-26B	165.00	157.00	161.00	24.90	23.10	24.00	3.00	3.00	3.00	79.00	81.50	(64.62)	80.25	(63.67)							
5	GI-35-15-15	225.00	238.50	231.75	23.90	21.60	22.75	2.50	3.00	2.75	97.00	80.16	(72.50)	84.75	(69.29)							
6	SPV-946	245.00	246.50	245.75	23.70	22.60	23.15	3.00	2.50	2.75	78.00	62.05	(53.73)	71.50	(57.89)							
7	SPV-104	200.00	218.00	209.00	21.10	21.20	21.15	4.00	3.50	3.75	69.50	56.48	(50.00)	59.75	(50.74)							
8	IS-2284	265.00	214.50	239.75	22.70	22.30	22.50	2.00	2.00	2.00	87.00	81.50	(68.87)	84.25	(66.95)							
9	IS-6335	245.00	241.50	243.25	24.80	23.20	24.00	2.00	2.00	2.00	94.00	94.00	(68.04)	90.00	(72.16)							
10	IS-9471	247.50	257.00	252.25	25.30	24.90	25.10	1.50	2.00	1.75	94.50	76.56	(76.26)	96.75	(69.80)							
11	SPV-1201 x ICSB-101B	217.00	299.50	258.25	24.40	23.70	24.05	3.00	2.75	2.88	67.00	67.50	(54.94)	85.00	(67.26)							
12	SPV-1201 x AKma-14 B	196.00	301.00	248.50	22.30	22.30	22.30	3.00	3.00	3.00	67.50	67.50	(55.26)	82.50	(52.25)							
13	SPV-1201 x SRT-26B	206.00	269.50	237.75	22.40	21.30	21.85	2.00	2.75	2.38	62.00	61.95	(65.00)	73.50	(59.79)							
14	SPV-1201 x GI-35-15-15	229.00	281.00	255.00	26.80	24.80	25.80	3.00	2.75	2.88	91.00	72.50	(64.50)	77.75	(63.02)							
15	SPV-1201 x SPV-946	201.00	233.50	217.25	27.00	25.80	26.40	2.50	2.50	2.50	90.50	90.50	(72.05)	57.00	(49.05)							
16	SPV-1201 x SPV-104	219.00	280.50	249.75	23.40	20.40	21.90	3.50	3.00	3.25	63.00	63.00	(52.54)	78.00	(62.12)							
17	SPV-1201 x IS-2284	243.00	293.00	268.00	23.00	22.90	22.95	2.00	2.00	2.00	91.00	72.56	(80.17)	94.50	(57.33)							
18	SPV-1201 x IS-6335	231.00	340.00	285.50	27.10	26.30	26.70	2.00	2.00	2.00	81.50	94.50	(64.55)	97.00	(72.36)							
19	SPV-1201 x IS-9471	194.00	343.00	268.50	28.40	27.30	27.85	2.00	2.00	2.00	81.00	94.00	(61.17)	89.25	(70.44)							
20	ICSB-101B x AKma 14 B	160.00	183.00	171.50	29.80	29.30	29.55	3.50	3.50	3.50	61.00	61.00	(51.35)	66.50	(54.70)							
21	ICSB-101B x SRT-26 B	188.50	222.00	205.25	28.50	28.90	28.70	3.00	3.00	3.00	52.00	46.15	(67.50)	59.75	(53.03)							
22	ICSB-101B x GI-35-15-15	178.00	266.00	222.00	29.70	28.50	29.10	3.00	3.00	3.00	57.00	49.02	(72.50)	64.75	(53.72)							
23	ICSB-101B x SPV-946	235.00	298.00	266.50	28.00	28.10	28.05	3.00	2.75	2.88	64.00	64.00	(53.13)	77.50	(61.75)							
24	ICSB-101B x SPV-104	150.50	217.50	184.00	23.10	24.10	23.60	4.00	3.00	3.50	68.00	68.00	(55.56)	58.50	(49.89)							
25	ICSB-101B x IS-2284	254.50	306.00	280.25	27.70	27.30	27.50	2.00	2.00	2.00	90.50	90.50	(77.14)	92.75	(74.60)							
26	ICSB-101B x IS-6335	263.50	312.00	287.75	29.90	28.50	29.20	2.00	2.00	2.00	91.00	91.00	(72.50)	91.00	(72.55)							
27	ICSB-101B x IS-9471	261.50	300.00	283.25	29.40	29.50	29.45	2.00	2.00	2.00	91.00	91.00	(72.56)	91.00	(72.56)							
28	AKma-14 B x SRT-26B	167.50	196.00	181.75	24.50	24.00	24.25	3.50	3.50	3.50	75.00	60.11	(62.50)	68.75	(56.17)							
29	AKma-14 B x GI-35-15-15	190.00	208.00	199.00	27.90	27.20	27.55	3.00	3.00	3.00	59.00	59.00	(50.18)	49.50	(44.71)							
30	AKma-14 B x SPV-946	227.50	286.00	256.75	27.00	26.80	26.90	3.50	3.50	3.50	78.00	62.05	(60.67)	54.25	(47.45)							
31	AKma-14 B x SPV-104	172.50	191.00	181.75	23.40	22.80	23.10	3.50	3.50	3.50	76.00	60.00	(54.93)	71.00	(58.49)							
32	AKma-14 B x IS-2284	220.00	279.00	249.50	26.00	24.80	25.40	2.00	2.00	2.00	78.00	62.08	(70.65)	83.50	(47.60)							
33	AKma-14 B x IS-6335	247.50	293.50	270.50	25.00	25.30	25.15	2.00	2.00	2.00	88.50	94.00	(76.28)	91.25	(66.36)							
34	AKma-14 B x IS-9471	230.00	299.00	264.50	28.00	27.80	27.90	2.00	2.00	2.00	86.00	83.00	(65.69)	84.50	(73.25)							

Contd.

Contd.

Table 19 Contd.

1	2	3	4	5	6	7	8	9	10	11	12	13	14
35	SRT - 26B x GI-35-15-15	155.00	206.50	180.75	27.00	26.70	26.85	3.50	3.00	3.25	92.00	88.50	90.25
36	SRT - 26B x SPV - 946	209.50	253.00	231.25	24.50	23.90	24.20	2.50	2.75	2.63	84.00	81.00	77.50
37	SRT - 26B x SPV - 104	177.50	213.50	195.50	22.10	22.10	22.10	3.50	3.00	3.25	91.00	88.50	85.00
38	SRT - 26B x IS - 2284	247.50	301.50	274.50	25.30	24.10	24.70	2.00	2.50	2.25	82.50	80.00	77.50
39	SRT - 26B x IS - 6335	250.00	274.50	262.25	23.50	23.40	23.45	2.00	2.50	2.25	78.00	75.00	72.50
40	SRT - 26B x IS - 9471	227.50	284.00	255.75	26.40	25.70	26.05	2.00	2.00	2.00	84.00	81.00	78.50
41	GI-35-15-15 x SPV - 946	207.50	283.50	245.50	25.70	22.80	24.25	2.50	2.00	2.25	85.00	82.00	79.50
42	GI-35-15-15 x SPV - 104	205.00	278.50	241.75	25.40	24.50	24.95	3.50	2.75	3.13	45.00	42.11	39.00
43	GI-35-15-15 x IS - 2284	277.50	325.00	301.25	22.90	23.00	22.95	2.00	2.00	2.00	82.00	80.00	78.50
44	GI-35-15-15 x IS - 6335	282.50	311.50	297.00	30.90	30.60	30.75	2.00	2.00	2.00	93.00	91.00	89.50
45	GI-35-15-15 x IS - 9471	280.00	342.00	311.00	29.60	29.60	29.60	2.00	2.00	2.00	78.00	75.00	72.50
46	SPV - 946 x SPV - 104	197.50	312.00	254.75	21.20	20.70	20.95	3.50	3.00	3.25	49.00	46.12	43.25
47	SPV - 946 x IS - 2284	242.50	250.50	246.50	23.60	23.50	23.55	2.00	2.00	2.00	92.00	90.00	88.50
48	SPV - 946 x IS - 6335	280.00	345.50	312.75	27.10	26.20	26.65	2.00	2.00	2.00	93.00	91.00	89.50
49	SPV - 946 x IS 9471	237.50	314.00	275.75	26.50	25.60	26.05	2.00	2.00	2.00	88.00	85.00	82.50
50	SPV - 104 x IS 2284	267.50	288.50	278.00	24.20	22.70	23.45	2.00	2.00	2.00	79.00	76.00	73.50
51	SPV - 104 x IS 6335	257.50	297.50	277.50	27.40	26.50	26.95	2.00	2.00	2.00	60.00	57.00	54.50
52	SPV - 104 x IS 9471	265.00	283.50	274.25	26.30	25.30	25.80	2.00	2.00	2.00	81.00	78.00	75.50
53	IS 2284 x IS-6335	262.50	280.50	271.50	29.80	25.60	27.70	2.00	2.00	2.00	88.00	85.00	82.50
54	IS 2284 x IS -9471	262.50	270.00	266.25	24.40	23.30	23.85	2.00	2.00	2.00	86.50	83.50	81.00
55	IS 6335 x IS 9471	245.00	287.50	266.25	29.60	29.40	29.50	2.00	2.00	2.00	91.00	89.00	87.50
Parental range													
		142.50	136.00	139.25	21.10	21.10	21.15	1.50	2.00	1.75	51.00	29.00	40.00
		to	to	to	to	To	to	to	to	to	to	to	To
		265.00	251.00	252.25	26.60	26.20	26.40	4.00	4.00	4.00	97.00	86.00	90.00
Hybrid range													
		150.50	183.00	171.50	21.20	20.40	20.95	2.00	2.00	2.00	45.00	49.50	54.25
		to	to	to	to	To	to	to	to	to	to	to	To
		282.50	345.50	312.75	30.90	30.60	30.75	3.50	3.50	3.50	93.00	98.00	94.50
SE (m)													
		5.113	3.708	5.147	0.690	1.634	1.273	0.263	0.207	0.470	2.170	3.606	5.092
CD at 5%													
		14.580	10.582	14.950	1.982	4.664	3.239	0.751	0.593	1.339	6.203	10.293	14.905
L ₁ = Akola 1995													
L ₂ = Akola 1996													

Table: 20 Mean infection % of parents and F₁ crosses for pathological characters in 10 x 10 diallel of sorghum, Akola, 1995 and 1996

S.C. No.	Parents / Crosses	<i>F. moniliforme</i> (%)				<i>F. pallidoreum</i> (%)				<i>C. furcata</i> (%)				Other fungi(%)			
		L1		L2		L1		L2		L1		L2		L1		L2	
		3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18
1	SPV-1201	8.89	20.67	14.78	0.71	0.71	0.71	0.71	1.46	56.00	28.73	16.67	23.33	20.00	20.00	20.00	20.00
2	ICSB-101B	31.11	26.67	28.89	0.71	0.71	0.71	0.71	43.34	58.00	25.56	15.33	20.45	20.00	20.00	20.00	20.00
3	AKms-14B	8.89	46.00	27.44	0.71	0.71	0.71	0.71	12.22	44.00	28.11	62.23	9.33	35.78	20.00	20.00	20.00
4	SRT-26B	34.45	26.01	30.23	0.71	2.36	1.54	0.71	31.11	56.00	43.56	21.11	16.00	18.56	20.00	20.00	20.00
5	GI-35-15-15	0.00	18.66	9.33	0.71	0.71	0.71	0.71	6.67	60.00	33.33	88.89	21.33	55.11	20.00	20.00	20.00
6	SPV-946	17.78	18.00	17.89	0.71	1.02	0.87	0.87	38.89	57.33	48.11	26.67	26.33	26.33	20.00	20.00	20.00
7	SPV-104	6.67	38.67	22.67	0.71	0.71	0.71	0.71	45.56	44.67	45.11	41.11	16.00	28.56	20.00	20.00	20.00
8	IS-2284	0.00	19.99	10.00	0.71	1.69	1.20	1.20	20.00	46.67	33.33	47.78	24.67	36.22	20.00	20.00	20.00
9	IS-6335	1.11	7.99	4.55	0.71	2.67	1.69	0.71	7.78	26.00	16.89	40.00	20.00	30.00	20.00	20.00	20.00
10	IS-9471	3.33	5.33	4.33	0.71	1.02	0.87	0.87	7.78	24.00	15.89	63.34	20.00	41.67	20.00	20.00	20.00
11	SPV-1201 x ICSB-101B	18.89	18.01	18.45	0.71	1.69	1.20	0.71	31.11	56.67	43.89	47.78	24.00	35.89	20.00	20.00	20.00
12	SPV-1201 x AKms-14 B	23.33	13.34	18.34	0.71	0.71	0.71	0.71	44.45	61.33	52.89	32.22	25.33	28.78	20.00	20.00	20.00
13	SPV-1201 x SRT - 26B	17.78	15.34	16.56	0.71	0.71	0.71	0.71	57.78	50.67	54.22	22.22	34.00	28.11	20.00	20.00	20.00
14	SPV-1201 x GI-35-15-15	11.11	29.33	20.22	0.71	1.69	1.20	1.20	51.11	50.67	50.89	37.78	19.33	28.56	20.00	20.00	20.00
15	SPV-1201 x SPV-946	24.45	35.99	30.22	0.71	0.71	0.71	0.71	40.00	50.00	43.00	32.22	16.00	24.11	20.00	20.00	20.00
16	SPV-1201 x SPV-104	21.11	14.67	17.89	0.71	0.71	0.71	0.71	51.11	55.33	53.22	28.89	32.67	30.78	20.00	20.00	20.00
17	SPV-1201 x IS-2284	5.56	8.01	6.78	2.22	2.35	2.29	2.29	14.45	28.00	21.22	35.56	21.33	28.45	20.00	20.00	20.00
18	SPV-1201 x IS-6335	11.11	7.34	9.23	0.71	0.71	0.71	0.71	22.22	18.00	20.11	37.78	18.67	28.22	20.00	20.00	20.00
19	SPV-1201 x IS - 9471	14.45	8.66	11.55	0.71	2.00	1.36	1.36	20.00	18.00	19.00	43.34	17.33	30.33	20.00	20.00	20.00
20	ICSB - 101B x AKms 14 B	23.33	19.33	21.33	0.71	1.69	1.20	0.71	50.00	59.33	54.67	25.56	19.33	22.45	20.00	20.00	20.00
21	ICSB - 101B x SRT-26 B	23.33	11.99	17.66	0.71	0.71	0.71	0.71	58.89	68.00	63.45	16.67	21.33	19.00	20.00	20.00	20.00
22	ICSB - 101B x GI-35-15-15	22.22	9.33	15.78	0.71	0.71	0.71	0.71	52.22	67.33	59.78	26.67	23.33	25.00	20.00	20.00	20.00
23	ICSB - 101B x SPV - 946	14.45	14.67	14.56	0.71	3.02	1.87	1.87	55.56	49.33	52.45	26.67	34.00	30.33	20.00	20.00	20.00
24	ICSB - 101B x SPV - 104	35.56	34.01	34.78	0.71	0.71	0.71	0.71	42.22	49.33	45.78	22.22	17.33	19.78	20.00	20.00	20.00
25	ICSB - 101B x IS - 2284	10.00	14.66	12.33	0.71	0.71	0.71	0.71	38.89	38.00	38.45	40.00	33.33	36.67	20.00	20.00	20.00
26	ICSB - 101B x IS - 6335	13.33	9.99	11.66	0.71	1.02	0.87	0.87	32.22	26.67	29.45	32.22	34.00	33.11	20.00	20.00	20.00
27	ICSB - 101B x IS - 9471	10.67	7.56	7.56	0.71	1.69	1.20	0.71	20.00	27.33	33.67	37.78	26.67	32.22	20.00	20.00	20.00
28	AKms - 14 B x SRT - 26B	73.34	28.67	51.01	0.71	1.69	1.20	0.71	17.78	50.67	34.22	8.89	19.33	14.11	20.00	20.00	20.00
29	AKms - 14 B x GI-35-15-15	24.45	23.33	23.89	0.71	4.00	2.36	2.36	56.67	52.67	54.67	18.89	19.33	19.11	20.00	20.00	20.00
30	AKms - 14 B x SPV - 946	16.67	17.33	17.00	0.71	0.71	0.71	0.71	54.45	59.33	56.89	30.00	23.33	26.67	20.00	20.00	20.00
31	AKms - 14 B x SPV - 104	17.78	35.33	26.55	0.71	4.67	2.69	2.69	57.78	48.00	52.89	22.22	12.00	17.11	20.00	20.00	20.00
32	AKms - 14 B x IS - 2284	23.33	15.34	19.34	0.71	6.00	3.36	3.36	38.89	47.33	43.11	23.33	27.33	25.33	20.00	20.00	20.00
33	AKms - 14 B x IS - 6335	8.89	14.01	11.45	0.71	3.33	2.02	2.02	33.33	30.67	37.00	17.78	37.33	27.56	20.00	20.00	20.00
34	AKms - 14 B x IS - 9471	11.11	16.67	13.89	0.71	2.35	1.53	1.53	35.56	39.33	37.45	32.22	22.00	27.11	20.00	20.00	20.00

Contd.....

Table 20. Contd.

	1	2	3	4	5	6	7	8	9	10	11	12	13	14
15	SRT-26B x GI-35-15-15	41.11	12.67	26.89	0.71	0.71	0.71	0.71	46.67	55.33	51.00	12.22	34.00	23.11
16	SRT-26B x SPV-946	35.56	26.01	30.78	0.71	0.71	0.71	0.71	44.45	54.00	49.22	17.78	20.67	19.22
17	SRT-26B x SPV-104	10.00	30.67	20.33	0.71	1.69	1.69	1.20	15.56	53.33	34.44	45.56	15.33	30.45
18	SRT-26B x IS-2284	5.56	11.33	8.44	0.71	1.69	1.69	1.20	18.89	46.00	32.44	31.11	34.00	32.56
19	SRT-26B x IS-6335	11.11	10.01	10.56	0.71	3.34	3.34	2.03	24.45	36.67	30.56	17.78	25.33	21.56
20	SRT-26B x IS-9471	8.89	9.34	9.11	0.71	3.02	3.02	1.87	31.11	31.33	31.22	27.78	17.33	22.56
21	GI-35-15-15 x SPV-946	15.56	20.00	17.78	0.71	1.02	1.02	0.87	31.11	56.00	43.56	46.67	24.67	35.67
22	GI-35-15-15 x SPV-104	43.34	22.67	33.00	0.71	2.67	2.67	1.69	35.56	50.00	42.78	20.00	18.67	19.33
23	GI-35-15-15 x IS-2284	7.78	9.34	8.56	0.71	4.67	4.67	2.69	26.67	38.67	32.67	43.34	18.67	31.00
24	GI-35-15-15 x IS-6335	3.33	13.99	8.66	0.71	4.67	4.67	2.69	15.56	32.00	23.78	24.45	22.00	23.22
25	GI-35-15-15 x IS-9471	8.89	7.99	8.44	0.71	1.33	1.33	1.02	30.00	32.00	31.00	30.00	22.67	26.33
26	SPV-946 x SPV-104	21.11	11.33	16.22	0.71	1.02	1.02	0.87	57.78	58.00	57.89	21.11	32.00	26.56
27	SPV-946 x IS-2284	6.67	15.34	11.00	0.71	4.67	4.67	2.69	24.45	30.67	27.56	28.89	23.33	26.11
28	SPV-946 x IS-6335	6.67	11.99	9.33	0.71	2.35	2.35	1.53	14.45	29.33	21.89	35.56	26.00	30.78
29	SPV-946 x IS-9471	11.11	11.99	11.55	0.71	3.33	3.33	2.02	13.33	30.67	22.00	34.45	23.33	28.89
30	SPV-104 x IS-2284	14.45	15.99	15.22	0.71	7.33	7.33	4.02	35.56	34.67	35.11	43.34	27.33	35.33
31	SPV-104 x IS-6335	37.78	5.99	21.89	0.71	1.69	1.69	1.20	35.56	36.00	35.78	25.56	29.33	27.45
32	SPV-104 x IS-9471	10.00	7.99	9.00	0.71	2.35	2.35	1.53	17.78	38.67	28.22	37.78	22.67	30.22
33	IS-2284 x IS-6335	8.89	6.01	7.45	0.71	3.02	3.02	1.87	22.22	31.33	26.78	18.89	24.67	21.78
34	IS-2284 x IS-9471	6.67	12.01	9.34	0.71	8.00	8.00	4.36	25.56	30.00	27.78	27.78	26.00	26.89
35	IS-6335 x IS-9471	3.34	6.01	4.67	0.71	2.35	2.35	1.53	26.67	23.33	25.00	32.22	17.33	24.78
Parental range														
	0.71	5.33	4.51	0.71	0.71	0.71	0.71	0.71	1.46	24.00	15.89	16.67	9.33	18.55
	to	to	to	to	to	to	to	to	to	to	to	to	to	to
	34.45	46.00	30.23	0.71	2.36	1.69	45.55	60.00	50.66	88.89	26.00	35.11		
Hybrid range														
	3.34	18.50	4.67	0.71	0.71	0.71	0.71	0.71	13.34	18.00	19.00	8.88	12.00	14.11
	to	to	to	to	to	to	to	to	to	to	to	to	to	to
	73.34	52.52	51.08	0.71	8.00	4.35	58.90	68.00	63.45	46.67	37.34	36.66		
SE (m)	1.103	1.203	1.644	0.148	0.163	0.220	1.430	1.630	2.132	1.111	1.213	1.657		
CD at 5%	2.185	2.383	3.256	0.293	0.322	0.435	2.832	3.229	4.223	2.200	2.402	3.282		
L ₁ = Akola, 1995 L ₂ = Akola, 1996														

Table 21 Analysis of variance for combining ability of F_1 crosses in 10×10 diallel of sorghum (Pooled) Akola and Patancheru, 1996

Sr. No.	Characters	Sources and degrees of freedom							
		GCA (9)	SCA (45)	Envior- ments (1)	GCA x Env. (9)	SCA x Env. (45)	Error (108)	δ^2 gca δ^2 sca	δ^2 gca/ δ^2 sca
1	100 grain weight	31.84**	2.33*	170.27**	2.63**	1.76**	0.04	0.032	0.016
2	Grain hardness	19.16**	1.89*	302.43**	2.88**	1.36	0.30	0.227	0.134
3	Endosperm texture	60.26**	3.86**	11.71**	1.87	1.80**	18.55	45.807	26.496
4	Electrical conductivity	53.24**	6.98**	419.70**	21.92**	3.53**	1055.92	2298.515	3156.782
5	DTF	104.18**	9.83**	122.67**	14.92**	3.95**	2.60	11.141	11.439
6	Plant height	120.33**	23.45**	0.26	6.22**	1.82**	103.22	513.178	1178.673
7	Cob length	16.73**	2.04*	17.43**	1.22	0.80	2.66	1.746	1.390
8	Glume covering	15.11**	1.91	19.99**	1.17	1.31	44.40	26.121	20.202
9	Mesocarp thickness	142.02**	37.08**	0.67	1.61	1.65*	12.73	74.811	229.710
10	TGMR	131.78**	6.55**	192.86**	32.92**	3.33**	0.06	0.333	0.170
11	Germination	89.27**	12.40**	726.47**	6.30**	4.33**	11.43	42.075	65.226
12	<i>F. moniliforme</i>	18.69**	1.94**	697.64**	2.45*	1.49*	38.39	28.293	17.981
13	<i>F. pallidorozeum</i>	1.48	1.92**	603.45**	2.02*	1.64*	5.09	0.101	2.327
14	<i>C. lunata</i>	14.56**	1.28	333.63**	10.13**	1.22	34.38	19.425	4.774
15	Other fungi	1.79	1.81**	426.93**	1.71	1.37	18.24	0.603	7.379

Figures in parenthesis indicates degree of freedom

* Significant at 5%

** Significant at 1%

Table 22 Analysis of variance for combining ability of F₂ progenies in 10 × 10 diallel of sorghum (Pooled) Akola and Patancheru, 1996

Sr No	Characters	Sources of degrees of freedom							δ^2 gca/ δ^2 sca
		GCA (9)	SCA (45)	Environ ments (1)	GCA × Env. (9)	SCA × Env. (45)	Error (108)	δ^2 gca	
1	Grain hardness	13.61**	3.78**	341.10**	7.07**	2.76**	0.25	0.131	0.349
2	Ele. conductivity	157.86**	8.61**	269.45**	11.43**	9.43**	721.54	4710.639	2744.384
3	Plant height	27.00**	2.77**	308.94**	5.49**	2.68**	229.47	248.582	202.530
4	Cob length	22.97**	1.93**	0.30	3.74	1.12	1.89	1.730	0.880
5	Glume covering	8.62**	2.32**	0.11	1.09	0.83	20.52	6.517	13.580
6	TGMR	41.70**	3.86**	140.15**	7.02**	1.42	0.07	0.117	0.099
7	Germination	26.83**	3.79**	215.10**	4.34**	2.21**	23.11	24.882	32.248
8	<i>F. moniliforme</i>	4.07**	1.35	130.25**	1.33	0.94	60.88	7.777	10.732
9	<i>F. pallidorozeum</i>	0.88	1.22	101.49**	1.02	1.49*	3.84	0.019	0.420
10	<i>C. lunata</i>	10.45**	1.02	116.14**	3.22**	1.12	47.21	18.588	0.373
11	Other fungi	2.15*	0.95	70.46**	0.71	1.69*	14.88	0.714	0.387

Figures in parenthesis indicate degrees of freedom

* Significant at 5%

** Significant at 1%

IV - 3
 Table 23 Analysis of variance for combining ability of F_1 crosses in 10×10 diallel of sorghum for biochemical characters Akola, 1995

Sr.No	Characters	Sources and degree of freedom					
		GCA (9)	SCA (44)	Error (54)	δ^2 gca	δ^2 sca	δ^2 gca / δ^2 sca
1	Protein	248.02**	204.51**	0.0005	0.0111	0.105	0.101
2	Soluble sugars	20.63**	6.23**	0.0016	0.003	0.008	0.313
3	Tannins	10938.26**	286.57**	0.0001	0.127	0.040	3.192
4	Flavan-4-ols	4668.30**	81.69**	0.0014	0.546	0.113	4.821

Figures in parenthesis indicate degree of freedom.

* Significant at 5%

** Significant at 1%

Table : 24 Estimates of general combining ability effects for parents from F₁ crosses (Pooled) Akola and Patancheru, 1996

Parents	100 grain weight	Grain hardness	Endosperm texture	Electrical conductivity	DTF	Plant height	Cob length	Gilane covering	Merocarp thickness	TGMR	Germination	F.m.	F.p.	C. Insect	Other fungi
SPV-1201	0.045	-0.151	-3.104**	39.208**	-8.375**	-68.667**	-0.466	3.220**	-2.605**	0.483**	-12.270**	7.593**	0.156	3.893**	-6.727**
ICSB-101B	-0.090**	0.051	-4.223**	81.792**	1.292**	-66.042**	-2.207**	10.511**	14.871**	0.420**	-17.589**	4.790**	-0.289	5.672**	-3.792**
AKMS-14B	-0.149**	0.784**	-17.316**	91.792**	2.875**	-57.167**	-2.191**	1.136	-3.499**	0.389**	-13.773**	10.955**	-3.555**	-1.328	-6.336**
SRT-26B	-0.282**	0.677**	-7.858**	112.000**	2.083**	-33.458**	-0.686*	0.095	-6.767**	0.004	-8.869**	2.238	-0.487	6.949**	-4.896**
QJ-35-15-15	0.132**	0.429**	-6.212**	44.042**	2.708**	-21.042**	-2.578**	-5.114**	4.185**	0.129**	-6.234**	-0.934	-1.412**	5.783**	-1.762*
SPV-946	-0.130**	-0.298**	-4.516**	65.792**	1.250**	-19.750**	-1.466**	-4.072**	-3.475**	-0.038	-3.858**	-0.708	-0.985*	3.727**	-1.897**
SPV-104	-0.132**	0.122	-8.400**	184.667**	5.917**	-36.875**	-2.349**	1.136	19.099**	0.316**	-10.361**	9.238**	-2.195**	-2.941**	-3.475**
IS-2284	-0.138**	0.020	-5.033**	67.792**	2.958**	-59.167**	-1.607**	8.428**	29.971**	0.587**	-13.293**	11.647**	1.164**	6.114**	-2.511**
IS-4335	-0.113**	0.464**	-6.225**	98.958**	3.333**	-55.208**	-2.641**	-5.114**	37.614**	0.795**	-13.705**	11.174**	0.431	4.171**	-3.777**
IS-M71	-0.266**	0.843**	-5.908**	85.208**	4.708**	-39.625**	-2.774**	7.386**	26.037**	0.629**	-13.676**	5.846**	-1.002**	4.895**	-0.066
SE (G)	0.030	0.106	0.834	6.293	0.312	1.967	0.316	1.290	0.691	0.048	0.655	1.200	0.437	1.135	0.927
SE(G-G)	0.045	0.158	1.243	9.380	0.465	2.933	0.471	1.924	1.030	0.071	0.976	1.789	0.651	1.693	1.232
CD # (G) 5% 0.060		0.210	1.652	12.466	0.618	3.897	0.626	2.556	1.369	0.095	1.297	2.377	0.865	2.249	1.638

* Significant at 5%

F.m. = *F. moniliforme*
F.p. = *F. pallidoroeseum*

** Significant at 1%

Table 25 Estimates of general combining ability effects for parents from F₂ progenies (Pooled), Akola and Patancheru, 1996

S. N	Parents	Grain hardness	Ele. Conductivity	Plant height	Cob length	Glume covering	TGMR	Germination	F. moniliforme	F. pallidoroseum	C. lunata	Other fungi
1	SPV-1201	1.301**	-101.451**	16.737**	0.072	5.576**	0.560**	-11.000**	-3.558*	2.216**	3.804**	-2.463**
2	ICSB-101B	0.610**	-32.617**	-15.434**	-1.458**	4.451**	0.714**	-14.845**	-1.614	2.518**	2.805*	-5.706**
3	AKms-14B	0.470**	19.049**	-34.722**	-1.349**	2.742**	1.014**	-16.211**	-2.364	1.372**	4.250**	-3.168**
4	SRT-26B	1.568**	-25.284**	-41.734**	-0.349	1.742*	0.619**	-7.298**	-1.316	2.555**	-1.061	-3.696**
5	GI-35-15-15	0.983**	-121.617**	11.808**	-3.499**	-0.258	0.777**	-9.063**	-7.828**	1.346**	4.102**	1.524*
6	SPV-946	0.534**	-78.284**	9.420**	-1.070**	-4.966**	0.648**	-11.000**	-2.989*	3.205**	1.975	-0.366
7	SPV-104	0.852**	-21.992**	2.028	-1.253**	-0.674	0.752**	-7.520**	-10.343**	0.344	4.200**	1.357
8	IS-2284	0.789**	-59.117**	-14.005**	-1.558**	-0.716	0.127**	-5.933**	-13.818**	-0.493	5.672**	-0.754
9	IS-6335	1.295**	-1.867	-1.551	-2.141**	11.701**	0.244**	-10.174**	-14.268**	-0.329	-0.508	-1.619*
10	IS-9471	1.388**	70.841**	12.312**	-1.453**	0.492	0.073	-6.538**	-15.954**	0.523	3.504**	0.760
	SE (G)	0.097	5.202	2.933	0.266	0.877	0.051	0.931	1.511	0.380	1.331	0.747
	SE (G-G)	0.145	7.754	4.373	0.397	1.308	0.076	1.388	2.252	0.566	1.983	1.114
	CD (G) at 5%	0.192	10.304	5.811	0.527	1.738	0.101	1.845	2.993	0.740	2.263	1.479

* Significant at 5%

** Significant at 1%

V-3

Table 26 Estimates of general combining ability effects of parents from F₁ crosses in 10 × 10 diallel of sorghum, biochemical character Akola, 1995

Sr.No.	Parents	Proteins	Soluble sugars	Tannins	Flavan-4-ols
1	SPV - 1201	0.372**	0.027*	-0.055**	-0.084**
2	ICSB-101B	0.637**	0.109**	0.018**	0.078**
3	AKms-14B	0.801**	0.280**	-0.120**	-0.210**
4	SRT - 26B	0.787**	0.197**	-0.124**	-0.247
5	GJ-35-15-15	0.619**	0.024*	0.033**	-0.050**
6	SPV-946	0.616**	0.112**	-0.002	-0.266**
7	SPV-104	0.274**	0.167**	-0.059**	-0.229**
8	IS - 2284	0.365**	0.092**	0.815**	0.469**
9	IS - 6335	0.334**	0.083**	0.796**	0.578**
10	IS - 9471	0.326**	0.037**	0.590**	1.858**
SE (Gi)		0.006	0.011	0.003	0.010
SE (Gi-Gj)		0.009	0.016	0.005	0.015
CD (Gi) 5%		0.012	0.022	0.006	0.021

* Significant at 5%

** Significant at 1%

VI - 1
 Table : 27 Estimates of specific combining ability effects for F_1 crosses in 10×10 diallel (Pooled) Akola and Patancheru, 1996

Sr. No.	Crosses	100 grain weight	Grain hardness	Endosperm texture	Electrical conductivity	DTF	Plant height	Cob length	Glume covering
		3	4	5	6	7	8	9	10
1	SPV-1201 x ICSB - 10B	0.125	0.238	-1.631	-17.375	1.458	41.521**	-2.161*	-5.634
2	SPV-1201 x AKms - 14 B	-0.132	-0.481	-3.935	11.125	4.125**	42.958**	-1.878	-4.072
3	SPV-1201 x SRT - 26B	0.336**	0.078	3.707	-28.771	4.104**	16.813**	-1.551	-4.593
4	SPV-1201 x GI-35-15-15	-0.024	0.948**	-6.283*	-32.750	0.292	10.021	1.428	-0.947
5	SPV-1201 x SPV - 946	-0.326**	0.274	0.715	21.500	-1.938	-54.208**	1.084	-6.676
6	SPV-1201 x SPV - 104	0.207*	0.435	1.511	-27.938	1.896	19.721**	-0.907	5.303
7	SPV-1201 x IS - 2284	-0.155	-0.413	5.832*	-9.625	-0.583	-0.042	0.989	-0.426
8	SPV-1201 x IS-6335	-0.176	-0.245	0.461	1.833	5.229**	26.188**	1.947	5.303
9	SPV-1201 x IS - 9471	0.054	-0.534	5.832*	3.583	2.167*	34.354**	2.280*	5.303
10	ICSB - 101B x AKms 14 B	-0.227*	0.163	4.038	0.542	0.708	-30.359**	1.926	-6.676
11	ICSB - 101B x SRT - 26 B	-0.094	-0.166	-2.935	-20.083	-0.375	4.500	2.103*	11.553**
12	ICSB - 101B x GI-35-15-15	0.009	0.460	3.063	2.667	1.646	37.479**	-0.411	-3.551
13	ICSB - 101B x SPV - 946	0.227*	0.016	-4.191	-21.771	0.229	-22.583**	-0.353	-3.030
14	ICSB - 101B x SPV - 104	0.198	-0.167	3.730	-25.708	-0.500	38.396**	0.643	-3.030
15	ICSB - 101B x IS - 2284	0.111	0.399	-0.041	-36.750	-4.188**	35.875**	0.701	2.699
16	ICSB - 101B x IS - 6335	0.054	-0.063	6.380**	-27.750	-0.500	28.292**	1.384	-3.551
17	ICSB - 101B x IS - 9471	-0.053	-1.420**	5.650	12.146	0.604	4.688	0.137	0.616
18	AKms - 14 B x SRT - 26B	-0.161	-1.345**	0.936	-12.083	-1.708	-13.354*	2.316*	10.511**
19	AKms - 14 B x GI-35-15-15	0.179	-0.056	9.484**	-11.583	-3.188**	36.667**	1.422	11.032**
20	AKms - 14 B x SPV - 946	0.065	-0.117	2.430	2.729	0.396	-15.896**	-0.220	-1.989
21	AKms - 14 B x SPV - 104	0.143	0.760*	5.000	-50.458**	0.167	25.333**	0.176	-1.718
22	AKms - 14 B x IS - 2284	0.296**	0.523	7.480**	-68.000**	-3.021**	24.313**	-0.416	-1.989
23	AKms - 14 B x IS - 6335	0.189	0.126	3.550	-68.000**	-3.833**	39.979**	0.918	-1.989
24	AKms - 14 B x IS - 9471								

Contd....

VI - 2

Table 27 Contd.

1	2	3	4	5	6	7	8	9	10
25	SRT - 26B x GJ-35-15-15	-0.033	0.382	0.002	29.771	-0.979	-34.000*	0.043	-2.509
26	SRT - 26B x SPV - 946	0.135	-0.124	4.100	-26.229	0.542	6.021	0.749	-1.989
27	SRT - 26B x SPV - 104	-0.025	0.652	0.546	-66.917**	-1.875	-4.792	0.257	-2.509
28	SRT - 26B x IS - 2284	0.149	0.144	-6.830	-50.604**	-3.854**	49.688**	1.303	-1.989
29	SRT - 26B x IS - 6335	0.192	-0.096	6.746**	-61.146**	-4.542**	9.167	-1.363	-2.509
30	SRT - 26B x IS - 9471	0.087	-0.268	-4.408	-14.896	2.896**	14.833*	-0.305	3.741
31	GJ-35-15-15 x SPV - 946	0.179	-0.112	0.186	16.792	1.229	-21.521**	-2.972**	1.657
32	GJ-35-15-15 x SPV 104	-0.122	0.459	0.582	-15.896	-0.688	26.667**	0.037	-5.114
33	GJ-35-15-15 x IS - 2284	-0.019	0.121	6.702**	-19.333	-1.667	30.896**	2.032	7.907
34	GJ-35-15-15 x IS - 6335	0.009	-0.843*	7.732**	-39.375	-2.354*	29.375**	1.041	1.936
35	GJ-35-15-15 x IS - 9471	0.002	-0.303	5.502	4.875	0.833	9.042	0.649	1.136
36	SPV - 946 x SPV - 104	-0.007	0.080	-3.045	5.854	0.833	35.688**	1.843	7.907
37	SPV - 946 x IS - 2284	0.191	-0.118	-3.150	-54.583**	-4.646**	-12.083	0.389	-10.322
38	SPV - 946 x IS - 6335	0.068	0.140	-4.070	-30.125	7.667**	11.396	0.197	1.677
39	SPV - 946 x IS 9471	-0.031	0.051	1.750	-55.875**	-4.646**	6.063	0.630	7.907
40	SPV - 104 x IS 2284	-0.121	-0.726*	4.521	-48.521**	-0.063	23.854**	-0.003	1.657
41	SPV - 104 x IS 6335	-0.140	-0.588	9.100**	-76.813**	-4.750**	14.333*	2.455**	7.356
42	SPV - 104 x IS 9471	0.180	-0.455	5.346	-120.063**	-7.813**	-3.250	1.589	-5.114
43	IS 2284 x IS - 6335	-0.052	-0.189	-7.354**	64.000**	1.521	-13.938*	0.151	7.907
44	IS 2284 x IS - 9471	-0.057	-0.179	-4.533	59.250**	3.708**	-23.771**	-2.166*	-10.843**
45	IS 6335 x IS 9471	0.054	-0.408	-7.604**	48.458**	-2.229*	-26.292**	-0.568	-11.364**
SE (Sij)									
SE (Sij - Sit)									
SE (Sij - Skm)									
CD (Sij) at 5%									
SE (Sij)									
SE (Sij - Sit)									
SE (Sij - Skm)									
CD (Sij) at 5%									

* Significant at 5%

** Significant at 1%

VI - 3

Table : 28 Estimates of specific combining ability effects for F_1 crosses in 10×10 diallel of sorghum (Pooled), Akola and Patancheru, 1996 (Mesocarp thickness, TGMR, Germination, F_1 moniliforme, F_1 pallidoroeseum, C , $lunata$, and other fungi)

SN	Crosses	Meso thickness	TGMR	Germination	F_1 moniliforme	F_1 pallidoroeseum	C lunata	Other fungi
1	2	3	4	5	6	7	8	9
1	SPV-1201 x ICSB - 101B	-17.545**	-0.298	5.235**	3.194	0.257	3.616	-2.013
2	SPV-1201 x AKms - 14 B	19.318**	-0.069	-1.442	-1.388	1.802	0.452	1.457
3	SPV-1201 x SRT - 26B	8.660**	-0.257	8.279**	-7.585	-1.077	-0.245	12.679**
4	SPV-1201 x GI-35-15-15	15.930**	-0.319*	-0.720	1.332	-0.472	-1.663	3.334
5	SPV-1201 x SPV - 946	0.231	0.098	-0.956	3.437	-1.823	-1.690	1.511
6	SPV-1201 x SPV - 104	10.385**	0.275	0.100	-6.584	-2.852	8.312*	5.637*
7	SPV-1201 x IS - 2284	-4.847*	-0.215	7.330**	-3.365	0.739	-5.329	-3.209
8	SPV-1201 x IS-6335	-11.506**	-0.111	2.629	0.390	2.137	-4.299	-3.288
9	SPV-1201 x IS - 9471	-15.447**	-0.069	4.135	-4.617	0.977	-6.940	-2.653
10	ICSB - 101B x AKms 14 B	3.139	0.525**	3.164	-0.460	0.901	-0.828	-0.319
11	ICSB - 101B x SRT - 26 B	1.328	0.462**	-6.025**	3.676	2.869	2.478	-2.768
12	ICSB - 101B x GI-35-15-15	17.438**	-0.100	2.476	-3.402	0.649	6.727	0.233
13	ICSB - 101B x SPV - 946	12.132**	-0.434**	6.534**	-1.625	1.941	-1.967	3.067
14	ICSB - 101B x SPV - 104	7.691**	-0.132	-3.614	12.017**	3.579**	-7.300	-4.122
15	ICSB - 101B x IS - 2284	-22.031**	-0.371*	9.225**	-5.284	-5.140**	-3.941	5.015
16	ICSB - 101B x IS - 6335	-22.195**	-0.267	7.657**	-10.017**	-2.761	-6.579	5.263
17	ICSB - 101B x IS - 9471	-9.699**	-0.225	10.526**	-7.678	-1.899	-3.549	3.238
18	AKms - 14 B x SRT - 26B	17.451**	0.316	-1.373	2.762	-0.608	-2.357	-0.295
19	AKms - 14 B x GI-35-15-15	-3.252	0.254	-1.344	-0.657	1.659	-1.105	-0.641
20	AKms - 14 B x SPV - 946	-0.971	0.545**	1.512	-7.875	-0.512	5.864	0.872
21	AKms - 14 B x SPV - 104	-10.666**	0.223	-7.261**	1.424	4.104**	-3.467	-2.672
22	AKms - 14 B x IS - 2284	-8.239**	-0.767**	6.336**	-5.861	1.205	4.562	4.155
23	AKms - 14 B x IS - 6335	-7.933**	-0.913**	15.185**	-9.259**	-0.907	-3.078	7.738**
24	AKms - 14 B x IS - 9471	-1.849	-0.871**	12.769**	-0.596	-0.535	1.614	2.376

Contd.

Table 28 Contd.

1	2	3	4	5	6	7	8	9
25	SRT - 26B x GI-35-15-15	-14.070**	0.441**	4.010	3.489	1.138	-5.802	4.760
26	SRT - 26B x SPV - 946	16.269**	-0.142	0.740	5.596	0.941	-2.493	-3.396
27	SRT - 26B x SPV - 104	7.131**	0.160	0.192	0.233	1.405	-2.841	-3.118
28	SRT - 26B x IS - 2284	-11.786**	-0.330*	2.866	-3.398	-2.162	-0.132	4.042
29	SRT - 26B x IS - 6335	-1.718	-0.225	4.188	-4.458	-1.118	-2.108	0.625
30	SRT - 26B x IS - 9471	-9.731**	-0.434**	4.912	-4.792	-0.413	-6.078	-2.737
31	GI-35-15-15 x SPV - 946	7.026**	0.420**	-3.291	1.180	-3.121*	2.086	0.082
32	GI-35-15-15 x SPV 104	6.343**	-0.027	0.666	-0.180	1.005	-1.123	-0.465
33	GI-35-15-15 x IS - 2284	-12.857**	-0.392**	3.068	-4.803	-1.224	-6.548	-4.303
34	GI-35-15-15 x IS - 6335	-14.928**	-0.288	7.987**	2.789	1.332	-2.188	-0.387
35	GI-35-15-15 x IS - 9471	-10.001**	-0.246	-0.384	2.120	1.859	-2.161	0.919
36	SPV - 946 x SPV - 104	-4.551	0.139	1.779	-3.403	-4.014**	3.727	3.559
37	SPV - 946 x IS - 2284	-4.893*	-0.100	1.078	0.973	1.915	-7.244	-1.791
38	SPV - 946 x IS - 6335	-5.755**	-0.246	4.207	3.233	4.313**	-2.215	-0.876
39	SPV - 946 x IS 9471	-12.538**	-0.205	-3.888	-0.101	2.330	-3.523	0.765
40	SPV - 104 x IS 2284	-26.604**	-0.173	9.925**	-8.392*	1.883	-1.245	2.664
41	SPV - 104 x IS 6335	-21.485**	-0.569**	8.824**	-9.795**	-0.381	3.782	4.248
42	SPV - 104 x IS 9471	-6.449**	-0.527**	10.111**	-3.794	-0.521	1.144	1.218
43	IS 2284 x IS - 6335	13.972**	0.441**	10.839**	1.919	-1.615	3.144	-2.263
44	IS 2284 x IS - 9471	17.312**	0.733**	-2.402	4.918	2.070	4.506	0.713
45	IS 6335 x IS 9471	-3.680	0.587**	-8.426**	2.850	-1.864	5.198	-3.707
SE (Sij)		2.324	0.161	2.203	0.436	1.459	3.819	2.782
SE (Sij - Sk)		3.416	0.237	3.238	5.932	2.159	5.614	4.089
SE (Sij - Skm)		3.257	0.226	3.088	5.656	2.059	5.352	3.898
CD (Sij) at 5%		4.604	0.319	4.364	7.994	2.910	7.566	5.510

* Significant at 5%

** Significant at 1%

Table: 29 Estimate of specific combining ability effects for F_2 progenies in 10×10 diallel (Pooled) Akola and Patancheru, 1996

Sr. No	Crosses	Grain hardness	Ele. conductivity	Plant height	Cob length	Glume covering	TGMR	Germination	F_m	F_p	C. <i>lunata</i>	Other lungi
		3	4	5	6	7	8	9	10	11	12	13
1	SPV-1201 x ICSB - 101B	-0.179	20.591**	-19.474	-0.643	0.638	-0.388**	5.635	0.119	-1.735	-4.738	1.994
2	SPV-1201 x AKms - 14 B	-0.107	23.424	23.583**	-1.214	4.159	-0.013	3.117	1.524	-0.345	-8.129	0.940
3	SPV-1201 x SRT - 26B	0.422	40.258**	-9.499	-0.114	1.159	-0.261	8.270**	4.273	0.110	-5.915	2.395
4	SPV-1201 x GJ-35-15-15	0.137	190.341**	20.222*	-0.189	-1.091	0.019	2.276	-0.575	0.954	1.925	-2.200
5	SPV-1201 x SPV - 946	-1.149**	-31.117	-5.547	-0.224	-5.945*	0.079	1.442	2.519	-0.162	-3.878	-1.983
6	SPV-1201 x SPV - 104	-0.691*	-7.847	-17.792	0.259	-6.299*	-0.219	6.478*	0.652	-2.149	-2.200	1.508
7	SPV-1201 x IS - 2284	-0.583	45.591**	-9.109	0.857	1.305	-0.081	0.372	-5.028	-1.684	4.828	3.115
8	SPV-1201 x IS-6335	-0.133	-49.909**	-5.582	1.690	-5.112	-0.148	-0.418	4.286	0.945	3.012	-1.102
9	SPV-1201 x IS - 9471	-0.318	-28.430	-0.716	-0.566	0.034	-0.109	-5.071	-0.656	-0.366	7.488	0.158
10	ICSB - 101B x AKms 14 B	-0.370	0.716	5.803	1.872*	0.472	-0.286	7.184**	-4.784	-0.606	5.159	-0.303
11	ICSB - 101B x SRT - 26 B	-0.269	20.799	11.041	0.772	-0.028	-0.259	2.565	1.785	-0.277	-2.887	3.423
12	ICSB - 101B x GJ-35-15-15	-0.291	9.133	3.667	0.222	-1.028	-0.304	5.941	-3.723	-1.493	1.434	3.888
13	ICSB - 101B x SPV - 946	-0.374	9.424	2.993	0.986	-3.383	-0.244	0.260	4.061	-0.433	-2.510	-1.146
14	ICSB - 101B x SPV - 104	-0.009	-20.305	-5.203	0.570	-1.237	-0.367*	-0.389	-0.246	2.675*	-4.862	0.135
15	ICSB - 101B x IS - 2284	-0.425	10.133	27.655**	0.767	-4.133	-0.004	1.152	6.264	-0.894	-1.804	1.312
16	ICSB - 101B x IS - 6335	0.254	13.633	4.333	-0.599	-3.549	0.154	2.649	-0.642	-2.567*	4.930	1.305
17	ICSB - 101B x IS - 9471	-0.426	2.112	11.689	-1.030	3.347	0.269	4.892	0.396	0.295	-0.334	0.705
18	AKms - 14 B x SRT - 26 B	0.424	-4.617	16.022	1.201	2.242	-0.059	-5.558	1.190	0.563	0.062	-3.482
19	AKms - 14 B x GJ-35-15-15	-0.209	-27.784	-7.332	0.751	1.242	-0.329	8.175**	-1.742	-1.635	-2.817	-2.701
20	AKms - 14 B x SPV - 946	-0.102	32.008	3.974	-0.060	3.138	-0.119	-1.693	-1.144	0.450	7.979	-2.701
21	AKms - 14 B x SPV - 104	-0.327	20.028	1.903	0.449	1.034	-0.217	0.660	7.759	-0.337	-9.273*	0.171
22	AKms - 14 B x IS - 2284	-0.426	-5.784	10.362	-0.078	-3.362	-0.279	4.647	2.689	0.370	-6.095	2.138
23	AKms - 14 B x IS - 6335	0.149	-76.284**	17.789	-0.270	-9.778	-0.346*	9.286**	-0.237	0.604	-1.791	4.770
24	AKms - 14 B x IS - 9471	0.029	0.195	9.045	0.049	-4.633	-0.381*	6.604*	-4.009	-1.807	1.745	7.320**

Contd.

Table 29 Condt.

1	2	3	4	5	6	7	8	9	10	11	12	13
25	SRT - 26B x GJ-35-15-15	-0.335	-50.451**	-31.463**	-0.124	0.748	-0.327	3.263	4.211	0.162	-10.513*	2.169
26	SRT - 26B x SPV - 946	-0.631	50.091**	-0.082	-0.910	-1.612	-0.292	3.800	-6.425	-0.931	5.103	2.215
27	SRT - 26B x SPV - 104	-0.450	-1.138	20.472*	0.299	2.534	-0.240	0.998	-6.582	-2.190	5.121	0.797
28	SRT - 26B x IS - 2284	-0.617	54.299**	24.505**	-2.628**	-3.862	0.223	-4.681	-5.342	-0.586	1.819	-0.346
29	SRT - 26B x IS - 6335	-1.112**	8.049	29.108**	0.880	-5.778	-0.194	5.327	3.042	-1.070	6.433	0.676
30	SRT - 26B x IS - 9471	-0.567*	-66.722**	23.364**	1.324	1.117	0.171	0.712	6.440	-0.891	2.899	-0.654
31	GJ-35-15-15 x SPV - 946	-0.268	112.424**	-11.086	0.765	-1.362	-0.138	1.301	-0.923	-2.512	-3.296	1.870
32	GJ-35-15-15 x SPV 104	0.135	64.195**	-2.507	0.424	-0.466	-0.011	-3.315	3.680	-0.414	3.581	-2.358
33	GJ-35-15-15 x IS - 2284	0.309	-45.867**	14.101	3.722**	8.388**	-0.248	5.247	1.710	2.391	-0.320	-1.941
34	GJ-35-15-15 x IS - 6335	-0.657*	-5.367	-9.997	0.105	-4.778	-0.090	-1.102	5.834	-0.403	-1.757	-1.259
35	GJ-35-15-15 x IS - 9471	-0.789	-3.388	0.810	1.224	-1.133	-0.125	-3.657	3.702	0.256	-1.451	-0.519
36	SPV - 946 x SPV - 104	0.047	8.987	7.974	-1.037	0.430	-0.300	7.053*	0.084	-0.809	-3.612	-2.322
37	SPV - 946 x IS - 2284	1.098**	29.674	-21.617*	0.036	4.284	-0.138	0.464	6.894	-0.105	0.886	1.875
38	SPV - 946 x IS - 6335	0.059	-16.576	50.985**	2.645**	11.617**	0.021	8.017**	-4.912	-1.608	-2.310	4.357
39	SPV - 946 x IS 9471	0.252	-38.347*	-11.434	-0.062	2.763	-0.165	1.357	5.826	-0.299	-2.264	-2.333
40	SPV - 104 x IS 2284	-0.535	-24.055	16.137	0.095	10.180**	0.139	3.796	-0.753	-0.482	4.194	-1.753
41	SPV - 104 x IS 6335	-0.058	51.695**	-10.386	0.853	-1.737	-0.177	-1.337	5.311	1.475	2.528	-0.001
42	SPV - 104 x IS 9471	0.165	-46.576**	-14.655	0.597	-3.091	-0.113	1.296	10.779*	1.544	-3.877	0.409
43	IS 2284 x IS - 6335	-0.609	46.633**	-19.453	-1.024	-8.133**	0.060	-4.031	13.791**	2.519	-6.984	-2.617
44	IS 2284 x IS - 9471	-0.639	7.612	-14.472	1.370	-3.237	0.075	4.944	7.409	-0.542	-7.868	-1.174
45	IS 6335 x IS 9471	-0.482	31.862	-18.695	0.003	3.847	0.233	1.999	2.023	0.565	-3.045	-2.392
SE (Sij)		0.326	17.496	9.867	0.895	2.951	0.171	3.132	5.082	1.277	4.475	2.513
SE (Sij - Sik)		0.480	25.718	14.503	1.316	4.337	0.252	4.604	7.471	1.877	6.578	3.694
SE (Sij - Skm)		0.457	24.521	13.828	1.255	4.135	0.240	4.389	7.123	1.790	6.272	3.522
CD (Sij) at 5%		0.646	34.659	19.546	1.774	5.845	0.339	6.204	10.068	2.530	8.822	4.978

* Significant at 5%

** Significant at 1%

F.m. = *F. moniliforme**F.p.* = *F. pallidoroseum*

Table : 30 Estimates of specific combining ability effects for F_1 crosses in 10×10 diallel of sorghum biochemical character Akola, 1995.

Sr.No.	Crosses	Protein	Soluble sugars	Tannins	Flavan-4-ols
1	2	3	4	5	6
1	SPV-1201 x ICSB - 101 B	-0.021	-0.038	-0.125**	-0.003
2	SPV-1201 x AKms - 14 B	-0.340**	-0.121**	-0.122**	-0.147**
3	SPV-1201 x SRT - 26B	-0.341**	-0.073	-0.106**	-0.166**
4	SPV-1201 x GJ-35-15-15	0.006	0.044	-0.031**	-0.067**
5	SPV-1201 x SPV - 946	-0.134**	0.017	-0.049**	-0.175**
6	SPV-1201 x SPV - 104	-0.158**	0.007	-0.071**	-0.157**
7	SPV-1201 x IS - 2284	0.181**	-0.025	0.058**	0.182**
8	SPV-1201 x IS-6335	-0.214**	0.089*	0.214**	0.072*
9	SPV-1201 x IS - 9471	0.278**	0.047	0.244**	0.063
10	ICSB - 101B x AKms 14 B	-0.125**	-0.127**	-0.058**	-0.066
11	ICSB - 101B x SRT - 26 B	-0.000	-0.112**	-0.063**	-0.085*
12	ICSB - 101B x GJ-35-15-15	-0.067**	0.013	0.005	0.014
13	ICSB - 101B x SPV - 946	-0.237**	0.017	0.008	-0.094**
14	ICSB - 101B x SPV - 104	-0.284**	0.004	-0.041**	-0.075*
15	ICSB - 101B x IS - 2284	-0.174**	0.018	0.108**	0.019
16	ICSB - 101B x IS - 6335	-0.179**	0.026	-0.015	0.225**
17	ICSB - 101B x IS - 9471	-0.186**	-0.017	0.045**	-0.091
18	AKms - 14 B x SRT - 26B	-0.179	-0.237*	-0.146**	-0.229**
19	AKms - 14 B x GJ-35-15-15	-0.132**	-0.149**	-0.071**	0.087*
20	AKms - 14 B x SPV - 946	-0.232	0.048	-0.081**	-0.238**
21	AKms - 14 B x SPV - 104	-0.317**	-0.041	-0.124**	-0.219**
22	AKms - 14 B x IS - 2284	0.188**	0.042	0.344**	0.635**
23	AKms - 14 B x IS - 6335	0.175**	-0.055	0.216**	0.192**
24	AKms - 14 B x IS - 9471	-0.264**	0.080*	0.283**	0.405**

Contd.

Table 30 Contd.

1	2	3	4	5	6
25	SRT - 26B x GJ-35-15-15	-0.154**	-0.057	-0.048**	-0.149**
26	SRT - 26B x SPV - 946	-0.130**	0.021	-0.080**	0.035
27	SRT - 26B x SPV - 104	-0.022	-0.067	-0.088**	-0.238**
28	SRT - 26B x IS - 2284	-0.290**	0.050	0.179**	0.299**
29	SRT - 26B x IS - 6335	-0.183**	0.026	0.299**	0.596**
30	SRT - 26B x IS - 9471	-0.274**	0.054	0.302**	0.431**
31	GJ-35-15-15 x SPV - 946	-0.001	-0.126**	0.009	0.184**
32	GJ-35-15-15 x SPV 104	-0.089**	0.095*	-0.040*	0.048
33	GJ-35-15-15 x IS - 2284	-0.187**	0.029	0.105**	0.032
34	GJ-35-15-15 x IS - 6335	-0.286**	0.037	-0.021	0.110**
35	GJ-35-15-15 x IS - 9471	-0.328**	0.066	0.028*	0.062
36	SPV - 946 x SPV - 104	0.070**	-0.062	-0.058**	-0.248**
37	SPV - 946 x IS - 2284	-0.141**	-0.074*	0.063**	0.246**
38	SPV - 946 x IS - 6335	-0.238**	0.009	0.163**	0.586**
39	SPV - 946 x IS 9471	-0.188**	-0.074*	0.028*	0.236**
40	SPV - 104 x IS 2284	0.119**	-0.048	0.074**	0.391**
41	SPV - 104 x IS 6335	0.170**	-0.129**	0.234**	0.530**
42	SPV - 104 x IS 9471	-0.037	-0.092*	0.233**	0.426**
43	IS 2284 x IS - 6335	-0.153**	-0.103**	0.763**	-0.749**
44	IS 2284 x IS - 9471	0.106**	-0.072	-0.408**	-0.492**
45	IS 6335 x IS 9471	0.240**	-0.065	-0.291**	-0.517**

SE (Sij)

SE (Sij - Sik)

CD (Sij) 5 %

* Significant at 5%

** Significant at 1%

0.035

0.051

0.069

Table 31. Protein fractions % of parents and F₁ crosses, Akola, 1995

Sr.No	Cross No.	Parents/Crosses	Albumin & Globulin	Profilamin	Cross link profilamin	Glutelin like	Glutelin	Residues
1	1	SPV-1201	14.41	6.96	10.41	2.51	28.74	10.68
2	2	ICSB-101B	15.84	5.49	8.57	3.01	27.90	8.82
3	3	AKms-14B	14.09	9.08	12.14	2.46	28.95	11.46
4	4	SRT-26B	20.63	5.88	10.18	3.22	32.47	6.78
5	5	GJ-35-15-15	15.60	6.47	13.39	3.33	35.11	9.12
6	6	SPV-946	13.62	6.36	13.62	3.18	37.06	9.78
7	7	SPV-104	12.62	4.88	14.92	3.98	32.51	8.64
8	8	IS-2284	1.81	0.70	5.24	12.39	6.27	7.26
9	9	IS-6335	1.96	2.79	10.53	12.79	57.25	10.20
10	10	IS-9471	8.73	1.90	6.83	9.07	56.43	8.04
11	13	SPV-1201 x SPV-946	14.19	3.65	12.89	2.86	23.81	11.14
12	12	SPV-1201 x SPV-104	14.94	3.15	13.03	4.04	30.67	20.70
13	11	SPV-1201 x IS-9471	9.00	1.17	8.90	10.88	38.48	4.41
14	16	ICSB-101B x SPV-946	15.13	2.42	12.10	3.33	39.18	9.96
15	15	ICSB-101B x SPV-104	10.91	1.71	13.40	3.81	37.58	6.30
16	14	ICSB-101 B x IS-9471	8.82	0.75	9.12	7.65	46.91	6.18
17	19	AKms-14B x SPV-946	15.89	3.97	10.19	3.80	35.58	9.30
18	18	AKms-14B x SPV-104	13.63	4.13	12.10	4.13	36.60	10.80
19	17	AKms-14B x IS-9471	8.63	0.54	5.94	7.73	45.32	5.40
20	22	SRT-26B x SPV-946	18.09	4.10	9.73	4.40	28.88	6.36
21	21	SRT-26B x SPV-104	14.37	3.53	9.78	5.54	26.62	7.80
22	20	SRT-26B x IS-9471	10.07	0.52	3.99	10.24	46.18	11.46
23	25	GJ-35-15-15 x SPV-946	15.88	4.72	10.59	3.00	32.76	3.42
24	24	GJ-35-15-15 x SPV-104	15.67	4.24	8.87	3.73	30.98	10.02
25	23	GJ-35-15-15 x IS-9471	15.30	1.34	5.54	7.65	51.05	4.56
26	27	SPV-946 x SPV-104	16.74	3.79	9.38	4.46	33.93	4.68
27	26	SPV-946 x IS-9471	13.89	1.80	7.19	7.02	51.96	3.90
28	28	SPV-104 x IS-9471	15.94	1.58	7.54	7.66	36.01	8.76
29	29	IS-2284 x IS-9471	~ 90	0.38	4.39	8.66	47.05	8.28
30	30	IS-6335 x IS-9471	9.77	2.55	6.58	0.96	49.68	10.80
SE (m)			0.217	0.035	0.048	0.070	0.279	0.133
C/D at 5%			0.620	0.101	0.139	0.210	0.798	0.381

Table : 32 Observation on glume colour, grain color and testa

Sr. No.	Parents/ Crosses	Glume color	Grain color	Testa	Sr. No.	Parents/ Crosses	Glume color	Grain color	Testa
1	SPV-1201	Light red	White	A	29	AKms - 14 B x GI-35-15-15	Light red	White	A
2	ICSB-101B	Light red	White	A	30	AKms - 14 B x SPV - 946	Light red	White	A
3	AKms-14B	Light red	White	A	31	AKms - 14 B x SPV - 104	Light red	White	A
4	SRT-26B	Red	White	A	32	AKms - 14 B x IS - 2284	Purple	Red	P
5	GI-35-15-15	Red	White	A	33	AKms - 14 B x IS - 6335	Black	Red	P
6	SPV-946	Light red	White	A	34	AKms - 14 B x IS - 9471	Purple	Red	P
7	SPV-104	Light red	White	A	35	SRT - 26B x GI-35-15-15	Light red	White	A
8	IS-2284	Purple	Light red	P	36	SRT - 26B x SPV - 946	Light red	White	A
9	IS-6335	Black	Dark red	P	37	SRT - 26B x SPV - 104	Light red	White	A
10	IS-9471	Purple	Dark red	P	38	SRT - 26B x IS - 2284	Purple	Light red	P
11	SPV-1201 x ICSB-101B	Tan	White	A	39	SRT - 26B x IS - 6335	Black	Light red	P
12	SPV-1201 x AKms-14 B	Tan	White	A	40	SRT - 26B x IS - 9471	Red	Red	P
13	SPV-1201 x SRT - 26B	Light red	White	A	41	GI-35-15-15 x SPV - 946	Light red	White	A
14	SPV-1201 x GI-35-15-15	Light red	White	A	42	GI-35-15-15 x SPV - 104	Light red	White	A
15	SPV-1201 x SPV - 946	Tan	White	A	43	GI-35-15-15 x IS - 2284	Purple	Dark red	P
16	SPV-1201 x SPV - 104	Tan	White	A	44	GI-35-15-15 x IS - 6335	Black	Dark red	P
17	SPV-1201 x IS - 2284	Purple	Light red	P	45	GI-35-15-15 x IS - 9471	Red	Red	P
18	SPV-1201 x IS-6335	Partly straw purple	Dark red	P	46	SPV - 946 x SPV - 104	Light red	White	A
19	SPV-1201 x IS - 9471	Red	Light red	P	47	SPV - 946 x IS - 2284	Purple	Red	P
20	ICSB - 101B x AKms 14 B	Light red	White	A	48	SPV - 946 x IS - 6335	Black	Light red	P
21	ICSB - 101B x SRT - 26 B	Red	White	A	49	SPV - 946 x IS 9471	Red	Red	P
22	ICSB - 101B x GI-35-15-15	Red	White	A	50	SPV - 104 x IS 2284	Purple	Red	P
23	ICSB - 101B x SPV - 946	Light Red	White	A	51	SPV - 104 x IS 6335	Black	Red	P
24	ICSB - 101B x SPV - 104	White	White	A	52	SPV - 104 x IS 9471	Black	Red	P
25	ICSB - 101B x IS - 2284	Purple	Light red	P	53	IS 2284 x IS-6335	Black	Light red	P
26	ICSB - 101B x IS - 6335	Black	Light red	P	54	IS 2284 x IS - 9471	Black	Red	P
27	ICSB - 101B x IS - 9471	Purple	Red	P	55	IS 6335 x IS 9471	Black	Dark red	P
28	AKms - 14 B x SRT - 26B	Light Red	White	A					

A = Absent

P = Present

VIII - 1
 Table 33. Effect of pre-treatment on germination (%) of parents, F_1 crosses at Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola		Mean	Patancheru		Mean	Pooled over locations		Mean
		Untreated	Treated		Untreated	Treated		Untreated	Treated	
1	2	3	4	5	6	7	8	9	10	11
1	SPV-1201	78.00	76.00	77.00	49.33	52.00	50.67	63.67	64.00	63.83
2	ICSB-101B	76.67	80.00	78.33	35.33	28.67	32.00	56.00	54.33	55.17
3	Akma-14B	54.00	68.67	86.00	4.67	5.33	5.00	29.33	37.00	33.17
4	SRT-26B	86.67	85.33	86.00	22.00	18.00	20.00	54.33	51.67	53.00
5	GI-35-15-15	81.33	77.33	79.33	24.67	12.00	18.33	53.00	44.67	48.83
6	SPV-946	87.33	81.33	84.33	32.67	22.67	27.67	60.00	52.00	56.00
7	SPV-104	61.33	58.00	59.67	4.00	8.67	6.33	32.67	33.33	33.00
8	IS-2284	93.33	90.67	92.00	66.67	82.00	74.33	80.00	86.33	83.17
9	IS-6335	93.33	93.33	93.33	78.00	83.33	80.67	85.67	88.33	87.00
10	IS-9471	93.33	94.00	93.67	71.33	76.67	74.00	82.33	85.33	83.83
11	SPV-1201 x ICSB-101B	85.33	88.00	86.67	52.67	52.67	52.67	69.00	70.33	69.67
12	SPV-1201 x Akma-14 B	82.67	85.33	84.00	52.67	61.33	57.00	67.67	73.33	70.50
13	SPV-1201 x SRT-26B	94.00	90.00	92.00	80.67	82.67	81.67	87.33	86.33	86.83
14	SPV-1201 x GI-35-15-15	60.67	94.67	77.67	71.33	84.00	77.67	66.00	89.33	77.67
15	SPV-1201 x SPV-946	65.33	74.67	70.00	68.00	81.33	74.67	66.67	78.00	72.33
16	SPV-1201 x SPV-104	84.67	87.33	86.00	43.33	62.00	52.67	64.00	74.67	69.33
17	SPV-1201 x IS-2284	91.33	96.00	93.67	74.67	79.33	77.00	83.00	87.67	85.33
18	SPV-1201 x IS-6335	96.00	93.33	94.67	70.00	68.67	69.33	83.00	81.00	82.00
19	SPV-1201 x IS-9471	95.33	92.67	94.00	76.67	72.67	74.67	86.00	82.67	84.33
20	ICSB-101B x Akma-14 B	78.67	76.00	77.33	45.33	42.67	44.00	62.00	59.33	60.67
21	ICSB-101B x SRT-26 B	88.00	91.33	89.67	39.33	41.33	40.33	63.67	66.33	65.00
22	ICSB-101B x GI-35-15-15	88.00	94.00	91.00	78.67	78.67	78.67	83.33	86.33	84.83
23	ICSB-101B x SPV-946	92.00	89.33	90.67	58.67	49.33	54.00	75.33	69.33	72.33
24	ICSB-101B x SPV-104	76.67	80.67	78.67	37.33	45.33	41.33	57.00	63.00	60.00
25	ICSB-101B x IS-2284	88.67	90.67	89.67	87.33	89.33	88.33	88.00	90.00	89.00
26	ICSB-101B x IS-6335	90.00	93.33	91.67	94.00	90.00	92.00	92.00	91.67	91.83
27	ICSB-101B x IS-9471	92.00	94.00	93.00	86.00	93.33	89.67	89.00	93.67	91.33
28	Akma-14 B x SRT-26B	72.00	82.00	77.00	26.00	28.67	27.33	49.00	55.33	52.17
29	Akma-14 B x GI-35-15-15	82.00	81.33	81.67	26.67	22.67	29.67	54.33	57.00	55.67
30	Akma-14 B x SPV-946	77.33	81.33	79.33	34.67	32.67	28.67	56.00	52.00	54.00
31	Akma-14 B x SPV-104	53.33	66.00	59.67	22.00	24.00	23.00	37.67	45.00	41.33
32	Akma-14 B x IS-2284	80.00	87.00	83.50	66.67	82.67	74.67	73.33	88.33	80.83
33	Akma-14 B x IS-6335	93.33	97.33	95.33	88.00	88.00	88.00	90.67	92.67	91.67
34	Akma-14 B x IS-9471	88.00	94.00	91.00	80.67	76.67	78.67	84.33	85.33	84.83

Contd....

VIII - 2

Table 33. Contd...

1	2	3	4	5	6	7	8	9	10	11
35	SRT- 26B x GI-35-15-15	92.67	90.00	91.33	40.67	40.67	40.67	56.67	56.33	66.00
36	SRT- 26B x SPV - 946	84.00	86.00	85.00	64.00	62.67	63.33	74.00	74.33	74.17
37	SRT- 26B x SPV -104	69.33	72.00	70.67	24.67	40.00	32.33	47.00	56.00	51.50
38	SRT- 26B x IS - 2284	88.00	78.67	83.33	80.67	95.33	88.00	84.33	87.00	85.67
39	SRT- 26B x IS - 6335	90.00	89.33	89.67	88.67	93.33	91.00	89.33	91.33	90.33
40	SRT- 26B x IS - 9471	90.00	95.33	92.67	86.00	90.67	88.33	88.00	93.00	90.50
41	GI-35-15-15 x SPV - 946	85.33	88.00	86.67	56.00	73.33	64.67	70.67	80.67	75.67
42	GI-35-15-15 x SPV - 104	83.33	86.00	84.67	50.67	42.67	46.67	67.00	64.33	65.67
43	GI-35-15-15 x IS - 2284	90.67	88.00	89.33	82.67	84.00	83.33	86.67	86.00	86.33
44	GI-35-15-15 x IS - 6335	92.00	90.00	91.00	83.33	88.00	85.67	87.67	89.00	88.33
45	GI-35-15-15 x IS - 9471	92.67	92.00	92.33	74.00	71.33	72.67	83.33	81.67	82.50
46	SPV - 946 x SPV - 104	84.67	80.00	82.33	27.33	42.67	35.00	56.00	61.33	58.67
47	SPV - 946 x IS - 2284	86.67	92.67	89.67	79.33	84.00	81.67	83.00	88.33	85.67
48	SPV - 946 x IS - 6335	93.33	95.33	94.33	77.33	83.33	80.33	85.33	89.33	87.33
49	SPV - 946 x IS 9471	92.00	89.33	90.67	78.00	82.00	80.00	85.00	85.67	85.33
50	SPV - 104 x IS 2284	88.67	89.33	89.00	80.67	87.33	84.00	84.67	88.33	86.50
51	SPV - 104 x IS 6335	92.67	93.33	93.00	84.67	93.33	89.00	88.67	93.33	91.00
52	SPV - 104 x IS 9471	92.00	94.00	93.00	78.00	92.00	85.00	85.00	93.00	89.00
53	IS 2284 x IS-6335	91.33	92.67	92.00	86.00	85.33	85.67	88.67	89.00	88.83
54	IS 2284 x IS -9471	88.00	84.00	86.00	77.33	92.67	85.00	82.67	88.33	85.50
55	IS 6335 x IS 9471	96.00	94.67	95.33	78.67	78.67	78.67	87.33	86.67	87.00
Mean		84.80	86.78	85.79	60.62	64.05	62.29	72.66	75.41	74.01
% increase (+)/decrease(-) over untreated				(+2.33)			(+5.51)			(+3.64)
% increase over location										

Patancheru.

27.39

SE (m)±	SE (m)±	SE (m)	CD 5%	CD 5%
0.430	0.639	0.578	1.557	1.681
2.254	3.324	3.030	9.213	8.847
3.389	4.700	4.286	13.029	12.471

A
B
A x B

VIII - 3

Table 34. Effect of pre-treatment on *F. moniliforme* (%) (GS) of parents and F_1 crosses at Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola		Patancheru		Mean		Pooled over locations		Mean
		Untreated	Treated	Untreated	Treated	5	8	Untreated	Treated	
1	2	3	4	5	6	7	8	9	10	11
1	SPV-1201	12.67	10.01	11.34	27.33	16.00	21.67	20.00	13.00	16.50
2	ICSB-101B	18.01	16.67	17.34	18.00	11.33	14.67	18.00	14.00	16.00
3	AKms-14B	24.67	17.99	21.33	2.67	2.67	2.67	13.67	10.33	12.00
4	SRT-26B	20.67	19.33	20.00	14.00	7.33	10.67	14.00	13.33	15.34
5	GI-35-15-15	11.99	18.67	15.33	16.67	7.33	12.00	14.33	13.00	13.67
6	SPV-946	20.00	22.67	17.34	18.00	8.67	13.33	15.00	15.67	15.34
7	SPV-104	20.01	27.33	23.67	4.00	4.67	4.33	12.00	16.00	14.00
8	IS-2284	15.99	8.67	12.33	28.00	16.00	22.00	22.00	12.34	17.17
9	IS-6335	5.33	3.33	4.33	32.00	12.67	22.33	18.66	8.00	13.33
10	IS-9471	3.33	3.33	3.33	20.67	15.33	18.00	12.00	9.33	10.67
11	SPV-1201 x ICSB-101B	12.67	9.34	11.01	28.67	11.33	20.00	20.67	10.34	15.50
12	SPV-1201 x AKms-14 B	8.67	10.01	9.34	26.00	12.67	19.33	17.34	11.34	14.34
13	SPV-1201 x SRT-26B	12.01	12.67	12.34	24.67	20.00	22.33	18.34	16.33	17.34
14	SPV-1201 x GI-35-15-15	12.67	11.34	12.00	25.33	16.67	21.00	19.00	14.00	16.50
15	SPV-1201 x SPV-946	16.66	13.99	15.33	21.33	20.67	21.00	19.00	17.33	18.16
16	SPV-1201 x SPV-104	10.00	9.99	10.00	22.67	8.67	15.67	16.33	9.33	12.83
17	SPV-1201 x IS-2284	5.34	3.33	4.34	25.33	9.33	17.33	15.34	6.33	10.84
18	SPV-1201 x IS-6335	5.34	3.99	4.67	26.00	10.00	18.00	15.67	7.00	11.33
19	SPV-1201 x IS-9471	6.66	4.67	5.67	17.33	12.00	14.67	12.00	8.34	10.17
20	ICSB-101B x AKms 14 B	13.33	9.99	11.66	24.00	13.33	18.67	18.67	11.66	15.17
21	ICSB-101B x SRT-26 B	7.99	5.34	6.67	26.67	12.67	19.67	17.33	9.00	13.17
22	ICSB-101B x SPV-946	5.33	14.01	9.67	38.00	16.00	27.00	21.67	15.00	18.34
23	ICSB-101B x SPV-104	11.33	12.01	11.67	33.33	10.00	21.67	22.33	11.00	16.67
24	ICSB-101B x IS-2284	22.67	10.67	16.67	27.33	14.00	20.67	25.00	12.34	18.67
25	ICSB-101B x IS-6335	8.66	3.99	6.33	24.00	10.67	17.33	16.33	7.33	11.83
26	ICSB-101B x IS-9471	6.66	5.34	6.00	16.67	12.00	14.33	11.66	8.67	10.17
27	AKms-14 B x SRT-26B	16.67	3.33	5.00	16.67	22.67	19.67	11.66	13.00	12.34
28	AKms-14 B x GI-35-15-15	4.67	14.67	14.67	16.00	8.00	12.00	15.34	11.33	13.34
29	AKms-14 B x SPV-946	15.99	11.33	13.66	16.67	10.67	13.67	16.33	11.00	13.67
30	AKms-14 B x SPV-104	9.33	9.99	9.66	14.00	8.67	11.33	11.67	9.33	10.50
31	AKms-14 B x IS-2284	17.99	12.67	15.33	10.00	8.67	9.33	14.00	10.67	12.33
32	AKms-14 B x IS-6335	10.01	5.34	7.34	22.67	9.33	16.00	16.34	7.00	11.67
33	AKms-14 B x IS-9471	10.67	4.67	8.01	18.00	8.67	13.33	14.34	7.00	10.67
34	AKms-14 B x IS-9471	12.01	4.67	8.34	23.33	8.67	16.00	17.67	6.67	12.17

Contd.

Table 35. Effect of pre-treatment on *F. pallidoseum* (%) (GS) of parents and F_1 crosses at Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola			Patancheru			Pooled over locations		
		Untreated	Treated	Mean	Untreated	Treated	Mean	Untreated	Treated	Mean
1	2	3	4	5	6	7	8	9	10	11
1	SPV-1201	0.00	0.67	0.33	6.00	5.33	5.67	3.00	3.00	300
2	ICSB-101B	0.00	4.67	2.33	3.33	3.33	3.33	1.67	4.00	283
3	AKms-14B	0.00	5.33	2.67	0.00	0.00	0.00	0.00	2.67	133
4	SRT-26B	0.00	4.00	2.00	1.33	4.00	2.67	1.67	4.00	283
5	GJ-35-15-15	0.00	6.67	3.33	4.00	0.00	2.00	2.00	3.33	267
6	SPV-946	0.67	6.00	3.33	6.00	3.33	4.67	3.33	4.67	400
7	SPV-104	0.00	6.67	3.33	0.00	0.00	0.00	0.00	3.33	167
8	IS-2284	1.33	2.67	2.00	12.00	6.00	9.00	6.67	4.33	550
9	IS-6335	2.67	2.67	2.67	7.33	5.33	6.33	5.00	4.00	450
10	IS-9471	0.67	0.67	0.67	6.00	6.00	6.00	3.33	3.33	333
11	SPV-1201 x ICSB-101B	1.33	0.00	0.67	4.00	4.67	4.33	2.67	2.33	250
12	SPV-1201 x AKms-14 B	0.00	2.67	1.33	8.67	2.67	5.67	4.33	2.67	350
13	SPV-1201 x SRT - 26B	0.00	2.00	1.00	8.00	6.67	7.33	4.00	4.33	417
14	SPV-1201 x GJ-35-15-15	0.00	2.67	1.33	6.67	4.67	5.67	3.33	3.67	350
15	SPV-1201 x SPV-946	0.00	0.67	0.33	5.33	6.00	5.67	2.67	3.33	380
16	SPV-1201 x SPV-104	0.00	0.67	0.33	4.00	2.67	3.33	2.00	1.67	183
17	SPV-1201 x IS-2284	2.00	0.00	1.00	8.67	1.33	5.00	5.33	0.67	300
18	SPV-1201 x IS-6335	0.00	2.00	1.00	8.67	2.67	5.67	4.33	2.33	333
19	SPV-1201 x IS-9471	2.00	0.67	1.33	6.67	2.00	4.33	4.33	1.33	283
20	ICSB - 101B x AKms 14 B	1.33	0.67	1.00	6.00	6.67	6.33	3.67	3.67	367
21	ICSB - 101B x SRT- 26 B	0.00	1.33	0.67	5.33	4.67	5.00	2.67	3.00	283
22	ICSB - 101B x GJ-35-15-15	0.00	0.00	0.00	14.67	6.00	10.33	7.33	3.00	517
23	ICSB - 101B x SPV - 946	2.67	1.33	2.00	8.00	4.00	6.00	5.33	2.67	400
24	ICSB - 101B x SPV - 104	0.00	2.67	1.33	6.67	4.67	5.67	3.33	3.67	350
25	ICSB - 101B x IS - 2284	0.00	2.67	1.33	5.33	4.67	5.00	2.67	3.67	317
26	ICSB - 101B x IS - 6335	0.67	0.67	0.67	6.67	3.33	5.00	3.67	2.00	283
27	ICSB - 101B x IS - 9471	1.33	0.00	0.67	6.00	6.00	6.00	3.67	3.00	333
28	AKms - 14 B x SRT - 26B	1.33	0.67	1.00	3.33	4.00	3.67	2.33	2.33	233
29	AKms - 14 B x GJ-35-15-15	3.33	1.33	2.33	4.00	6.00	5.00	3.67	3.67	367
30	AKms - 14 B x SPV - 946	0.00	0.67	0.33	3.33	2.00	2.67	1.67	1.33	150
31	AKms - 14 B x SPV - 104	4.00	2.00	3.00	4.00	3.33	3.67	4.00	2.67	333
32	AKms - 14 B x IS - 2284	4.00	1.33	2.67	10.67	3.33	7.00	7.33	2.33	483
33	AKms - 14 B x IS - 6335	3.33	2.00	2.67	8.00	4.00	6.00	5.67	3.00	433
34	AKms - 14 B x IS - 9471	1.33	0.67	1.00	7.33	5.33	6.33	4.33	3.00	367

Contd.

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Table 35. Contd. 2										
1	2	3	4	5	6	7	8	9	10	11
35	SRT- 26B x GJ-35-15-15	0.00	0.00	0.00	6.67	6.67	6.67	3.33	3.33	3.33
36	SRT - 26B x SPV - 946	0.00	6.00	3.00	10.67	4.67	7.67	5.33	5.33	5.33
37	SRT - 26B x SPV - 104	0.67	2.00	1.33	3.33	6.67	5.00	2.00	4.33	3.17
38	SRT - 26B x IS - 2284	1.33	1.33	1.33	6.67	5.33	6.00	4.00	3.33	3.67
39	SRT - 26B x IS - 6335	3.33	3.33	3.33	7.33	7.33	7.33	5.33	5.33	5.33
40	SRT - 26B x IS - 9471	1.33	2.00	1.67	7.33	7.33	7.33	4.33	4.67	4.50
41	GJ-35-15-15 x SPV - 946	0.67	0.00	0.33	4.67	3.33	3.67	2.67	1.33	2.00
42	GJ-35-15-15 x SPV - 104	1.33	1.33	1.33	8.67	3.33	6.00	5.00	2.33	3.67
43	GJ-35-15-15 x IS - 2284	3.33	2.00	2.67	7.33	3.33	5.33	5.33	2.67	4.00
44	GJ-35-15-15 x IS - 6335	3.33	2.00	2.67	10.67	4.67	7.67	7.00	3.33	5.17
45	GJ-35-15-15 x IS - 9471	1.33	2.00	1.67	12.67	5.33	9.00	7.00	3.33	5.33
46	SPV - 946 x SPV - 104	0.67	0.67	0.67	3.33	6.00	4.67	2.00	3.33	2.67
47	SPV - 946 x IS - 2284	2.67	2.67	2.67	13.33	6.00	9.67	8.00	4.33	6.17
48	SPV - 946 x IS - 6335	1.33	0.00	0.67	17.33	6.00	11.67	9.33	3.00	6.17
49	SPV - 946 x IS 9471	3.33	2.67	3.00	12.00	6.00	9.00	7.67	4.33	6.00
50	SPV - 104 x IS 2284	6.00	2.00	4.00	8.67	4.00	6.33	7.33	3.00	5.17
51	SPV - 104 x IS 6335	1.33	5.33	3.33	9.33	4.67	7.00	5.33	5.00	5.17
52	SPV - 104 x IS 9471	2.00	0.67	1.33	7.33	6.00	6.67	4.67	3.33	4.00
53	IS 2284 x IS-6335	1.33	4.67	3.00	8.00	6.00	7.00	4.67	5.33	5.00
54	IS 2284 x IS - 9471	4.67	6.00	5.33	8.00	4.00	6.00	6.33	5.00	5.67
55	IS 6335 x IS 9471	2.00	0.67	1.33	6.00	4.67	5.33	4.00	2.67	3.33
Mean		2.20	2.12	2.16	7.00	4.46	5.73	4.21	3.29	3.75
% increase (+)/decrease (-)				(-3.64)			(-36.28)			(-21.75)
over untreated										
% increase over location							165.27			
							Akola			
		SE (m)	CD 5%		SE (m)	CD 5%		SE (m)	CD 5%	
	A	0.168	0.468		0.220	0.610		0.216	0.628	
	B	0.885	2.454		1.154	3.200		1.131	3.291	
	A x B	1.252	3.470		1.632	4.526		1.599	4.656	

VIII - 7
 Table 36. Effect of pre-treatment on *C. lunata* (%) (GS) of parents and F_1 crosses at Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola		Patancheru		Mean		Pooled over locations		
		Untreated	Treated	Untreated	Treated	Mean	Mean	Untreated	Treated	Mean
		3	4	6	7	5	8	9	10	11
1	SPV-1201	45.33	45.36	45.35	15.33	45.35	17.00	30.33	32.01	31.17
2	ICSB-101B	45.33	39.32	42.33	11.33	42.33	12.67	28.33	26.66	27.50
3	AKma-14B	24.67	37.36	31.01	2.00	31.01	2.67	13.33	20.01	16.67
4	SRT-26B	48.67	34.00	41.33	6.00	41.33	6.00	27.33	20.00	23.67
5	GI-35-15-15	49.33	30.68	40.01	4.00	40.01	4.33	26.67	17.67	22.17
6	SPV-946	51.33	28.68	40.01	10.67	40.01	8.67	31.00	18.67	24.84
7	SPV-104	28.67	16.68	22.67	0.00	22.67	1.33	14.33	9.67	12.00
8	IS-2284	43.33	30.68	37.01	18.00	37.01	18.00	30.67	24.34	27.50
9	IS-6335	22.00	14.00	18.00	12.67	18.00	15.00	17.33	15.67	16.50
10	IS-9471	20.00	16.68	18.34	14.67	18.34	19.00	17.33	20.01	18.67
11	SPV-1201 x ICSB-101B	48.67	46.68	47.67	18.67	47.67	13.33	33.67	30.01	31.84
12	SPV-1201 x AKma-14 B	50.67	40.00	45.33	14.00	45.33	13.00	32.33	26.00	29.17
13	SPV-1201 x SRT-26B	48.00	44.68	46.34	24.67	46.34	24.67	36.33	34.67	35.50
14	SPV-1201 x GI-35-15-15	37.33	39.36	38.68	20.67	38.68	23.00	27.33	35.35	31.34
15	SPV-1201 x SPV-946	48.00	44.00	46.00	14.67	46.00	26.00	28.67	35.68	32.17
16	SPV-1201 x SPV-104	24.67	9.36	17.01	24.00	17.01	18.33	31.33	31.33	17.67
17	SPV-1201 x IS-2284	16.67	8.68	12.67	23.33	12.67	16.67	20.00	8.34	14.67
18	SPV-1201 x IS-6335	16.00	9.36	12.68	19.33	12.68	15.33	17.67	10.35	14.01
19	SPV-1201 x IS-9471	47.33	32.68	40.01	11.33	40.01	14.67	29.33	25.34	27.34
20	ICSB-101B x AKma 14 B	62.00	28.68	45.34	6.67	45.34	9.33	34.33	20.34	27.34
21	ICSB-101B x SRT-26 B	60.67	46.00	53.33	20.00	53.33	19.67	40.33	32.67	36.50
22	ICSB-101B x GI-35-15-15	45.33	26.68	36.01	15.33	36.01	16.67	30.33	22.34	26.34
23	ICSB-101B x SPV-946	38.67	39.32	38.99	3.33	38.99	9.33	21.00	27.33	24.16
24	ICSB-101B x SPV-104	34.00	19.36	26.68	16.67	26.68	14.00	25.33	16.68	21.01
25	ICSB-101B x IS-2284	23.33	16.00	19.67	12.67	19.67	12.67	18.00	17.33	18.50
26	ICSB-101B x IS-6335	24.00	16.00	20.00	15.33	20.00	17.00	19.67	17.33	18.50
27	ICSB-101B x IS-9471	41.33	48.68	45.01	6.67	45.01	11.33	24.00	32.34	28.17
28	AKma-14 B x SRT-26B	47.33	50.00	47.33	5.33	47.33	8.67	25.00	31.00	28.00
29	AKma-14 B x GI-35-15-15	44.67	52.00	49.67	13.33	49.67	13.33	30.33	32.67	31.50
30	AKma-14 B x SPV-946	47.33	39.36	31.68	6.00	31.68	8.67	15.00	25.35	20.17
31	AKma-14 B x SPV-104	24.00	22.00	30.00	17.33	30.00	16.67	27.67	23.33	25.50
32	AKma-14 B x IS-2284	38.00	27.33	32.68	16.00	32.68	14.33	21.67	14.33	18.00
33	AKma-14 B x IS-6335	27.33	16.00	21.67	20.67	21.67	16.67	27.33	17.67	22.50
34	AKma-14 B x IS-9471	34.00	22.68	28.34	20.67	28.34	16.67	27.33	17.67	22.50

Contd.,

Table 36. Contd.....

1	2	3	4	5	6	7	8	9	10	11
35	SRT - 26B x GI-35-15-15	50.67	42.00	46.33	4.67	15.33	10.00	27.67	28.67	28.17
36	SRT - 26B x SPV - 946	44.67	48.00	46.33	11.33	26.67	19.00	28.00	37.33	32.67
37	SRT - 26B x SPV - 104	36.67	36.68	36.67	8.00	10.67	9.33	22.33	23.67	23.00
38	SRT - 26B x IS - 2284	38.00	17.36	27.68	15.33	22.00	18.67	26.67	19.68	23.17
39	SRT - 26B x IS - 6335	31.33	16.68	24.01	12.00	22.67	17.33	21.67	19.67	20.67
40	SRT - 26B x IS - 9471	27.33	20.00	23.67	12.00	21.33	16.67	19.67	20.67	20.17
41	GI-35-15-15 x SPV - 946	51.33	36.68	44.01	17.33	37.33	27.33	34.33	37.01	35.67
42	GI-35-15-15 x SPV - 104	41.33	48.00	44.67	10.67	13.33	12.00	26.00	30.67	28.33
43	GI-35-15-15 x IS - 2284	34.67	21.36	28.01	14.00	30.67	22.33	24.33	26.01	25.17
44	GI-35-15-15 x IS - 6335	29.33	22.00	25.67	15.33	38.67	27.00	22.33	30.33	26.33
45	GI-35-15-15 x IS - 9471	28.67	34.00	31.33	18.00	26.00	22.00	23.33	30.00	26.67
46	SPV - 946 x SPV - 104	47.33	36.68	42.01	6.67	17.33	12.00	27.00	27.01	27.00
47	SPV - 946 x IS - 2284	26.00	14.00	20.00	17.33	22.67	20.00	21.67	18.33	20.00
48	SPV - 946 x IS - 6335	28.00	14.00	21.00	16.67	30.67	23.67	22.33	22.33	22.33
49	SPV - 946 x IS 9471	26.00	20.00	23.00	14.67	22.67	18.67	20.33	21.33	20.83
50	SPV - 104 x IS 2284	31.33	18.00	24.67	20.67	16.00	18.33	26.00	17.00	21.50
51	SPV - 104 x IS 6335	30.00	20.68	25.34	18.67	18.00	18.33	24.33	19.34	21.84
52	SPV - 104 x IS 9471	34.00	20.00	27.00	11.33	18.00	14.67	22.67	19.00	20.83
53	IS 2284 x IS-6335	25.33	16.68	21.01	16.00	22.00	19.00	20.67	19.34	20.00
54	IS 2284 x IS-9471	26.00	17.36	21.68	20.67	15.33	18.00	23.33	16.35	19.84
55	IS 6335 x IS 9471	21.33	13.36	17.35	12.67	18.67	15.67	17.00	16.01	16.51
Mean		36.43	28.89	32.66	13.72	17.53	15.63	25.08	23.21	24.15
% increase (+)/decrease (-)				(-)20.69			(+)21.73			(-)17.44
over untreated										
% increase over location										

Palancheru

SE (m)	CD 5%	SE (m)	CD 5%	SE (m)	CD 5%
0.504	1.455	0.056	0.164	0.616	1.792
2.643	7.633	0.297	0.860	3.232	9.405
3.738	10.795	0.421	1.216	4.571	13.302

A
B
A x B

Table 37. Effect of pre-treatment on other fungi (%) (GS) of parents and F_1 crosses at Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola		Mean		Patancheru		Mean		Pooled over locations		Mean
		Untreated	Treated	5	6	Untreated	Treated	7	8	Untreated	Treated	
1	SPV-1201	20.00	18.67	19.33	0.67	2.00	1.33	10.33	10.33	10.33	10.33	11
2	ICSB-101B	13.33	12.67	13.00	2.67	0.00	1.33	6.33	8.00	6.33	7.17	7.17
3	AKms-14B	4.67	8.00	6.33	0.00	0.00	0.00	2.33	4.00	2.33	3.17	3.17
4	SRT-26B	15.33	16.67	16.00	0.00	0.67	0.33	7.67	8.67	7.67	8.17	8.17
5	GI-35-15-15	20.00	14.67	17.33	0.00	0.00	0.00	10.00	7.33	10.00	8.67	8.67
6	SPV-946	23.33	14.00	18.67	0.00	2.00	1.00	11.67	8.00	11.67	9.83	9.83
7	SPV-104	12.67	7.33	10.00	0.00	1.33	0.67	6.33	4.33	6.33	5.33	5.33
8	IS-2284	24.67	10.00	17.33	5.33	4.00	4.67	15.00	7.00	15.00	11.00	11.00
9	IS-6335	19.33	6.67	13.33	4.67	6.00	5.33	12.33	6.33	12.33	9.33	9.33
10	IS-9471	20.00	18.67	19.33	4.00	4.67	4.33	11.67	5.33	11.67	8.50	8.50
11	SPV-1201 x ICSB-101B	22.67	18.67	20.67	1.33	3.33	2.33	12.00	11.00	12.00	11.50	11.50
12	SPV-1201 x AKms-14 B	23.33	18.00	20.67	5.33	4.67	5.00	14.33	11.33	14.33	12.83	12.83
13	SPV-1201 x SRT - 26B	34.00	20.67	27.33	23.33	13.33	16.00	16.33	15.33	16.33	15.83	15.83
14	SPV-1201 x GI-35-15-15	14.00	17.33	15.67	18.67	21.33	22.00	21.67	20.00	21.67	18.17	18.17
15	SPV-1201 x SPV-946	11.33	18.00	14.67	21.33	2.67	3.33	15.33	15.00	15.33	15.17	15.17
16	SPV-1201 x SPV-104	26.67	27.33	27.00	4.00	2.00	2.00	14.00	13.67	14.00	13.67	13.67
17	SPV-1201 x IS-2284	18.67	9.33	14.00	8.67	2.00	5.33	13.67	3.33	13.67	9.67	9.67
18	SPV-1201 x IS-6335	18.00	5.33	11.67	9.33	1.33	5.33	13.67	3.33	13.67	8.50	8.50
19	SPV-1201 x IS-9471	16.67	5.33	11.00	10.00	0.67	5.33	13.33	3.00	13.33	8.17	8.17
20	ICSB-101B x AKms 14 B	16.67	10.67	13.67	4.00	6.00	5.00	10.33	8.33	10.33	9.33	9.33
21	ICSB-101B x SRT-26 B	18.00	10.00	14.00	1.33	3.33	2.33	9.67	6.67	9.67	8.17	8.17
22	ICSB-101B x GI-35-15-15	22.00	8.67	15.33	6.00	1.33	3.67	14.00	5.00	14.00	9.50	9.50
23	ICSB-101B x SPV-946	33.33	10.67	22.00	2.00	0.00	1.00	17.67	5.33	17.67	11.50	11.50
24	ICSB-101B x SPV-104	16.00	11.33	13.67	0.00	0.00	0.00	8.00	5.67	8.00	6.83	6.83
25	ICSB-101B x IS-2284	32.00	10.00	21.00	8.00	0.00	4.00	20.00	12.50	20.00	12.50	12.50
26	ICSB-101B x IS-6335	31.33	8.00	19.67	7.33	3.33	5.33	19.33	5.67	19.33	10.83	10.83
27	ICSB-101B x IS-9471	26.00	9.33	17.67	8.00	0.00	4.00	17.00	4.67	17.00	10.83	10.83
28	AKms-14 B x SRT-26B	14.67	18.00	16.33	0.00	0.67	0.33	7.33	9.33	7.33	8.33	8.33
29	AKms-14 B x GI-35-15-15	18.00	18.67	18.33	0.67	4.00	2.33	9.33	10.33	9.33	10.33	10.33
30	AKms-14 B x SPV-946	20.67	18.67	19.67	4.00	0.00	2.33	12.33	9.33	12.33	10.83	10.83
31	AKms-14 B x SPV-104	8.00	12.67	10.33	2.00	0.67	1.33	5.00	6.67	5.00	5.83	5.83
32	AKms-14 B x IS-2284	24.00	6.00	15.00	7.33	4.00	5.67	15.67	5.00	15.67	10.33	10.33
33	AKms-14 B x IS-6335	37.33	7.33	22.33	6.00	2.00	4.00	21.67	13.17	21.67	13.17	13.17
34	AKms-14 B x IS-9471	20.67	10.67	15.67	8.00	2.00	5.00	14.33	6.33	14.33	10.33	10.33

Contd.

VIII - 10

Table 37. Contd.		1	2	3	4	5	6	7	8	9	10	11
35	SRT- 26B x GI-35-15-15	34.00	19.33	26.67	0.00	3.33	1.67	17.00	11.33	14.17		
36	SRT - 26B x SPV - 946	20.00	21.33	20.67	1.33	6.00	3.67	10.67	13.67	12.17		
37	SRT - 26B x SPV - 104	13.33	18.67	16.00	1.33	0.67	1.00	7.33	9.67	8.50		
38	SRT - 26B x IS - 2284	32.00	11.33	21.67	5.33	6.67	6.00	18.67	9.00	13.83		
39	SRT - 26B x IS - 6335	24.00	13.33	18.67	6.00	4.67	5.33	15.00	9.00	12.00		
40	SRT - 26B x IS - 9471	16.67	14.00	15.33	4.67	4.00	4.33	10.67	9.00	9.83		
41	GI-35-15-15 x SPV - 946	24.00	16.00	20.00	4.67	4.00	5.67	14.33	11.33	12.83		
42	GI-35-15-15 x SPV - 104	17.33	24.00	20.67	4.00	4.00	4.00	10.67	14.00	12.33		
43	GI-35-15-15 x IS - 2284	17.33	14.00	15.67	4.67	7.33	6.00	11.00	10.67	10.83		
44	GI-35-15-15 x IS - 6335	22.00	13.33	17.67	7.33	6.67	7.00	14.67	10.00	12.33		
45	GI-35-15-15 x IS - 9471	22.67	15.33	19.00	7.33	7.33	7.33	15.00	11.33	13.17		
46	SPV - 946 x SPV - 104	30.67	22.67	26.67	0.67	6.67	3.67	15.67	14.67	15.17		
47	SPV - 946 x IS - 2284	23.33	7.33	15.33	6.00	4.67	5.33	14.67	6.00	10.33		
48	SPV - 946 x IS - 6335	26.00	12.00	19.00	4.00	5.33	4.67	15.00	8.67	11.83		
49	SPV - 946 x IS 9471	23.33	12.67	18.00	5.33	8.00	6.67	14.33	10.33	12.33		
50	SPV - 104 x IS 2284	25.33	12.67	19.00	7.33	5.33	6.33	16.33	9.00	12.67		
51	SPV - 104 x IS 6335	29.33	10.67	20.00	6.67	6.67	6.67	18.00	8.67	13.33		
52	SPV - 104 x IS 9471	22.67	10.00	16.33	5.33	4.67	5.00	14.00	7.33	10.67		
53	IS 2284 x IS-6335	24.67	10.67	17.67	4.67	4.00	4.33	14.67	7.33	11.00		
54	IS 2284 x IS - 9471	25.33	11.33	18.33	5.33	3.33	4.33	15.33	7.33	11.33		
55	IS 6335 x IS 9471	16.67	6.67	11.67	5.33	5.33	5.33	11.00	6.00	8.50		
Mean		21.60	13.14	17.34	5.11	3.94	4.53	13.36	8.54	10.95		
% increase (-)/decrease (-)				(-)39.16			(-)22.89			(-)36.09		
% increase over location												
A		SE (m)	CD 5%	Palancheru		SE (m)	CD 5%	SE (m)		CD 5%		
B		0.675	1.871			0.400	1.108			0.666		
A x B		3.539	9.810			2.097	5.813			3.494		
		5.005	13.873			2.966	8.220			4.942		

Table 38. Effect of pre-treatment on score (GS) of parents and F₁ crosses at Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola		Patancheru		Mean		Pooled over locations		Mean
		Untreated	Treated	Untreated	Treated	Untreated	Treated	Untreated	Treated	
1	2	3	4	5	6	7	8	9	10	11
1	SPV-1201	4.00	3.00	3.50	4.00	4.00	4.00	4.00	3.50	3.75
2	ICSB-101B	4.00	3.00	3.50	4.00	4.00	4.00	4.00	3.50	3.75
3	AKms-14B	5.00	3.50	4.25	3.00	4.33	3.67	4.00	3.92	3.96
4	SRT-26B	3.83	3.00	3.42	3.50	4.00	3.75	3.67	3.50	3.58
5	GI-35-15-15	3.50	3.00	3.25	3.50	3.33	3.42	3.50	3.17	3.33
6	SPV-946	3.83	2.50	3.17	3.50	3.00	3.25	3.67	2.75	3.21
7	SPV-104	4.50	3.50	4.00	3.33	3.33	3.33	3.92	3.42	3.67
8	IS-2284	2.00	1.00	1.50	2.00	1.00	1.50	2.00	1.00	1.50
9	IS-6335	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00
10	IS-9471	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00
11	SPV-1201 x ICSB-101B	3.00	2.50	2.75	3.00	3.00	3.00	3.00	2.75	2.88
12	SPV-1201 x AKms-14 B	3.00	2.50	2.75	3.67	3.00	3.33	3.33	2.75	3.04
13	SPV-1201 x SRT-26B	2.83	3.00	2.92	3.33	3.00	3.17	3.08	3.00	3.04
14	SPV-1201 x GI-35-15-15	2.50	3.00	2.75	3.50	3.00	3.25	3.00	3.00	3.00
15	SPV-1201 x SPV-946	2.83	3.00	2.92	3.50	3.00	3.25	3.17	3.00	3.08
16	SPV-1201 x SPV-104	3.17	2.83	3.00	3.50	3.00	3.25	3.33	2.92	3.13
17	SPV-1201 x IS-2284	1.00	1.00	1.00	2.50	1.00	1.75	1.75	1.00	1.38
18	SPV-1201 x IS-6335	1.00	1.00	1.00	1.50	1.00	1.25	1.25	1.00	1.13
19	SPV-1201 x IS-9471	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00
20	ICSB-101B x AKms-14 B	3.00	3.00	3.00	3.00	3.50	3.25	3.00	3.25	3.13
21	ICSB-101B x SRT-26 B	3.00	2.00	2.50	3.00	3.17	3.08	3.00	2.58	2.79
22	ICSB-101B x GI-35-15-15	2.67	2.00	2.33	3.00	3.00	3.00	2.83	2.50	2.67
23	ICSB-101B x SPV-946	2.67	3.00	2.83	3.00	3.00	3.00	2.83	3.00	2.92
24	ICSB-101B x SPV-104	3.33	2.00	2.67	3.50	2.00	2.75	3.42	2.00	2.71
25	ICSB-101B x IS-2284	1.33	1.00	1.17	1.50	1.00	1.25	1.42	1.00	1.21
26	ICSB-101B x IS-6335	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00
27	ICSB-101B x IS-9471	1.50	1.00	1.25	1.00	1.00	1.00	1.25	1.00	1.13
28	AKms-14 B x SRT-26B	3.00	2.83	2.92	3.50	3.33	3.42	3.25	3.08	3.17
29	AKms-14 B x GI-35-15-15	3.00	3.00	3.00	3.50	2.50	3.00	3.25	2.75	3.00
30	AKms-14 B x SPV-946	3.00	3.00	3.00	3.50	4.00	3.75	3.25	3.50	3.38
31	AKms-14 B x SPV-104	3.50	4.00	3.75	3.67	4.00	3.83	3.58	4.00	3.79
32	AKms-14 B x IS-2284	2.00	1.00	1.50	1.50	1.00	1.25	1.75	1.00	1.38
33	AKms-14 B x IS-6335	1.50	1.00	1.25	1.00	1.00	1.00	1.25	1.00	1.13
34	AKms-14 B x IS-9471	2.17	1.00	1.58	1.00	1.00	1.00	1.58	1.00	1.29

Contd.

	1	2	3	4	5	6	7	8	9	10	11
35	SRT-26B x GJ-35-15-15	2.83	2.50	2.67	3.50	2.83	3.17	3.17	3.17	2.67	2.92
36	SRT-26B x SPV - 946	3.00	2.50	2.75	3.50	3.17	3.33	3.33	3.25	2.83	3.04
37	SRT-26B x SPV - 104	3.00	3.00	3.00	3.00	2.67	2.83	3.00	3.00	2.83	2.92
38	SRT-26B x IS - 2284	1.00	1.00	1.00	1.50	2.00	1.75	1.75	1.25	1.50	1.38
39	SRT-26B x IS - 6335	1.00	1.00	1.00	1.50	1.67	1.58	1.25	1.25	1.33	1.29
40	SRT-26B x IS - 9471	1.00	1.00	1.00	1.50	1.00	1.25	1.25	1.25	1.00	1.13
41	GJ-35-15-15 x SPV - 946	3.00	2.33	2.67	3.50	3.83	3.67	3.25	3.25	3.08	3.17
42	GJ-35-15-15 x SPV - 104	2.33	2.50	2.42	3.50	4.00	3.75	2.92	2.92	3.25	3.08
43	GJ-35-15-15 x IS - 2284	1.00	1.00	1.00	1.50	2.00	1.75	1.25	1.25	1.50	1.38
44	GJ-35-15-15 x IS - 6335	1.00	1.00	1.00	1.50	2.17	1.83	1.25	1.25	1.58	1.42
45	GJ-35-15-15 x IS - 9471	1.00	2.00	1.50	1.67	2.00	1.83	1.33	1.33	2.00	1.67
46	SPV - 946 x SPV - 104	3.33	3.00	3.17	3.67	4.00	3.83	3.50	3.50	3.50	3.50
47	SPV - 946 x IS - 2284	1.17	1.00	1.08	1.50	4.00	2.75	1.33	1.33	1.92	2.50
48	SPV - 946 x IS - 6335	1.00	1.00	1.00	1.50	2.00	2.00	1.50	1.50	1.50	1.50
49	SPV - 946 x IS 9471	1.00	1.00	1.00	1.50	2.00	1.75	1.25	1.25	1.50	1.38
50	SPV - 104 x IS 2284	1.50	1.00	1.25	1.17	1.50	1.33	1.33	1.33	1.25	1.29
51	SPV - 104 x IS 6335	1.50	1.00	1.25	1.00	1.00	1.00	1.00	1.25	1.00	1.13
52	SPV - 104 x IS 9471	1.50	2.00	1.75	1.50	2.00	1.75	1.50	1.50	2.00	1.75
53	IS 2284 x IS-6335	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00
54	IS 2284 x IS-9471	1.50	1.00	1.25	1.00	1.00	1.00	1.00	1.25	1.00	1.13
55	IS 6335 x IS 9471	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00
Mean		2.26	1.96	2.11	2.41	2.39	2.40	2.34	2.18	2.26	(-)-6.83
% increase (+)/decrease (-) over untreated				(-)13.27			(-)0.83				
% increase over location							12.74 (Aroclor)				
A	SE (m)	0.039	CD 5%		SE (m)	CD 5%	SE (m)	CD 5%			
B	0.209	0.110	0.041	0.115	0.041	0.115	0.043	0.125			
A x B	0.296	0.581	0.603	0.658	0.218	0.658	0.226	0.658			
		0.821	0.821	0.821	0.307	0.821	0.310	0.821			

Table 39. Effect of pre-treatment on Ungermminated seed of parents and F₁ crosses at Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola		Patancheru		Mean		Pooled over locations		
		Untreated	Treated	Untreated	Treated	5	8	Untreated	Treated	Mean
1	SPV-1201	22.00	24.00	23.00	50.67	23.00	48.00	36.33	36.00	36.17
2	ICSB-101B	23.33	20.00	21.67	64.67	21.67	71.33	44.00	45.67	44.83
3	AKma-14B	46.00	31.33	38.67	96.00	38.67	94.67	71.00	67.00	69.00
4	SRT-26B	13.33	14.67	14.00	78.00	14.00	82.00	45.67	48.33	47.00
5	GI-35-15-15	18.67	22.67	20.67	75.33	20.67	88.00	47.00	55.33	51.17
6	SPV-946	12.67	18.67	15.67	68.67	15.67	77.33	40.67	44.33	42.33
7	SPV-104	38.67	42.00	40.33	96.00	40.33	91.33	67.33	66.67	67.00
8	IS-2284	6.67	9.33	8.00	33.33	8.00	18.00	20.00	13.67	16.83
9	IS-6335	6.67	6.67	6.67	22.00	6.67	16.67	14.33	11.67	13.00
10	IS-9471	6.67	6.00	6.33	28.67	6.33	23.33	17.67	14.67	16.17
11	SPV-1201 x ICSB-101B	14.67	12.00	13.33	47.33	13.33	47.33	31.00	29.67	30.33
12	SPV-1201 x AKma-14 B	17.33	14.67	16.00	47.33	16.00	38.67	32.33	26.67	29.50
13	SPV-1201 x SRT-26B	6.00	10.00	8.00	19.33	10.00	17.33	12.67	13.67	13.17
14	SPV-1201 x GI-35-15-15	39.33	5.33	22.33	28.00	22.33	16.00	33.67	10.67	22.17
15	SPV-1201 x SPV-946	34.67	25.33	30.00	32.00	30.00	18.67	33.33	22.00	27.67
16	SPV-1201 x SPV-104	15.33	12.67	14.00	56.67	14.00	38.00	36.00	25.33	30.67
17	SPV-1201 x IS-2284	8.67	4.00	6.33	25.33	6.33	20.67	17.00	12.33	14.67
18	SPV-1201 x IS-6335	4.00	6.67	5.33	30.00	5.33	31.33	17.00	19.00	18.00
19	SPV-1201 x IS-9471	4.67	7.33	6.00	23.33	6.00	27.32	14.00	17.33	15.66
20	ICSB-101B x AKma 14 B	21.33	24.00	22.67	54.67	22.67	57.33	38.00	40.67	39.33
21	ICSB-101B x SRT-26 B	12.00	8.67	10.33	60.67	10.33	58.67	36.33	33.67	35.00
22	ICSB-101B x GI-35-15-15	12.00	6.00	9.00	21.33	9.00	21.33	16.67	15.17	15.17
23	ICSB-101B x SPV-946	8.00	10.67	9.33	41.33	9.33	50.67	24.67	30.67	27.67
24	ICSB-101B x SPV-104	23.33	19.33	21.33	62.67	21.33	54.67	43.00	37.00	40.00
25	ICSB-101B x IS-2284	11.33	9.33	10.33	12.67	10.33	10.67	12.00	10.00	11.00
26	ICSB-101B x IS-6335	10.00	6.67	8.33	6.00	8.33	6.67	8.00	8.33	8.17
27	ICSB-101B x IS-9471	8.00	6.00	7.00	14.00	7.00	10.00	11.00	6.33	8.67
28	AKma-14 B x SRT-26B	28.00	18.00	23.00	74.00	23.00	71.33	51.00	44.67	47.83
29	AKma-14 B x GI-35-15-15	18.00	18.67	18.33	73.33	18.33	67.33	45.67	43.00	44.33
30	AKma-14 B x SPV-946	22.67	19.33	21.00	65.33	21.00	77.33	44.00	48.33	46.17
31	AKma-14 B x SPV-104	46.00	34.00	40.00	76.67	40.00	76.00	61.33	55.00	58.17
32	AKma-14 B x IS-2284	20.00	6.00	13.00	33.33	13.00	17.33	26.67	11.67	19.17
33	AKma-14 B x IS-6335	6.67	2.67	4.67	12.00	4.67	12.00	9.33	7.33	8.33
34	AKma-14 B x IS-9471	12.00	6.00	9.00	19.33	9.00	23.33	15.67	14.67	15.17

Contd.

Table 39. Contd.

	1	2	3	4	5	6	7	8	9	10	11
35	SRT-26B x GI-35-15-15	7.33	10.00	8.67	59.33	59.33	59.33	59.33	33.33	34.67	34.00
36	SRT-26B x SPV-946	16.00	14.00	15.00	36.00	37.33	37.33	36.67	26.00	25.67	25.83
37	SRT-26B x SPV-104	30.67	28.00	29.33	75.33	60.00	60.00	67.67	53.00	44.00	48.50
38	SRT-26B x IS-2284	12.00	21.33	16.67	19.33	4.67	4.67	12.00	15.67	13.00	14.33
39	SRT-26B x IS-6335	10.00	10.67	10.33	11.33	6.67	6.67	9.00	10.67	8.67	9.67
40	SRT-26B x IS-9471	10.00	4.67	7.33	14.00	9.33	9.33	11.67	7.00	7.00	9.50
41	GI-35-15-15 x SPV-946	14.67	12.00	13.33	44.00	26.67	26.67	35.33	29.33	19.33	24.33
42	GI-35-15-15 x SPV-104	16.67	14.00	15.33	49.33	57.33	57.33	53.33	33.00	35.67	34.33
43	GI-35-15-15 x IS-2284	9.33	12.00	10.67	17.33	16.00	16.00	16.67	13.33	14.00	13.67
44	GI-35-15-15 x IS-6335	8.00	10.00	9.00	16.67	12.00	12.00	14.33	12.33	11.00	11.67
45	GI-35-15-15 x IS-9471	7.33	8.00	7.67	26.00	28.67	28.67	27.33	16.67	18.33	17.50
46	SPV-946 x SPV-104	15.33	20.00	17.67	72.67	54.67	54.67	63.67	44.00	37.33	40.67
47	SPV-946 x IS-2284	13.33	7.33	10.33	20.67	16.00	16.00	18.33	17.00	11.67	14.33
48	SPV-946 x IS-6335	6.67	4.67	5.67	22.67	16.67	16.67	19.67	14.67	10.67	12.67
49	SPV-946 x IS-9471	8.00	10.67	9.33	22.00	18.00	18.00	20.00	15.00	14.33	14.67
50	SPV-104 x IS-2284	11.33	10.67	11.00	19.33	12.67	12.67	16.00	15.33	11.67	13.50
51	SPV-104 x IS-6335	7.33	7.00	7.00	15.33	6.67	6.67	11.00	11.33	6.67	9.00
52	SPV-104 x IS-9471	8.00	6.00	7.00	22.00	8.00	8.00	15.00	15.00	7.00	11.00
53	IS-2284 x IS-6335	8.67	7.33	8.00	14.00	14.67	14.67	14.33	11.33	11.00	11.17
54	IS-2284 x IS-9471	12.00	16.00	14.00	22.67	7.33	7.33	15.00	17.33	11.67	14.50
55	IS-6335 x IS-9471	4.00	5.33	4.67	21.33	21.33	21.33	21.33	12.67	13.33	13.00
Mean		15.18	13.24	14.21	39.48	35.90	35.90	37.69	27.33	24.57	25.95
% increase (+)/decrease(-) over untreated				(-)12.77				(-)9.06			(-)10.09
% increase over location								165.24 (Akola)			
		SE (m)	CD 5%		SE (m)	CD 5%			SE (m)	CD 5%	
	A	0.430	0.193		0.631	1.750			0.575	1.673	
	B	2.258	6.258		3.311	9.178			3.019	8.785	
	A x B	3.193	8.850		4.683	12.980			4.270	12.426	

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 Table 40. Effect of pre-treatment on *F. montiforme* (%) (UGS) of parents and F_1 crosses at Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola		Mean		Patancheru		Mean		Pooled over locations	
		Untreated	Treated	3	4	Untreated	Treated	5	6	Untreated	Treated
1	SPV-1201	8.00	8.00			27.33	22.00	24.67	17.67	15.00	16.33
2	ICSB-101B	8.67	8.67	8.00		34.67	22.00	31.67	21.67	15.00	16.33
3	AKms-14B	21.33	16.00	18.67	8.67	56.00	28.00	42.00	38.67	20.67	21.17
4	SRT-26B	5.33	8.00	6.67	6.67	37.33	38.67	38.00	21.33	23.33	30.33
5	GI-35-15-15	6.67	8.67	7.67	7.67	36.00	34.67	35.33	21.33	21.67	21.50
6	SPV-946	6.00	8.00	7.00	7.00	44.67	37.33	41.00	25.33	22.67	24.00
7	SPV-104	18.67	14.67	16.67	16.67	67.33	40.67	54.00	43.00	27.67	35.33
8	IS-2284	4.00	3.33	3.67	3.67	20.00	9.33	14.67	12.00	6.33	9.17
9	IS-6335	2.67	4.00	3.33	3.33	15.33	8.00	11.67	9.00	6.00	7.50
10	IS-9471	2.00	2.67	2.33	2.33	13.33	10.67	12.00	7.67	6.67	7.17
11	SPV-1201 x ICSB-101B	5.33	4.00	4.67	4.67	24.67	16.00	20.33	15.00	10.00	12.50
12	SPV-1201 x AKms-14 B	4.67	5.33	5.00	5.00	29.33	16.00	22.67	17.00	10.67	13.83
13	SPV-1201 x SRT-26B	3.33	4.67	4.00	4.00	11.33	11.33	11.33	7.33	8.00	7.67
14	SPV-1201 x GI-35-15-15	16.67	2.67	9.67	9.67	14.67	10.00	12.33	15.67	6.33	11.00
15	SPV-1201 x SPV-946	19.33	14.67	17.00	17.00	20.67	20.00	22.67	15.00	12.00	17.00
16	SPV-1201 x SPV-104	4.67	4.00	4.33	4.33	25.33	13.33	17.00	20.00	14.00	13.50
17	SPV-1201 x IS-2284	2.67	2.00	2.33	2.33	12.00	10.67	11.33	7.33	6.33	6.83
18	SPV-1201 x IS-6335	2.00	4.00	3.00	3.00	14.00	14.00	14.00	8.00	9.00	8.50
19	SPV-1201 x IS-9471	2.00	2.67	2.33	2.33	8.67	12.00	10.33	5.33	7.33	6.33
20	ICSB-101B x AKms 14 B	6.00	5.33	5.67	5.67	32.00	32.67	32.33	19.00	19.00	19.00
21	ICSB-101B x SRT-26 B	4.00	0.67	2.33	2.33	40.00	32.00	36.00	22.00	16.33	19.17
22	ICSB-101B x GI-35-15-15	4.00	2.67	3.33	3.33	17.33	12.67	15.00	10.67	7.67	9.17
23	ICSB-101B x SPV-946	3.33	2.67	3.00	3.00	24.67	29.33	27.00	14.00	16.00	15.00
24	ICSB-101B x SPV-104	11.33	8.00	9.67	9.67	43.33	32.00	37.67	27.33	20.00	23.67
25	ICSB-101B x IS-2284	6.00	1.33	3.67	3.67	8.00	6.67	7.33	7.00	4.00	5.50
26	ICSB-101B x IS-6335	3.33	2.67	3.00	3.00	4.67	6.00	5.33	4.00	4.33	4.17
27	ICSB-101B x IS-9471	4.00	1.33	2.67	2.67	38.67	33.33	36.00	26.33	19.33	3.83
28	AKms-14 B x SRT-26B	14.00	5.33	9.67	9.67	36.67	24.00	30.33	22.00	14.67	22.83
29	AKms-14 B x GI-35-15-15	7.33	5.33	6.33	6.33	35.33	26.67	31.00	21.67	16.33	18.33
30	AKms-14 B x SPV-946	17.00	6.00	7.00	7.00	44.67	28.67	36.67	31.00	19.67	25.33
31	AKms-14 B x SPV-104	8.33	10.67	14.00	14.00	14.00	9.33	11.67	9.67	6.33	8.00
32	AKms-14 B x IS-2284	5.33	3.33	4.33	4.33	7.33	7.33	7.33	5.33	3.67	4.50
33	AKms-14 B x IS-6335	3.33	0.00	1.67	1.67	14.00	13.33	13.67	9.33	7.67	8.50
34	AKms-14 B x IS-9471	4.67	2.00	3.33	3.33	14.00	13.33	13.67	9.33	7.67	8.50

Contd.

Table 40. Contd.

	1	2	3	4	5	6	7	8	9	10	11
35	SRT- 26B x GI-35-15-15	2.67	6.00	4.33	38.00	26.00	32.00	20.33	16.00	18.17	
36	SRT- 26B x SPV - 946	6.67	6.00	6.33	22.00	21.33	21.67	14.33	13.67	14.00	
37	SRT- 26B x SPV -104	11.33	14.00	12.67	40.00	31.33	35.67	25.67	22.67	24.17	
38	SRT- 26B x IS - 2284	5.33	8.00	6.67	11.33	4.00	7.67	8.33	6.00	7.17	
39	SRT- 26B x IS - 6335	3.33	4.00	3.67	8.67	2.67	5.67	6.00	3.33	4.67	
40	SRT- 26B x IS - 9471	4.00	1.33	2.67	8.67	4.67	6.67	6.33	3.00	4.67	
41	GI-35-15-15 x SPV - 946	9.33	5.33	7.33	30.00	12.00	21.00	19.67	8.67	14.17	
42	GI-35-15-15 x SPV - 104	5.33	5.33	5.33	31.33	28.00	29.67	18.33	16.67	17.50	
43	GI-35-15-15 x IS - 2284	2.67	5.33	4.00	11.33	7.33	9.33	7.00	6.33	6.67	
44	GI-35-15-15 x IS - 6335	4.00	4.67	4.33	10.00	6.00	8.00	7.00	5.33	6.17	
45	GI-35-15-15 x IS - 9471	4.00	4.00	4.00	18.00	8.67	13.33	11.00	6.33	8.67	
46	SPV - 946 x SPV - 104	4.00	7.33	5.67	52.00	36.67	44.33	28.00	22.00	25.00	
47	SPV - 946 x IS - 2284	6.67	4.67	5.67	12.00	8.00	10.00	9.33	6.33	7.83	
48	SPV - 946 x IS- 6335	4.67	1.33	3.00	14.67	9.33	12.00	9.67	5.33	7.50	
49	SPV - 946 x IS 9471	3.33	6.00	4.67	11.33	8.67	10.00	7.33	7.33	7.33	
50	SPV - 104 x IS 2284	4.67	6.00	5.33	8.67	7.33	8.00	6.67	6.67	6.67	
51	SPV - 104 x IS 6335	1.33	2.67	2.00	8.67	4.67	6.67	5.00	3.67	4.33	
52	SPV - 104 x IS 9471	3.33	4.00	3.67	12.67	4.67	8.67	8.00	4.33	6.17	
53	IS 2284 x IS-6335	1.33	4.00	2.67	8.00	8.00	8.00	4.67	6.00	5.33	
54	IS 2284 x IS -9471	4.00	7.33	5.67	12.00	4.00	8.00	8.00	5.67	6.83	
55	IS 6335 x IS 9471	1.33	2.00	1.67	11.33	10.00	10.67	6.33	6.00	6.17	

Table 41. Effect of pre-treatment on *F. pallidorostrum* (%) (UGS) of parents and F₁ crosses at Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola		Patancheru		Mean		Untreated		Treated		Mean		Untreated		Treated		Mean	
		Untreated	Treated	Untreated	Treated	3	4	5	6	7	8	9	10	11	12	13	14	15	16
1	SPV-1201	0.00	1.33	0.67	6.00	6.00	6.00	6.00	6.00	6.00	6.00	6.00	6.00	3.00	3.67	3.33	3.33	3.33	3.33
2	ICSB-101B	0.00	0.00	0.00	9.33	9.33	9.33	9.33	9.33	10.00	9.67	4.67	5.00	4.83	4.83	4.83	4.83	4.83	4.83
3	AKms-14B	0.00	0.67	0.33	8.00	8.00	8.00	8.00	8.00	7.33	7.67	3.67	4.33	4.00	4.00	4.00	4.00	4.00	4.00
4	SRT-26B	0.00	0.00	0.00	8.67	8.67	8.67	8.67	8.67	7.33	8.00	4.33	3.67	4.00	4.00	4.00	4.00	4.00	4.00
5	GI-35-15-15	0.00	0.00	0.00	9.33	9.33	9.33	9.33	9.33	9.00	9.67	4.67	4.33	4.50	4.50	4.50	4.50	4.50	4.50
6	SPV-946	0.00	0.00	0.00	8.67	8.67	8.67	8.67	8.67	12.00	10.33	4.33	6.00	5.17	5.17	5.17	5.17	5.17	5.17
7	SPV-104	0.00	8.00	4.00	10.00	10.00	10.00	10.00	10.00	10.67	10.33	5.00	9.33	7.17	7.17	7.17	7.17	7.17	7.17
8	IS-2284	0.00	0.00	0.00	7.33	7.33	7.33	7.33	7.33	0.00	3.67	3.67	0.00	1.83	1.83	1.83	1.83	1.83	1.83
9	IS-6335	0.00	0.00	0.00	3.33	3.33	3.33	3.33	3.33	0.67	2.00	1.67	0.33	1.00	1.00	1.00	1.00	1.00	1.00
10	IS-9471	0.00	0.67	0.33	4.00	4.00	4.00	4.00	4.00	4.00	4.00	2.00	2.33	2.17	2.17	2.17	2.17	2.17	2.17
11	SPV-1201 x ICSB-101B	0.00	0.00	0.00	8.00	8.00	8.00	8.00	8.00	6.00	7.00	4.00	3.00	3.50	3.50	3.50	3.50	3.50	3.50
12	SPV-1201 x AKms-14 B	0.00	0.00	0.00	8.00	8.00	8.00	8.00	8.00	5.33	6.67	4.00	2.67	3.33	3.33	3.33	3.33	3.33	3.33
13	SPV-1201 x SRT - 26B	0.00	0.67	0.33	2.00	2.00	2.00	2.00	2.00	2.00	2.00	1.00	1.33	1.17	1.17	1.17	1.17	1.17	1.17
14	SPV-1201 x GI-35-15-15	1.33	0.00	0.67	5.33	5.33	5.33	5.33	5.33	2.00	3.67	3.33	1.00	1.33	1.33	1.33	1.33	1.33	1.33
15	SPV-1201 x SPV-946	0.00	0.00	0.00	3.33	3.33	3.33	3.33	3.33	4.00	2.67	2.67	0.00	1.33	1.33	1.33	1.33	1.33	1.33
16	SPV-1201 x SPV-104	0.00	0.00	0.00	6.00	6.00	6.00	6.00	6.00	4.00	4.00	3.00	1.33	2.17	2.17	2.17	2.17	2.17	2.17
17	SPV-1201 x IS-2284	0.00	0.67	0.33	7.33	7.33	7.33	7.33	7.33	3.33	5.33	3.67	1.67	2.00	2.00	2.00	2.00	2.00	2.00
18	SPV-1201 x IS-6335	0.00	0.00	0.00	6.00	6.00	6.00	6.00	6.00	0.67	3.33	3.00	0.33	1.67	1.67	1.67	1.67	1.67	1.67
19	SPV-1201 x IS-9471	0.00	0.00	0.00	8.67	8.67	8.67	8.67	8.67	6.00	7.33	4.33	3.00	3.67	3.67	3.67	3.67	3.67	3.67
20	ICSB - 101B x AKms 14 B	0.00	0.00	0.00	13.33	13.33	13.33	13.33	13.33	6.67	10.00	6.67	3.33	3.67	3.67	3.67	3.67	3.67	3.67
21	ICSB - 101B x SRT-26 B	0.00	0.00	0.00	8.67	8.67	8.67	8.67	8.67	6.00	7.33	4.33	3.00	3.67	3.67	3.67	3.67	3.67	3.67
22	ICSB - 101B x GI-35-15-15	0.00	0.00	0.00	13.33	13.33	13.33	13.33	13.33	6.67	10.00	6.67	3.33	3.67	3.67	3.67	3.67	3.67	3.67
23	ICSB - 101B x SPV - 946	0.00	0.00	0.00	8.67	8.67	8.67	8.67	8.67	7.33	8.00	0.67	0.67	5.00	5.00	5.00	5.00	5.00	5.00
24	ICSB - 101B x SPV - 104	0.00	0.00	0.00	14.67	14.67	14.67	14.67	14.67	7.33	8.00	4.33	3.67	4.00	4.00	4.00	4.00	4.00	4.00
25	ICSB - 101B x IS-2284	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.67	3.33	3.00	0.33	1.67	1.67	1.67	1.67	1.67	1.67
26	ICSB - 101B x IS-6335	0.00	0.00	0.00	2.00	2.00	2.00	2.00	2.00	0.00	0.33	0.00	0.33	0.17	0.17	0.17	0.17	0.17	0.17
27	ICSB - 101B x IS-9471	0.00	0.00	0.00	8.00	8.00	8.00	8.00	8.00	0.00	1.00	1.00	0.00	0.50	0.50	0.50	0.50	0.50	0.50
28	AKms - 14 B x SRT - 26B	0.00	2.00	1.00	11.33	11.33	11.33	11.33	11.33	8.00	9.67	6.00	4.33	5.17	5.17	5.17	5.17	5.17	5.17
29	AKms - 14 B x GI-35-15-15	0.67	0.67	0.67	11.33	11.33	11.33	11.33	11.33	11.33	11.33	5.67	6.33	7.00	7.00	7.00	7.00	7.00	7.00
30	AKms - 14 B x SPV - 946	0.00	1.33	0.67	14.67	14.67	14.67	14.67	14.67	8.00	11.33	7.67	6.33	7.00	7.00	7.00	7.00	7.00	7.00
31	AKms - 14 B x SPV - 104	0.67	0.67	0.67	2.67	2.67	2.67	2.67	2.67	3.33	3.00	2.33	1.67	2.00	2.00	2.00	2.00	2.00	2.00
32	AKms - 14 B x IS-2284	0.00	0.00	0.00	0.67	0.67	0.67	0.67	0.67	0.00	0.33	0.33	0.00	0.17	0.17	0.17	0.17	0.17	0.17
33	AKms - 14 B x IS-6335	0.00	0.00	0.00	3.33	3.33	3.33	3.33	3.33	2.67	3.00	2.00	1.33	1.67	1.67	1.67	1.67	1.67	1.67
34	AKms - 14 B x IS-9471	0.67	0.00	0.33	3.33	3.33	3.33	3.33	3.33	2.67	3.00	2.00	1.33	1.67	1.67	1.67	1.67	1.67	1.67

Concl.

Table 41. Contd.

1	2	3	4	5	6	7	8	9	10	11
35	SRT-26B x GJ-35-15-15	0.00	0.00	0.00	10.00	6.00	8.00	5.00	3.00	4.00
36	SRT-26B x SPV-946	0.00	0.67	0.33	6.00	4.00	5.00	3.00	2.33	2.67
37	SRT-26B x SPV-104	0.67	1.33	1.00	12.00	7.33	9.67	6.33	4.33	5.33
38	SRT-26B x IS-2284	0.00	0.67	0.33	3.33	0.00	1.67	1.67	0.33	1.00
39	SRT-26B x IS-6335	0.00	0.67	0.33	0.00	0.00	0.00	0.00	0.33	0.17
40	SRT-26B x IS-9471	1.33	0.67	1.00	2.00	0.67	1.33	1.67	0.67	1.17
41	GJ-35-15-15 x SPV-946	0.00	0.00	0.00	5.33	2.00	3.67	2.67	1.00	1.83
42	GJ-35-15-15 x SPV-104	1.33	0.00	0.67	6.67	4.67	5.67	4.00	2.33	3.17
43	GJ-35-15-15 x IS-2284	1.33	0.00	0.67	3.33	1.33	2.33	2.33	0.67	1.50
44	GJ-35-15-15 x IS-6335	1.33	0.67	1.00	2.00	0.00	1.00	1.67	0.33	1.00
45	GJ-35-15-15 x IS-9471	0.00	0.00	0.00	4.67	2.67	3.67	2.33	1.33	1.83
46	SPV-946 x SPV-104	0.00	1.33	0.67	4.00	5.33	4.67	2.00	3.33	2.67
47	SPV-946 x IS-2284	2.00	0.67	1.33	4.00	0.67	2.33	3.00	0.67	1.83
48	SPV-946 x IS-6335	0.67	0.00	0.33	4.00	1.33	2.67	2.33	0.67	1.50
49	SPV-946 x IS-9471	0.00	0.00	0.00	4.67	2.67	3.67	2.33	1.33	1.83
50	SPV-104 x IS-2284	1.33	0.67	1.00	4.67	1.33	3.00	3.00	1.00	2.00
51	SPV-104 x IS-6335	0.00	1.33	0.67	2.00	0.67	1.33	1.00	1.00	1.00
52	SPV-104 x IS-9471	0.00	0.00	0.00	3.33	0.00	1.67	1.67	0.00	0.83
53	IS-2284 x IS-6335	1.33	1.33	1.33	1.33	2.00	1.67	1.33	1.67	1.50
54	IS-2284 x IS-9471	3.33	0.67	2.00	4.00	0.00	2.00	3.67	0.33	2.00
55	IS-6335 x IS-9471	0.00	0.00	0.00	0.67	3.33	2.00	0.33	1.67	1.00
Mean		0.36	0.57	0.47	5.71	3.82	4.77	3.04	2.20	2.62
% increase (+)/decrease (-)				(+) 58.33			(-) 33.09			(-) 27.68
over untreated										
% increase over location										
		SE (m)	CD 5%		SE (m)	CD 5%	918.89 (A) (k)	SE (m)	CD 5%	
	A	0.104	0.288		0.306	0.849		0.161	0.869	
	B	0.545	1.511		1.607	4.454		0.842	2.450	
	A x B	0.771	2.136		2.272	6.299		1.191	3.466	

Table 42. Effect of pre-treatment on *C. lunata* (%) (UGS) of parents and F₁ crosses at Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola		Patancheru		Pooled over localities		Mean
		Untreated 3	Treated 4	Untreated 6	Treated 7	Untreated 9	Treated 10	
1								11
1	SPV-1201	10.67	13.33	19.33	19.33	15.00	16.33	15.67
2	ICSB-101B	12.67	8.00	16.67	27.33	14.67	17.67	16.17
3	AKms-14B	19.33	12.67	32.00	54.00	25.67	33.33	29.50
4	SRT-26B	7.33	6.00	31.33	32.67	19.33	19.33	19.33
5	GI-35-15-15	10.67	11.33	26.00	36.00	18.33	23.67	21.00
6	SPV-946	6.00	10.67	15.33	24.67	10.67	17.67	14.17
7	SPV-104	16.00	16.00	18.67	36.00	17.33	26.00	21.67
8	IS-2284	3.33	6.00	6.00	7.33	4.67	6.67	5.67
9	IS-6335	4.00	2.67	2.67	7.33	3.33	5.00	4.17
10	IS-9471	4.00	2.67	3.33	8.00	6.00	6.00	6.00
11	SPV-1201 x ICSB-101B	8.00	8.00	10.67	21.33	9.33	14.67	12.00
12	SPV-1201 x AKms-14 B	10.67	7.33	7.33	17.33	9.00	12.33	10.67
13	SPV-1201 x SRT-26B	2.67	4.67	5.33	4.00	4.00	4.33	4.17
14	SPV-1201 x GI-35-15-15	16.67	2.67	6.00	4.00	11.33	3.33	7.33
15	SPV-1201 x SPV-946	12.67	10.00	6.00	4.67	9.33	7.33	8.33
16	SPV-1201 x SPV-104	7.33	8.00	22.67	12.67	15.00	10.33	12.67
17	SPV-1201 x IS-2284	3.33	2.00	8.00	7.33	5.67	4.67	5.17
18	SPV-1201 x IS-6335	1.33	2.00	8.00	13.33	10.67	7.67	6.17
19	SPV-1201 x IS-9471	2.00	4.00	8.67	12.67	5.33	8.33	6.83
20	ICSB-101B x AKms-14 B	12.00	16.67	5.33	16.67	8.67	12.33	12.67
21	ICSB-101B x SRT-26 B	6.00	6.67	6.33	18.00	12.67	16.67	9.50
22	ICSB-101B x GI-35-15-15	6.67	3.33	2.67	7.33	5.00	5.33	5.00
23	ICSB-101B x SPV-946	4.00	7.33	6.67	13.33	5.33	10.33	7.83
24	ICSB-101B x SPV-104	10.67	10.00	4.67	14.00	7.67	12.00	9.83
25	ICSB-101B x IS-2284	4.00	7.33	4.00	3.33	4.00	5.33	4.67
26	ICSB-101B x IS-6335	3.33	2.67	3.33	2.33	2.33	3.00	2.67
27	ICSB-101B x IS-9471	3.33	4.00	6.00	2.00	4.67	3.00	3.83
28	AKms-14 B x SRT-26B	9.33	9.33	18.00	26.67	13.67	18.00	15.83
29	AKms-14 B x GI-35-15-15	8.00	12.67	21.33	31.33	14.67	22.00	18.33
30	AKms-14 B x SPV-946	12.00	10.00	17.33	34.00	14.67	25.67	18.33
31	AKms-14 B x SPV-104	24.00	18.00	14.00	33.33	19.00	25.67	22.33
32	AKms-14 B x IS-2284	9.33	2.67	14.00	4.67	11.67	3.67	7.67
33	AKms-14 B x IS-6335	3.33	2.67	4.00	4.67	3.67	3.67	3.67
34	AKms-14 B x IS-9471	5.33	4.00	2.00	7.33	3.67	5.67	4.67

Contd.

Table 42 . Contd.

	1	2	3	4	5	6	7	8	9	10	11
SRT - 26B x GJ-35-15-15	4.67	3.33	4.00	7.33	23.33	15.33	6.00	13.33	9.67		
SRT - 26B x SPV - 946	9.33	6.67	8.00	7.33	11.33	9.33	8.33	9.00	8.67		
SRT - 26B x SPV - 104	16.67	11.33	14.00	18.67	20.00	19.33	17.67	15.67	16.67		
SRT - 26B x IS - 2284	8.00	10.67	9.33	7.33	0.67	4.00	7.67	5.67	6.67		
SRT - 26B x IS - 6335	5.33	6.00	5.67	3.33	4.00	3.67	4.33	5.00	4.67		
SRT - 26B x IS - 9471	4.00	2.00	3.00	2.67	4.67	3.67	3.33	3.33	3.33		
GJ-35-15-15 x SPV - 946	4.67	6.67	5.67	8.00	12.00	10.00	6.33	9.33	7.83		
GJ-35-15-15 x SPV - 104	8.67	8.00	8.33	9.33	20.67	15.00	9.00	14.33	11.67		
GJ-35-15-15 x IS - 2284	4.00	6.67	5.33	2.67	6.67	4.67	3.33	6.67	5.00		
GJ-35-15-15 x IS - 6335	2.67	4.67	3.67	4.00	5.33	4.67	3.33	5.00	4.17		
GJ-35-15-15 x IS - 9471	3.33	4.00	3.67	3.33	16.67	10.67	3.33	10.33	6.83		
SPV - 946 x SPV - 104	10.67	10.00	10.33	15.33	14.00	14.67	13.00	12.00	12.50		
SPV - 946 x IS - 2284	4.67	2.00	3.33	4.67	7.33	6.00	4.67	4.67	4.17		
SPV - 946 x IS - 6335	1.33	3.33	2.33	4.00	5.33	4.67	2.67	4.33	3.50		
SPV - 946 x IS 9471	4.67	4.67	4.67	4.00	6.00	5.00	4.33	5.33	4.83		
SPV - 104 x IS 2284	3.33	3.33	3.33	6.00	4.00	5.00	4.67	3.67	4.17		
SPV - 104 x IS 6335	6.00	3.33	4.67	4.00	1.33	2.67	5.00	2.33	3.67		
SPV - 104 x IS 9471	4.67	2.00	3.33	5.33	2.00	3.67	5.00	2.00	3.50		
IS 2284 x IS-6335	6.00	2.00	4.00	4.67	4.00	4.33	5.33	3.00	4.17		
IS 2284 x IS - 9471	4.00	6.67	5.33	6.00	2.67	4.33	5.00	4.67	4.83		
IS 6335 x IS 9471	2.00	3.33	2.67	9.33	8.67	9.00	5.67	6.00	5.83		
Mean	7.25	6.85	6.95	9.53	13.67	11.60	8.39	10.16	9.28		
% increase (+)/decrease (-) over untreated			-18.27			(+)-43.44			(+)-21.09		
% increase over location						66.91 (Akola)					
SE (m)	0.282	CD 5%		SE (m)	CD 5%		SE (m)	CD 5%			
A	1.781			0.428			0.396				
B	4.096			2.250			2.080				
A x B	2.090			3.181			2.941				

Table 43. Effect of pre-treatment on other fungi (%) (UGS) of parents and F₁ crosses at Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola				Patancheru				Pooled over locations			
		Untreated	Treated	Mean	5	Untreated	Treated	Mean	8	Untreated	Treated	Mean	11
1	SPV-1201	3.33	1.33	2.33	2.33	0.00	1.33	0.67	1.67	1.67	1.33	1.50	
2	ICSB-101B	2.00	2.00	2.00	2.00	3.33	1.33	2.33	2.67	2.67	1.67	2.17	
3	AKma-14B	4.67	2.00	3.33	3.33	0.67	4.67	2.67	2.67	2.67	3.33	3.00	
4	SRT-26B	0.67	0.67	0.67	0.67	1.33	3.33	2.33	1.00	1.00	2.00	1.50	
5	GI-35-15-15	1.33	2.67	2.00	2.00	4.00	8.67	6.33	2.67	2.67	5.67	4.17	
6	SPV-946	2.67	0.00	1.33	1.33	0.67	3.33	2.00	1.67	1.67	1.67	1.67	
7	SPV-104	3.33	4.00	3.67	3.67	0.00	4.00	2.00	1.67	1.67	4.00	2.83	
8	IS-2284	0.00	0.00	0.00	0.00	0.00	1.33	0.67	0.67	0.00	0.67	0.33	
9	IS-6335	0.00	0.00	0.00	0.00	0.67	0.67	0.67	0.33	0.33	0.33	0.33	
10	IS-9471	0.67	0.00	0.33	0.33	3.33	0.00	1.67	2.00	2.00	0.00	1.00	
11	SPV-1201 x ICSB-101B	1.33	0.00	0.67	0.67	4.00	4.00	4.00	2.67	2.67	2.00	2.33	
12	SPV-1201 x AKma-14 B	2.00	1.33	1.67	1.67	2.67	0.00	1.33	2.33	2.33	0.67	1.50	
13	SPV-1201 x SRT - 26B	0.00	0.67	0.33	0.33	0.67	0.00	0.33	0.33	0.33	0.33	0.33	
14	SPV-1201 x GI-35-15-15	5.33	0.00	2.67	2.67	2.00	0.00	1.00	3.67	3.67	0.00	1.83	
15	SPV-1201 x SPV-946	4.67	0.67	2.67	2.67	0.00	0.67	0.33	2.33	2.33	0.67	1.50	
16	SPV-1201 x SPV-104	6.00	0.00	3.00	3.00	5.33	1.33	3.33	5.67	5.67	3.33	3.17	
17	SPV-1201 x IS-2284	2.67	0.00	1.33	1.33	0.33	0.67	0.67	0.33	0.33	0.33	0.33	
18	SPV-1201 x IS-6335	0.67	0.00	0.33	0.33	0.67	0.67	0.67	0.67	0.67	0.33	0.50	
19	SPV-1201 x IS - 9471	0.67	0.00	0.33	0.33	0.00	2.00	1.00	0.33	1.00	0.33	0.50	
20	ICSB - 101B x AKma 14 B	2.67	2.67	2.67	2.67	2.00	2.00	2.00	2.33	2.33	2.33	2.33	
21	ICSB - 101B x SRT - 26 B	3.33	2.00	2.67	2.67	0.00	2.00	1.00	1.67	1.67	2.00	0.33	
22	ICSB - 101B x GI-35-15-15	1.33	0.00	0.67	0.67	0.00	0.00	0.00	0.67	0.67	0.00	0.33	
23	ICSB - 101B x SPV - 946	0.67	0.67	0.67	0.67	0.00	0.67	0.33	0.33	0.33	0.67	0.50	
24	ICSB - 101B x SPV - 104	1.33	1.33	1.33	1.33	0.00	1.33	0.67	0.67	0.67	1.33	1.00	
25	ICSB - 101B x IS - 2284	1.33	0.67	1.00	1.00	0.67	0.00	0.33	1.00	1.00	0.33	0.67	
26	ICSB - 101B x IS - 6335	2.67	0.00	1.33	1.33	0.00	0.00	0.00	1.33	1.33	0.00	0.67	
27	ICSB - 101B x IS - 9471	0.67	0.67	0.67	0.67	0.00	0.67	0.33	0.33	0.33	0.67	0.50	
28	AKma - 14 B x SRT - 26B	4.67	1.33	3.00	3.00	5.33	3.33	4.33	5.00	5.00	2.33	3.67	
29	AKma - 14 B x GI-35-15-15	1.33	0.00	0.67	0.67	4.67	4.00	4.33	3.00	3.00	2.00	2.50	
30	AKma - 14 B x SPV - 946	2.67	2.00	2.33	2.33	1.33	5.33	3.33	2.00	2.00	3.67	2.83	
31	AKma - 14 B x SPV - 104	4.00	0.67	2.33	2.33	4.00	6.67	3.33	4.00	4.00	3.67	3.83	
32	AKma - 14 B x IS - 2284	3.33	0.00	1.67	1.67	2.67	0.00	1.33	3.00	3.00	0.00	1.50	
33	AKma - 14 B x IS - 6335	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	
34	AKma - 14 B x IS - 9471	1.33	0.00	0.67	0.67	0.00	0.00	0.00	0.67	0.67	0.00	0.33	

Contd.

Table 43. Contd.

	1	2	3	4	5	6	7	8	9	10	11
35	SRT-26B x GJ-35-15-15	0.00	0.67	0.67	0.33	3.33	4.00	3.67	1.67	2.33	2.00
36	SRT-26B x SPV-946	0.67	0.67	0.67	0.67	0.00	0.67	0.33	0.33	0.67	0.50
37	SRT-26B x SPV-104	2.00	1.33	1.33	1.67	2.67	2.67	2.67	2.33	2.00	2.17
38	SRT-26B x IS-2284	2.00	2.00	2.00	2.00	0.00	0.00	0.00	1.00	1.00	1.00
39	SRT-26B x IS-6335	1.33	0.00	0.67	0.67	0.00	0.00	0.00	0.67	0.00	0.33
40	SRT-26B x IS-9471	0.67	0.67	0.67	0.33	0.00	0.00	0.00	0.33	0.00	0.17
41	GJ-35-15-15 x SPV-946	0.67	0.00	0.00	0.33	0.67	0.67	0.67	0.67	0.33	0.50
42	GJ-35-15-15 x SPV-104	1.33	0.67	0.67	1.00	2.67	4.00	3.33	2.00	2.33	2.17
43	GJ-35-15-15 x IS-2284	1.33	0.00	0.00	0.67	0.00	0.67	0.33	0.67	0.33	0.50
44	GJ-35-15-15 x IS-6335	0.00	0.67	0.67	0.33	0.67	0.67	0.67	0.33	0.67	0.50
45	GJ-35-15-15 x IS-9471	0.00	0.00	0.00	0.00	0.00	0.67	0.33	0.00	0.33	0.17
46	SPV-946 x SPV-104	1.33	0.67	1.00	1.00	1.33	1.33	1.33	1.33	1.00	1.17
47	SPV-946 x IS-2284	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
48	SPV-946 x IS-6335	0.00	0.00	0.00	0.00	0.00	0.67	0.33	0.00	0.33	0.17
49	SPV-946 x IS-9471	0.00	0.00	0.00	0.00	2.00	0.67	1.33	1.00	0.33	0.67
50	SPV-104 x IS-2284	2.00	0.00	0.00	1.00	0.00	0.00	0.00	1.00	0.00	0.50
51	SPV-104 x IS-6335	0.00	0.00	0.00	0.00	0.67	0.00	0.33	0.33	0.00	0.17
52	SPV-104 x IS-9471	0.00	0.00	0.00	0.00	0.00	1.33	0.67	0.00	0.67	0.33
53	IS-2284 x IS-6335	0.00	0.00	0.00	0.00	0.00	0.00	0.33	0.00	0.33	0.17
54	IS-2284 x IS-9471	0.67	1.33	1.33	1.00	1.33	0.67	1.00	1.00	1.00	1.00
55	IS-6335 x IS-9471	0.67	0.00	0.00	0.33	0.00	0.00	0.00	0.33	0.00	0.17
Mean		1.60	0.64	1.12	1.19	1.51	1.35	1.39	1.08	1.24	
% increase (+)/decrease (-) over untreated				(-60.00)			(+26.89)			(-22.30)	
% increase over location							20.53 (Akola)				
		SE (m)	CD 5%		SE (m)	CD 5%		SE (m)		CD 5%	
A		0.120	0.333		0.146	0.405		0.126		0.367	
B		0.631	1.749		0.766	2.125		0.659		1.917	
A x B		0.892	2.473		1.084	3.006		0.932		2.712	

Table 44. Effect of pre-treatment on score (UGS) of parents and F_1 crosses at Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola				Patancheru				Pooled over locations			
		Untreated		Treated		Untreated		Treated		Untreated		Treated	
		3	4	5	6	7	8	9	10	11	12	13	14
1	SPV-1201	4.50	4.00	4.25	5.00	5.00	5.00	4.75	4.75	4.50	4.83	4.50	4.83
2	ICSB-101B	4.67	4.00	4.33	4.83	5.00	4.92	4.75	4.75	4.50	4.83	4.50	4.83
3	AKms-14B	5.00	4.50	4.75	5.00	5.00	5.00	5.00	5.00	4.75	4.88	4.50	4.88
4	SRT-26B	4.00	4.00	4.00	4.50	5.00	4.75	4.25	4.25	4.50	4.38	4.50	4.38
5	GJ-35-15-15	4.00	4.00	4.00	4.50	5.00	4.75	4.25	4.25	4.50	4.38	4.50	4.38
6	SPV-946	4.00	3.50	3.75	4.83	5.00	4.92	4.42	4.42	4.25	4.33	4.50	4.33
7	SPV-104	4.50	4.00	4.25	5.00	5.00	5.00	4.75	4.75	4.50	4.83	4.50	4.83
8	IS-2284	2.00	2.17	2.08	3.50	2.17	2.83	2.75	2.17	2.17	2.46	2.17	2.46
9	IS-6335	1.83	1.50	1.67	2.33	2.83	2.50	2.08	2.17	2.17	2.13	2.17	2.13
10	IS-9471	1.33	1.00	1.17	2.00	3.00	2.50	1.67	2.00	2.00	1.83	2.00	1.83
11	SPV-1201 x ICSB-101B	3.50	3.50	3.50	4.50	4.00	4.25	4.00	4.00	3.75	3.88	4.00	3.88
12	SPV-1201 x AKms-14 B	4.33	4.00	4.17	4.67	4.00	4.33	4.50	4.50	4.00	4.25	4.00	4.25
13	SPV-1201 x SRT-26B	3.17	4.00	3.58	4.50	4.00	4.25	3.83	4.00	4.00	3.92	4.00	3.92
14	SPV-1201 x GJ-35-15-15	4.50	4.00	4.25	4.50	4.00	4.25	4.50	4.50	4.00	4.25	4.00	4.25
15	SPV-1201 x SPV-946	4.50	4.00	4.25	4.50	4.00	4.25	4.50	4.50	4.00	4.25	4.00	4.25
16	SPV-1201 x SPV-104	4.00	3.50	3.75	5.00	4.00	4.50	4.50	4.50	3.75	4.13	4.00	4.13
17	SPV-1201 x IS-2284	0.67	1.17	0.92	3.00	2.00	2.50	1.83	1.71	1.58	1.71	1.58	1.71
18	SPV-1201 x IS-6335	1.17	1.33	1.25	2.83	2.00	2.42	2.00	1.67	1.67	1.83	1.67	1.83
19	SPV-1201 x IS-9471	1.50	1.00	1.25	2.50	2.00	2.25	2.00	1.50	1.50	1.75	1.50	1.75
20	ICSB-101B x AKms 14 B	4.00	4.00	4.00	4.17	5.00	4.58	4.08	4.08	4.50	4.29	4.50	4.29
21	ICSB-101B x SRT-26 B	4.00	3.50	3.75	4.50	5.00	4.75	4.25	4.25	4.25	4.25	4.25	4.25
22	ICSB-101B x GJ-35-15-15	3.67	3.00	3.33	4.00	4.00	4.00	3.83	3.83	3.50	3.67	3.50	3.67
23	ICSB-101B x SPV-946	3.00	4.00	3.50	4.00	4.00	4.00	3.75	4.00	4.00	3.87	4.00	3.87
24	ICSB-101B x SPV-104	4.00	3.00	3.50	4.50	4.00	4.25	4.25	3.50	3.50	3.88	3.50	3.88
25	ICSB-101B x IS-2284	1.83	1.83	1.83	1.50	2.00	1.75	1.67	1.67	1.92	1.79	1.92	1.79
26	ICSB-101B x IS-6335	1.50	1.00	1.25	1.00	2.00	1.50	1.25	1.50	1.50	1.38	1.50	1.38
27	ICSB-101B x IS-9471	2.00	1.50	1.75	1.83	1.00	1.50	1.42	1.92	1.25	1.58	1.25	1.58
28	AKms-14 B x SRT-26B	3.50	3.50	3.50	4.50	4.50	4.50	4.00	4.00	4.00	4.00	4.00	4.00
29	AKms-14 B x GJ-35-15-15	3.50	4.00	3.75	4.50	4.00	4.25	4.00	4.00	4.00	4.00	4.00	4.00
30	AKms-14 B x SPV-946	4.17	4.00	4.08	4.83	5.00	4.92	4.50	4.50	4.50	4.50	4.50	4.50
31	AKms-14 B x SPV-104	4.83	4.33	4.58	4.67	5.00	4.83	4.75	4.75	4.67	4.71	4.67	4.71
32	AKms-14 B x IS-2284	2.00	1.67	1.83	2.00	2.00	2.00	2.00	2.00	1.83	2.00	1.83	2.00
33	AKms-14 B x IS-6335	2.00	0.83	1.42	2.00	2.00	2.00	2.00	2.00	1.42	2.00	1.42	2.00
34	AKms-14 B x IS-9471	3.17	1.50	2.33	2.50	2.00	2.25	2.83	2.83	1.75	2.29	1.75	2.29

Contd.

Table 44 . Contd..

1	2	3	4	5	6	7	8	9	10	11
35	SRT- 26B x GI-35-15-15	3.83	3.50	3.67	4.50	4.00	4.25	4.17	3.75	3.96
36	SRT- 26B x SPV - 946	4.00	3.50	3.75	4.50	4.00	4.25	4.25	3.75	4.00
37	SRT- 26B x SPV - 104	4.00	4.00	4.00	4.50	4.00	4.25	4.25	4.13	4.25
38	SRT- 26B x IS - 2284	2.00	2.00	2.00	2.33	2.33	2.17	2.00	2.17	2.08
39	SRT- 26B x IS - 6335	2.00	2.00	2.00	2.00	2.50	2.25	2.00	2.25	2.13
40	SRT- 26B x IS - 9471	1.83	1.67	1.75	2.17	2.33	2.25	2.00	2.00	2.00
41	GI-35-15-15 x SPV - 946	3.33	2.83	3.08	4.50	5.00	4.75	3.92	3.92	3.92
42	GI-35-15-15 x SPV - 104	3.17	3.00	3.08	4.50	5.00	4.75	3.83	4.00	3.92
43	GI-35-15-15 x IS - 2284	1.00	2.00	1.50	2.33	3.00	2.67	1.67	2.50	2.08
44	GI-35-15-15 x IS - 6335	1.17	2.00	1.58	2.00	3.00	2.50	1.58	2.50	2.08
45	GI-35-15-15 x IS - 9471	1.00	3.00	2.00	2.50	3.00	2.75	1.75	3.00	2.38
46	SPV - 946 x SPV - 104	4.00	3.83	3.92	5.00	5.00	5.00	4.50	4.42	4.46
47	SPV - 946 x IS - 2284	2.00	2.00	2.00	2.00	5.00	3.50	2.00	3.50	2.75
48	SPV - 946 x IS - 6335	1.67	1.67	1.67	2.17	4.00	3.08	1.92	2.83	2.38
49	SPV - 946 x IS 9471	2.00	1.83	1.92	2.00	3.00	2.50	2.00	2.42	2.21
50	SPV - 104 x IS 2284	2.00	2.00	2.00	2.50	3.00	2.75	2.25	2.50	2.38
51	SPV - 104 x IS 6335	2.00	2.00	2.00	2.50	3.00	2.75	2.25	2.50	2.38
52	SPV - 104 x IS 9471	2.50	3.00	2.75	2.50	3.00	2.75	2.50	3.00	2.75
53	IS 2284 x IS-6335	1.67	2.00	1.83	2.00	2.50	2.25	1.83	2.25	2.04
54	IS 2284 x IS -9471	2.00	2.00	2.00	2.50	2.00	2.25	2.25	2.00	2.13
55	IS 6335 x IS 9471	1.17	2.00	1.58	2.00	2.50	2.25	1.58	2.25	1.92
Mean		2.88	2.77	2.83	3.43	3.55	3.49	3.18	3.16	3.17
% increase (+)/decrease(-)				(-33.97)			(+33.38)			(-10.63)

Table 4.5 : Effect of pre-treatment on germination (%) of parents and F₂ progenies at Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola		Patancheru		Pooled over locations				
		Untreated	Treated	Mean	Untreated	Treated	Mean	Untreated	Treated	Mean
1	2	3	4	5	6	7	8	9	10	11
1	SPV-1201	78.00	76.00	77.00	52.00	49.33	50.67	65.00	62.67	63.83
2	ICSB-101B	76.67	80.00	78.33	28.67	35.33	32.00	52.67	57.67	55.17
3	AKms-14B	54.00	68.67	61.33	5.33	4.67	5.00	29.67	36.67	33.17
4	SRT-26B	86.67	85.33	86.00	18.00	22.00	20.00	52.33	53.67	53.00
5	GI-35-15-15	81.33	77.33	79.33	12.00	24.67	18.33	46.67	51.00	48.83
6	SPV-946	87.33	81.33	84.33	22.67	32.67	27.67	55.00	57.00	56.00
7	SPV-104	61.33	58.00	59.67	8.67	4.00	6.33	35.00	31.00	33.00
8	IS-2284	93.33	90.67	92.00	82.00	66.67	74.33	87.67	78.67	83.17
9	IS-6335	93.33	93.33	93.33	83.33	83.33	80.67	88.33	85.67	87.00
10	IS-9471	93.33	94.00	93.67	76.67	71.33	74.00	85.00	82.67	83.83
11	SPV-1201 x ICSB-101B	87.00	91.40	89.20	76.20	79.40	77.80	81.60	85.40	83.50
12	SPV-1201 x AKms-14 B	74.40	79.40	76.90	45.66	51.34	48.50	60.03	65.37	62.70
13	SPV-1201 x SRT-26B	86.80	87.40	87.10	68.66	72.50	70.58	77.73	79.96	78.84
14	SPV-1201 x GI-35-15-15	84.40	84.80	84.60	74.26	71.50	72.88	79.33	78.16	78.74
15	SPV-1201 x SPV-946	83.00	86.00	84.50	63.90	69.50	66.70	73.45	77.76	75.60
16	SPV-1201 x SPV-104	79.20	81.40	80.30	53.34	58.60	55.97	66.27	70.00	68.14
17	SPV-1201 x IS-2284	85.40	90.40	87.90	75.26	84.50	81.96	87.45	88.36	85.39
18	SPV-1201 x IS-6335	89.60	92.20	90.90	75.26	84.50	79.88	82.43	88.36	85.39
19	SPV-1201 x IS-9471	87.80	89.00	88.40	59.50	67.50	63.50	73.65	78.25	75.95
20	ICSB-101B x AKms-14 B	74.34	80.00	77.17	58.34	60.84	59.59	66.34	70.42	68.38
21	ICSB-101B x SRT-26 B	85.00	89.60	87.30	59.16	64.66	61.91	72.08	77.13	74.61
22	ICSB-101B x GI-35-15-15	83.80	89.40	86.60	75.74	77.74	76.74	79.77	83.57	81.67
23	ICSB-101B x SPV-946	75.00	78.80	76.90	65.74	72.76	69.25	70.37	75.78	73.08
24	ICSB-101B x SPV-104	80.80	84.80	82.80	30.00	32.00	31.00	55.40	58.40	56.90
25	ICSB-101B x IS-2284	86.20	91.60	88.90	69.84	77.42	73.63	78.02	84.51	81.27
26	ICSB-101B x IS-6335	87.80	92.20	90.00	79.26	82.60	80.88	83.53	87.35	85.44
27	ICSB-101B x IS-9471	90.00	93.80	91.90	69.00	74.50	71.75	79.50	84.15	81.83
28	AKms-14 B x SRT-26B	75.00	81.60	78.30	18.50	22.50	20.50	46.75	52.05	49.40
29	AKms-14 B x GI-35-15-15	72.00	80.80	76.40	54.50	58.66	56.58	63.25	69.73	66.49
30	AKms-14 B x SPV-946	69.60	77.80	73.70	39.50	38.00	38.75	54.55	57.90	56.23
31	AKms-14 B x SPV-104	52.00	60.56	56.28	30.50	34.00	32.25	41.25	47.28	44.27
32	AKms-14 B x IS-2284	83.60	90.20	86.90	69.50	74.00	71.75	76.55	82.10	79.33
33	AKms-14 B x IS-6335	87.60	92.40	90.00	73.00	76.00	74.50	80.30	84.20	82.25
34	AKms-14 B x IS-9471	85.40	88.80	87.10	60.50	69.00	64.75	72.96	78.90	76.93

Contd.

Table 45. Contd.....

1	2	3	4	5	6	7	8	9	10	11
35	SRT- 26B x GJ-35-15-15	80.20	85.00	82.60	62.00	67.50	64.75	71.10	76.25	73.68
36	SRT- 26B x SPV - 946	81.60	88.60	85.10	70.16	78.60	74.38	75.88	83.60	79.74
37	SRT- 26B x SPV - 104	68.00	73.00	70.50	50.50	55.00	52.75	59.25	64.00	61.63
38	SRT- 26B x IS - 2284	83.80	89.40	86.60	79.50	84.00	81.75	81.65	86.70	84.18
39	SRT- 26B x IS - 6335	78.60	86.60	82.60	78.00	84.26	81.13	78.30	85.43	81.87
40	SRT- 26B x IS - 9471	82.00	87.60	84.80	71.00	79.00	75.00	76.50	83.30	79.90
41	GJ-35-15-15 x SPV - 946	74.20	77.00	75.60	66.00	69.84	67.92	70.10	73.42	71.76
42	GJ-35-15-15 x SPV - 104	71.40	72.20	71.80	32.50	38.00	35.25	51.95	55.10	53.53
43	GJ-35-15-15 x IS - 2284	84.80	89.60	87.20	91.00	92.00	91.50	87.90	90.80	89.35
44	GJ-35-15-15 x IS - 6335	85.40	89.60	87.50	73.84	78.30	76.07	79.62	83.95	81.79
45	GJ-35-15-15 x IS - 9471	72.00	76.74	74.37	71.50	76.60	74.05	71.75	76.67	74.21
46	SPV - 946 x SPV - 104	73.40	75.00	74.20	57.50	63.00	60.25	65.45	69.00	67.23
47	SPV - 946 x IS - 2284	81.00	84.80	82.90	73.16	79.90	76.53	77.08	82.35	79.72
48	SPV - 946 x IS - 6335	90.00	92.80	91.40	88.66	90.34	89.50	89.33	91.57	90.45
49	SPV - 946 x IS 9471	83.00	88.16	85.58	67.20	72.20	69.70	75.10	80.18	77.64
50	SPV - 104 x IS 2284	73.80	77.60	75.70	64.66	76.00	70.33	69.23	76.80	73.02
51	SPV - 104 x IS 6335	70.60	74.10	72.35	58.00	63.50	60.75	64.30	68.80	66.55
52	SPV - 104 x IS 9471	65.40	72.00	68.70	65.00	71.50	68.25	65.20	71.75	68.48
53	IS 2284 x IS-6335	90.00	94.26	92.13	69.00	71.00	70.00	79.50	82.63	81.07
54	IS 2284 x IS -9471	90.50	90.14	90.32	89.50	92.00	90.75	90.00	91.07	90.54
55	IS 6335 x IS 9471	89.64	93.94	91.79	78.00	85.34	81.67	83.82	89.64	86.73
Mean		80.37	83.95	82.16	59.390	63.82	61.87	70.14	73.89	72.02
% increase (+)/decrease(-) over untreated				(+/-) 4.45			(+/-) 6.56			(+/-) 5.35
% increase over localton										
		SE (m)	CD 5%	Palancheru		SE (m)	CD 5%	SE (m)	CD 5%	
		0.833	2.310			1.113	3.086	0.462	1.349	
		A				5.838	16.184	2.425	7.078	
		B				8.257	12.888	3.429	9.978	
		A x B								

Table 46: Effect of pre-treatment on *F. moniliforme* (%) (GS) of parents and F_2 progenies Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola				Patancheru				Pooled over locations			
		Untreated		Treated		Untreated		Treated		Untreated		Treated	
		3	4	5	6	7	8	9	10	11	12	13	14
1	SPV-1201	12.67	10.01	11.34	27.33	16.00	21.67	20.00	13.00	16.50			
2	ICSB-101B	18.01	16.67	17.34	18.00	11.33	14.67	18.00	14.00	16.00			
3	AKma-14B	24.67	17.99	21.33	2.67	2.67	2.67	13.67	10.33	12.00			
4	SRT-26B	20.67	19.33	20.00	14.00	7.33	10.67	17.34	13.33	15.34			
5	GI-35-15-15	11.99	18.67	15.33	16.67	7.33	12.00	14.33	13.00	13.67			
6	SPV-946	12.00	22.67	17.34	18.00	8.67	13.33	15.00	15.67	15.34			
7	SPV-104	20.01	27.33	23.67	4.00	4.67	4.33	12.00	16.00	14.00			
8	IS-2284	15.99	8.67	12.33	28.00	16.00	22.00	22.00	12.34	17.17			
9	IS-6335	5.33	3.33	4.33	32.00	12.67	22.33	18.66	8.00	13.33			
10	IS-9471	3.33	3.33	3.33	20.67	15.33	18.00	12.00	9.33	10.67			
11	SPV-1201 x ICSB-101B	25.86	16.54	21.20	25.66	20.40	23.03	25.76	18.47	22.12			
12	SPV-1201 x AKma-14 B	22.66	17.50	20.08	15.66	16.34	16.00	19.16	16.92	18.04			
13	SPV-1201 x SRT - 26B	26.50	20.50	23.50	22.66	14.16	18.41	24.58	17.33	20.96			
14	SPV-1201 x GI-35-15-15	15.70	11.40	13.55	32.50	17.50	25.00	24.10	14.45	19.28			
15	SPV-1201 x SPV - 946	14.96	10.26	12.61	27.20	19.40	23.30	21.08	14.83	17.96			
16	SPV-1201 x IS-2284	19.90	13.46	16.68	28.34	16.60	22.47	24.12	15.03	19.58			
17	SPV-1201 x IS-9471	21.36	13.10	17.23	20.50	13.50	17.00	20.93	13.30	17.12			
18	SPV-1201 x IS-6335	25.00	18.16	21.58	25.50	17.26	21.38	25.25	17.71	21.48			
19	SPV-1201 x IS - 9471	15.40	9.86	12.63	27.00	17.00	22.00	21.20	13.43	17.32			
20	ICSB - 101B x AKma 14 B	14.16	10.42	12.29	24.00	22.84	23.42	19.08	16.63	17.86			
21	ICSB - 101B x SRT - 26 B	15.20	12.00	13.60	33.50	25.66	29.58	24.35	18.83	21.59			
22	ICSB - 101B x GI-35-15-15	22.20	11.60	16.90	36.50	20.76	23.63	24.35	16.18	20.27			
23	ICSB - 101B x SPV - 946	19.40	14.00	16.70	30.26	22.50	26.38	24.83	18.25	21.54			
24	ICSB - 101B x SPV - 104	21.70	15.30	18.50	18.00	13.00	15.50	19.85	14.15	17.00			
25	ICSB - 101B x IS - 2284	29.40	12.76	21.08	30.10	23.10	26.60	29.75	17.93	23.84			
26	ICSB - 101B x IS - 6335	15.20	7.66	11.43	34.00	35.76	34.88	24.60	21.71	23.16			
27	ICSB - 101B x IS - 9471	23.40	16.80	20.10	24.50	30.00	27.25	23.95	23.40	23.68			
28	AKma - 14 B x SRT - 26B	19.20	14.60	16.90	13.50	17.50	15.50	16.35	16.05	16.20			
29	AKma - 14 B x GI-35-15-15	17.00	13.60	15.30	29.34	20.34	24.84	23.17	16.97	20.07			
30	AKma - 14 B x SPV - 946	15.60	11.40	13.50	18.00	18.00	18.00	16.80	14.70	15.75			
31	AKma - 14 B x SPV - 104	18.60	14.20	16.40	32.50	17.50	18.00	18.56	15.85	17.20			
32	AKma - 14 B x IS - 2284	21.26	14.56	17.91	32.50	21.00	26.75	26.88	17.78	22.33			
33	AKma - 14 B x IS - 6335	22.16	14.00	18.08	23.50	17.00	20.25	22.83	15.50	19.17			
34	AKma - 14 B x IS - 9471	19.80	13.80	16.80	19.00	19.00	19.00	19.40	16.40	17.90			

Contd.

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Table 46. Condit.

	1	2	3	4	5	6	7	8	9	10	11
35	SRT-26B x GI-35-15-15	24.00	18.60	21.30	30.00	28.50	29.25	27.00	23.55	25.28	
36	SRT-26B x SPV-946	12.20	9.66	10.93	28.20	19.20	23.70	20.20	14.43	17.32	
37	SRT-26B x SPV-104	12.00	12.80	12.40	26.00	16.00	21.00	19.00	14.40	16.70	
38	SRT-26B x IS-2284	17.96	13.06	15.51	28.00	23.50	25.75	22.98	18.28	20.63	
39	SRT-26B x IS-6335	15.80	11.74	13.77	36.76	25.00	30.88	26.28	18.37	22.33	
40	SRT-26B x IS-9471	24.60	19.20	21.90	30.66	19.00	24.83	27.63	19.10	23.37	
41	GI-35-15-15 x SPV-946	15.50	11.48	13.49	26.42	17.16	21.79	20.96	14.32	17.64	
42	GI-35-15-15 x SPV-104	18.76	14.66	16.71	21.00	21.00	21.00	19.88	17.83	18.86	
43	GI-35-15-15 x IS-2284	27.00	16.80	21.90	38.00	15.00	26.50	32.50	15.90	24.20	
44	GI-35-15-15 x IS-6335	23.86	13.54	18.70	32.50	26.50	29.55	28.18	20.07	24.13	
45	GI-35-15-15 x IS-9471	22.40	15.06	18.73	24.00	16.00	20.00	23.20	15.53	19.37	
46	SPV-946 x SPV-104	16.40	11.60	14.00	28.00	26.00	27.00	27.00	18.80	20.50	
47	SPV-946 x IS-2284	27.00	13.80	20.40	35.16	20.70	27.93	31.08	17.25	24.17	
48	SPV-946 x IS-6335	15.84	10.16	13.00	26.34	18.00	22.17	21.09	14.08	17.59	
49	SPV-946 x IS-9471	25.60	18.86	22.23	26.00	16.80	21.40	25.80	17.83	21.82	
50	SPV-104 x IS-2284	21.36	14.80	18.08	31.34	23.50	27.42	26.35	19.15	22.75	
51	SPV-104 x IS-6335	29.46	20.16	24.81	27.50	17.00	22.25	28.48	18.58	23.53	
52	SPV-104 x IS-9471	30.26	17.86	24.06	33.50	18.50	26.00	31.88	18.18	25.03	
53	IS-2284 x IS-6335	26.30	18.76	22.53	39.00	28.00	33.50	32.65	23.38	28.02	
54	IS-2284 x IS-9471	27.92	18.34	23.13	39.50	26.50	33.00	33.71	22.42	28.07	
55	IS-6335 x IS-9471	21.50	19.60	20.55	17.00	12.00	14.50	24.25	15.80	20.03	
Mean		19.85	14.47	17.16	24.19	19.84	22.02	22.02	16.96	19.49	
% increase (+) / decrease (-) over untreated				(-)27.10			(-)17.98			(-)22.98	
% increase over location							28.32 (Akola)				
		SE (m)	CD 5%	SE (m)	CD 5%	SE (m)	CD 5%	SE (m)	CD 5%		
A		0.824	2.284	0.734	2.034		2.034	1.348	3.923		
B		4.320	11.975	3.849	10.669		10.669	7.076	20.591		
A x B		6.110	16.936	5.443	15.068		15.068	10.008	29.124		

Table 47 : Effect of pre-treatment on *F. pallidorozeum* (%) (GS) of parents and F₂ progenies Akola and Patancheru, 1996

S.N.	Parent/Crosses	Akola				Patancheru				Pooled over locations			
		Untreated	Treated	Mean	Mean	Untreated	Treated	Mean	Mean	Untreated	Treated	Mean	Mean
1	2	3	4	5	5	6	7	8	8	9	10	11	11
1	SPV-1201	0.00	0.67	0.33	0.33	5.33	6.00	5.67	5.67	2.67	3.33	3.00	3.00
2	ICSD-101B	0.00	4.67	2.33	2.33	3.33	3.33	3.33	3.33	1.67	4.00	2.83	2.83
3	AKms-14B	0.00	5.33	2.67	2.67	0.00	0.00	0.00	0.00	0.00	2.67	1.33	1.33
4	SRT-26B	2.00	4.00	3.00	3.00	4.00	1.33	2.67	2.67	3.00	2.67	2.67	2.67
5	GI-35-15-15	0.00	6.67	3.33	3.33	0.00	4.00	2.00	2.00	0.00	5.33	2.67	2.67
6	SPV-946	0.67	6.00	3.33	3.33	3.33	6.00	4.67	4.67	2.00	6.00	4.00	4.00
7	SPV-104	0.00	6.67	3.33	3.33	0.00	0.00	0.00	0.00	0.00	3.33	1.67	1.67
8	IS-2284	1.33	2.67	2.00	2.00	6.00	12.00	9.00	9.00	3.67	7.33	5.50	5.50
9	IS-6335	2.67	2.67	2.67	2.67	5.33	7.33	6.33	6.33	4.00	5.00	4.50	4.50
10	IS-9471	0.67	0.67	0.67	0.67	6.00	6.00	6.00	6.00	3.33	3.33	3.33	3.33
11	SPV-1201 x ICSD-101B	1.50	2.56	2.03	2.03	2.14	1.46	1.80	1.80	1.82	2.01	1.92	1.92
12	SPV-1201 x AKms-14 B	1.50	0.60	1.05	1.05	1.34	0.50	0.92	0.92	1.42	0.55	0.99	0.99
13	SPV-1201 x SRT - 26B	1.80	1.40	1.60	1.60	5.50	0.84	3.17	3.17	3.65	1.12	2.39	2.39
14	SPV-1201 x GI-35-15-15	1.20	0.40	0.80	0.80	6.50	3.26	4.88	4.88	3.85	1.83	2.20	2.20
15	SPV-1201 x SPV - 946	1.00	0.20	0.60	0.60	5.20	2.40	2.66	2.66	3.10	1.30	2.20	2.20
16	SPV-1201 x SPV - 104	2.50	0.40	1.45	1.45	3.00	2.66	2.83	2.83	2.75	1.53	2.14	2.14
17	SPV-1201 x IS - 2284	1.20	0.50	0.85	0.85	2.96	2.00	2.48	2.48	2.08	1.25	1.67	1.67
18	SPV-1201 x IS - 6335	3.36	1.96	2.66	2.66	6.00	2.00	4.00	4.00	4.68	1.98	3.33	3.33
19	SPV-1201 x IS - 9471	1.56	0.50	1.03	1.03	4.00	3.50	3.75	3.75	2.78	2.00	2.39	2.39
20	ICSD - 101B x AKms 14 B	1.26	0.50	1.63	1.63	1.66	1.66	1.66	1.66	1.46	1.83	1.65	1.65
21	ICSD - 101B x SRT - 26 B	3.40	2.40	2.90	2.90	2.34	0.34	1.34	1.34	2.87	1.37	2.12	2.12
22	ICSD - 101B x GI-35-15-15	1.60	1.60	1.60	1.60	3.00	2.00	2.50	2.50	2.30	1.80	2.05	2.05
23	ICSD - 101B x SPV - 946	2.20	1.80	2.00	2.00	3.50	1.00	2.25	2.25	2.85	1.40	2.13	2.13
24	ICSD - 101B x SPV - 104	3.80	3.70	3.75	3.75	1.50	0.00	0.75	0.75	2.65	1.85	2.25	2.25
25	ICSD - 101B x IS - 2284	3.20	2.16	2.68	2.68	3.10	2.26	2.68	2.68	3.15	2.21	2.68	2.68
26	ICSD - 101B x IS - 6335	1.66	0.70	1.18	1.18	1.76	4.26	3.01	3.01	1.71	2.48	2.10	2.10
27	ICSD - 101B x IS - 9471	3.86	1.10	2.48	2.48	4.16	4.00	4.08	4.08	4.01	2.55	3.28	3.28
28	AKms - 14 B x SRT - 26B	2.20	1.20	1.70	1.70	1.00	0.00	0.50	0.50	1.60	0.60	1.10	1.10
29	AKms - 14 B x GI-35-15-15	0.80	1.60	1.20	1.20	1.50	0.34	0.92	0.92	1.15	0.97	1.06	1.06
30	AKms - 14 B x SPV - 946	0.80	0.40	0.60	0.60	6.50	1.50	4.00	4.00	3.65	0.95	2.30	2.30
31	AKms - 14 B x SPV - 104	1.00	1.00	1.30	1.30	2.50	2.50	2.50	2.50	1.75	2.05	1.90	1.90
32	AKms - 14 B x IS - 2284	1.56	1.86	1.71	1.71	5.00	3.00	4.00	4.00	3.28	2.43	2.86	2.86
33	AKms - 14 B x IS - 6335	1.00	2.34	1.67	1.67	6.00	5.50	5.75	5.75	3.50	3.92	3.71	3.71
34	AKms - 14 B x IS - 9471	1.00	1.50	1.25	1.25	3.00	4.50	3.75	3.75	2.00	3.00	2.50	2.50

Contd.

Table 47. Contd..

1	2	3	4	5	6	7	8	9	10	11
35	SRT- 26B x GJ-35-15-15	3.60	2.60	3.10	3.00	2.00	2.50	3.30	2.30	2.80
36	SRT - 26B x SPV - 946	1.10	1.50	1.30	6.10	1.40	3.75	3.60	1.45	2.53
37	SRT - 26B x SPV -104	1.20	1.20	1.20	3.00	1.50	2.25	2.10	1.35	1.73
38	SRT - 26B x IS - 2284	0.60	1.00	0.80	5.00	0.50	2.75	2.80	0.75	1.78
39	SRT - 26B x IS - 6335	1.80	2.20	2.00	4.00	1.00	2.50	2.90	1.60	2.25
40	SRT - 26B x IS - 9471	2.40	2.20	2.30	3.50	1.00	2.25	2.95	1.60	2.28
41	GJ-35-15-15 x SPV - 946	0.34	0.34	0.34	1.50	1.00	1.25	0.92	0.67	0.80
42	GJ-35-15-15 x SPV- 104	2.86	1.96	2.41	1.00	1.50	1.25	1.93	1.73	1.83
43	GJ-35-15-15 x IS - 2284	4.46	2.14	3.30	8.00	3.00	5.50	6.23	2.57	4.40
44	GJ-35-15-15 x IS - 6335	2.20	0.20	1.20	5.00	3.00	4.00	3.60	1.60	2.60
45	GJ-35-15-15 x IS - 9471	2.92	2.80	2.86	4.76	1.00	2.88	3.84	1.90	2.87
46	SPV - 946 x SPV- 104	2.20	1.40	1.80	5.00	3.00	4.00	3.60	2.20	2.90
47	SPV - 946 x IS - 2284	4.40	2.20	3.30	4.00	2.50	3.25	4.20	2.35	3.28
48	SPV - 946 x IS- 6335	3.42	2.16	2.79	2.66	1.34	2.00	3.04	1.75	2.40
49	SPV - 946 x IS 9471	3.76	3.00	3.38	4.80	1.60	3.20	4.28	2.30	3.29
50	SPV - 104 x IS 2284	2.30	2.40	2.35	3.66	1.00	2.33	2.98	1.70	2.34
51	SPV - 104 x IS 6335	3.70	5.10	4.40	5.00	1.50	3.25	4.35	3.30	3.83
52	SPV - 104 x IS 9471	4.60	2.86	3.73	4.50	1.00	2.75	4.55	1.93	3.24
53	IS 2284 x IS-6335	6.04	3.52	4.78	4.00	3.00	3.50	5.02	3.26	4.14
54	IS 2284 x IS -9471	3.74	0.92	2.33	4.50	0.50	2.50	4.12	0.71	2.42
55	IS 6335 x IS 9471	5.34	4.10	4.72	5.34	2.34	3.84	5.34	3.22	4.28
Mean		2.29	2.18	2.24	3.72	2.40	3.06	2.91	2.29	2.60
% increase (+)/decrease(-) over untreated				(-)4.80			(-)35.48			(-)21.30
% increase over location							36.60 (Akola)			
		SE (m)	CD 5%		SE (m)	CD 5%		SE (m)	CD 5%	
A		0.148	0.409		0.205	0.569		0.216	0.598	
B		0.775	2.147		1.076	2.983		1.133	3.139	
A x B		1.095	3.036		1.522	4.220		1.602	4.441	

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 Table 48. Effect of pre-treatment on *C. lunata* (%) (GS) of parents and F₂ progenies at Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola				Patancheru				Pooled over locations			
		Untreated		Treated		Untreated		Treated		Untreated		Treated	
		3	4	5	6	7	8	9	10	11	12	13	14
1	SPV-1201	45.33	45.36	45.35	18.67	15.33	17.00	32.00	30.35	31.17			
2	ICSB-101B	45.33	39.32	42.33	14.00	11.33	12.67	29.67	25.33	27.50			
3	AKme-14B	24.67	37.36	31.01	2.67	2.00	2.33	13.67	19.68	16.67			
4	SRT-26B	48.67	34.00	41.33	6.00	6.00	6.00	27.33	20.00	23.67			
5	GI-35-15-15	49.33	30.68	40.01	4.67	4.00	4.33	27.00	17.34	22.17			
6	SPV-946	51.33	28.68	40.01	8.67	10.67	9.67	30.00	19.67	24.84			
7	SPV-104	28.67	16.68	22.67	2.67	0.00	1.33	15.67	8.34	12.00			
8	IS-2284	43.33	30.68	37.01	18.00	18.00	18.00	30.67	24.34	27.50			
9	IS-6335	22.00	14.00	18.00	17.33	12.67	15.00	19.67	13.33	16.50			
10	IS-9471	20.00	16.68	18.34	23.33	14.67	19.00	21.67	15.67	18.67			
11	SPV-1201 x ICSB-101B	40.76	29.10	34.93	25.80	29.94	27.87	33.28	29.52	31.40			
12	SPV-1201 x AKme-14 B	31.16	34.30	32.73	14.00	24.16	19.08	22.58	29.23	25.91			
13	SPV-1201 x SRT - 26B	34.90	30.00	32.45	22.16	23.84	23.00	28.53	26.92	27.73			
14	SPV-1201 x GI-35-15-15	51.76	35.60	43.68	26.00	24.50	25.25	38.88	30.05	34.47			
15	SPV-1201 x SPV -946	39.10	35.05	37.08	21.40	23.00	22.20	30.25	29.03	29.64			
16	SPV-1201 x SPV - 104	37.26	33.66	35.46	16.34	25.00	20.67	26.80	29.33	28.07			
17	SPV-1201 x IS -2284	39.90	34.46	37.18	29.26	21.50	25.38	34.58	27.98	31.28			
18	SPV-1201 x IS-6335	32.16	27.05	29.61	30.50	25.26	27.88	31.33	26.16	28.75			
19	SPV-1201 x IS - 9471	36.50	35.60	36.05	20.50	16.00	18.25	28.90	25.80	27.15			
20	ICSB - 101B x AKme 14 B	46.16	40.84	43.50	25.34	18.84	22.09	35.75	29.84	32.80			
21	ICSB - 101B x SRT - 26 B	51.00	37.20	44.10	16.84	14.24	15.54	33.92	25.72	29.82			
22	ICSB - 101B x GI-35-15-15	43.40	29.20	36.30	32.26	27.26	29.76	37.83	28.23	33.03			
23	ICSB - 101B x SPV - 946	40.40	33.60	37.00	20.50	18.50	19.50	30.45	26.05	28.25			
24	ICSB - 101B x SPV - 104	36.60	36.40	36.50	8.50	10.50	9.50	22.55	23.45	23.00			
25	ICSB - 101B x IS - 2284	34.26	31.20	32.73	21.76	20.66	21.21	28.01	25.93	26.97			
26	ICSB - 101B x IS - 6335	51.30	37.36	44.33	16.26	18.76	17.51	33.78	28.06	30.92			
27	ICSB - 101B x IS - 9471	40.16	32.36	36.26	16.34	15.34	15.84	28.25	23.85	26.05			
28	AKme - 14 B x SRT - 26B	43.60	35.60	39.60	3.50	4.50	4.00	23.55	20.05	21.80			
29	AKme - 14 B x GI-35-15-15	44.80	37.60	41.20	20.00	15.00	22.50	32.40	31.30	31.85			
30	AKme - 14 B x SPV - 946	42.60	36.20	39.40	18.00	15.00	16.50	30.30	25.60	27.95			
31	AKme - 14 B x SPV - 104	25.30	25.23	25.23	8.50	8.50	8.50	16.90	16.83	16.87			
32	AKme - 14 B x IS - 2284	32.36	27.66	30.01	18.00	16.00	17.00	25.18	21.83	23.51			
33	AKme - 14 B x IS - 6335	36.50	21.16	28.83	18.00	18.50	18.25	27.25	19.83	23.54			
34	AKme - 14 B x IS - 9471	32.70	20.40	26.55	20.00	21.00	20.50	26.35	20.70	23.53			

Contd.

Table 48. Conid.

1	2	3	4	5	6	7	8	9	10	11
35	SRT-26B x GJ-35-15-15	33.80	29.60	31.70	18.00	18.00	18.00	25.90	23.80	24.85
36	SRT-26B x SPV-946	52.20	41.86	47.03	25.10	20.10	22.60	38.65	30.98	34.82
37	SRT-26B x SPV-104	41.00	34.60	37.80	17.50	16.50	17.00	29.25	25.55	27.40
38	SRT-26B x IS-2284	39.46	32.16	35.81	24.00	19.50	21.75	31.73	25.83	28.78
39	SRT-26B x IS-6335	44.80	36.54	40.67	20.26	17.00	18.63	32.53	26.77	29.65
40	SRT-26B x IS-9471	36.00	27.00	31.50	25.66	24.00	24.83	30.83	25.50	28.17
41	GJ-35-15-15 x SPV-946	35.30	25.74	30.52	22.34	23.52	22.93	28.82	24.63	26.73
42	GJ-35-15-15 x SPV-104	51.16	24.70	37.93	9.00	12.00	10.50	30.08	18.35	24.22
43	GJ-35-15-15 x IS-2284	39.20	27.60	33.40	26.00	32.00	29.00	32.60	29.80	31.20
44	GJ-35-15-15 x IS-6335	31.86	25.80	28.83	23.00	19.50	21.25	27.43	22.65	25.04
45	GJ-35-15-15 x IS-9471	29.40	25.80	27.60	18.76	21.00	19.88	24.08	23.40	23.74
46	SPV-946 x SPV-104	38.20	34.20	36.20	11.50	18.00	14.75	24.85	26.10	25.48
47	SPV-946 x IS-2284	31.80	29.80	30.80	24.84	28.00	26.42	28.32	28.90	28.61
48	SPV-946 x IS-6335	37.26	28.26	32.76	18.66	21.00	19.83	27.96	24.63	26.30
49	SPV-946 x IS-9471	23.80	22.36	23.08	19.20	18.60	18.90	21.50	20.48	20.99
50	SPV-104 x IS-2284	33.56	27.26	30.41	19.00	17.50	18.25	26.28	22.38	24.33
51	SPV-104 x IS-6335	23.70	21.40	22.55	17.00	18.00	17.50	20.35	19.70	20.03
52	SPV-104 x IS-9471	18.52	17.46	17.99	15.00	15.50	15.25	16.76	16.62	16.48
53	IS-2284 x IS-6335	19.60	16.26	17.93	14.00	13.00	13.50	16.80	14.63	15.72
54	IS-2284 x IS-9471	18.50	16.92	17.71	17.00	18.50	17.75	17.75	17.71	17.73
55	IS-6335 x IS-9471	17.00	13.34	15.17	17.66	14.00	15.83	17.33	13.67	15.50
Mean		36.72	29.61	33.17	17.99	17.55	17.77	27.36	23.58	25.47
% increase (+)/decrease (-) over untreated				(-)19.36			(-)12.44			(-)13.81
% increase over location				46.43						
				Palancheru						
		SE (m)	CD 5%	SE (m)	CD 5%	SE (m)	CD 5%	SE (m)	CD 5%	
A		1.092	3.028	0.601	1.666	0.500	1.457	0.500	1.457	
B		5.729	15.890	3.152	8.736	3.710	7.638	3.710	7.638	
A x B		8.102	22.458	4.457	12.354					

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 Table 49. Effect of pre-treatment on other fungi (%) (GS) of parents and F₂ progenies at Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola			Patancheru			Pooled over locations		
		Untreated	Treated	Mean	Untreated	Treated	Mean	Untreated	Treated	Mean
1	SPV-1201	20.00	18.67	19.33	2.00	0.67	1.33	11.00	9.67	10.33
2	ICSB-101B	13.33	12.67	13.00	0.00	2.67	1.33	6.67	7.67	7.17
3	AKms-14B	4.67	8.00	6.33	0.00	0.00	0.00	2.33	4.00	3.17
4	SRT-26B	15.33	16.67	16.00	0.67	0.00	0.33	8.00	8.33	8.17
5	GI-35-15-15	20.00	14.67	17.33	2.00	0.00	1.00	10.00	7.33	8.67
6	SPV-946	23.33	14.00	18.67	2.00	0.00	1.00	12.67	7.00	9.83
7	SPV-104	12.67	7.33	10.00	1.33	0.00	0.67	7.00	3.67	5.33
8	IS-2284	24.67	10.00	17.33	4.00	5.33	4.67	14.33	7.67	11.00
9	IS-6335	20.00	6.67	13.33	6.00	4.67	5.33	13.00	5.67	9.33
10	IS-9471	19.33	6.00	12.67	4.67	4.00	4.33	12.00	5.00	8.50
11	SPV-1201 x ICSB-101B	16.70	16.00	16.35	12.46	19.60	16.03	14.58	17.80	16.19
12	SPV-1201 x AKms-14 B	15.60	21.30	18.45	5.34	7.84	6.59	10.47	14.57	12.52
13	SPV-1201 x SRT-26B	20.10	14.60	17.35	9.32	10.00	9.66	14.71	12.30	13.51
14	SPV-1201 x GI-35-15-15	15.90	13.76	14.83	6.26	11.50	8.88	11.08	12.63	11.86
15	SPV-1201 x SPV-946	17.86	14.10	15.98	10.10	13.98	10.65	13.98	12.65	13.32
16	SPV-1201 x SPV-104	17.00	17.96	17.48	6.00	6.64	6.32	11.50	12.30	11.90
17	SPV-1201 x IS-2284	16.56	16.30	16.43	17.76	11.50	14.63	17.16	13.90	15.53
18	SPV-1201 x IS-6335	16.36	16.30	16.33	10.26	12.76	11.51	13.31	14.53	13.92
19	SPV-1201 x IS-9471	16.20	18.10	17.15	10.00	7.34	8.50	13.10	13.30	13.20
20	ICSB-101B x AKms 14 B	12.26	18.50	15.38	7.34	9.34	8.34	9.80	13.92	11.86
21	ICSB-101B x SRT-26 B	26.60	16.40	21.50	6.00	12.66	9.33	16.30	14.53	15.42
22	ICSB-101B x GI-35-15-15	16.80	19.40	18.10	14.00	12.50	13.25	15.40	15.95	15.68
23	ICSB-101B x SPV-946	11.60	19.40	15.50	13.00	13.76	13.38	12.30	16.58	14.44
24	ICSB-101B x SPV-104	18.50	16.40	17.45	2.00	7.00	4.50	10.25	11.70	10.98
25	ICSB-101B x IS-2284	18.90	20.50	19.70	10.76	8.34	9.56	14.83	14.63	14.73
26	ICSB-101B x IS-6335	14.96	15.60	15.28	14.26	8.00	11.13	14.61	11.80	13.21
27	ICSB-101B x IS-9471	14.40	12.26	13.33	11.66	10.00	10.83	13.03	11.13	12.08
28	AKms-14 B x SRT-26B	9.80	16.60	13.20	3.00	0.50	0.50	5.15	6.85	6.50
29	AKms-14 B x GI-35-15-15	9.80	16.20	13.00	3.00	7.66	5.33	6.40	11.93	9.17
30	AKms-14 B x SPV-946	10.80	15.80	13.30	5.00	5.00	5.00	7.90	10.90	9.40
31	AKms-14 B x SPV-104	7.86	15.16	11.51	1.00	5.00	3.00	4.43	10.08	7.26
32	AKms-14 B x IS-2284	20.10	14.46	17.28	9.00	11.00	10.00	14.56	12.73	13.64
33	AKms-14 B x IS-6335	25.16	14.00	19.58	7.50	7.00	7.25	16.33	10.50	13.42
34	AKms-14 B x IS-9471	17.90	13.50	15.70	13.50	8.00	10.75	15.70	13.23	13.23

Contd.....

Table 50. Effect of pre-treatment on score (GS) of parents and F₂ progenies at Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola				Patancheru				Pooled over locations			
		4		5		6		7		8		9	
		Untreated	Treated	Mean	Mean	Untreated	Treated	Untreated	Treated	Mean	Mean	Untreated	Treated
1	SPV-1201	4.00	3.00	3.50	4.00	4.00	4.00	4.00	4.00	4.00	4.00	4.00	3.75
2	ICSB-101B	4.00	3.00	3.50	4.00	4.00	4.00	4.00	4.00	4.00	4.00	4.00	3.75
3	AKms-14B	5.00	3.50	4.25	4.00	4.33	3.00	4.67	3.00	3.67	4.67	3.25	3.96
4	SRT-26B	3.83	3.00	3.42	4.00	3.33	3.50	3.92	3.50	3.75	3.92	3.25	3.58
5	GI-35-15-15	3.50	3.00	3.25	3.33	3.00	3.50	3.42	3.50	3.42	3.42	3.25	3.33
6	SPV-946	3.83	2.50	3.17	3.00	3.00	3.50	3.25	3.50	3.25	3.42	3.00	3.21
7	SPV-104	4.50	3.50	4.00	4.00	3.33	3.33	3.92	3.33	3.92	3.42	3.42	3.67
8	IS-2284	2.00	1.00	1.50	1.50	1.00	2.00	1.50	2.00	1.50	1.50	1.50	1.50
9	IS-6335	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00
10	IS-9471	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00
11	SPV-1201 x ICSB-101B	3.20	3.00	3.10	3.45	3.45	3.15	3.30	3.33	3.30	3.33	3.08	3.20
12	SPV-1201 x AKms-14B	3.70	3.40	3.55	3.50	3.50	3.92	3.71	3.60	3.66	3.66	3.63	3.63
13	SPV-1201 x SRT-26B	3.60	3.10	3.35	3.40	3.40	2.79	3.10	3.50	2.95	3.22	3.37	3.37
14	SPV-1201 x GI-35-15-15	3.73	3.13	3.43	3.50	3.50	3.13	3.31	3.61	3.45	3.38	3.38	3.16
15	SPV-1201 x SPV-946	3.05	2.70	2.88	3.70	3.70	3.20	3.45	3.33	3.38	3.38	2.88	3.38
16	SPV-1201 x SPV-104	3.75	3.10	3.43	4.00	4.00	2.67	3.00	3.06	3.11	2.79	2.95	2.95
17	SPV-1201 x IS-2284	3.10	2.58	2.84	3.13	3.13	3.00	3.15	2.85	3.13	2.85	2.85	2.85
18	SPV-1201 x IS-6335	3.10	2.88	2.99	3.19	3.19	2.50	2.50	2.50	2.50	2.50	2.50	2.50
19	SPV-1201 x IS-9471	2.98	2.73	2.85	3.75	3.75	2.50	2.50	2.50	2.50	2.50	2.50	2.50
20	ICSB-101B x AKms-14B	3.75	3.00	3.38	3.75	3.75	3.42	3.58	3.75	3.58	3.75	3.58	3.58
21	ICSB-101B x SRT-26B	3.65	2.90	3.28	3.84	3.84	3.00	3.07	3.07	3.07	3.07	3.07	3.07
22	ICSB-101B x GI-35-15-15	3.65	3.00	3.33	3.50	3.50	3.07	3.28	3.58	3.58	3.58	3.58	3.58
23	ICSB-101B x SPV-946	3.85	3.20	3.53	3.50	3.50	3.25	3.38	3.58	3.58	3.58	3.58	3.58
24	ICSB-101B x SPV-104	3.40	3.10	3.25	3.75	3.75	3.00	3.38	3.58	3.58	3.58	3.58	3.58
25	ICSB-101B x IS-2284	3.05	2.85	2.95	2.96	2.96	2.63	2.79	2.79	2.79	2.79	2.79	2.79
26	ICSB-101B x IS-6335	3.05	2.98	3.01	3.07	3.07	2.69	2.88	2.88	2.88	2.88	2.88	2.88
27	ICSB-101B x IS-9471	3.00	2.08	2.54	3.25	3.25	3.25	3.25	3.25	3.25	3.25	3.25	3.25
28	AKms-14B x SRT-26B	3.65	3.00	3.33	3.13	3.13	3.50	3.31	3.39	3.39	3.39	3.39	3.39
29	AKms-14B x GI-35-15-15	3.75	3.00	3.38	3.84	3.84	3.34	3.79	3.79	3.79	3.79	3.79	3.79
30	AKms-14B x SPV-946	4.00	3.30	3.65	3.75	3.75	3.50	3.88	3.88	3.88	3.88	3.88	3.88
31	AKms-14B x SPV-104	4.00	3.40	3.70	3.88	3.88	3.75	3.81	3.81	3.81	3.81	3.81	3.81
32	AKms-14B x IS-2284	2.78	2.17	2.48	3.00	3.00	2.75	2.89	2.89	2.89	2.89	2.89	2.89
33	AKms-14B x IS-6335	2.34	2.17	2.25	3.13	3.13	2.75	2.94	2.94	2.94	2.94	2.94	2.94
34	AKms-14B x IS-9471	2.70	2.40	2.55	3.75	3.75	3.25	3.50	3.50	3.50	3.50	3.50	3.50

Contd.

Table 50. Contd.

1	2	3	4	5	6	7	8	9	10	11
35	SRT- 26B x GJ-35-15-15	3.30	2.80	3.05	3.50	3.10	3.30	3.40	2.95	3.18
36	SRT - 26B x SPV - 946	3.70	2.90	3.30	3.50	2.85	3.18	3.60	2.88	3.24
37	SRT - 26B x SPV - 104	4.00	3.40	3.70	3.88	3.25	3.56	3.94	3.33	3.63
38	SRT - 26B x IS - 2284	3.23	2.58	2.90	3.50	2.38	2.94	3.36	2.48	2.92
39	SRT - 26B x IS - 6335	3.60	2.80	3.20	2.94	2.50	2.72	3.27	2.65	2.96
40	SRT - 26B x IS - 9471	3.00	2.60	2.80	3.25	2.92	3.08	3.13	2.76	2.94
41	GJ-35-15-15 x SPV - 946	3.55	2.92	3.23	4.00	3.23	3.62	3.78	3.07	3.42
42	GJ-35-15-15 x SPV - 104	3.80	3.35	3.58	4.00	3.00	3.50	3.90	3.18	3.54
43	GJ-35-15-15 x IS - 2284	3.38	2.74	3.06	3.50	2.50	3.00	3.44	2.62	3.03
44	GJ-35-15-15 x IS - 6335	2.67	2.40	2.53	3.38	3.05	3.21	3.02	2.73	2.87
45	GJ-35-15-15 x IS - 9471	3.19	2.54	2.86	3.00	2.55	2.78	3.09	2.54	2.82
46	SPV - 946 x SPV - 104	3.80	3.60	3.70	3.63	2.75	3.19	3.71	3.18	3.44
47	SPV - 946 x IS - 2284	3.50	2.70	3.10	3.88	3.15	3.51	3.69	2.93	3.31
48	SPV - 946 x IS - 6335	2.92	2.34	2.63	2.75	3.00	2.88	2.83	2.67	2.75
49	SPV - 946 x IS 9471	2.94	2.30	2.62	3.20	3.00	3.10	3.07	2.65	2.86
50	SPV - 104 x IS 2284	3.87	3.07	3.47	3.84	2.75	3.29	3.85	2.91	3.38
51	SPV - 104 x IS 6335	3.90	3.03	3.46	3.50	3.00	3.25	3.70	3.01	3.36
52	SPV - 104 x IS 9471	3.79	2.97	3.38	2.75	2.75	2.75	3.27	2.86	3.06
53	IS 2284 x IS-6335	2.70	1.88	2.29	3.25	3.25	3.25	2.98	2.56	2.77
54	IS 2284 x IS -9471	2.44	1.75	2.10	2.75	2.50	2.63	2.60	2.13	2.36
55	IS 6335 x IS 9471	2.69	2.13	2.41	1.84	1.67	1.75	2.26	1.90	2.08
Mean		3.33	2.74	3.03	3.32	2.95	3.14	3.33	2.85	3.09
% increase (+)/decrease(-) over untreated				(-)17.71			(-)11.14			(-)14.41
% increase over location						3.63 (Akola)				
	SE (m)	CD 5%		SE (m)	CD 5%		SE (m)	CD 5%		
A	0.041	0.113		0.047	0.131		0.052	0.151		
B	0.214	0.595		0.248	0.688		0.275	0.800		
A x B	0.304	0.841		0.351	0.972		0.388	1.128		

Table 51. Effect of pre-treatment on ungerminated (%) seed of parents and F_2 progenies at Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola		Mean	Patancheru		Mean	Pooled over locations		
		Untreated	Treated		Untreated	Treated		Untreated	Treated	Mean
1	SPV-1201	22.00	24.00	23.00	48.00	50.67	49.33	35.00	37.33	36.17
2	ICSB-101B	23.33	20.00	21.67	71.33	64.67	68.00	47.33	42.33	44.83
3	AKma-14B	46.00	31.33	38.67	94.67	96.00	95.33	70.33	63.67	67.00
4	SRT-26B	13.33	14.67	14.00	82.00	78.00	80.00	47.67	46.33	47.00
5	GI-35-15-15	18.67	22.67	20.67	88.00	75.33	81.67	53.33	49.00	51.17
6	SPV-946	12.67	18.67	15.67	77.33	68.67	73.00	45.00	44.33	44.67
7	SPV-104	38.67	42.00	40.33	91.33	96.00	93.67	65.00	69.00	67.00
8	IS-2284	6.67	9.33	8.00	18.00	33.33	25.67	18.00	21.33	19.67
9	IS-6335	6.67	6.67	6.67	16.67	22.00	19.33	11.67	14.33	13.00
10	IS-9471	6.67	6.00	6.33	23.33	28.67	26.00	15.00	17.33	16.17
11	SPV-1201 x ICSB-101B	13.00	8.60	10.80	19.80	20.60	20.20	16.40	14.60	15.50
12	SPV-1201 x AKma-14 B	25.60	20.60	23.10	54.34	48.74	51.54	39.97	34.67	37.32
13	SPV-1201 x SRT-26B	13.20	12.60	12.90	31.34	27.50	29.42	22.27	20.06	21.16
14	SPV-1201 x GI-35-15-15	15.60	15.20	15.40	25.74	28.50	27.12	20.67	21.85	21.26
15	SPV-1201 x SPV-946	16.80	18.00	17.40	36.10	30.50	33.30	26.45	22.25	24.35
16	SPV-1201 x SPV-104	20.80	14.60	19.70	46.66	41.40	44.03	33.73	30.00	31.87
17	SPV-1201 x IS-2284	14.80	9.60	12.10	21.50	15.50	18.50	12.56	12.56	15.30
18	SPV-1201 x IS-6335	10.40	7.80	9.10	24.74	15.50	20.12	11.66	11.66	14.61
19	SPV-1201 x IS-9471	12.20	11.00	11.60	40.50	32.50	36.50	26.36	21.75	24.05
20	ICSB-101B x AKma-14 B	25.66	20.00	22.83	41.66	39.16	40.41	33.66	29.58	31.62
21	ICSB-101B x SRT-26 B	15.00	10.40	12.70	40.84	35.34	38.09	27.92	22.87	25.40
22	ICSB-101B x GI-35-15-15	16.20	10.60	13.40	24.26	22.26	23.26	20.23	16.43	18.33
23	ICSB-101B x SPV-946	25.00	21.20	23.10	34.26	27.24	30.75	29.63	24.22	26.93
24	ICSB-101B x SPV-104	19.20	15.20	17.20	70.00	68.00	69.00	44.60	41.60	43.10
25	ICSB-101B x IS-2284	13.80	8.40	11.10	30.16	22.58	26.37	21.98	15.49	18.74
26	ICSB-101B x IS-6335	12.20	7.80	10.00	20.74	17.50	19.12	16.47	12.65	14.56
27	ICSB-101B x IS-9471	10.00	6.20	8.10	31.00	25.50	28.25	20.50	15.85	18.18
28	AKma-14 B x SRT-26B	25.00	18.40	21.70	81.50	77.50	79.50	53.25	47.96	50.60
29	AKma-14 B x GI-35-15-15	28.00	19.20	23.60	45.50	41.34	43.42	36.75	30.27	33.51
30	AKma-14 B x SPV-946	30.40	22.20	26.30	60.50	62.00	61.25	45.45	42.10	43.78
31	AKma-14 B x SPV-104	48.00	39.44	43.72	69.50	66.00	67.75	58.75	52.72	55.74
32	AKma-14 B x IS-2284	16.40	11.80	14.10	30.50	26.00	28.25	23.45	18.90	21.18
33	AKma-14 B x IS-6335	12.40	7.60	10.00	27.00	24.00	25.50	19.70	15.80	17.75
34	AKma-14 B x IS-9471	14.60	11.20	12.90	39.50	31.00	35.25	27.05	21.10	24.08

Contd.

	1	2	3	4	5	6	7	8	9	10	11
35	SRT-26B x GI-35-15-15	19.80	15.00	17.40	38.00	33.50	35.75	28.90	24.25	26.58	
36	SRT-26B x SPV-946	18.40	11.40	14.90	29.84	21.40	25.62	24.12	16.40	20.26	
37	SRT-26B x SPV-104	33.60	27.00	30.30	49.50	45.00	47.25	41.55	36.00	38.78	
38	SRT-26B x IS-2284	16.20	10.60	13.40	20.50	16.00	18.25	18.35	13.30	15.83	
39	SRT-26B x IS-6335	21.40	12.40	17.40	22.00	15.74	18.87	21.70	14.57	18.14	
40	SRT-26B x IS-9471	18.00	13.40	15.20	29.00	21.00	25.00	23.50	16.70	20.10	
41	GI-35-15-15 x SPV-946	25.80	23.00	24.40	34.00	30.84	32.42	29.90	26.92	28.41	
42	GI-35-15-15 x SPV-104	28.60	27.80	28.20	67.50	62.00	64.75	48.05	44.90	46.48	
43	GI-35-15-15 x IS-2284	15.20	10.40	12.80	9.00	8.00	8.50	12.10	9.20	10.65	
44	GI-35-15-15 x IS-6335	14.60	10.40	12.50	26.16	21.70	23.93	20.38	16.05	18.22	
45	GI-35-15-15 x IS-9471	28.00	23.26	25.63	28.50	23.40	25.95	28.25	23.33	25.79	
46	SPV-946 x SPV-104	28.60	25.00	26.80	43.50	37.00	40.25	36.05	31.00	33.53	
47	SPV-946 x IS-2284	19.00	15.20	17.10	26.84	20.10	23.47	22.92	17.65	20.29	
48	SPV-946 x IS-6335	10.00	7.20	8.60	11.34	9.66	10.50	10.67	8.43	9.55	
49	SPV-946 x IS 9471	17.00	11.84	14.42	32.80	27.80	30.30	24.90	19.82	22.36	
50	SPV-104 x IS 2284	26.20	22.40	24.30	35.34	24.00	29.67	30.77	23.20	26.99	
51	SPV-104 x IS 6335	29.40	25.90	27.65	42.00	36.50	39.25	35.70	31.20	33.45	
52	SPV-104 x IS 9471	34.60	28.00	31.30	35.00	28.50	31.75	34.80	28.25	31.53	
53	IS 2284 x IS-6335	10.00	5.74	7.87	31.00	29.00	30.00	20.50	17.37	18.94	
54	IS 2284 x IS-9471	9.40	5.86	7.63	10.50	8.00	9.25	9.95	6.93	8.44	
55	IS 6335 x IS 9471	10.40	6.06	8.23	22.00	14.66	18.33	16.20	10.36	13.28	
Mean		19.69	16.02	17.86 (-)18.63	40.05	36.24	38.15 (-)9.51	29.87	26.13	28.02 (-)12.50	
% increase (+)/decrease(-) over untreated											
% increase over location							113.60 (Akola)				
	SE (m)		CD 5%		SE (m)		CD 5%		SE (m)		CD 5%
A	0.830		2.301		1.112		3.081		0.460		1.338
B	4.354		12.067		5.829		16.157		2.418		7.059
A x B	3.156		17.065		8.243		22.860		3.413		9.931

Table 52. Effect of pre-treatment on *F. moniliforme* (%) (UGS) of parents and F₂ progenies at Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola			Patancheru			Pooled over locations		
		Untreated	Treated	Mean	Untreated	Treated	Mean	Untreated	Treated	Mean
1	2	3	4	5	6	7	8	9	10	11
1	SPV-1201	8.00	8.00	8.00	22.00	27.33	24.67	15.00	17.67	16.33
2	ICSB-101B	8.67	8.67	8.67	32.67	34.67	33.67	20.67	21.67	21.17
3	AKms-14B	21.33	16.00	18.67	28.00	56.00	42.00	24.67	36.00	30.33
4	SRT-26B	5.33	8.00	6.67	38.67	37.33	38.00	22.00	22.67	22.33
5	GJ-35-15-15	6.67	8.67	7.67	34.67	36.00	35.33	20.67	22.33	21.50
6	SPV-946	6.00	8.00	7.00	37.33	44.67	41.00	21.67	26.33	24.00
7	SPV-104	18.67	14.67	16.67	40.67	67.33	54.00	29.67	41.00	35.33
8	IS-2284	4.00	3.33	3.67	9.33	20.00	14.67	6.67	11.67	9.17
9	IS-6335	2.67	4.00	3.33	8.00	15.33	11.67	5.33	9.67	7.50
10	IS-9471	2.00	2.67	2.33	10.67	13.33	12.00	6.33	8.00	7.17
11	SPV-1201 x ICSB-101B	6.36	4.40	5.38	12.20	10.34	11.27	9.28	7.37	8.33
12	SPV-1201 x AKms-14 B	9.16	7.60	8.38	29.16	19.50	24.33	19.16	13.55	16.36
13	SPV-1201 x SRT - 26B	6.10	6.30	6.20	23.50	14.16	18.83	14.80	10.23	12.52
14	SPV-1201 x GJ-35-15-15	5.70	4.90	5.30	16.00	13.00	14.50	10.85	8.95	9.90
15	SPV-1201 x SPV - 946	7.76	6.26	7.01	23.00	17.70	20.35	15.38	11.98	13.68
16	SPV-1201 x SPV - 104	8.30	7.86	8.08	30.00	29.26	29.63	19.15	18.56	18.86
17	SPV-1201 x IS - 2284	6.96	5.50	6.23	10.50	7.26	8.88	8.73	6.38	7.56
18	SPV-1201 x IS-6335	5.56	4.16	4.86	14.00	7.76	10.88	9.78	5.96	7.87
19	SPV-1201 x IS - 9471	6.30	4.60	5.45	14.50	12.50	13.50	10.40	8.55	9.48
20	ICSB - 101B x AKms 14 B	7.42	5.50	6.46	22.50	22.34	22.42	14.96	13.92	14.44
21	ICSB - 101B x SRT- 26 B	3.80	2.80	3.30	25.34	24.34	24.84	14.57	13.57	14.07
22	ICSB - 101B x GJ-35-15-15	7.20	4.40	5.80	11.76	10.76	11.26	9.48	7.58	8.53
23	ICSB - 101B x SPV - 946	10.40	7.40	8.90	20.00	13.26	16.63	15.20	10.33	12.77
24	ICSB - 101B x SPV - 104	9.60	7.00	8.30	39.50	39.00	39.25	24.55	23.00	23.78
25	ICSB - 101B x IS - 2284	5.70	4.20	4.95	20.76	15.00	17.88	13.23	9.60	11.42
26	ICSB - 101B x IS - 6335	3.30	2.16	2.73	11.76	12.00	11.88	7.53	7.08	7.31
27	ICSB-101B x IS - 9471	3.96	3.66	3.81	17.50	16.00	16.75	10.73	9.83	10.28
28	AKms - 14 B x SRT - 26B	10.20	8.00	9.10	37.50	33.50	35.50	23.85	20.75	22.30
29	AKms - 14 B x GJ-35-15-15	9.00	7.20	8.10	27.00	24.66	25.83	18.00	15.93	16.97
30	AKms - 14 B x SPV - 946	7.80	6.20	7.00	32.00	40.50	36.25	19.90	23.35	21.63
31	AKms - 14 B x SPV - 104	22.46	16.30	19.38	49.00	38.50	43.75	35.73	27.40	31.57
32	AKms - 14 B x IS - 2284	7.80	6.00	6.90	21.00	17.50	19.25	14.40	11.75	13.08
33	AKms - 14 B x IS - 6335	5.66	3.66	4.66	17.50	14.50	16.00	11.58	9.08	10.33
34	AKms - 14 B x IS - 9471	6.50	5.80	6.15	19.00	15.00	17.00	12.75	10.40	11.58

Contd.

Table 53. Effect of pre-treatment on *F. pallidorozeum* (%)(UGS) of parents and F_2 progenies at Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola			Patancheru			Pooled over locations		
		Untreated	Treated	Mean	Untreated	Treated	Mean	Untreated	Treated	Mean
1	2	3	4	5	6	7	8	9	10	11
1	SPV-1201	0.00	1.33	0.67	6.00	6.00	6.00	3.00	3.67	3.33
2	ICSB-101B	0.00	0.00	0.00	10.00	9.33	9.67	5.00	4.67	4.83
3	AKms-14B	0.00	0.67	0.33	8.00	7.33	7.67	4.00	4.00	4.00
4	SRT-26B	0.00	0.00	0.00	7.33	8.67	8.00	3.67	4.33	4.00
5	GI-35-15-15	0.00	0.00	0.00	8.67	9.33	9.00	4.33	4.67	4.50
6	SPV-946	0.00	0.00	0.00	12.00	8.67	10.33	6.00	4.33	5.17
7	SPV-104	0.00	8.00	4.00	10.67	10.00	10.33	5.33	9.00	7.17
8	IS-2284	0.00	0.00	0.00	0.00	7.33	3.67	0.00	3.67	1.83
9	IS-6335	0.00	0.00	0.00	0.67	3.33	2.00	0.33	1.67	1.00
10	IS-9471	0.00	0.67	0.33	4.00	4.00	4.00	2.00	2.33	2.17
11	SPV-1201 x ICSB-101B	0.00	0.00	0.00	1.20	0.00	0.60	0.60	0.00	0.30
12	SPV-1201 x AKms-14 B	0.26	0.00	0.13	3.00	1.84	2.42	1.63	0.92	1.28
13	SPV-1201 x SRT - 26B	0.00	0.00	0.00	0.66	0.16	0.41	0.33	0.08	0.21
14	SPV-1201 x GI-35-15-15	0.00	0.00	0.00	1.00	0.26	0.63	0.50	0.13	0.32
15	SPV-1201 x SPV -946	0.00	0.00	0.00	1.90	0.50	1.20	0.95	0.25	0.60
16	SPV-1201 x SPV - 104	0.00	0.00	0.00	1.34	3.00	2.17	0.67	1.50	1.09
17	SPV-1201 x IS -2284	0.00	0.00	0.00	0.26	0.00	0.13	0.13	0.00	0.07
18	SPV-1201 x IS-6335	0.00	0.00	0.00	1.00	0.00	0.50	0.50	0.00	0.25
19	SPV-1201 x IS - 9471	0.00	0.00	0.00	2.50	1.00	1.75	1.25	0.50	0.88
20	ICSB - 101B x AKms 14 B	0.34	0.00	0.17	2.66	2.00	2.33	1.50	1.00	1.25
21	ICSB - 101B x SRT - 26 B	0.00	0.00	0.00	2.50	1.00	1.75	1.25	0.50	0.88
22	ICSB - 101B x GI-35-15-15	0.00	0.40	0.20	0.26	0.00	0.13	0.13	0.20	0.17
23	ICSB - 101B x SPV - 946	0.20	0.00	0.10	2.00	0.00	1.00	1.10	0.00	0.55
24	ICSB - 101B x SPV - 104	0.40	0.20	0.30	9.50	3.50	6.50	4.95	1.85	3.40
25	ICSB - 101B x IS - 2284	0.00	0.00	0.00	0.60	0.00	0.30	0.30	0.00	0.15
26	ICSB - 101B x IS - 6335	0.00	0.00	0.00	0.26	0.00	0.13	0.13	0.00	0.07
27	ICSB - 101B x IS - 9471	0.40	0.00	0.20	1.66	0.00	0.83	1.03	0.00	0.52
28	AKms - 14 B x SRT - 26B	0.20	0.00	0.10	5.00	4.00	4.50	2.60	2.00	2.30
29	AKms - 14 B x GI-35-15-15	0.40	0.00	0.20	1.00	2.00	1.50	0.20	1.00	0.60
30	AKms - 14 B x SPV - 946	0.80	0.00	0.40	1.00	2.50	1.75	0.70	1.25	0.98
31	AKms - 14 B x SPV - 104	0.80	0.80	0.80	3.00	4.50	3.75	1.90	2.65	2.28
32	AKms - 14 B x IS - 2284	0.00	0.00	0.00	1.50	0.50	1.00	0.75	0.35	0.55
33	AKms - 14 B x IS - 6335	0.00	0.34	0.17	1.50	1.00	1.25	0.75	0.67	0.71
34	AKms - 14 B x IS - 9471	0.00	0.00	0.00	2.00	2.00	1.00	0.00	1.00	0.50

Contd.

	1	2	3	4	5	6	7	8	9	10	11
35	SRT- 26B x GJ-35-15-15	0.00	0.00	0.00	0.00	1.00	1.00	1.00	0.50	0.50	0.50
36	SRT- 26B x SPV - 946	0.00	0.00	0.00	0.00	0.20	0.00	0.10	0.10	0.00	0.05
37	SRT - 26B x SPV -104	0.20	0.00	0.00	0.10	0.50	0.50	0.50	0.35	0.25	0.30
38	SRT - 26B x IS - 2284	0.00	0.00	0.00	0.00	1.50	0.00	0.75	0.75	0.00	0.38
39	SRT - 26B x IS - 6335	0.40	0.00	0.00	0.20	0.26	0.00	0.13	0.33	0.00	0.17
40	SRT - 26B x IS - 9471	0.40	0.00	0.00	0.20	0.84	0.00	0.42	0.62	0.00	0.31
41	GJ-35-15-15 x SPV - 946	0.28	0.42	0.35	0.66	0.50	0.50	0.58	0.47	0.46	0.47
42	GJ-35-15-15 x SPV - 104	0.00	0.00	0.00	4.00	2.00	2.00	3.00	2.00	1.00	1.50
43	GJ-35-15-15 x IS - 2284	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
44	GJ-35-15-15 x IS - 6335	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
45	GJ-35-15-15 x IS - 9471	0.66	1.86	1.26	0.50	0.50	0.00	0.25	0.58	0.93	0.76
46	SPV - 946 x SPV - 104	0.00	0.00	0.00	1.50	0.50	0.50	1.00	0.75	0.25	0.50
47	SPV - 946 x IS - 2284	0.20	0.00	0.10	0.50	0.50	0.00	0.25	0.35	0.00	0.18
48	SPV - 946 x IS - 6335	0.00	0.00	0.00	0.34	0.00	0.00	0.17	0.17	0.00	0.09
49	SPV - 946 x IS 9471	0.00	0.00	0.00	0.80	0.00	0.00	0.40	0.40	0.00	0.20
50	SPV - 104 x IS 2284	0.26	0.00	0.13	2.50	0.00	0.00	1.25	1.38	0.00	0.89
51	SPV - 104 x IS 6335	1.26	0.50	0.88	3.00	1.50	1.50	2.25	2.13	1.00	1.57
52	SPV - 104 x IS 9471	1.32	1.34	1.33	3.00	3.00	2.00	1.50	2.16	0.67	1.42
53	IS 2284 x IS-6335	0.00	0.00	0.00	4.00	2.00	2.00	3.00	2.00	1.00	1.50
54	IS 2284 x IS -9471	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
55	IS 6335 x IS 9471	0.50	0.00	0.25	0.00	0.00	0.00	0.00	0.25	0.00	0.13
Mean		0.16	0.30	0.23	2.48	2.032	2.26	1.32	1.67	1.50	
% increase (+)/decrease(-)				(+)/87.50			(+)/88.15			(+)/2652	
% increase over location							882.60 (Akola)				
		SE (m)	CD 5%		SE (m)	CD 5%		SE (m)	CD 5%		
A		0.059	0.164		0.228	0.631		1.00	0.291		
B		0.311	0.861		1.193	3.308		0.526	1.535		
A x B		0.439	1.218		1.688	4.678		0.742	2.156		

Table 34. Effect of pre-treatment on *C. lunata* (%) (UGS) of parents and F₂ progenies Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola		Patancheru		Mean		Pooled over locations	
		Untreated	Treated	Untreated	Treated	Untreated	Treated	Untreated	Treated
1	2	3	4	5	6	7	8	9	10
1	SPV-1201	10.67	13.33	12.00	19.33	19.33	19.33	15.00	16.33
2	ICSB-101B	12.67	8.00	10.33	27.33	16.67	22.00	20.00	12.33
3	Akma-14B	19.33	12.67	16.00	54.00	32.00	43.00	36.67	22.33
4	SRT-26B	7.33	6.00	6.67	31.33	31.33	32.00	20.00	18.67
5	GJ-35-15-15	10.67	11.33	11.00	36.00	26.00	31.00	23.33	18.67
6	SPV-946	6.00	10.67	8.33	24.67	15.33	20.00	15.33	21.00
7	SPV-104	16.00	16.00	16.00	36.00	18.67	27.33	26.00	14.17
8	IS-2284	3.33	6.00	4.67	7.33	6.00	6.67	5.33	21.67
9	IS-6335	4.00	2.67	3.33	7.33	2.67	5.00	5.67	6.00
10	IS-9471	4.00	2.67	3.33	9.33	8.00	8.67	5.67	2.67
11	SPV-1201 x ICSB-101B	5.76	3.86	4.81	8.26	9.94	9.10	6.67	5.33
12	SPV-1201 x Akma-14 B	14.20	11.26	12.73	13.66	22.66	18.16	13.93	6.96
13	SPV-1201 x SRT-26B	6.20	5.40	5.80	16.50	10.66	13.58	11.35	16.96
14	SPV-1201 x GJ-35-15-15	9.26	9.70	9.48	6.26	12.76	9.51	7.76	8.03
15	SPV-1201 x SPV-946	8.10	6.66	7.38	10.20	11.50	10.85	7.76	11.23
16	SPV-1201 x SPV-104	11.00	9.50	10.25	11.66	7.00	9.33	11.33	9.08
17	SPV-1201 x IS-2284	6.76	3.40	5.08	7.26	6.26	6.76	7.01	8.25
18	SPV-1201 x IS-6335	4.40	3.36	3.88	8.00	6.76	7.38	6.20	4.83
19	SPV-1201 x IS-9471	5.50	5.76	5.63	20.50	16.00	18.25	13.00	5.06
20	ICSB-101B x Akma-14 B	18.10	13.50	15.80	13.66	14.00	13.83	10.88	5.63
21	ICSB-101B x SRT-26 B	10.80	14.80	12.80	10.84	8.66	9.75	16.88	11.94
22	ICSB-101B x GJ-35-15-15	7.80	5.20	6.50	12.50	11.76	12.13	10.82	13.76
23	ICSB-101B x SPV-946	13.80	13.20	13.50	10.50	11.00	12.13	10.15	11.73
24	ICSB-101B x SPV-104	9.00	7.80	8.40	20.50	11.00	10.75	12.15	9.32
25	ICSB-101B x IS-2284	7.56	4.20	5.88	10.00	7.10	21.25	14.75	12.13
26	ICSB-101B x IS-6335	8.00	5.66	6.83	7.00	7.10	8.55	8.78	14.83
27	ICSB-101B x IS-9471	5.36	2.66	4.01	9.16	5.50	6.25	7.50	5.65
28	Akma-14 B x SRT-26B	13.00	10.00	11.50	34.50	8.34	8.75	7.26	6.54
29	Akma-14 B x GJ-35-15-15	16.40	11.80	14.10	15.50	30.50	32.50	15.50	5.50
30	Akma-14 B x SPV-946	19.20	14.80	17.00	25.50	13.16	14.33	15.25	20.25
31	Akma-14 B x SPV-104	19.70	17.56	18.63	11.50	9.00	21.25	22.35	12.48
32	Akma-14 B x IS-2284	7.36	3.40	5.38	6.50	7.00	10.25	15.60	15.90
33	Akma-14 B x IS-6335	5.84	3.34	4.59	8.00	7.50	6.75	6.92	13.28
34	Akma-14 B x IS-9471	6.60	5.40	6.00	14.50	13.00	13.75	10.55	5.42
									9.20

Contd.

Table 54. Contd.....

1	2	3	4	5	6	7	8	9	10	11
35	SRT-26B x GI-35-15-15	8.80	6.60	7.70	13.00	11.40	12.20	10.90	9.00	9.95
36	SRT-26B x SPV-946	11.10	7.36	9.23	13.56	10.60	12.08	12.33	8.98	10.66
37	SRT-26B x SPV-104	17.60	14.00	15.80	20.00	12.50	16.25	18.80	13.26	16.03
38	SRT-26B x IS-2284	7.90	4.66	6.28	11.00	7.00	9.00	9.45	5.83	7.84
39	SRT-26B x IS-6335	14.54	8.06	11.30	7.50	3.76	5.63	11.02	5.91	8.47
40	SRT-26B x IS-9471	7.20	4.20	5.70	10.16	8.66	9.41	8.68	6.43	7.56
41	GI-35-15-15 x SPV-946	13.20	13.14	13.17	12.16	13.92	13.04	12.68	13.53	13.11
42	GI-35-15-15 x SPV-104	15.20	16.06	15.63	15.50	13.00	14.25	15.35	14.53	14.94
43	GI-35-15-15 x IS-2284	8.72	2.40	5.66	2.00	4.00	3.00	5.36	3.20	4.28
44	GI-35-15-15 x IS-6335	5.20	4.86	5.03	8.50	10.10	9.30	6.85	7.48	7.17
45	GI-35-15-15 x IS-9471	8.50	8.14	8.32	11.50	11.50	11.50	10.00	9.82	9.91
46	SPV-946 x SPV-104	16.40	13.40	14.90	7.50	8.50	8.00	11.95	10.95	11.45
47	SPV-946 x IS-2284	7.00	5.80	6.40	11.82	11.60	11.71	9.41	8.70	9.06
48	SPV-946 x IS-6335	5.16	4.26	4.71	3.50	3.00	3.25	4.33	3.63	3.98
49	SPV-946 x IS-9471	6.46	4.36	5.41	14.20	12.60	13.40	10.33	8.48	9.41
50	SPV-104 x IS-2284	13.46	11.46	12.46	10.16	4.50	7.33	11.81	7.98	9.90
51	SPV-104 x IS-6335	11.16	10.96	11.06	16.50	19.00	17.75	13.83	14.98	14.41
52	SPV-104 x IS-9471	9.52	8.00	8.76	11.50	11.00	11.25	10.51	9.50	10.01
53	IS-2284 x IS-6335	2.60	1.26	1.93	6.00	6.00	6.00	4.30	3.63	3.97
54	IS-2284 x IS-9471	2.42	1.10	1.76	1.50	2.00	1.75	1.96	1.55	1.76
55	IS-6335 x IS-9471	2.26	1.10	1.68	7.66	5.66	6.66	4.96	3.38	4.17
Mean		8.8	7.25	7.77	6.22	4.98	5.60	7.25	6.12	6.69
% increase (+)/decrease(-) over untreated				(-12.44)			(-19.93)			(-15.58)
% increase over location				27.92						
				Palancheru						
		SE (m)	CD 5%		SE (m)	CD 5%		SE (m)	CD 5%	
A		0.534	1.480		0.989	2.740		0.330	0.960	
B		2.799	7.79		5.184	14.370		1.731	5.051	
A x B		3.958	10.972		7.331	20.323		2.449	7.126	

S.N.	Parents/Crosses	Akola				Palamcham				Pooled over locations			
		3		4		5		6		7		8	
		Untreated	Treated	Untreated	Treated	Mean	Untreated	Untreated	Treated	Mean	Untreated	Treated	Mean
1	SPV-1201	3.33	1.33	2.33	1.33	0.00	0.67	2.33	0.67	1.50	2.33	0.67	1.50
2	ICSB-101B	2.00	2.00	2.00	1.33	0.00	0.67	2.33	0.67	1.50	2.33	0.67	1.50
3	AKms-14B	4.67	2.00	3.33	4.67	0.67	3.33	4.67	0.67	3.00	4.67	0.67	3.00
4	SRT-26B	0.67	0.67	0.67	0.67	0.67	0.67	0.67	0.67	0.67	0.67	0.67	0.67
5	GI-35-15-15	1.33	2.67	2.00	8.67	0.67	4.00	8.67	0.67	4.17	8.67	0.67	4.17
6	SPV-946	2.67	0.00	1.33	3.33	0.67	0.67	3.33	0.67	2.00	3.33	0.67	2.00
7	SPV-104	3.33	4.00	3.67	3.33	0.67	2.00	3.67	0.67	2.83	3.67	0.67	2.83
8	IS-2284	0.00	0.00	0.00	1.33	0.00	0.67	1.33	0.00	0.67	1.33	0.00	0.67
9	IS-6335	0.00	0.00	0.00	0.00	0.00	0.67	0.00	0.67	0.33	0.67	0.00	0.33
10	IS-9471	0.67	0.00	0.33	0.00	0.33	0.33	0.00	0.33	0.33	0.33	0.33	0.33
11	SPV-1201 x ICSB-101B	0.70	0.46	0.58	2.34	1.34	1.84	2.34	1.34	1.21	1.84	1.34	1.21
12	SPV-1201 x AKms-14 B	1.90	2.00	1.95	5.00	4.66	4.83	5.00	4.66	3.39	4.83	3.33	3.39
13	SPV-1201 x SRT-26B	0.70	0.90	0.80	4.16	2.16	3.16	4.16	2.16	1.98	3.16	1.53	1.98
14	SPV-1201 x GI-35-15-15	0.70	0.70	0.70	2.00	2.50	2.25	2.00	2.50	1.48	2.25	1.60	1.48
15	SPV-1201 x SPV-946	0.70	0.76	0.73	1.00	0.80	0.90	1.00	0.80	0.82	0.90	0.78	0.82
16	SPV-1201 x SPV-104	1.46	4.00	2.73	3.66	2.00	2.83	3.66	2.00	2.78	3.66	2.00	2.78
17	SPV-1201 x IS-2284	0.90	0.20	0.55	2.76	1.76	2.26	2.76	1.76	1.41	2.26	0.98	1.41
18	SPV-1201 x IS-9471	0.66	0.46	0.56	1.26	1.00	1.13	1.26	1.00	0.85	1.13	0.73	0.85
19	SPV-1201 x IS-9471	0.50	0.70	0.60	2.50	3.00	2.75	2.50	3.00	1.68	2.75	1.85	1.68
20	ICSB-101B x AKms 14 B	0.34	0.50	0.42	3.84	1.16	2.50	3.84	1.16	1.46	2.50	0.83	1.46
21	ICSB-101B x SRT-26 B	0.40	0.40	0.20	2.16	1.34	1.75	2.16	1.34	0.98	1.75	0.67	0.98
22	ICSB-101B x GI-35-15-15	4.60	0.40	2.50	0.26	0.76	0.51	0.26	0.76	0.51	0.51	0.58	0.51
23	ICSB-101B x SPV-946	1.60	0.60	1.10	1.76	1.50	1.63	1.76	1.50	1.37	1.63	1.05	1.37
24	ICSB-101B x SPV-104	0.20	1.20	0.70	4.50	4.00	2.35	4.50	4.00	2.05	2.35	1.75	2.05
25	ICSB-101B x IS-6335	0.80	0.00	0.40	2.34	1.67	1.47	2.34	1.67	1.29	1.47	1.10	1.29
26	ICSB-101B x IS-9471	0.80	0.00	0.40	1.76	1.38	1.28	1.76	1.38	0.89	1.28	0.50	0.89
27	ICSB-101B x IS-9471	0.00	0.00	0.00	2.66	0.34	1.50	2.66	0.34	0.75	1.50	0.17	0.75
28	AKms-14 B x SRT-26B	1.60	0.40	1.00	5.00	7.25	3.30	5.00	7.25	4.13	3.30	4.95	4.13
29	AKms-14 B x GI-35-15-15	2.20	0.20	1.20	3.00	1.83	2.60	3.00	1.83	1.52	2.60	0.43	1.52
30	AKms-14 B x SPV-946	3.00	0.00	2.00	2.00	3.50	5.48	2.00	3.50	2.00	5.48	1.50	2.00
31	AKms-14 B x SPV-104	4.96	3.60	4.28	6.00	1.00	1.25	6.00	1.00	3.89	1.25	2.30	3.89
32	AKms-14 B x IS-2284	1.20	0.00	0.60	1.50	1.00	1.35	1.50	1.00	0.93	1.35	0.50	0.93
33	AKms-14 B x IS-6335	1.00	0.34	0.67	0.00	1.00	0.50	0.00	1.00	0.50	0.50	0.67	0.50
34	AKms-14 B x IS-9471	0.50	0.00	0.25	6.00	1.00	3.50	6.00	1.00	1.88	3.50	0.50	1.88

Contd.

Table 56. Effect of pre-treatment on score (UGS) of parents and F₂ progenies at Akola and Patancheru, 1996

S.N.	Parents/Crosses	Akola-96				Patancheru				Pooled over locations			
		3		4		5		6		7		8	
		Untreated	Treated	Untreated	Treated	Mean	Mean	Untreated	Treated	Untreated	Treated	Mean	Mean
1	SPV-1201	4.50	4.00	4.25	5.00	5.00	5.00	4.75	4.50	4.75	4.50	4.63	4.63
2	ICSB-101B	4.67	4.00	4.33	5.00	4.83	4.83	4.92	4.42	4.83	4.42	4.63	4.63
3	AKms-14B	5.00	4.50	4.75	5.00	5.00	5.00	5.00	4.75	5.00	4.75	4.88	4.88
4	SRT-26B	4.00	4.00	4.00	5.00	4.50	4.50	4.75	4.25	4.50	4.25	4.38	4.38
5	GI-35-15-15	4.00	4.00	4.00	5.00	4.50	4.50	4.75	4.25	4.50	4.25	4.38	4.38
6	SPV-946	4.00	3.50	3.75	5.00	4.83	4.83	4.92	4.17	4.33	4.17	4.63	4.63
7	SPV-104	4.50	4.25	4.25	5.00	5.00	5.00	4.75	4.50	4.75	4.50	4.63	4.63
8	IS-2284	2.00	2.17	2.08	2.17	2.83	2.83	2.08	2.83	2.08	2.83	2.46	2.46
9	IS-6335	1.83	1.50	1.67	2.83	2.33	2.33	2.58	1.92	2.33	1.92	2.13	2.13
10	IS-9471	1.33	1.00	1.17	3.00	2.00	2.00	2.17	1.50	2.17	1.50	1.83	1.83
11	SPV-1201 x ICSB-101B	4.05	3.80	3.93	4.54	4.49	4.49	4.29	4.14	4.29	4.14	4.22	4.22
12	SPV-1201 x AKms-14 B	4.75	4.40	4.58	5.00	5.00	5.00	4.88	4.70	4.88	4.70	4.79	4.79
13	SPV-1201 x SRT-26B	4.50	4.00	4.25	4.63	4.25	4.25	4.44	4.13	4.56	4.13	4.34	4.34
14	SPV-1201 x GI-35-15-15	4.83	4.18	4.50	4.75	4.38	4.38	4.56	4.28	4.79	4.28	4.53	4.53
15	SPV-1201 x SPV-946	4.00	3.60	3.80	4.65	4.60	4.60	4.67	4.10	4.63	4.10	4.21	4.21
16	SPV-1201 x SPV-104	4.70	4.25	4.48	5.00	4.34	4.34	4.85	4.29	4.85	4.29	4.57	4.57
17	SPV-1201 x IS-2284	3.98	3.08	3.53	3.75	3.75	3.75	3.86	3.60	3.86	3.60	3.73	3.73
18	SPV-1201 x IS-6335	4.00	3.68	3.84	4.00	3.44	3.44	3.72	3.56	3.72	3.56	3.78	3.78
19	SPV-1201 x IS-9471	3.90	3.58	3.74	4.75	4.00	4.00	4.38	3.79	4.33	3.79	4.06	4.06
20	ICSB-101B x AKms-14 B	4.75	3.84	4.29	5.00	4.34	4.34	4.67	4.09	4.88	4.09	4.48	4.48
21	ICSB-101B x SRT-26 B	3.90	3.90	4.28	5.00	4.25	4.25	4.63	4.08	4.83	4.08	4.45	4.45
22	ICSB-101B x GI-35-15-15	4.65	4.60	4.30	5.00	4.13	4.13	4.56	4.06	4.80	4.06	4.43	4.43
23	ICSB-101B x SPV-946	4.65	4.20	4.43	5.00	4.25	4.25	4.63	4.23	4.83	4.23	4.53	4.53
24	ICSB-101B x SPV-104	4.40	4.10	4.25	4.88	4.31	4.31	4.64	3.93	4.28	3.93	4.28	4.28
25	ICSB-101B x IS-2284	3.85	3.85	3.98	4.15	3.88	3.88	4.01	3.86	4.13	3.86	3.99	3.99
26	ICSB-101B x IS-6335	4.10	3.98	4.04	4.25	4.00	4.00	4.18	3.86	4.10	3.86	4.02	4.02
27	ICSB-101B x IS-9471	3.95	2.99	3.47	4.25	4.34	4.34	4.29	3.66	4.10	3.66	3.88	3.88
28	AKms-14 B x SRT-26B	5.00	4.00	4.50	5.00	5.00	5.00	5.00	4.75	5.00	4.75	4.88	4.88
29	AKms-14 B x GI-35-15-15	4.70	4.00	4.35	4.84	4.50	4.50	4.67	4.25	4.77	4.25	4.51	4.51
30	AKms-14 B x SPV-946	5.00	4.40	4.70	5.00	5.00	5.00	5.00	4.70	5.00	4.70	4.85	4.85
31	AKms-14 B x SPV-104	5.00	4.40	4.70	5.00	4.75	4.75	4.88	4.58	5.00	4.58	4.79	4.79
32	AKms-14 B x IS-2284	3.78	3.20	3.49	4.50	4.00	4.00	4.25	3.60	4.14	3.60	3.87	3.87
33	AKms-14 B x IS-6335	3.17	2.84	3.00	4.50	4.25	4.25	4.38	3.42	3.83	3.42	3.63	3.63
34	AKms-14 B x IS-9471	3.70	3.65	3.68	4.75	4.25	4.25	4.50	3.95	4.23	3.95	4.09	4.09

Table 20. Contd.

1	2	3	4	5	6	7	8	9	10	11
35	SRT-26B x GI-35-15-15	4.30	3.80	4.05	4.50	4.60	4.55	4.40	4.20	4.30
36	SRT-26B x SPV-946	4.70	3.90	4.30	4.75	4.20	4.48	4.73	4.05	4.39
37	SRT-26B x SPV-104	5.00	4.40	4.70	5.00	4.25	4.63	5.00	4.33	4.66
38	SRT-26B x IS-2284	4.23	3.58	3.90	4.25	3.75	4.00	4.24	3.66	3.95
39	SRT-26B x IS-6335	4.40	3.80	4.10	4.07	3.50	3.78	4.23	3.65	3.94
40	SRT-26B x IS-9471	4.05	3.60	3.83	4.34	4.59	4.46	4.19	4.09	4.14
41	GI-35-15-15 x SPV-946	4.60	3.80	4.20	5.00	4.53	4.76	4.80	4.16	4.48
42	GI-35-15-15 x SPV-104	4.70	3.90	4.30	5.00	4.50	4.75	4.85	4.20	4.53
43	GI-35-15-15 x IS-2284	4.23	3.39	3.81	4.50	3.50	4.00	4.37	3.44	3.90
44	GI-35-15-15 x IS-6335	3.12	3.10	3.11	4.46	4.50	4.48	3.79	3.80	3.79
45	GI-35-15-15 x IS-9471	4.18	3.39	3.78	4.07	4.25	4.16	4.12	3.82	3.97
46	SPV-946 x SPV-104	4.85	4.55	4.70	4.50	4.00	4.25	4.68	4.28	4.48
47	SPV-946 x IS-2284	4.50	3.55	4.03	4.58	4.80	4.69	4.54	4.18	4.36
48	SPV-946 x IS-6335	3.67	2.65	3.16	3.84	4.00	3.92	3.75	3.33	3.54
49	SPV-946 x IS-9471	3.99	2.95	3.47	4.05	4.10	4.08	4.02	3.53	3.77
50	SPV-104 x IS-2284	4.59	4.25	4.42	4.84	3.88	4.36	4.71	4.06	4.39
51	SPV-104 x IS-6335	4.33	4.13	4.23	4.50	4.00	4.25	4.41	4.06	4.24
52	SPV-104 x IS-9471	4.78	3.97	4.37	4.13	4.00	4.06	4.45	3.98	4.22
53	IS-2284 x IS-6335	3.50	2.88	3.19	4.25	4.50	4.38	3.88	3.69	3.78
54	IS-2284 x IS-9471	3.44	2.38	2.91	3.75	3.50	3.63	3.59	2.94	3.27
55	IS-6335 x IS-9471	3.69	2.57	3.13	3.17	2.67	2.92	3.43	2.62	3.02
Mean		4.16	3.61	3.89	4.48	4.18	4.33	4.32	3.90	4.11
% increase (+)/decrease (-) over untreated				(-)/13.22			(-)/6.96			(-)/9.72
% increase over location							11.31 (Akola)			
		SE (m)	CD 5%		SE (m)	CD 5%		SE (m)	CD 5%	
A		0.050	0.139		0.039	0.110		0.047	0.248	
B		0.263	0.729		0.209	0.579		0.248	0.723	
A x B		0.372	1.216		0.296	0.750		0.350	1.020	

Table 57. Effect of pre-treatment on germination (%) of parents and F₁ crosses at Akola 1995 and Akola 1996

S.N.	Parents/Crosses	Akola-1995		Akola-1996		Mean		Mean		Pooled over locations	
		Untreated	Treated	Untreated	Treated	3	4	5	6	Untreated	Treated
1	SPV-1201	97.78	95.56	96.67	78.00	96.67	76.00	77.00	87.89	85.78	86.84
2	ICSB-101B	74.45	90.00	82.23	76.67	82.23	80.00	78.33	75.56	85.00	80.28
3	AKma-14B	51.11	56.56	53.34	54.00	53.34	54.00	61.33	52.56	62.11	57.33
4	SRT-26B	73.34	68.89	71.11	86.67	71.11	85.33	86.00	80.00	77.11	78.56
5	CU-35-15-15	97.78	92.23	95.00	81.33	95.00	77.33	79.33	89.56	84.78	87.17
6	SPV-946	71.11	67.78	72.78	87.33	72.78	81.33	79.33	82.56	74.56	78.56
7	SPV-104	71.11	82.23	76.67	61.33	76.67	58.00	59.67	66.22	70.11	68.17
8	IS-2284	87.78	91.12	89.45	93.33	89.45	90.67	92.00	90.56	90.89	90.72
9	IS-6335	97.78	96.67	97.23	93.33	97.23	93.33	93.33	95.56	95.00	95.28
10	IS-9471	97.78	96.67	97.23	93.33	97.23	93.33	93.33	95.56	95.00	95.28
11	SPV-1201 x ICSB-101B	67.78	78.89	73.34	82.67	73.34	85.33	86.67	76.56	89.56	83.06
12	SPV-1201 x AKma-14 B	64.45	73.89	73.89	82.67	73.89	85.33	84.00	75.22	82.11	78.67
13	SPV-1201 x SRT-26B	96.67	97.78	97.23	94.67	97.23	94.67	77.67	78.67	86.67	82.96
14	SPV-1201 x CU-35-15-15	92.23	94.45	93.34	65.33	93.34	74.67	70.00	78.78	96.22	87.45
15	SPV-1201 x SPV-946	67.78	75.56	71.67	84.67	71.67	87.33	86.00	76.22	81.45	78.84
16	SPV-1201 x IS-2284	82.23	96.67	96.12	91.33	96.12	96.00	93.67	93.45	96.34	94.89
17	SPV-1201 x IS-6335	80.00	88.89	84.45	95.33	84.45	92.67	94.00	89.11	92.78	90.96
18	SPV-1201 x IS-9471	85.56	78.89	72.23	78.67	72.23	76.00	77.33	72.11	77.45	74.78
19	ICSB-101B x AKma-14 B	35.56	62.23	48.89	88.00	48.89	91.33	89.67	61.78	76.78	69.28
20	ICSB-101B x SRT-26 B	58.89	78.89	68.89	88.00	68.89	94.00	91.00	73.45	86.45	79.96
21	ICSB-101B x CU-35-15-15	52.23	80.00	71.11	76.67	71.11	89.33	90.67	77.11	84.67	80.89
22	ICSB-101B x SPV-946	71.11	72.23	71.67	76.67	71.67	80.67	78.67	73.89	76.45	75.17
23	ICSB-101B x SPV-104	94.45	95.56	95.00	88.67	95.00	90.67	89.67	91.56	93.11	92.34
24	ICSB-101B x IS-2284	92.23	94.45	93.34	90.00	93.34	93.33	91.67	93.89	93.11	92.50
25	ICSB-101B x IS-6335	92.23	94.45	93.34	92.00	93.34	92.00	93.00	92.11	94.22	93.17
26	AKma-14 B x SRT-26B	76.67	81.12	78.89	82.00	78.89	81.33	77.00	74.34	81.56	77.96
27	AKma-14 B x SPV-946	61.11	74.45	67.78	77.33	67.78	81.33	81.67	71.56	77.89	74.72
28	AKma-14 B x CU-35-15-15	78.89	86.67	82.78	53.33	82.78	66.00	59.67	65.00	84.00	81.06
29	AKma-14 B x SPV-104	76.67	83.34	80.00	80.00	80.00	81.33	79.33	79.45	90.34	84.89
30	AKma-14 B x IS-2284	90.00	93.34	91.67	93.33	91.67	97.33	96.33	91.67	95.34	93.50
31	AKma-14 B x IS-6335	84.45	90.00	87.23	88.00	87.23	94.00	91.00	86.22	92.00	89.11
32	AKma-14 B x IS-9471										
33											
34											

Contd.

Table 57. Contd.

	1	2	3	4	5	6	7	8	9	10	11
35	SRT- 26B x GI-35-15-15		93.34	95.56	94.45	92.67	90.00	91.33	93.00	92.78	92.89
36	SRT- 26B x SPV- 946		84.45	87.78	86.12	84.00	86.00	85.00	84.22	86.89	85.56
37	SRT- 26B x SPV- 104		92.23	96.67	94.45	69.33	72.00	70.67	80.78	84.34	82.56
38	SRT- 26B x IS - 2284		84.45	82.23	83.34	88.00	78.67	83.33	86.22	80.45	83.34
39	SRT- 26B x IS - 6335		78.89	84.45	81.67	90.00	89.33	89.67	84.45	86.89	85.67
40	SRT- 26B x IS - 9471		82.23	84.45	83.34	90.00	95.33	92.67	86.11	89.89	88.00
41	GI-35-15-15 x SPV- 946		86.67	94.45	90.56	85.33	88.00	86.67	86.00	91.22	88.61
42	GI-35-15-15 x SPV- 104		45.56	54.45	50.00	83.33	86.00	84.67	64.45	70.22	67.33
43	GI-35-15-15 x IS - 2284		85.56	85.56	85.56	90.67	88.00	89.33	88.11	86.78	87.45
44	GI-35-15-15 x IS - 6335		96.67	98.89	97.78	92.00	90.00	91.00	94.34	94.45	94.39
45	GI-35-15-15 x IS - 9471		78.89	82.23	80.56	92.67	92.00	92.33	85.78	87.11	86.45
46	SPV- 946 x SPV- 104		51.11	55.56	53.34	84.67	80.00	82.33	67.89	67.78	67.83
47	SPV- 946 x IS - 2284		96.67	98.89	97.78	86.67	92.67	89.67	91.67	95.78	93.72
48	SPV- 946 x IS- 6335		97.78	97.78	97.78	93.33	95.33	94.33	95.56	96.56	96.06
49	SPV- 946 x IS 9471		90.00	90.00	90.00	92.00	89.33	90.67	91.00	89.67	90.34
50	SPV- 104 x IS 2284		81.12	86.67	83.89	88.67	89.33	89.00	84.89	88.00	86.45
51	SPV - 104 x IS 6335		61.11	70.00	65.56	92.67	93.33	93.00	76.89	81.67	79.28
52	SPV - 104 x IS 9471		83.34	85.56	84.45	92.00	94.00	93.00	87.67	89.78	88.72
53	IS 2284 x IS-6335		90.00	98.89	94.45	91.33	92.67	92.00	90.67	95.78	93.22
54	IS 2284 x IS -9471		91.12	97.78	94.45	88.00	84.00	86.00	89.56	90.89	90.22
55	IS 6335 x IS 9471		93.34	97.78	95.56	96.00	94.67	95.33	94.67	96.72	95.45

VIII

Table 58. Effect of pre-treatment on *F. moniliforme* (%) (GS) of parents and F_1 crosses at Akola 1995 and Akola 1996

S.N.	Parents/Crosses	Akola-1995				Akola-1996				Pooled over years			
		3		4		5		6		7		8	
		Untreated	Treated	Untreated	Treated	Mean	Mean	Untreated	Treated	Untreated	Treated	Mean	Mean
1	SPV-1201	6.67	12.61	9.74	12.67	10.01	11.34	9.67	11.41	10.54	11		
2	ICSB-101B	24.45	22.22	23.33	18.01	16.67	17.34	16.67	19.45	20.34			
3	AKma-14B	4.44	4.44	4.44	24.67	17.99	21.33	14.56	11.22	12.89			
4	SRT-26B	21.11	13.33	17.22	20.67	19.33	20.00	20.89	16.33	18.61			
5	GJ-35-15-15	0.00	2.22	1.11	11.99	18.67	15.33	6.00	10.45	8.22			
6	SPV-946	12.22	11.11	11.67	12.00	22.67	17.34	12.11	16.89	14.50			
7	SPV-104	2.22	5.56	3.89	20.01	27.33	23.67	11.11	16.44	13.78			
8	IS-2284	0.00	3.33	1.67	15.99	8.67	12.33	8.00	6.00	7.00			
9	IS-6355	1.11	1.11	1.11	5.33	3.33	4.33	3.22	2.22	2.72			
10	IS-9471	2.22	1.11	1.67	3.33	3.33	3.33	2.78	2.22	2.50			
11	SPV-1201 x ICSB-101B	12.22	6.67	9.44	12.67	9.34	11.01	12.45	8.00	10.23			
12	SPV-1201 x AKma-14 B	12.22	8.89	10.56	8.67	10.01	9.34	10.45	9.45	9.95			
13	SPV-1201 x SRT-26B	7.78	7.78	7.78	12.01	12.67	12.34	9.89	10.22	10.06			
14	SPV-1201 x GJ-35-15-15	10.00	11.11	10.56	12.67	11.34	15.33	11.33	11.23	11.28			
15	SPV-1201 x SPV-946	20.00	7.78	13.89	16.66	13.99	15.33	18.33	10.89	14.61			
16	SPV-1201 x SPV-104	12.22	10.00	11.11	10.00	9.99	10.00	11.11	10.00	10.55			
17	SPV-1201 x IS-2284	3.33	4.44	3.89	5.34	3.33	4.34	4.34	3.89	4.11			
18	SPV-1201 x IS-6355	4.44	4.44	4.44	5.34	3.99	4.67	4.89	4.22	4.56			
19	SPV-1201 x IS-9471	6.67	5.56	6.11	6.66	4.67	5.67	6.66	5.11	5.89			
20	ICSB-101B x AKma 14 B	10.00	7.78	8.89	13.33	9.99	11.66	11.67	8.89	10.28			
21	ICSB-101B x SRT-26 B	6.67	8.89	7.78	7.99	5.34	6.67	7.33	7.11	7.22			
22	ICSB-101B x GJ-35-15-15	10.00	6.67	8.33	5.33	14.01	9.67	7.67	10.34	9.00			
23	ICSB-101B x SPV-946	5.56	11.11	8.33	11.33	12.01	11.67	8.44	11.56	10.00			
24	ICSB-101B x SPV-104	23.33	15.56	19.45	22.67	10.67	16.67	23.00	13.11	18.06			
25	ICSB-101B x IS-2284	5.56	4.44	5.00	8.66	3.99	6.33	7.11	4.22	5.66			
26	ICSB-101B x IS-6355	10.00	5.56	7.78	6.66	5.34	5.00	8.33	5.45	6.89			
27	ICSB-101B x IS-9471	3.33	3.33	1.67	6.67	3.33	5.00	5.00	1.67	3.34			
28	AKma-14 B x SRT-26B	52.22	26.67	39.45	14.67	14.67	14.67	33.45	20.67	27.06			
29	AKma-14 B x GJ-35-15-15	15.56	8.89	12.22	15.99	11.33	13.66	15.77	10.11	12.94			
30	AKma-14 B x SPV-946	10.00	5.56	7.78	9.33	9.99	9.66	9.67	7.77	8.72			
31	AKma-14 B x SPV-104	15.56	13.33	14.45	17.99	12.67	15.33	16.77	13.00	14.89			
32	AKma-14 B x IS-2284	10.00	5.56	7.78	10.67	4.67	7.34	10.00	5.11	7.56			
33	AKma-14 B x IS-6355	7.78	5.56	6.67	10.67	5.34	8.01	9.23	5.45	7.34			
34	AKma-14 B x IS-9471	5.56	2.22	3.89	12.01	4.67	8.34	8.78	3.45	6.11			

Contd.

Table 59. Effect of pre-treatment on *F. pallidorostrum* (%) (GS) of parents and F₁ crosses at Akola 1995 and Akola 1996

S.N.	Parents/Crosses	Akola-1995			Akola -1996			Pooled over locations		
		Untreated	Treated	Mean	Untreated	Treated	Mean	Untreated	Treated	Mean
1	2	3	4	5	6	7	8	9	10	11
1	SPV-1201	0.00	0.00	0.00	0.00	0.67	0.33	0.00	0.33	0.17
2	ICSB-101B	0.00	0.00	0.00	0.00	4.67	2.33	0.00	2.33	1.17
3	AKma-14B	0.00	0.00	0.00	0.00	5.33	2.67	0.00	2.67	1.33
4	SRT-26B	0.00	0.00	0.00	0.00	4.00	3.00	1.00	2.00	1.50
5	GJ-35-15-15	0.00	0.00	0.00	0.00	6.67	3.33	0.00	3.33	1.67
6	SPV-946	0.00	0.00	0.00	0.67	6.00	3.33	0.33	3.00	1.87
7	SPV-104	0.00	0.00	0.00	0.00	6.67	3.33	0.00	3.33	1.67
8	IS-2284	0.00	0.00	0.00	1.33	2.67	2.00	0.67	1.33	1.00
9	IS-6335	0.00	0.00	0.00	2.67	2.67	2.67	1.33	1.33	1.33
10	IS-9471	0.00	0.00	0.00	0.67	0.67	0.67	0.33	0.33	0.33
11	SPV-1201 x ICSB-101B	0.00	0.00	0.00	1.33	0.00	0.67	0.67	0.00	0.33
12	SPV-1201 x AKma-14 B	0.00	0.00	0.00	0.00	2.67	1.33	0.00	1.33	0.67
13	SPV-1201 x SRT-26B	0.00	0.00	0.00	0.00	2.00	1.00	0.00	1.00	0.50
14	SPV-1201 x GJ-35-15-15	0.00	0.00	0.00	0.00	2.67	1.33	0.00	1.33	0.67
15	SPV-1201 x SPV-946	0.00	0.00	0.00	0.00	0.67	0.33	0.00	0.33	0.17
16	SPV-1201 x SPV-104	0.00	0.00	0.00	0.00	0.67	0.33	0.00	0.33	0.17
17	SPV-1201 x IS-2284	1.11	0.00	0.00	0.00	0.67	0.33	0.00	0.33	0.17
18	SPV-1201 x IS-6335	0.00	0.00	0.00	0.96	2.00	1.00	1.56	1.00	0.78
19	SPV-1201 x IS-9471	0.00	0.00	0.00	0.00	2.00	1.00	1.00	1.00	0.50
20	ICSB-101B x AKma 14 B	0.00	0.00	0.00	2.00	0.67	1.33	1.00	0.33	0.67
21	ICSB-101B x SRT-26 B	0.00	0.00	0.00	1.33	0.67	1.00	0.67	0.33	0.50
22	ICSB-101B x GJ-35-15-15	0.00	0.00	0.00	0.00	1.33	0.67	0.00	0.67	0.33
23	ICSB-101B x SPV-946	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
24	ICSB-101B x SPV-104	0.00	0.00	0.00	2.67	1.33	2.00	1.33	0.67	1.00
25	ICSB-101B x IS-2284	0.00	0.00	0.00	0.00	2.67	1.33	0.00	1.33	0.67
26	ICSB-101B x IS-6335	0.00	0.00	0.00	0.00	2.67	1.33	0.00	1.33	0.67
27	ICSB-101B x IS-9471	0.00	0.00	0.00	0.67	0.67	0.67	0.33	0.33	0.33
28	AKma-14 B x SRT-26B	0.00	0.00	0.00	1.33	0.00	0.67	0.67	0.00	0.33
29	AKma-14 B x GJ-35-15-15	0.00	0.00	0.00	1.33	0.67	1.00	0.67	0.33	0.50
30	AKma-14 B x SPV-946	0.00	0.00	0.00	3.33	1.33	2.33	1.67	0.67	1.17
31	AKma-14 B x SPV-104	0.00	0.00	0.00	0.00	0.67	0.33	0.00	0.33	0.17
32	AKma-14 B x IS-2284	0.00	0.00	0.00	4.00	2.00	3.00	2.00	1.00	1.50
33	AKma-14 B x IS-6335	0.00	0.00	0.00	4.00	1.33	2.67	2.00	0.67	1.33
34	AKma-14 B x IS-9471	0.00	0.00	0.00	3.33	2.00	2.67	1.67	1.00	1.33
					1.33	0.67	1.00	0.67	0.33	0.50

Contd.

Table 99. Contd.

1	2	3	4	5	6	7	8	9	10	11
35	SRT - 26B x GI-35-15-15	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
36	SRT - 26B x SPV - 946	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
37	SRT - 26B x SPV - 104	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
38	SRT - 26B x IS - 2284	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
39	SRT - 26B x IS - 6335	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
40	SRT - 26B x IS - 9471	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
41	GI-35-15-15 x SPV - 946	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
42	GI-35-15-15 x SPV - 104	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
43	GI-35-15-15 x IS - 2284	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
44	GI-35-15-15 x IS - 6335	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
45	GI-35-15-15 x IS - 9471	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
46	SPV - 946 x SPV - 104	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
47	SPV - 946 x IS - 2284	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
48	SPV - 946 x IS - 6335	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
49	SPV - 946 x IS 9471	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
50	SPV - 104 x IS 2284	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
51	SPV - 104 x IS 6335	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
52	SPV - 104 x IS 9471	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
53	IS 2284 x IS-6335	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
54	IS 2284 x IS-9471	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
55	IS 6335 x IS 9471	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
Mean		0.02	0.00	0.01	1.42	2.12	1.77	0.72	1.06	0.89
% increase (+)/decrease (-)				(-)/100.00			(+49.29			(+47.00
over untreated										
% increase over location							176.00			
							(Alole 95)			
		SE (m)	CD 5%	SE (m)	CD 5%	SE (m)	CD 5%	SE (m)	CD 5%	
		0.014	0.039	0.169	0.168	0.118	0.343	0.118	0.343	
		0.075	0.208	0.885	2.454	0.623	1.818	0.623	1.818	
		0.106	0.294	1.252	3.471	0.881	2.564	0.881	2.564	
		A								
		B								
		A x B								

Table 60. Effect of pre-treatment on *C. lunata* (%) (GS) of parents and F₁ crosses at Akola 1995 and Akola 1996

S.N.	Parents/Crosses	Akola-1995				Akola-1996				Pooled over locations			
		4		5		6		7		8		9	
		Untreated	Treated	Mean	Mean	Untreated	Treated	Mean	Mean	Untreated	Treated	Mean	Mean
1	SPV-1201	1.11	7.78	4.44	45.33	45.36	45.36	45.36	45.36	23.22	26.57	24.90	24.90
2	ICSB-101B	28.89	27.78	28.33	45.33	39.32	37.11	33.53	37.11	33.53	33.53	33.53	33.53
3	AKms-14B	6.67	20.00	13.33	24.67	37.36	31.01	34.33	31.01	15.67	28.68	22.17	22.17
4	SRT-26B	20.00	16.67	18.33	48.67	34.00	41.33	40.01	41.33	28.00	25.33	26.11	26.11
5	GI-35-15-15	6.67	17.78	12.22	49.33	30.68	40.01	37.33	40.01	22.12	25.01	23.56	23.56
6	SPV-946	23.33	15.56	19.45	51.33	28.68	40.01	37.33	40.01	22.12	25.01	23.56	23.56
7	SPV-104	35.56	33.34	34.45	28.67	16.68	32.11	25.01	32.11	26.45	27.11	26.78	26.78
8	IS-2284	12.22	22.22	17.22	43.33	30.68	37.01	33.84	37.01	13.78	15.89	14.83	14.83
9	IS-6335	5.56	17.78	11.67	22.00	14.00	18.00	16.00	18.00	13.89	15.56	14.73	14.73
10	IS-9471	7.78	14.45	11.11	20.00	48.67	47.67	48.00	47.67	33.22	33.90	33.56	33.56
11	SPV-1201 x ICSB-101B	17.78	21.11	19.45	48.67	40.00	45.33	41.45	45.33	26.67	34.06	30.36	30.36
12	SPV-1201 x AKms-14 B	32.22	13.33	22.78	48.00	34.00	46.34	42.33	46.34	32.90	37.62	35.26	35.26
13	SPV-1201 x SRT - 26B	36.67	21.11	28.89	48.00	34.00	46.34	42.33	46.34	32.90	37.62	35.26	35.26
14	SPV-1201 x GI-35-15-15	48.89	26.67	37.78	37.33	37.33	39.36	38.35	39.36	37.56	38.73	38.15	38.15
15	SPV-1201 x SPV - 946	37.78	26.67	32.22	48.00	44.00	46.00	45.00	46.00	32.56	36.89	34.73	34.73
16	SPV-1201 x SPV - 104	34.45	21.11	27.78	48.00	44.00	46.00	45.00	46.00	32.56	36.89	34.73	34.73
17	SPV-1201 x IS - 2284	13.33	8.89	11.11	24.67	16.67	18.00	17.33	18.00	17.22	18.46	17.84	17.84
18	SPV-1201 x IS - 6335	17.78	12.22	15.00	16.67	62.00	62.00	62.00	62.00	40.33	40.33	40.33	40.33
19	SPV-1201 x IS - 9471	15.56	11.11	13.33	16.00	36.67	47.33	32.68	40.01	40.33	36.34	38.34	38.34
20	ICSB - 101B x AKms 14 B	33.34	40.00	36.67	47.33	62.00	62.00	62.00	62.00	40.33	40.33	40.33	40.33
21	ICSB - 101B x SRT - 26 B	20.00	32.22	26.11	60.00	62.00	62.00	62.00	62.00	40.33	40.33	40.33	40.33
22	ICSB - 101B x GI-35-15-15	31.11	45.56	38.34	60.67	46.00	53.33	45.89	53.33	45.89	45.89	45.89	45.89
23	ICSB - 101B x SPV - 946	38.89	32.22	35.56	45.33	26.68	36.01	31.01	36.01	42.11	42.11	42.11	42.11
24	ICSB - 101B x SPV - 104	27.78	25.56	26.67	38.67	38.67	39.32	39.32	39.32	33.22	32.44	32.83	32.83
25	ICSB - 101B x IS - 2284	37.78	18.89	28.33	34.00	19.36	26.68	23.02	26.68	19.12	27.51	23.33	23.33
26	ICSB - 101B x IS - 6335	30.00	18.89	24.45	23.33	16.00	16.00	16.00	16.00	20.89	22.06	19.48	19.48
27	ICSB - 101B x IS - 9471	17.78	10.00	13.89	24.00	20.00	20.00	20.00	20.00	28.44	31.56	25.00	25.00
28	AKms - 14 B x SRT - 26B	15.56	14.45	15.00	41.33	48.68	50.00	49.33	48.68	39.56	40.56	40.06	40.06
29	AKms - 14 B x GI-35-15-15	34.45	31.11	32.78	47.33	52.00	49.67	44.22	49.67	33.67	35.79	34.73	34.73
30	AKms - 14 B x SPV - 946	41.11	33.34	37.22	47.33	24.00	39.36	32.00	39.36	22.56	23.22	22.89	22.89
31	AKms - 14 B x SPV - 104	43.34	32.22	37.78	38.00	27.33	25.00	26.17	25.00	30.89	26.90	28.90	28.90
32	AKms - 14 B x IS - 2284	33.34	24.45	28.89	38.00	27.33	25.00	26.17	25.00	30.89	26.90	28.90	28.90
33	AKms - 14 B x IS - 6335	17.78	32.22	25.00	27.33	34.00	27.33	30.89	27.33	26.90	28.90	28.90	28.90
34	AKms - 14 B x IS - 9471	27.78	31.11	29.45	34.00	22.56	22.56	22.56	22.56	30.89	26.90	28.90	28.90

Table ou Contd.

	1	2	3	4	5	6	7	8	9	10	11
35	SRT- 26B x GJ-35-15-15	45.56	24.45	35.00	50.67	42.00	46.33	48.11	33.22	40.67	
36	SRT- 26B x SPV - 946	42.22	21.11	31.67	44.67	48.00	46.33	43.45	34.56	39.00	
37	SRT- 26B x SPV - 104	14.45	13.33	13.89	36.67	36.68	36.67	25.56	25.01	25.28	
38	SRT- 26B x IS - 2284	12.22	16.67	14.45	38.00	17.36	27.68	25.11	17.01	21.06	
39	SRT- 26B x IS - 6335	14.45	24.45	19.45	31.33	16.68	24.01	22.89	20.56	21.73	
40	SRT- 26B x IS - 9471	23.33	12.22	17.78	27.33	20.00	23.67	25.33	16.11	20.72	
41	GJ-35-15-15 x SPV - 946	23.33	23.33	23.33	51.33	36.68	44.01	37.33	30.01	33.67	
42	GJ-35-15-15 x SPV- 104	21.11	20.00	20.56	41.33	48.00	44.67	31.22	34.00	32.61	
43	GJ-35-15-15 x IS - 2284	22.22	13.33	17.78	34.67	21.36	28.01	28.45	17.35	22.90	
44	GJ-35-15-15 x IS - 6335	15.56	14.45	15.00	29.33	22.00	25.67	22.44	18.22	20.33	
45	GJ-35-15-15 x IS - 9471	20.00	16.67	18.33	28.67	34.00	31.33	24.33	25.33	24.83	
46	SPV - 946 x SPV - 104	28.89	18.89	23.89	47.33	36.68	42.01	38.11	27.78	32.95	
47	SPV - 946 x IS - 2284	23.33	12.22	17.78	26.00	14.00	20.00	24.67	13.11	18.89	
48	SPV - 946 x IS- 6335	14.45	5.56	10.00	28.00	14.00	21.00	21.22	9.78	15.50	
49	SPV - 946 x IS 9471	10.00	12.22	11.11	26.00	20.00	23.00	18.00	16.11	17.06	
50	SPV - 104 x IS 2284	28.89	22.22	25.56	31.33	18.00	24.67	30.11	20.11	25.11	
51	SPV - 104 x IS 6335	22.22	21.11	21.67	30.00	20.68	25.34	26.11	20.90	23.50	
52	SPV - 104 x IS 9471	8.89	11.11	10.00	34.00	20.00	27.00	21.44	15.56	18.50	
53	IS 2284 x IS-6335	20.00	18.89	19.45	25.33	16.68	21.01	22.67	17.78	20.23	
54	IS 2284 x IS -9471	22.22	17.78	20.00	26.00	17.36	21.68	24.11	17.57	20.84	
55	IS 6335 x IS 9471	25.56	17.78	21.67	21.33	13.36	17.35	23.45	15.57	19.51	

Mean

% increase (+)/decrease (-)

over untreated

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Table 61. Effect of pre-treatment on other fungi (%) (GS) of parents and F₁ crosses at Akola 1995 and Akola 1996

S.N.	Parents/Crosses	Akola-1995				Akola-1996				Pooled over localities			
		2		3		4		5		6		7	
		Untreated	Treated	Untreated	Treated	Untreated	Treated	Untreated	Treated	Untreated	Treated	Untreated	Treated
1	SPV-1201	16.67	13.33	15.00	20.00	18.67	19.33	18.33	16.00	18.33	10	11	17.17
2	ICSB-101B	21.11	33.34	27.22	13.33	12.67	13.00	17.22	20.11	17.22	10	11	20.11
3	AKma-14B	25.56	25.56	25.56	4.67	8.00	6.33	15.11	15.95	15.11	10	11	15.95
4	SRT-26B	20.00	26.67	23.33	15.33	16.67	16.00	17.67	21.67	17.67	10	11	19.67
5	GI-35-15-15	86.67	42.22	64.45	20.00	14.67	17.33	53.34	28.45	53.34	10	11	40.89
6	SPV-946	24.45	25.56	25.00	23.33	14.00	18.67	23.89	19.78	23.89	10	11	21.83
7	SPV-104	30.00	23.33	26.67	12.67	7.33	10.00	21.33	18.33	21.33	10	11	18.33
8	IS-2284	44.45	26.67	35.56	24.67	10.00	17.33	34.56	26.45	34.56	10	11	26.45
9	IS-6335	40.00	22.22	31.11	20.00	6.67	13.33	30.00	14.45	30.00	10	11	22.22
10	IS-9471	62.23	33.34	47.78	19.33	6.00	12.67	40.78	19.67	40.78	10	11	30.22
11	SPV-1201 x ICSB-101B	35.56	14.45	25.00	22.67	18.67	20.67	29.11	16.56	29.11	10	11	22.83
12	SPV-1201 x AKma-14 B	23.33	18.89	21.11	23.33	18.00	20.67	23.33	18.44	23.33	10	11	20.89
13	SPV-1201 x SRT-26B	17.78	12.22	15.00	34.00	20.67	27.33	25.89	16.44	27.33	10	11	21.17
14	SPV-1201 x GI-35-15-15	37.78	16.67	27.22	14.00	17.33	15.67	25.89	17.00	25.89	10	11	21.45
15	SPV-1201 x SPV-946	31.11	14.45	22.78	11.33	18.00	14.67	21.22	16.22	21.22	10	11	18.72
16	SPV-1201 x SPV-104	22.22	22.22	22.22	26.67	27.33	27.00	24.45	24.78	24.45	10	11	24.61
17	SPV-1201 x IS-2284	34.45	15.56	25.00	18.67	9.33	14.00	26.56	12.44	26.56	10	11	19.50
18	SPV-1201 x IS-6335	31.11	15.56	23.33	18.00	5.33	11.67	24.56	10.44	24.56	10	11	17.50
19	SPV-1201 x IS-9471	35.56	15.56	25.56	16.67	5.33	11.00	26.11	10.44	26.11	10	11	18.28
20	ICSB-101B x AKma-14 B	21.11	20.00	20.56	16.67	10.67	13.67	18.89	15.33	18.89	10	11	17.11
21	ICSB-101B x SRT-26 B	10.00	11.11	10.56	18.00	10.00	14.00	14.00	10.56	14.00	10	11	12.28
22	ICSB-101B x GI-35-15-15	17.78	18.89	18.33	22.00	8.67	15.33	19.89	13.78	19.89	10	11	16.83
23	ICSB-101B x SPV-946	24.45	24.45	21.11	33.33	10.67	22.00	25.56	17.56	25.56	10	11	21.68
24	ICSB-101B x SPV-104	20.00	24.45	22.22	16.00	11.33	13.67	18.00	17.89	18.00	10	11	17.96
25	ICSB-101B x IS-2284	40.00	7.78	23.89	32.00	10.00	21.00	36.00	8.89	36.00	10	11	22.45
26	ICSB-101B x IS-6335	30.00	15.56	22.78	31.33	8.00	19.67	30.67	11.78	30.67	10	11	21.22
27	ICSB-101B x IS-9471	33.34	15.56	24.45	26.00	9.33	17.67	29.67	12.44	29.67	10	11	21.06
28	AKma-14 B x SRT-26B	8.89	11.11	10.00	14.67	18.00	16.33	11.78	14.56	16.33	10	11	13.17
29	AKma-14 B x GI-35-15-15	11.11	14.45	12.78	18.00	18.67	18.33	14.56	15.66	18.33	10	11	15.66
30	AKma-14 B x SPV-946	28.89	16.67	22.78	20.67	18.67	19.67	24.78	17.67	24.78	10	11	21.22
31	AKma-14 B x SPV-104	17.78	22.22	20.00	8.00	12.67	10.33	12.89	17.45	12.89	10	11	15.17
32	AKma-14 B x IS-2284	22.22	14.45	18.33	24.00	6.00	15.00	10.22	10.22	15.00	10	11	16.67
33	AKma-14 B x IS-6335	17.78	14.45	16.11	37.33	7.33	10.67	22.33	10.89	22.33	10	11	19.22
34	AKma-14 B x IS-9471	30.00	12.22	21.11	20.67	10.67	15.67	25.33	11.44	25.33	10	11	18.39

Contd.

Table 62. Effect of pre-treatment on score (GS) of parents and F₁ crosses at Akola 1995 and Akola 1996

S.N.	Parents/Crosses	Akola-1995				Akola-1996				Pooled over locations			
		Untreated		Treated		Untreated		Treated		Untreated		Treated	
		3	Mean	4	5	6	Mean	7	8	9	Mean	10	11
1	SPV-1201	1.50	1.50	1.50	1.50	4.00	3.00	3.50	3.50	2.75	2.25	2.25	2.50
2	ICSB-101B	4.17	3.83	3.50	3.83	5.00	3.00	3.50	4.08	4.08	3.25	3.25	3.67
3	AKma-14B	2.83	3.50	2.83	3.17	5.00	3.50	3.50	4.25	3.92	3.50	3.71	3.50
4	SRT-26B	3.50	2.83	2.83	3.17	3.83	3.00	3.42	3.42	3.67	2.92	2.92	3.29
5	GI-35-15-15	2.50	2.67	2.67	2.58	3.50	3.00	3.25	3.00	3.00	2.83	2.83	2.92
6	SPV-946	3.67	3.33	3.33	3.50	3.83	3.17	3.17	3.17	3.75	2.92	2.92	3.33
7	SPV-104	3.50	2.50	2.50	3.00	4.50	4.00	3.50	4.00	4.00	3.00	3.00	3.50
8	IS-2284	2.00	2.00	2.00	2.00	2.00	1.00	1.00	1.50	2.00	1.50	1.50	1.75
9	IS-6335	2.00	2.00	2.00	2.00	2.00	1.00	1.00	1.00	1.50	1.50	1.50	1.50
10	IS-9471	2.00	2.00	2.00	2.00	2.00	1.00	1.00	1.00	1.50	1.50	1.50	1.50
11	SPV-1201 x ICSB-101B	2.83	2.83	1.83	2.33	3.00	2.50	2.50	2.75	2.92	2.17	2.17	2.54
12	SPV-1201 x AKma-14 B	2.83	2.83	2.00	2.42	3.00	2.50	2.50	2.75	2.92	2.25	2.25	2.58
13	SPV-1201 x SRT - 26B	2.67	2.33	2.33	2.50	2.83	3.00	3.00	2.92	2.75	2.67	2.67	2.71
14	SPV-1201 x GI-35-15-15	3.00	2.50	2.50	2.75	2.50	3.00	3.00	2.75	2.75	2.75	2.75	2.75
15	SPV-1201 x SPV - 946	2.83	2.17	2.17	2.50	2.83	2.92	2.92	2.83	3.00	2.58	2.58	2.71
16	SPV-1201 x SPV - 104	3.33	3.00	3.00	3.17	3.17	3.00	3.25	3.00	3.25	2.92	2.92	3.08
17	SPV-1201 x IS - 2284	2.00	1.50	1.50	1.75	1.00	1.00	1.00	1.00	1.50	1.25	1.25	1.38
18	SPV-1201 x IS - 6335	2.00	1.50	1.50	1.75	1.00	1.00	1.00	1.00	1.50	1.25	1.25	1.38
19	SPV-1201 x IS - 9471	2.00	2.00	2.00	2.00	1.00	1.00	1.00	1.00	1.50	1.50	1.50	1.50
20	ICSB - 101B x AKma 14 B	4.00	2.50	2.50	3.25	3.00	3.00	3.00	3.00	3.50	2.75	2.75	3.13
21	ICSB - 101B x SRT - 26 B	4.17	2.67	2.67	3.42	3.00	2.50	2.50	2.50	3.58	2.33	2.33	2.86
22	ICSB - 101B x GI-35-15-15	4.17	2.67	2.67	3.42	2.67	2.00	2.00	2.33	3.42	2.33	2.33	2.86
23	ICSB - 101B x SPV - 946	4.33	3.50	3.50	3.92	2.67	2.00	2.00	2.83	3.50	3.25	3.25	3.38
24	ICSB - 101B x SPV - 104	4.83	3.67	3.67	4.25	3.33	3.00	3.00	2.67	4.08	2.83	2.83	3.46
25	ICSB - 101B x IS - 2284	2.00	1.50	1.50	1.75	1.33	1.00	1.00	1.17	1.67	1.25	1.25	1.46
26	ICSB - 101B x IS - 6335	2.00	1.50	1.50	1.75	1.00	1.00	1.00	1.00	1.50	1.25	1.25	1.38
27	ICSB - 101B x IS - 9471	2.00	2.00	2.00	2.00	1.50	1.00	1.00	1.25	1.75	1.50	1.50	1.63
28	AKma - 14 B x SRT - 26B	4.00	3.67	3.67	3.83	3.00	2.83	2.83	2.92	3.50	3.25	3.25	3.38
29	AKma - 14 B x GI-35-15-15	4.00	3.33	3.33	3.75	3.00	3.00	3.00	3.00	3.17	2.92	2.92	3.04
30	AKma - 14 B x SPV - 946	3.33	2.83	2.83	3.08	3.00	3.00	3.00	3.00	3.75	3.50	3.50	3.63
31	AKma - 14 B x SPV - 104	4.00	3.00	3.00	3.50	3.50	4.00	4.00	1.50	2.00	1.42	1.42	1.71
32	AKma - 14 B x IS - 2284	2.00	1.83	1.83	1.92	2.00	1.00	1.00	1.25	1.75	1.25	1.25	1.50
33	AKma - 14 B x IS - 6335	2.00	1.50	1.50	1.75	1.50	1.00	1.00	1.58	1.83	1.50	1.50	1.67
34	AKma - 14 B x IS - 9471	1.50	2.00	2.00	1.75	2.17	1.00	1.00	1.58	1.83	1.50	1.50	1.67

Contd.

TABLE 26

	1	2	3	4	5	6	7	8	9	10	11
35	SRT- 26B x GI-35-15-15		3.00	2.33	2.67	2.83	2.50	2.67	2.92	2.42	2.67
36	SRT-26B x SPV - 946		2.83	2.00	2.42	3.00	2.50	2.75	2.92	2.25	2.58
37	SRT- 26B x SPV -104		1.50	1.50	1.50	3.00	3.00	3.00	2.25	2.25	2.25
38	SRT- 26B x IS - 2284		1.50	1.50	1.50	1.00	1.00	1.00	1.25	1.25	1.25
39	SRT- 26B x IS - 6335		1.50	1.50	1.50	1.00	1.00	1.00	1.25	1.25	1.25
40	SRT- 26B x IS - 9471		1.50	1.50	1.50	1.00	1.00	1.00	1.25	1.25	1.25
41	GI-35-15-15 x SPV -946		2.17	1.83	2.00	3.00	2.33	2.67	2.58	2.08	2.33
42	GI-35-15-15 x SPV- 104		2.83	2.50	2.67	2.33	2.50	2.42	2.58	2.50	2.54
43	GI-35-15-15 x IS -2284		1.50	1.50	1.50	1.00	1.00	1.00	1.25	1.25	1.25
44	GI-35-15-15 x IS -6335		1.50	1.50	1.50	1.00	1.00	1.00	1.25	1.25	1.25
45	GI-35-15-15 x IS - 9471		1.50	1.50	1.50	1.00	2.00	1.50	1.25	1.75	1.50
46	SPV- 946 x SPV- 104		4.00	3.00	3.50	3.33	3.00	3.17	3.67	3.00	3.33
47	SPV- 946 x IS - 2284		1.50	1.50	1.50	1.17	1.00	1.08	1.33	1.25	1.29
48	SPV- 946 x IS- 6335		1.50	1.00	1.25	1.00	1.00	1.00	1.25	1.00	1.13
49	SPV- 946 x IS 9471		1.50	1.50	1.50	1.00	1.00	1.00	1.25	1.25	1.25
50	SPV - 104 x IS 2284		2.00	1.50	1.75	1.50	1.00	1.25	1.75	1.25	1.50
51	SPV - 104 x IS 6335		2.50	2.33	2.42	1.50	1.00	1.25	2.00	1.67	1.83
52	SPV - 104 x IS 9471		1.50	2.00	1.75	1.50	2.00	1.75	1.50	2.00	1.75
53	IS 2284 x IS-6335		1.50	1.50	1.50	1.00	1.00	1.00	1.25	1.25	1.25
54	IS 2284 x IS -9471		1.50	1.50	1.50	1.50	1.00	1.25	1.50	1.25	1.38
55	IS 6335 x IS 9471		1.50	1.50	1.50	1.00	1.00	1.00	1.25	1.25	1.25
Mean			2.54	2.17	2.36	2.26	1.96	2.11	2.40	2.07	2.24
% increase (+)/decrease(-)					(-)14.56			(-)13.27			(-)13.75

Table 63. Effect of pre-treatment on ungerminated (%) seed of parents and F₁ crosses of AKOLA-1995 and AKOLA-1996

S.N.	Parental/Crosses	Akola-1995				Akola-1996				Pooled over localities				Mean
		Untreated		Treated		Untreated		Treated		Untreated		Treated		
		3	4	5	6	7	8	9	10	11				
1	SPV-1201	2.20	4.44	3.32	22.00	24.00	23.00	12.10	14.22	13.16				
2	ICSB-101B	25.56	10.00	17.78	23.33	20.00	21.67	24.45	15.00	19.72				
3	AKma-141	48.89	41.11	45.00	46.00	31.33	38.67	47.45	36.22	41.83				
4	SRT-26B	26.67	31.11	28.89	13.33	14.67	14.00	20.00	22.89	21.46				
5	GI-35-15-15	2.22	7.78	5.00	18.67	22.67	20.67	10.44	15.22	12.83				
6	SPV-946	22.22	32.22	27.22	12.67	18.67	15.67	17.45	25.45	21.45				
7	SPV-104	28.89	17.78	23.33	38.67	42.00	40.33	33.78	29.89	31.83				
8	IS-2284	12.22	8.89	10.56	6.67	9.33	8.00	9.44	9.11	9.28				
9	IS-6335	2.22	3.33	2.78	6.67	6.67	6.67	4.44	5.00	4.72				
10	IS-9471	1.11	3.33	2.22	6.67	6.00	6.33	3.89	4.67	4.28				
11	SPV-1201 x ICSB-101B	32.22	8.89	20.56	14.67	12.00	13.33	23.45	10.44	16.94				
12	SPV-1201 x AKma-14 B	32.22	21.11	26.67	17.33	14.67	16.00	24.78	17.89	21.33				
13	SPV-1201 x SRT-26B	35.56	16.67	26.11	6.00	10.00	8.00	20.78	13.33	17.06				
14	SPV-1201 x GI-35-15-15	3.33	2.22	2.78	39.33	5.33	22.33	21.33	3.78	12.56				
15	SPV-1201 x SPV-946	7.78	5.56	6.67	34.67	25.33	30.00	21.22	15.44	18.33				
16	SPV-1201 x SPV-104	32.22	24.45	28.33	15.33	12.67	14.00	23.78	18.56	21.17				
17	SPV-1201 x IS-2284	4.44	3.33	3.89	8.67	4.00	6.33	6.56	3.67	5.11				
18	SPV-1201 x IS-6335	17.78	7.78	12.78	4.00	6.67	5.33	10.89	7.22	9.06				
19	SPV-1201 x IS-9471	20.00	11.11	15.66	4.67	7.33	6.00	12.33	9.22	10.78				
20	ICSB-101B x AKma-14 B	34.45	21.11	27.78	21.33	24.00	22.67	27.89	22.56	25.22				
21	ICSB-101B x SRT-26 B	64.45	37.78	51.11	12.00	8.67	10.33	38.22	23.22	30.72				
22	ICSB-101B x GI-35-15-15	41.11	21.11	31.11	12.00	6.00	9.00	26.56	13.56	20.06				
23	ICSB-101B x SPV-946	37.78	20.00	28.89	8.00	10.67	9.33	22.89	15.33	19.11				
24	ICSB-101B x SPV-104	28.89	27.78	28.33	23.33	19.33	21.33	26.11	23.56	24.83				
25	ICSB-101B x IS-2284	5.56	4.44	5.00	11.33	9.33	10.33	8.44	6.89	7.67				
26	ICSB-101B x IS-6335	7.78	5.56	6.67	10.00	6.67	8.33	8.89	6.11	7.50				
27	ICSB-101B x IS-9471	7.78	5.56	6.67	8.00	6.00	7.00	7.89	5.78	6.83				
28	AKma-14 B x SRT-26B	23.33	18.89	21.11	28.00	18.00	23.00	25.67	18.44	22.06				
29	AKma-14 B x GI-35-15-15	38.89	25.56	32.22	18.00	18.67	18.33	28.45	22.11	25.28				
30	AKma-14 B x SPV-946	21.11	13.33	17.22	22.67	19.33	21.00	21.89	16.33	19.11				
31	AKma-14 B x SPV-104	23.33	16.67	20.00	46.00	34.00	40.00	34.67	25.33	30.00				
32	AKma-14 B x IS-2284	21.11	13.33	17.22	20.00	6.00	13.00	20.56	9.67	15.11				
33	AKma-14 B x IS-6335	10.00	6.67	8.33	6.67	2.67	4.67	8.33	4.67	6.50				
34	AKma-14 B x IS-9471	15.56	10.00	12.78	12.00	6.00	9.00	13.78	8.00	10.89				

Count

Contd.

Table 63. Contd.

1	2	3	4	5	6	7	8	9	10	11
35	SRT- 26B x GJ-35-15-15	6.67	4.44	5.56	7.33	10.00	8.67	7.00	7.22	7.11
36	SRT-26B x SPV - 946	15.56	12.22	13.89	16.00	14.00	15.00	15.78	13.11	14.44
37	SRT - 26B x SPV -104	7.78	3.33	5.56	30.67	28.00	29.33	19.22	15.67	17.44
38	SRT - 26B x IS - 2284	15.56	17.78	16.67	12.00	21.33	16.67	13.78	19.56	16.67
39	SRT - 26B x IS - 6335	21.11	15.56	18.33	10.00	10.67	10.33	15.56	13.11	14.33
40	SRT - 26B x IS - 9471	17.78	15.56	16.67	10.00	4.67	7.33	13.89	10.11	12.00
41	GJ-35-15-15 x SPV - 946	13.33	5.56	9.44	14.67	12.00	13.33	14.00	8.78	11.39
42	GJ-35-15-15 x SPV- 104	54.45	45.56	50.00	16.67	14.00	15.33	35.56	29.78	32.67
43	GJ-35-15-15 x IS - 2284	14.45	14.45	14.45	9.33	12.00	10.67	11.89	13.22	12.56
44	GJ-35-15-15 x IS - 6335	3.33	1.11	2.22	8.00	10.00	9.00	5.67	5.56	5.61
45	GJ-35-15-15 x IS - 9471	21.11	17.78	19.45	7.33	8.00	7.67	14.22	12.89	13.56
46	SPV - 946 x SPV- 104	48.89	44.45	46.67	15.33	20.00	17.67	32.11	32.22	32.17
47	SPV - 946 x IS - 2284	3.33	1.11	2.22	13.33	7.33	10.33	8.33	4.22	6.28
48	SPV - 946 x IS- 6335	5.56	2.22	3.89	6.67	4.67	5.67	6.11	3.44	4.78
49	SPV - 946 x IS 9471	10.00	10.00	10.00	8.00	10.67	9.33	9.00	10.33	9.67
50	SPV - 104 x IS 2284	18.89	13.33	16.11	11.33	10.67	11.00	15.11	12.00	13.56
51	SPV - 104 x IS 6335	37.78	30.00	33.89	7.33	6.67	7.00	22.56	18.33	20.45
52	SPV - 104 x IS 9471	16.67	14.45	15.56	8.00	6.00	7.00	12.33	10.22	11.28
53	IS 2284 x IS-6335	10.00	1.11	5.56	8.67	7.33	8.00	9.33	4.22	6.78
54	IS 2284 x IS -9471	8.89	2.22	5.56	12.00	16.00	14.00	10.44	9.11	9.78
55	IS 6335 x IS 9471	5.56	2.22	3.89	4.00	5.33	4.67	4.78	3.78	4.28
Mean		19.91	14.24	17.08	15.19	13.24	14.22	17.55	13.74	15.65
% increase (+)/decrease(-) over untreated				(-)28.47			(-)12.83			(-)21.71
% increase over location				16.74 (Akola 96)						
	SE (m)	CD 5%			SE (m)	CD 5%		SE (m)	CD 5%	
A	0.452	1.252			0.430	1.193		0.589	1.714	
B	2.368	6.563			2.258	3.258		3.093	9.029	
A x B	3.348	9.281			3.193	8.8580		4.375	12.731	

Table 64. Effect of pre-treatment on *F. moniliforme* (%) (UGS) of parents and F_1 crosses at Akola 1995 and Akola, 1996

S.N.	Parents/Crosses	Akola-1995		Akola-1996		Pooled over locations				
		Untreated	Treated	Mean	Untreated	Treated	Mean	Untreated	Treated	Mean
1	2	3	4	5	6	7	8	9	10	11
1	SPV-1201	2.22	1.11	1.67	8.00	8.00	8.00	5.11	4.56	483
2	ICSB-101B	6.67	5.56	6.11	8.67	8.67	8.67	7.67	7.11	739
3	AKms-14B	4.44	8.89	6.67	21.33	16.00	18.67	12.89	12.44	1267
4	SRT-26B	13.33	15.56	14.45	5.33	8.00	6.67	9.33	11.78	1056
5	GI-35-15-15	0.00	2.22	1.11	6.67	8.67	7.67	3.33	5.44	439
6	SPV-946	5.56	5.56	5.56	6.00	8.00	7.00	5.78	15.78	628
7	SPV-104	4.44	10.00	7.22	18.67	14.67	16.67	11.56	12.33	1194
8	IS-2284	0.00	2.22	1.11	4.00	3.33	3.67	2.00	2.78	239
9	IS-6335	0.00	1.11	1.11	2.67	4.00	3.33	1.33	3.11	222
10	IS-9471	1.11	1.11	1.11	2.00	2.67	2.33	1.56	1.89	172
11	SPV-1201 x ICSB-101B	6.67	2.22	4.44	5.33	4.00	4.67	6.00	3.11	456
12	SPV-1201 x AKms-14 B	11.11	6.67	8.89	4.67	5.33	5.00	7.89	15.00	694
13	SPV-1201 x SRT-26B	10.00	7.78	8.89	3.33	4.67	4.00	6.67	15.22	644
14	SPV-1201 x GI-35-15-15	1.11	1.11	1.11	16.67	2.67	9.67	8.89	1.89	539
15	SPV-1201 x SPV-946	4.44	3.33	3.89	19.33	14.67	17.00	11.89	19.00	1044
16	SPV-1201 x SPV-104	8.89	7.78	8.33	4.67	4.00	4.33	6.78	15.89	633
17	SPV-1201 x IS-2284	2.22	2.22	2.22	2.67	2.00	2.33	2.44	2.11	228
18	SPV-1201 x IS-6335	6.67	4.44	5.56	2.00	4.00	3.00	4.33	4.22	428
19	SPV-1201 x IS-9471	7.78	4.44	6.11	2.00	2.67	2.33	4.89	3.56	422
20	ICSB-101B x AKms-14 B	13.33	4.44	8.89	6.00	5.33	5.67	9.67	4.89	728
21	ICSB-101B x SRT-26B	16.67	7.78	12.22	4.00	0.67	2.33	10.33	4.22	728
22	ICSB-101B x GI-35-15-15	12.22	5.56	8.89	4.00	0.67	2.33	10.33	4.22	728
23	ICSB-101B x SPV-946	8.89	6.67	7.78	3.33	2.67	3.00	6.11	4.11	611
24	ICSB-101B x SPV-104	12.22	7.78	10.00	11.33	8.00	9.67	11.78	7.89	983
25	ICSB-101B x IS-2284	4.44	2.22	3.33	6.00	1.33	3.67	5.22	1.78	350
26	ICSB-101B x IS-6335	3.33	5.56	4.44	3.33	2.67	3.00	3.33	4.11	372
27	ICSB-101B x IS-9471	1.11	1.11	1.11	4.00	1.33	2.67	2.56	1.22	189
28	AKms-14 B x SRT-26B	21.11	16.67	18.89	14.00	5.33	9.67	17.56	11.00	1428
29	AKms-14 B x GI-35-15-15	8.89	10.00	9.44	7.33	5.33	6.33	8.11	7.67	789
30	AKms-14 B x SPV-946	6.67	4.44	5.56	8.00	6.00	7.00	7.33	5.22	628
31	AKms-14 B x SPV-104	2.22	2.22	2.22	17.33	10.67	14.00	9.78	6.44	811
32	AKms-14 B x IS-2284	13.33	4.44	8.89	5.33	3.33	4.33	9.33	3.89	651
33	AKms-14 B x IS-6335	1.11	2.22	1.67	3.33	0.00	1.67	2.22	1.11	157
34	AKms-14 B x IS-9471	5.56	2.22	3.89	4.67	2.00	3.33	5.11	2.11	351

Contd...

Concl.

Table 64 . Contd.

1	2	3	4	5	6	7	8	9	10	11
35	SRT- 26B x GI-35-15	5.56	4.44	5.00	2.67	6.00	4.33	4.11	5.22	4.67
36	SRT- 26B x SPV- 946	8.89	6.67	7.78	6.67	6.00	6.33	7.78	6.33	7.06
37	SRT- 26B x SPV-104	6.67	2.22	4.44	11.33	14.00	12.67	9.00	8.11	8.56
38	SRT- 26B x IS - 2284	4.44	4.44	4.44	5.33	8.00	6.67	4.89	6.22	5.56
39	SRT- 26B x IS - 6335	10.00	6.67	8.33	3.33	4.00	3.67	6.67	5.33	6.00
40	SRT- 26B x IS - 9471	6.67	8.89	7.78	4.00	1.33	2.67	5.33	5.11	5.22
41	GI-35-15 x SPV- 946	5.56	2.22	3.89	9.33	5.33	7.33	7.44	3.78	5.61
42	GI-35-15 x SPV- 104	32.22	28.89	30.56	5.33	5.33	5.33	18.78	17.11	17.96
43	GI-35-15 x IS - 2284	4.44	3.33	3.89	2.67	5.33	4.00	3.56	4.33	3.94
44	GI-35-15 x IS - 6335	2.22	1.11	1.67	4.00	4.67	4.33	3.11	2.89	3.00
45	GI-35-15 x IS - 9471	6.67	4.44	5.56	4.00	4.00	4.00	5.33	4.22	4.78
46	SPV- 946 x SPV- 104	12.22	13.33	12.78	4.00	7.33	5.67	8.11	10.33	9.22
47	SPV- 946 x IS - 2284	2.22	1.11	1.67	6.67	4.67	5.67	4.44	2.89	3.67
48	SPV- 946 x IS - 6335	1.11	1.11	1.11	4.67	1.33	3.00	2.89	1.22	2.06
49	SPV- 946 x IS 9471	3.33	5.56	4.44	3.33	6.00	4.67	3.33	5.78	4.56
50	SPV- 104 x IS 2284	7.78	5.56	6.67	4.67	6.00	5.33	6.22	5.78	6.00
51	SPV- 104 x IS 6335	20.00	15.56	17.78	1.33	2.67	2.00	10.67	9.11	9.89
52	SPV- 104 x IS 9471	4.44	7.78	6.11	3.33	4.00	3.67	3.89	5.89	4.89
53	IS 2284 x IS-6335	4.44	1.11	2.78	1.33	4.00	2.67	2.89	2.56	2.72
54	IS 2284 x IS-9471	3.33	2.22	2.78	4.00	7.33	5.67	3.67	4.78	4.22
55	IS 6335 x IS 9471	1.11	1.11	1.11	1.33	2.00	1.67	1.22	1.56	1.39
Mean		6.93	5.56	6.25	6.16	5.32	5.74	6.55	5.44	5.99
% increase (+)/decrease(-) over untreated				(-)/19.76			(-)/13.64			(-)/16.95
% increase over location				14.88 (Akola 96)						
		SE (m)	CD 5%		SE (m)	CD 5%		SE (m)	CD 5%	
		0.439	1.217		0.353	0.979		0.239	0.695	
		2.301	6.380		1.853	5.136		1.253	3.656	
		3.256	9.022		3.706	7.264		1.520	5.004	
		A								
		B								
		A x B								

Table 65. Contd.

	1	2	3	4	5	6	7	8	9	10	11
35	SRT- 26B x GI-35-15-15		0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
36	SRT- 26B x SPV - 946		0.00	0.00	0.00	0.00	0.67	0.33	0.00	0.33	0.17
37	SRT- 26B x SPV - 104		0.00	0.00	0.00	0.67	1.33	1.00	0.33	0.67	0.50
38	SRT- 26B x IS - 2284		0.00	0.00	0.00	0.00	0.67	0.33	0.00	0.33	0.17
39	SRT- 26B x IS - 6335		0.00	0.00	0.00	0.00	0.67	0.33	0.00	0.33	0.17
40	SRT- 26B x IS - 9471		0.00	0.00	0.00	1.33	0.67	1.00	0.67	0.33	0.50
41	GI-35-15-15 x SPV - 946		0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
42	GI-35-15-15 x SPV - 104		0.00	0.00	0.00	1.33	0.00	0.67	0.67	0.00	0.33
43	GI-35-15-15 x IS - 2284		0.00	0.00	0.00	1.33	0.67	1.00	0.67	0.33	0.50
44	GI-35-15-15 x IS - 6335		0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
45	GI-35-15-15 x IS - 9471		0.00	0.00	0.00	0.00	1.33	0.67	0.00	0.67	0.33
46	SPV - 946 x SPV - 104		0.00	0.00	0.00	0.00	0.67	1.33	1.00	0.33	0.67
47	SPV - 946 x IS - 2284		0.00	0.00	0.00	0.67	0.00	0.33	0.33	0.00	0.17
48	SPV - 946 x IS - 6335		0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
49	SPV - 946 x IS 9471		0.00	0.00	0.00	1.33	0.67	1.00	0.67	0.33	0.50
50	SPV - 104 x IS 2284		0.00	0.00	0.00	0.00	1.33	0.67	0.00	0.67	0.33
51	SPV - 104 x IS 6335		0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
52	SPV - 104 x IS 9471		0.00	0.00	0.00	0.00	1.33	0.67	0.67	0.33	0.50
53	IS 2284 x IS-6335		0.00	0.00	0.00	1.33	0.67	2.00	1.67	0.33	1.00
54	IS 2284 x IS -9471		0.00	0.00	0.00	3.33	0.67	2.00	1.67	0.33	1.00
55	IS 6335 x IS 9471		0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00

Mean

% increase (+)/decrease(-)
over untreated

% increase over location

SE (m)

CD 5%

SE (m)

CD 5%

SE (m)

CD 5%

SE (m)

CD 5%

SE (m)

CD 5%

SE (m)

CD 5%

SE (m)

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SE (m)

CD 5%

SE (m)

CD 5%

SE (m)

CD 5%

SE (m)

CD 5%

SE (m)

CD 5%

SE (m)

CD 5%

SE (m)

Table 66. Effect of pre-treatment on *C. lunata* (%) (UGS) of parents and F₁ crosses at Akola 1995 and Akola 1996

S.N.	Parents/Crosses	Akola-1995			Akola-1996			Pooled over locations		
		Untreated	Treated	Mean	Untreated	Treated	Mean	Untreated	Treated	Mean
1	2	3	4	5	6	7	8	9	10	11
1	SPV-1201	0.00	0.00	0.00	10.67	13.33	12.00	5.33	6.67	6.00
2	ICSB-101B	14.45	4.44	9.44	12.67	8.00	10.33	13.56	6.22	9.89
3	AKma-14B	5.56	14.45	10.00	19.33	12.67	16.00	12.44	13.56	13.00
4	SRT-26B	11.11	14.45	12.78	7.33	6.00	6.67	9.22	10.22	9.72
5	GJ-35-15-15	0.00	4.44	2.22	10.67	11.33	11.00	5.33	7.89	6.61
6	SPV-946	15.56	17.78	16.67	6.00	10.67	8.33	10.78	14.22	12.50
7	SPV-104	10.00	6.67	8.33	16.00	16.00	16.00	13.00	11.33	12.17
8	IS-2284	7.78	5.56	6.67	3.33	6.00	4.67	5.56	5.78	5.67
9	IS-6335	2.22	0.00	1.11	4.00	2.67	3.33	3.11	1.33	2.22
10	IS-9471	0.00	0.00	0.00	4.00	2.67	3.33	2.00	1.33	1.67
11	SPV-1201 x ICSB-101B	13.33	4.44	8.89	8.00	8.00	8.00	10.67	6.22	8.44
12	SPV-1201 x AKma-14 B	12.22	5.56	8.89	10.67	7.33	9.00	11.44	6.44	8.94
13	SPV-1201 x SRT - 26B	21.11	6.67	13.89	2.67	4.67	3.67	11.89	5.67	8.78
14	SPV-1201 x GJ-35-15-15	2.22	0.00	1.11	16.67	2.67	9.67	9.44	1.33	5.39
15	SPV-1201 x SPV -946	2.22	2.22	2.22	12.67	10.00	11.33	7.44	6.11	6.78
16	SPV-1201 x SPV- 104	16.67	11.11	13.89	7.33	8.00	7.67	12.00	9.56	10.78
17	SPV-1201 x IS -2284	1.11	0.00	0.56	3.33	2.00	2.67	2.22	1.00	1.61
18	SPV-1201 x IS-6335	4.44	2.22	3.33	1.33	2.00	1.67	2.89	2.11	2.50
19	SPV-1201 x IS - 9471	4.44	4.44	4.44	2.00	4.00	3.00	3.22	4.22	3.72
20	ICSB - 101B x AKma 14 B	16.67	15.56	16.11	12.00	16.67	14.33	14.33	16.11	15.22
21	ICSB - 101B x SRT- 26 B	38.89	24.45	31.67	6.00	6.67	6.33	22.45	15.56	19.00
22	ICSB - 101B x GJ-35-15-15	21.11	15.56	18.33	6.67	3.33	5.00	13.89	9.44	11.67
23	ICSB - 101B x SPV - 946	16.67	12.22	14.45	4.00	7.33	5.67	10.33	9.78	10.06
24	ICSB - 101B x SPV - 104	14.45	11.11	12.78	10.67	10.00	10.33	12.56	10.56	11.56
25	ICSB - 101B x IS - 2284	1.11	0.00	0.56	4.00	7.33	5.67	2.56	3.67	3.11
26	ICSB - 101B x IS - 6335	2.22	0.00	1.11	3.33	2.67	3.00	2.78	1.33	2.06
27	ICSB - 101B x IS - 9471	2.22	2.22	2.22	3.33	4.00	3.67	2.78	3.11	2.94
28	AKma - 14 B x SRT - 26B	2.22	2.22	2.22	9.33	9.33	9.33	5.78	5.78	5.78
29	AKma - 14 B x GJ-35-15-15	22.22	11.11	16.67	8.00	12.67	10.33	15.11	11.89	13.50
30	AKma - 14 B x SPV - 946	13.33	7.78	10.56	12.00	10.00	11.00	12.67	8.89	10.78
31	AKma - 14 B x SPV - 104	14.45	11.11	12.78	24.00	18.00	21.00	19.22	14.56	16.89
32	AKma - 14 B x IS - 2284	5.56	8.89	7.22	9.33	2.67	6.00	7.44	5.78	6.61
33	AKma - 14 B x IS - 6335	5.56	1.11	3.33	3.33	2.67	3.00	4.44	1.89	3.17
34	AKma - 14 B x IS - 9471	7.78	5.56	6.67	5.33	4.00	4.67	6.56	4.78	5.67

Contd.

	1	2	3	4	5	6	7	8	9	10	11
35	SRT- 26B x GJ-35-15-15										
36	SRT- 26B x SPV - 946										
37	SRT- 26B x SPV - 104										
38	SRT- 26B x IS - 2284										
39	SRT- 26B x IS - 6335										
40	SRT- 26B x IS - 9471										
41	GJ-35-15-15 x SPV - 946										
42	GJ-35-15-15 x SPV - 104										
43	GJ-35-15-15 x IS - 2284										
44	GJ-35-15-15 x IS - 6335										
45	GJ-35-15-15 x IS - 9471										
46	SPV - 946 x SPV - 104										
47	SPV - 946 x IS - 2284										
48	SPV - 946 x IS - 6335										
49	SPV - 946 x IS 9471										
50	SPV - 104 x IS 2284										
51	SPV - 104 x IS 6335										
52	SPV - 104 x IS 9471										
53	IS 2284 x IS-6335										
54	IS 2284 x IS - 9471										
55	IS 6335 x IS 9471										
Mean			8.42	6.10	7.26	7.25	6.65	6.95	7.78	6.38	7.08
% increase (+)/decrease (-)					(-)27.55			(-)8.27			(-)17.99

Table 67. Effect of pre-treatment on other fungi (%) (UGS) of parents and F_1 crosses at Akola 1995 and Akola 1996

N.	Parents/Crosses	Akola-1995			Akola-1996			Pooled over locations		
		Untreated	Treated	Mean	Untreated	Treated	Mean	Untreated	Treated	Mean
1										
1	SPV-1201	0.00	0.00	0.00	3.33	1.33	2.33	1.67	0.67	1.17
2	ICSB-101B	4.44	0.00	2.22	2.00	2.00	2.00	3.22	1.00	2.11
3	AKms-14B	36.67	15.56	26.11	4.67	2.00	3.33	20.67	3.78	14.72
4	SRT-26B	1.11	2.22	1.67	0.67	0.67	0.67	0.89	1.44	1.17
5	GI-35-15-15	2.22	1.11	1.67	1.33	2.67	2.00	1.78	1.89	1.83
6	SPV-946	2.22	8.89	5.56	2.67	4.00	3.33	2.44	4.44	3.44
7	SPV-104	11.11	1.11	6.11	3.33	4.00	3.67	7.22	2.56	4.89
8	IS-2284	3.33	2.22	2.78	0.00	0.00	0.00	1.67	1.11	1.39
9	IS-6335	0.00	1.11	0.56	0.00	0.00	0.00	0.00	0.56	0.28
10	IS-9471	1.11	2.22	1.67	0.67	0.00	0.33	0.89	1.11	1.00
11	SPV-1201 x ICSB-101B	12.22	2.22	7.22	1.33	0.00	0.67	6.78	1.11	3.94
12	SPV-1201 x AKms-14 B	8.89	8.89	8.89	2.00	1.33	1.67	5.44	5.11	5.28
13	SPV-1201 x SRT-26B	4.44	2.22	3.33	0.00	0.67	0.33	2.22	1.44	1.83
14	SPV-1201 x GI-35-15-15	0.00	1.11	0.56	5.33	0.00	2.67	2.67	0.56	1.61
15	SPV-1201 x SPV-946	1.11	0.00	0.56	4.67	0.67	2.67	2.89	0.33	1.61
16	SPV-1201 x SPV-104	6.67	4.44	5.56	6.00	0.00	3.00	6.33	2.22	4.28
17	SPV-1201 x IS-2284	1.11	0.00	0.56	2.67	0.00	1.33	1.89	0.00	0.94
18	SPV-1201 x IS-6335	6.67	0.00	3.33	0.67	0.00	0.33	3.67	0.00	1.83
19	SPV-1201 x IS-9471	7.78	1.11	4.44	0.67	0.00	0.33	4.22	0.56	2.39
20	ICSB-101B x AKms 14 B	4.44	1.11	2.78	2.67	2.67	2.67	3.56	2.72	3.17
21	ICSB-101B x SRT-26 B	6.67	5.56	6.11	3.33	2.00	2.67	5.00	3.78	4.39
22	ICSB-101B x GI-35-15-15	8.89	0.00	4.44	1.33	0.00	0.67	5.11	0.00	2.56
23	ICSB-101B x SPV-946	8.89	1.11	5.00	0.67	0.67	0.67	4.78	0.89	2.83
24	ICSB-101B x SPV-104	2.22	7.78	5.00	1.33	1.33	1.33	1.78	4.56	3.17
25	ICSB-101B x IS-2284	0.00	2.22	1.11	1.33	0.67	1.00	0.67	1.44	1.06
26	ICSB-101B x IS-6335	2.22	0.00	1.11	2.67	0.00	1.33	2.44	0.00	1.22
27	ICSB-101B x IS-9471	4.44	2.22	3.33	0.67	0.67	0.67	2.56	1.44	2.00
28	AKms-14 B x SRT-26B	0.00	0.00	0.00	4.67	1.33	3.00	2.33	0.67	1.50
29	AKms-14 B x GI-35-15-15	7.78	4.44	6.11	1.33	0.00	0.67	4.56	2.22	3.39
30	AKms-14 B x SPV-946	1.11	0.00	0.56	2.67	2.00	2.33	1.89	1.00	1.44
31	AKms-14 B x SPV-104	4.44	3.33	3.89	4.00	0.67	2.33	2.00	4.22	3.11
32	AKms-14 B x IS-2284	1.11	0.00	0.56	3.33	0.00	1.67	2.22	0.00	1.11
33	AKms-14 B x IS-6335	0.00	3.33	1.67	0.00	0.00	0.00	0.00	1.67	0.83
34	AKms-14 B x IS-9471	2.22	2.22	2.22	1.33	0.00	0.67	1.78	1.11	1.44

Contd.

VIII - 70

Table 67. Contd.

1	2	3	4	5	6	7	8	9	10	11
35	SRT-26B x GJ-35-15-15	0.00	0.00	0.00	0.00	0.67	0.33	0.00	0.33	0.17
36	SRT-26B x SPV-946	3.33	1.11	2.22	0.67	0.67	0.67	2.00	0.89	1.44
37	SRT-26B x SPV-104	0.00	0.00	0.00	2.00	1.33	1.67	1.00	0.87	0.83
38	SRT-26B x IS-2284	4.44	0.00	2.22	2.00	2.00	2.00	2.11	1.00	2.11
39	SRT-26B x IS-6335	1.11	1.11	1.11	1.33	0.00	0.67	1.22	0.56	0.89
40	SRT-26B x IS-9471	3.33	1.11	2.22	0.67	0.00	0.33	2.00	0.56	1.28
41	GJ-35-15-15 x SPV-946	2.22	2.22	2.22	0.67	0.00	0.33	1.44	1.11	1.28
42	GJ-35-15-15 x SPV-104	6.67	3.33	5.00	1.33	0.67	1.00	4.00	2.00	3.00
43	GJ-35-15-15 x IS-2284	5.56	0.00	2.78	1.33	0.00	0.67	3.44	1.72	3.44
44	GJ-35-15-15 x IS-6335	1.11	0.00	0.56	0.00	0.67	0.33	0.56	0.33	0.44
45	GJ-35-15-15 x IS-9471	4.44	2.22	3.33	0.00	0.00	0.00	2.22	1.11	1.67
46	SPV-946 x SPV-104	7.78	7.78	7.78	1.33	0.67	1.00	4.56	4.22	4.39
47	SPV-946 x IS-2284	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
48	SPV-946 x IS-6335	1.11	1.11	1.11	0.00	0.00	0.00	0.56	0.56	0.56
49	SPV-946 x IS-9471	3.33	1.11	2.22	0.00	0.00	0.00	1.67	0.56	1.11
50	SPV-104 x IS-2284	4.44	3.33	3.89	2.00	0.00	1.00	3.22	1.87	2.44
51	SPV-104 x IS-6335	5.56	4.44	5.00	0.00	0.00	0.00	2.78	2.22	2.50
52	SPV-104 x IS-9471	3.33	0.00	1.67	0.00	0.00	0.00	1.67	0.00	0.83
53	IS-2284 x IS-6335	2.22	0.00	1.11	0.00	0.00	0.00	1.11	0.00	0.56
54	IS-2284 x IS-9471	2.22	0.00	1.11	0.67	1.33	1.00	1.44	0.67	1.06
55	IS-6335 x IS-9471	3.33	0.00	1.67	0.67	0.00	0.33	2.00	0.00	1.00

Mean

% increase (+)/decrease (-)
over untreated

% increase over location

SE (m)

CD 5%

SE (m)

CD 5%

3.16

(-)49.52

1.60

0.64

1.12

(-)60.00

2.9

1.38

2.14

(-)52.41

0.170

0.494

64.56

(Akola 96)

0.892

1.261

CD 5%

0.494

2.421

3.670

A

B

A x B

VIII - 70
 Table 68. Effect of pre-treatment on score (UGS) of parents and F₁ crosses at Akola 1995 and Akola 1996

S.N.	Parents/Crosses	Akola-1995				Akola-1996				Pooled over locations			
		Untreated		Treated		Untreated		Treated		Untreated		Treated	
		3	4	5	6	7	8	9	10	Mean	Mean	Mean	Mean
1	SPV-1201	1.00	0.67	0.83	4.50	4.00	4.25	2.75	2.33	2.33	11	2.33	254
2	ICSD-101B	4.17	4.00	4.08	4.67	4.00	4.33	4.42	4.00	4.00	421	4.00	421
3	AKma-14B	3.00	4.00	3.50	5.00	4.50	4.75	4.00	4.25	4.25	413	4.25	413
4	SRT-26B	4.00	3.33	3.67	4.00	4.00	4.00	4.00	3.67	3.67	383	3.67	383
5	GI-35-15-15	0.67	2.83	1.75	4.00	4.00	4.00	2.33	3.42	3.42	288	3.42	288
6	SPV-946	4.17	3.83	4.00	4.00	3.50	3.75	4.08	3.67	3.67	388	3.67	388
7	SPV-104	4.00	3.50	3.75	4.50	4.00	4.25	3.75	3.75	3.75	400	3.75	400
8	IS-2284	2.50	2.17	2.33	2.00	2.17	2.08	2.25	2.17	2.17	221	2.17	221
9	IS-6335	0.67	1.33	1.00	1.83	1.50	1.67	1.25	1.42	1.42	133	1.42	133
10	IS-9471	0.67	1.17	0.92	1.33	1.00	1.17	1.00	1.08	1.08	194	1.08	194
11	SPV-1201 x ICSD-101B	3.67	1.50	2.58	3.50	3.50	3.50	3.58	2.50	2.50	304	2.50	304
12	SPV-1201 x AKma-14 B	3.50	2.67	3.08	4.33	4.00	4.17	3.92	3.33	3.33	363	3.33	363
13	SPV-1201 x SRT-26B	3.33	2.83	3.08	3.17	4.00	3.58	3.25	3.42	3.42	333	3.42	333
14	SPV-1201 x GI-35-15-15	2.00	0.67	1.33	4.50	4.00	4.25	3.25	2.33	2.33	279	2.33	279
15	SPV-1201 x SPV-946	2.83	3.00	2.92	4.50	4.00	4.25	3.67	3.50	3.50	358	3.50	358
16	SPV-1201 x SPV-104	4.00	3.67	3.83	4.00	3.50	3.75	4.00	3.58	3.58	379	3.58	379
17	SPV-1201 x IS-2284	1.33	1.50	1.42	0.67	1.17	0.92	1.00	1.33	1.33	117	1.33	117
18	SPV-1201 x IS-6335	2.50	2.00	2.25	1.17	1.33	1.25	1.83	1.67	1.67	176	1.67	176
19	SPV-1201 x IS-9471	2.33	2.50	2.42	1.50	1.50	1.75	1.92	1.83	1.83	183	1.83	183
20	ICSD-101B x AKma-14 B	5.00	3.00	4.00	4.00	4.00	4.00	4.50	3.50	3.50	400	3.50	400
21	ICSD-101B x SRT-26 B	4.83	3.67	4.25	4.00	4.00	3.75	4.42	3.58	3.58	400	3.58	400
22	ICSD-101B x GI-35-15-15	5.00	3.17	4.08	3.67	3.00	3.33	3.33	3.08	3.08	371	3.08	371
23	ICSD-101B x SPV-946	5.00	4.00	4.50	3.00	3.00	3.50	4.00	4.00	4.00	400	4.00	400
24	ICSD-101B x SPV-104	5.00	4.00	4.50	3.00	3.00	3.50	4.00	4.00	4.00	400	4.00	400
25	ICSD-101B x IS-2284	1.33	1.33	1.33	1.83	1.83	1.83	1.58	1.58	1.58	158	1.58	158
26	ICSD-101B x IS-6335	2.17	1.63	2.00	1.50	1.50	1.25	1.83	1.42	1.42	183	1.42	183
27	ICSD-101B x IS-9471	2.17	1.67	1.92	2.00	1.50	1.75	2.08	1.58	1.58	183	1.58	183
28	AKma-14 B x SRT-26B	5.00	4.50	4.75	3.50	3.50	3.50	4.25	4.00	4.00	413	4.00	413
29	AKma-14 B x GI-35-15-15	4.67	4.33	4.50	3.50	4.33	3.75	4.08	4.17	4.17	413	4.17	413
30	AKma-14 B x SPV-946	4.93	4.00	4.17	4.17	4.00	4.08	4.25	4.00	4.00	413	4.00	413
31	AKma-14 B x SPV-104	4.33	4.00	4.50	4.83	4.33	4.58	4.92	4.17	4.17	454	4.17	454
32	AKma-14 B x IS-2284	3.00	2.67	2.83	2.00	1.67	1.83	2.50	2.17	2.17	233	2.17	233
33	AKma-14 B x IS-6335	1.67	1.67	1.67	2.00	0.83	1.25	1.83	1.25	1.25	154	1.25	154
34	AKma-14 B x IS-9471	2.17	2.50	2.33	3.17	1.50	2.33	2.67	2.00	2.00	233	2.00	233

Contd.

VIII - 72

1	2	3	4	5	6	7	8	9	10	11
1	SRT - 26B x GI-35-15-15	1.83	2.00	1.92	3.83	3.50	3.67	2.83	2.75	2.79
5	SRT - 26B x SPV - 946	3.50	3.00	3.25	4.00	3.50	3.75	3.75	3.25	3.50
7	SRT - 26B x SPV - 104	2.17	1.33	1.75	4.00	4.00	4.00	3.08	2.67	2.88
8	SRT - 26B x IS - 2284	2.50	3.00	2.75	2.00	2.00	2.00	2.25	2.50	2.38
9	SRT - 26B x IS - 6335	3.00	2.50	2.75	2.00	2.00	2.00	2.25	2.25	2.38
0	SRT - 26B x IS - 9471	2.33	2.33	2.08	1.83	1.67	1.75	1.83	2.00	1.92
1	GI-35-15-15 x SPV - 946	3.33	1.83	2.58	3.33	2.83	3.08	3.33	2.83	2.83
2	GI-35-15-15 x SPV - 104	4.00	3.50	3.75	3.17	3.00	3.08	3.58	3.25	3.42
3	GI-35-15-15 x IS - 2284	2.33	2.67	2.50	1.00	2.00	1.50	1.67	2.33	2.00
4	GI-35-15-15 x IS - 6335	1.17	1.50	1.33	1.17	2.00	1.58	1.17	1.75	1.46
5	GI-35-15-15 x IS - 9471	2.33	2.17	2.25	1.00	3.00	2.00	1.67	2.58	2.13
16	SPV - 946 x SPV - 104	4.67	4.33	4.50	4.00	3.83	3.92	4.33	4.08	4.21
17	SPV - 946 x IS - 2284	1.17	0.50	0.83	2.00	2.00	2.00	1.58	1.25	1.42
18	SPV - 946 x IS - 6335	0.67	0.50	0.58	1.67	1.67	1.67	1.17	1.08	1.13
19	SPV - 946 x IS 9471	2.00	2.17	2.08	2.00	1.83	1.92	2.00	2.00	2.00
50	SPV - 104 x IS 2284	3.00	2.50	2.75	2.00	2.00	2.00	2.50	2.25	2.38
51	SPV - 104 x IS 6335	3.50	3.00	3.25	2.00	2.00	2.00	2.75	2.50	2.63
52	SPV - 104 x IS 9471	3.00	3.00	2.75	2.50	3.00	2.75	2.50	3.00	2.75
53	IS 2284 x IS-6335	2.50	0.00	1.25	1.67	2.00	1.83	2.08	1.50	1.54
54	IS 2284 x IS - 9471	2.17	1.00	1.58	2.00	2.00	2.00	2.08	1.50	1.79
55	IS 6335 x IS 9471	1.17	0.83	1.00	1.17	2.00	1.58	1.17	1.42	1.29
Mean		2.87	2.49	2.68	2.88	2.76	2.82	2.88	2.63	2.76
% increase (+)/decrease (-)				(+)13.24			(+)4.16			(+)8.68
% increase over location										
		SE (m)	CD 5%		SE (m)	CD 5%		SE (m)	CD 5%	
	A	0.092	0.255		.052	0.145		0.045	0.131	
	B	0.484	1.342		0.274	0.760		0.235	0.638	
	A x B	0.685	1.898		0.388	1.075		0.332	0.967	

(Akola 95)

CD 5%

SE (m)

CD 5%

SE (m)

CD 5%

SE (m)

CD 5%

SE (m)

Table : 69 Correlation coefficient between fungal load of germinated and ungerminated seed (Untreated), Akola and Patancheru, 1996

Sr. No	Characters	Loca- tion	Germin- ated Seed		F. m.		F. p.		C. lunata		Score		Other fungi		Score		TGM/R
			GS	UGS	GS	UGS	GS	UGS	GS	UGS	GS	UGS	GS	UGS	GS	UGS	
1	Germinated Seed	A	-	-1.0**	-0.71**	0.58**	0.21	-0.11	0.66**	-0.69**	-0.95**	-0.04	-0.94**	-0.74**	-0.74**	-0.72**	-0.87**
		P	-	-0.99**	-0.71*	0.58**	0.75**	0.11	0.58*	-0.77**	-0.96**	-0.80**	-0.82**	-0.52**	-0.83**	-0.52**	-0.87**
2	Ungerminated seed	A	-	-	-0.71*	-0.21	0.11	-0.66**	-0.58**	0.69**	0.95**	0.04	0.95**	0.75**	0.74**	0.72**	0.72**
		P	-	-	-0.58**	-0.66**	-0.75**	-0.58**	-0.49**	0.77**	0.96**	0.80**	0.82**	0.51**	0.83**	0.83**	0.87**
3	<i>F. moniliforme</i> (GS)	A	-	-	-0.18	0.28*	0.28*	0.48**	0.17	-0.20	0.72**	-0.06	0.67**	0.38**	0.71**	0.65**	0.65**
		P	-	-	0.78**	-	0.48**	0.11	-0.45**	-0.25	-0.52**	-0.31*	-0.64**	-0.29*	-0.30*	-0.38	-0.38
4	<i>F. pallidorostrum</i> (GS)	A	-	-	-	-	-0.38**	-0.56**	0.13	-0.41**	-0.25	-0.40**	-0.25**	-0.25	-0.47**	-0.40**	-0.40**
		P	-	-	-	-	0.28*	0.13	0.62**	-0.43**	0.05	-0.25	-0.60**	-0.36**	-0.50**	-0.56**	-0.56**
5	<i>C. lunata</i> (GS)	A	-	-	-	-	-	-	0.71**	-0.38**	-0.63**	-0.05	-0.56**	-0.32*	-0.43**	-0.61**	-0.61**
		P	-	-	-	-	-	-	-	-0.24	-0.57**	-0.45**	-0.46**	-0.28*	-0.27*	-0.38**	-0.38**
6	Other fungi (GS)	A	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
		P	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
7	Score (GS)	A	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
		P	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
8	<i>F. moniliforme</i> (UGS)	A	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
		P	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
9	<i>F. pallidorostrum</i> (UGS)	A	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
		P	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
10	<i>C. lunata</i> (UGS)	A	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
		P	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
11	Other fungi (UGS)	A	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
		P	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
12	Score (UGS)	A	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
		P	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-

F. m. = *F. moniliforme*F. p. = *F. pallidorostrum*

GS = Germinated Seed

UGS = Ungerminated seed

* Significant at 5%

** Significant at 1%

A = Akola

P = Patancheru

Table : 70 Correlation coefficient between fungal load of germinated and ungerminated seed (treated), Akola and Patancheru, 1996

Sr. No	Characters	Loca- tion	Germi- nated seed	Ungermi- nated seed	F. m.		F. p.		C. lunata		Other fungi		Score		TGMR
			GS	UGS	GS	UGS	GS	UGS	GS	UGS	GS	UGS	Score	UGS	
1	Germinated Seed	A	-	-0.99**	-0.72**	-0.36**	-0.40**	-0.25	-0.72**	-0.89**	-0.63**	-0.92**	-0.75**	-0.69**	-0.75**
2	Ungerminated seed	P	-	-0.99**	0.45**	0.46**	0.60**	0.43**	-0.72**	-0.94**	-0.94**	-0.95**	-0.83**	-0.77**	-0.83**
3	<i>F. moniliforme</i> (GS)	A	-	-	0.72**	0.36**	0.40**	0.26*	0.72**	0.89**	0.63**	0.92**	0.75**	0.69**	0.75**
3	<i>F. moniliforme</i> (UGS)	P	-	-	-0.45**	-0.47**	-0.60**	-0.43**	0.71**	0.94**	0.94**	0.95**	0.83**	0.76**	0.83**
4	<i>P. pallidorozeum</i> (GS)	A	-	-	-	0.49**	0.51**	0.33	0.75**	0.70**	0.44**	0.62**	0.54**	0.73**	0.69**
4	<i>P. pallidorozeum</i> (UGS)	P	-	-	-	0.60**	0.59**	0.46**	-0.04	-0.32*	-0.44**	-0.52**	-0.39**	-0.09	-0.08
5	<i>C. lunata</i> (GS)	A	-	-	-	-	-0.07	-0.08	0.13	0.41**	0.30**	0.21	0.36**	0.19	0.14
5	<i>C. lunata</i> (UGS)	P	-	-	-	-	0.48**	0.36**	-0.22	-0.34**	-0.41**	-0.53**	-0.45**	-0.21	-0.28
6	Other fungi (GS)	A	-	-	-	-	-	0.76**	0.77**	0.33*	0.01	0.48**	0.23	0.80**	0.71**
6	Other fungi (UGS)	P	-	-	-	-	-	0.65**	-0.10	-0.53**	-0.56**	-0.60**	-0.54**	-0.14	-0.33**
7	Score (GS)	A	-	-	-	-	-	-	0.56**	0.23	-0.04	0.31*	0.06	0.60**	0.59**
7	Score (UGS)	P	-	-	-	-	-	-	-0.03	-0.36**	-0.44**	-0.43**	-0.38**	-0.03	-0.15
8	<i>F. moniliforme</i> (UGS)	A	-	-	-	-	-	-	-	0.62**	0.36	0.73**	0.53**	0.95**	0.87**
8	<i>F. moniliforme</i> (UGS)	P	-	-	-	-	-	-	-	0.71**	0.64**	0.70**	0.54**	0.94**	0.83**
9	<i>P. pallidorozeum</i> (UGS)	A	-	-	-	-	-	-	-	-	-	0.69**	0.58**	0.61**	0.59**
9	<i>P. pallidorozeum</i> (UGS)	P	-	-	-	-	-	-	-	-	-	0.80**	0.69**	0.78**	0.85**
10	<i>C. lunata</i> (UGS)	A	-	-	-	-	-	-	-	-	-	0.50**	0.48**	0.26*	0.39**
10	<i>C. lunata</i> (UGS)	P	-	-	-	-	-	-	-	-	-	0.86**	0.71**	0.77**	0.76**
11	Other fungi (UGS)	A	-	-	-	-	-	-	-	-	-	-	0.69**	0.70**	0.76**
11	Other fungi (UGS)	P	-	-	-	-	-	-	-	-	-	-	0.84**	0.68**	0.74**
12	Score (UGS)	A	-	-	-	-	-	-	-	-	-	-	-	0.54**	0.68**
12	Score (UGS)	P	-	-	-	-	-	-	-	-	-	-	-	0.57**	0.65**
															0.85**

A= Akola

P= Patancheru

* Significant at 5%

** Significant at 1%

GS = Germinated Seed

UGS = Ungerminated seed

F. m. = *F. moniliforme*F. p. = *F. pallidorozeum*

Table : 71 Correlation coefficient between physical characters, Akola and Patancheru, 1996

Sr. No	Characters	100 grain weight	Grain hardness	End. texture	Ele. Conductivity	DTF	Gluco covering	Meso. thickness	<i>F. moniforme</i>	<i>F. pallidiorosum</i>	<i>C. lunata</i>	Other fungi	TOMR	Germination
1	100 grain weight	A	0.09	-0.02	0.11	0.22	-0.26**	0.21	0.28*	-0.26*	0.25	0.07	0.40**	-0.22
2	Grain hardness	P	0.03	0.09	0.14	-0.02	-0.25	-0.02	0.02	-0.11	-0.12	0.03	0.08	0.01
3	End. texture	P	-0.33*	-0.61**	0.15	0.42**	0.19	0.13	0.10	-0.32*	0.17	0.10	0.05	-0.07
4	Ele. conductivity	P	-0.54**	-0.51**	-0.19	0.39**	-0.05	0.07	-0.26*	-0.11	0.25	0.50**	-0.23	0.30*
5	DTF	P	0.54**	0.54**	0.06	-0.51**	0.01	-0.39**	-0.35**	0.41**	-0.52**	0.14	-0.36**	-0.41**
6	Glu. covering	P	-0.33*	-0.12	-0.07	0.56**	-0.07	0.56**	0.67**	-0.17	-0.24	0.06	-0.87**	0.52*
7	Meso. thickness	P	0.40**	-0.12	0.40**	0.59**	0.56**	0.72**	0.08	-0.33*	0.61**	-0.44**	0.64**	-0.66**
8	<i>F. moniforme</i>	P	0.31*	-0.33*	-0.05	0.41**	0.39**	0.37**	-0.32*	0.26*	0.23	-0.31**	0.82**	-0.82**
9	<i>F. pallidiorosum</i>	P	0.51**	-0.06	-0.13	-0.14	-0.05	0.31*	0.18	-0.12	0.46**	0.24	0.29*	-0.38**
10	<i>C. lunata</i>	P	0.58**	-0.38**	-0.13	-0.05	-0.05	-0.13	0.18	0.02	-0.04	-0.04	-0.10	0.06
11	Other fungi	P	0.65**	-0.20	-0.07	0.68**	0.14	0.14	0.18	0.02	0.68**	-0.20	0.65**	0.11
12	TOMR	P	-0.57**	-0.53**	-0.53**	-0.57**	-0.53**	-0.53**	-0.53**	-0.53**	-0.53**	-0.53**	-0.53**	-0.53**

* = Significant at 5%
 ** = Significant at 1%

X - IV

Table 72 Correlation coefficient between biochemical characters, Akola, 1995

Sr. No	Characters	Protein	Soluble sugars	Tannins	Flavan-4-ols	Grain hardness	TGMR
1	Protein	-	0.27*	0.01	-0.15	0.29*	0.16
2	Soluble sugars			0.35**	0.24	0.26*	-0.26*
3	Tannins				0.73**	-0.37**	-0.75**
4	Flavan-4-ols					-0.59**	-0.78**
5	Grain hardness						-0.30*

* Significant at 5%

** Significant at 1%

Table :73 Correlation coefficient between protein fractions and physical characters Akola and Patancheru, 1996

Sr No	Characters	Albumin and Globulin	Prolamin	Cross link Protein	Glutelin like	Glutelin	Residues	Meso cap thickness	Endo- perm texture	Germination	Grain hardness
1	Alb. & Glob.	-	0.55*	0.36*	-0.73**	-0.78**	-0.10	0.09	-0.45**	-0.37	0.24
2	Prolamin			0.64**	-0.69**	-0.66**	0.34	0.35	-0.73**	-0.69**	0.63**
3	Cross link Prol.				-0.58**	-0.59**	0.11	0.42*	-0.57**	-0.60**	0.55**
4	Glutelin like					0.72**	-0.18	-0.20	0.62**	0.57**	-0.47**
5	Glutelin						-0.07	-0.34	0.52**	0.56**	-0.42*
6	Residues							0.10	-0.10	-0.32	0.02
7	Meso thickness								-0.33	-0.48**	0.41*
8	End. texture									0.44*	0.77**
9	Germination										-0.38*

*

Significant of 5%

**

Significant at 1%

X - 1

Table 74 GCA effects in parents of 10×10 diallel of F_1 crosses and F_2 progenies

Sr. No	Characters	SPV-1201		ICSB-101B		AKms-14B		SRT-26B		GJ-35-15-15		SPV-946		SPV-104		IS-2284		IS-4335		IS-9471	
		F_1	F_2	F_1	F_2	F_1	F_2	F_1	F_2	F_1	F_2	F_1	F_2	F_1	F_2	F_1	F_2	F_1	F_2	F_1	F_2
1	100 grain weight	-	-	-	-	-	-	-	-	+	+	-	-	-	-	-	-	-	-	-	-
2	Grain hardness	-	+	-	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+
3	Endosperm texture	+	-	+	-	+	-	+	-	+	-	+	-	+	-	+	-	+	-	+	-
4	Electrical conductivity	-	+	-	+	-	+	-	+	-	+	-	+	-	+	-	+	-	+	-	+
5	DTF	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
6	Plant height	+	-	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+
7	Cob length	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
8	Glume covering	-	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+
9	Mesocarp thickness	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
10	TOMR	+	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
11	Oermination	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
12	<i>F. moniliforme</i>	-	+	-	-	-	-	-	-	+	+	-	+	+	+	-	+	-	+	-	+
13	<i>F. pallidirostrum</i>	-	-	-	-	+	+	-	-	+	+	-	+	+	+	-	+	-	+	-	+
14	<i>C. brevis</i>	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
15	Other fungi	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+
16	Proteins	+	-	+	-	+	-	+	-	+	-	+	-	+	-	+	-	+	-	+	-
17	Soluble sugars	+	-	+	-	+	-	+	-	+	-	+	-	+	-	+	-	+	-	+	-
18	Tannins	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
19	Flavan-4-ols	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-

+ Significant and desirable general combining ability estimates

- Non-significant and undesirable general combining ability estimates

X-2

Table : 75 Superior combinations showing significant sea effects for germination (%) with their desirable sea effects for various components

Sr. No	Desirable crosses for germination (%) on the basis of sea effects of F_1 crosses	Sea estimates for germination (%) in F_1 crosses	Sea estimates for germination (%) in F_2 progenies	Significant sea effects for components characters in F_1 crosses		Significant sea effects for components characters in F_2 progenies	
				100 grain weight, electrical conductivity, DTF, mesocarp thickness, TGM, <i>F. moniliforme</i> , protein, tannins and flavan-4-ols		Electrical conductivity and TGM, <i>F. moniliforme</i> , protein, tannins and flavan-4-ols	
1	AKms-14B x IS-6335	15.785**	9.286**				
2	AKms-14B x IS-9471	12.769**	6.604**				
3	IS-2284 x IS-6335	10.839**	-4.031				
4	ICSB-101B x IS-9471	10.526**	4.832				
5	SPV-104 x IS-9471	10.111**	1.296				
6	SPV-104 x IS-2284	9.925**	3.796				
7	ICSB-101B x IS-2284	9.225**	1.152				
8	SPV-104 x IS-6335	8.824**	-1.337				
9	SPV-1201 x SRT-26B	8.229**	8.270**				
10	GI-35-15-15 x IS-6335	7.987**	-1.102				

X - 3

Table : 76 Gene action of various parameters related to grain mold resistance in 10×10 diallel (F_1 - F_2) progenies of sorghum lines

Sr.No.	Characters	Gene action in	
		F_1 diallel	F_2 diallel
1	100 grain weight	Additive	-
2	Grain hardness	Additive	Non-additive
3	Endosperm texture	Additive	-
4	Electrical conductivity	Non-additive	Additive
5	DTF	Non-additive	-
6	Plant height	Non-additive	Additive
7	Cob length	Additive	Additive
8	Glume covering	Additive	Non-additive
9	Mesocarp thickness	Non-additive	-
10	TQMR	Additive	Additive
11	Germination	Non-additive	Non-additive
12	<i>F. moniliforme</i>	Additive	Additive/ Non-additive
13	<i>F. pallidovirium</i>	Non-additive	Non-additive
14	<i>C. lunata</i>	Additive	Additive
15	Other fungi	Non-additive	Additive
16	Proteins	Non-additive	-
17	Soluble sugars	Non-additive	-
18	Tannins	Additive	-
19	Flavan-4-ols	Additive	-

Table: 77 Heterotic crosses for germination (%) showing heterosis and heterobelliosis for other characters

Sr. No	Superior crosses for germination(%) on basis of mean performance	Mean germination of (%) of crosses	Mean performance of P ₁	Mean performance of P ₂	Heterosis for germination (%) over mid parent H ₁	Heterosis for germination (%) over better parent H ₂	Significant heterosis for component characters	
							Over mid parent H ₁ (%)	Over better parent H ₂ (%)
1	ICSB-101B x IS-2284	90.00	36.75	71.75	51.76**	24.28**	Electrical conductivity, mesocarp thickness, TGMAR, <i>P. pallidirostrum</i> , tannins and flavan-4-ols	Electrical conductivity, glume covering, mesocarp thickness, TGMAR and flavan-4-ols
2	ICSB-101B x IS-6335	89.75	36.75	69.50	51.44**	24.69**	Electrical conductivity, DTF, mesocarp thickness, TGMAR, <i>P. moniliforme</i> and flavan-4-ols	Electrical conductivity, DTF, mesocarp thickness, TGMAR, <i>P. moniliforme</i> and <i>C. laevis</i>
3	ICSB-101B x IS-9471	88.25	36.75	63.75	57.68**	33.45**	Electrical conductivity, mesocarp thickness, TGMAR, tannins and flavan-4-ols	Electrical conductivity, glume covering, mesocarp thickness, TGMAR, <i>P. moniliforme</i> C. <i>laevis</i>
4	SPV-1201 x IS-2284	86.75	48.75	71.75	39.30**	22.50**	Mesocarp thickness, TGMAR and flavan-4-ols	Electrical conductivity, mesocarp thickness and TGMAR
5	AKma-14B x IS-6335	85.50	15.75	69.50	76.76**	20.08*	100 grain weight, electrical conductivity, DTF, mesocarp thickness, TGMAR, <i>P. moniliforme</i> , tannins and flavan-4-ols	Electrical conductivity, DTF, mesocarp thickness, TGMAR and <i>P. moniliforme</i>
6	CU-35-15-15 x IS-6335	84.75	51.00	69.50	38.09**	19.76*	Electrical conductivity, DTF, mesocarp thickness, TGMAR and flavan-4-ols	Electrical conductivity, DTF, mesocarp thickness, TGMAR and <i>C. laevis</i>
7	SRT-26B x IS-9471	84.25	46.75	63.75	33.73**	19.74*	Electrical conductivity, mesocarp thickness, TGMAR, tannins and flavan-4-ols	Electrical conductivity, mesocarp thickness, TGMAR, <i>P. moniliforme</i> and <i>C. laevis</i>
8	IS-2284 x IS-9471	84.00	71.75	63.75	19.83**	15.09	Glume covering and mesocarp thickness	Mesocarp thickness and glume covering
9	SRT-26B x IS-6335	82.75	46.75	69.50	33.03**	15.63	100 grain weight, electrical conductivity, DTF, mesocarp thickness, TGMAR C. <i>laevis</i> , tannins and flavan-4-ols	Electrical conductivity, DTF, mesocarp thickness, TGMAR and <i>C. laevis</i>
10	SPV-104 x IS-6335	81.75	28.00	69.50	50.32**	13.79	Electrical conductivity, DTF, cob length, mesocarp thickness, <i>P. moniliforme</i> , soluble sugars, tannins and flavan-4-ols	Electrical conductivity, DTF, mesocarp thickness, TGMAR and <i>P. moniliforme</i>

APPENDIX - XI-1

Table 78. Composition of stain

Mac Grunwald's stain

Methylene blue 0.5%

Eosin Y 0.5% in methanol

Planeze III b

Malachite green 0.5 g

Acid fuschin 0.1 g

Martin gleb 0.01 g

Water 150 ml

Ethanol (95%) 50 ml

Host tissues staining green and fungal mycelium deep pink in lignified and non-lignified tissue.

APPENDIX – XI-1

Table 78. Composition of stain

Mac Grunwald's stain

Methylene blue 0.5%

Eosin Y 0.5% in methanol

Pianeze III b

Malachite green 0.5 g

Acid fuschin 0.1 g

Martin gleb 0.01 g

Water 150 ml

Ethanol (95%) 50 ml

Host tissues staining green and fungal mycelium deep pink in lignified and non-lignified tissue.

VITA

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VITA

Gulab Daultrao Agarkar was borne on August 30th 1947 at Loni-District Amravati. He completed his higher secondary school certificate examination in 1964 from Smt. Ramparibai Chandak High School, Loni, in first division, B.Sc. (Agri.) from Shri Shivaji Agriculture College, Amravati in 1969, and stood 10th in order of merit and M.Sc. (Agri.) in Plant Pathology in 1971 in first division from College of Agriculture, Nagpur, Dr. PDKV, Akola.



He was appointed as Agricultural Officer in 1971 at Agriculture. Research Station, Washim. He was selected as Asstt. Prof. of Plant Pathology in 1977 and till 1993 he was at College of Agriculture Akola. In 1993 he was selected for Ph.D. in service training and served for three years in Sorghum Research Unit as a Sorghum Pathologist. In 1998 he was again posted at College of Agriculture Akola and presently working at same place.

He has published five research articles in National Journals, participated and presented nearly 10 papers in National symposia and workshops. He has guided two students for thesis in M.Sc. (Agri.). He is associated with the release of sorghum hybrids SPH-840 and pre-release hybrids SPH-792, SPH-1010, and W-815 and W-2019 at the State level.

THESIS ABSTRACT

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THESIS ABSTRACT

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- b) Full name of student : Gulab Daulatrao Agarkar
- c) Name and Address of major Advisor : Dr. R.B. Somani, Asso. Prof. of Plant Pathology & Head Deptt. of Agro-Product Processing Centre, Dr.P.D.K. V., Akola - 444 104
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- j) Signature, Name and Address of the forwarding authority : 
(Dr. N.L. Peshney)
Head,
Uni. Deptt. of Plant Pathology
Dr. PDKV, Akola

ABSTRACT

Grain mold of sorghum is a complex problem involving several fungal species. Resistance to grain mold in sorghum is known to be imparted by various factors associated with host genotypes, fungal species and their interactions. The objectives of the present investigations were to : determine infection sites and colonization by major mold fungi (*Fusarium moniliforme*, *F. pallidoroseum* and *Curvularia lunata*); determine physical, physiological and biochemical mechanism of resistance and determine genetics

and heritability of various host factors contributing to grain mold resistance in selected sorghum lines.

The experimental comprised of 10 divergent parents and their 45 F_1 crosses and 45 F_2 , progenies of partial diallel. Experiments were conducted at two locations, Akola under natural condition during 1995 and 1996 and at Patancheru under controlled condition during 1996. Data were analyzed using model I, method 2 of Griffings (1956b) further extended by Singh (1973a, 1973b).

Data were recorded on agronomic, physical, pathological and biochemical parameters. Superior combination showing significant sca effects for germination with their desirable sca effects were observed in crosses Akms 14B x IS-6335, Akms 14B x IS-9471 and SPV-1201 x SRT - 26B. On the basis of superior mean performance in germination, four crosses viz. ICSB-101B x IS-2284, ICSB-101B x IS-6335, ICSB-101B x IS-9471 and SPV-1201 x IS 2284, were selected to understand significant heterosis for component characters. The results revealed that electrical conductivity, mesocarp thickness, thresh grain mold rating (TGM), glume covering, tannins and flavan-4-ols are the major components imparting resistance. As regards to gene action governing inheritance, it was noticed that in F_1 diallel the gene action was non-additive for electrical conductivity, mesocarp thickness, germination, proteins, and it was additive for grain hardness, glume covering, TGM, *F. moniliforme*, *C. lunata*, tannins and flavan-4-ols, whereas, in F_2 diallel, gene action changed for grain hardness and electrical conductivity.

Over all there was decrease in 100-grain weight, grain hardness, germination, *C. lunata* and fungal load; and increase in electrical conductivity, TGM, *F. moniliforme* and *F. pallidoroseum* load at Patancheru compared with Akola location. *F. moniliforme*, *C. lunata*, and *F. pallidoroseum* were important mold fungi, at Patancheru, while *C. lunata* and *F. moniliforme* were important at Akola. *F. pallidoroseum* remained a minor mold fungi at Akola. In general more fungal load was recorded at Akola during 1996 than in 1995. Pre-treatment with $HgCl_2$ reduced fungal load of grain deteriorating fungi and improved seed germination, except *C. lunata* at Patancheru. However, fungal load of all fungi was reduced at Akola in both seasons.

Soluble sugars could not show any effect on grain mold resistance. However, tannins and flavan-4-ols were important biochemical parameters conferring resistance to grain mold singles or in combination. Prolamin and cross-link prolamin were recorded low in colored grain parents and crosses, have resistance to grain mold in colored grain could be attributed to tannins and flavan-4-ols. Prolamin and cross-link prolamin were more in white grain thus contribute for mold resistance. *Fusarial* infection (both species) takes place through hilar areas, however, *Curvularia* infection takes place from both ends i.e. hilar and stylar or directly through pericarp.

Germinated seed (treated with $HgCl_2$) showed significant negative association with fungal load of all fungi at Akola, while it was positive at Patancheru. Ungerminated seed exhibited positive with fungal load of all fungi and score, negative with germination and positive with TGMR. Grain hardness had significantly negative correlation with endosperm texture, however, endosperm texture showed negative association with electrical conductivity, mesocarp thickness, *F. moniliforme*, *C. lunata* and TGMR. Mesocarp thickness exhibited positive correlation with *F. moniliforme*, *C. lunata* and TGMR and negative with germination. Prolamin and cross-link prolamin had showed positive correlation with grain hardness, however it was negative with endosperm texture and germination.

Considering the above results it is concluded that :

1. Parental lines IS-9471, SPV-1201, IS-6335, GJ-35-35-15 and IS-2284 with high gca effect for most of the characters contributing towards resistance to grain mold may be utilized in hybrid breeding program.
2. The cross : Akms 14B x IS-6335, ICSB-101B x IS-9471, ICSB-101B x IS-2284, SPV-104 x IS-6335, GJ-35-35-15 x IS-335 and SPV-1201 x SRT-26B exhibited high gca effects, desirable heterosis for most of the characters and additive gene action for important traits (agronomic, physical and biochemical) related to grain mold resistance. Therefore it is suggested that these crosses and their progenies may be utilized to generate better tolerance to grain mold infection.
3. Important traits imparting resistance to grain mold fungi are : low electrical conductivity, thin mesocarp, low TGMR, grain hardness, more glume covering, more tannins, more flavan-4-ols, more prolamin and cross-link prolamin.