Plant Disease Problems in Central India



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Pathometry of Fusarium Wilt of Pigeonpea

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One of the reasons for low seed yield of pigeonpea in the SemiArid Tropics is the occurrence of diseases, particularly Fusorium wilt (Fusarium udum Butler) which causes substantial losses (Kannaiyan et al. 1984). In recent years, much progress has been made in measuring and analyzing the epidemics of foliar diseases but our understanding about the measurement and quantitative aspects of soil-borne diseases is rudimentary. Analysis of disease variables in meaningful units help in better understanding of diseases and as well help in taking policy decisions about disease management. Hence, we used disease progress curves, area under the disease progress curve (AUDPC), apparent rate of infection (r), and inoculum density and disease relationship (ID-D) to study the epidemiology of fusarium wilt of pigeonpea.

MATERIALS AND METHODS

The experiment was conducted in a well established Fusarium wilt sick plot in vertisol at Patancheru, India. The sick plot was developed during 1975 to 1977 by growing wilt susceptible lines such as ICP 6997 and incorporating the wilted plants into soil, Sickness of the plot was subsequently maintained by inter-planting susceptible lines (1:2 or 1:4) with germplasm accessions and breeding material. The sick plot contined 4945 colony forming units (cfu per gram of soil). Sowing were made lune in 1991 with the inter- and intra-row spacings of 60 cm and 20 cm. respectively. The trial was conducted under rainfed conditions (annual rainfall 718.9 mm), F, udum population in wilt sick soil was estimated at sowing using melachite

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green medium (Singh and Chaube 1970). Three cultivars, ICP 8863 (resistant), C-11 (moderately susceptible) and ICP 2376 (susceptible) were used in the study. The experimental design used was R8D with 2 replications. The wilted plants in different cultivars were recorded at 15 days interval starting from 30 days after sowing.

Computation of apparent rate of infection (r)

Procedures or equations to calculate the apparent rate of infection at different stages have been established by Vanderplank (1963). Hence, the following formula with scope for introduction of correction factor was directly used for computing infection rate:

$$r = \frac{1}{(t_2 - t_1)} \times [\log e \frac{1}{(1 - Y_2)} - \log e \frac{1}{(1 - Y_1)}]$$

where y, and y_2 = disease incidence levels at time t, and t₂ (1-y) = correction factor, which allows for a decreasing proportion of healthy plant left for infection.

Estimation of area under disease progress curves (AUDPC)

The following formula given by Wilcoxson et al. (1975) was used.

AUDPC =
$$\sum_{i=1}^{k} \frac{1/2}{2} (y_i + y_{i+1}) \times d$$

where, $y_i = disease$ incidence at the end of week i; k = number of successive evaluationsof disease; and d = interval between twoevaluations.

Plotting the disease progress curves

Disease incidence was plotted in a cartesian plane taking log e (1/1-y) on the ordinate (Y axis) and the time on the abscissa

(X axis) where (1-y) is the correction factor for disease level.

Inoculum density and disease relation (ID-D)

Eight soil dilutions from wilt sick plot at ICRISAT (4945 cfu/g soil) were prepared in 8⁻diameter pots by mixing the sick soil with sterilized soil. Populations of *F. udum* in different dilutions were estimated by using aselective medium (malachite green). Each soil sample was plated on five petridishes for estimating the inoculum. These inoculum dilutions were used to measure wilt incidence in the three genotype ((CP 2370, C-11) and (CP 8863).

RESULTS AND DISCUSSION

Log transformation as advocated by Vanderplank (1963) was used for linearization of disease progress curve. Initially the progress was conspicuously logarithmic both in susceptible and moderately-susceptible genotypes owing to low rate of infection and thereafter the disease progress switched over to nonlogarithmic phase due to higher infection rate. The incidence level reached a plateau at the termination of the season. The term "logarithmic" usually refers to low infection rate of the disease in disease progress curve. In case of pigeonpea wilt though infection occurs early (with in a week after sowing) disease symptoms appear only after a month. Disease progress is slow during the vegetative phase (4 months) and accelerates only in the reproductive phase. The reasons for such disease pattern are not clear. Some of the possible explanations for higher incidence of wilt in the reproductive phase include: increased susceptability of pigeonpea to wilt in the reproductive stage compared to vegetative stage; slow colonization of F. udum in the host due to woody nature of the plant: and availability

| Days after sowing | ICP 2376 (Suspectable) | | C-11 (Moderately susceptible) | | ICP 8863 (Resistant) | |
|----------------------|------------------------|-------|-------------------------------|-------|----------------------|-------|
| | 1 | AUDPC | 1 | AUDPC | ' | AUDPC |
| 35 | | | | | | |
| 50 | 0.0037 | 110 | 0.0045 | 147 | 0 | 0 |
| 65 | 0.0089 | 232 | 0.0055 | 245 | 0.00670 | 7.01 |
| 80 | 0.0265 | 510 | 0.0180 | 438 | 0.00013 | 16.61 |
| 95 | 0.0271 | 837 | 0.0136 | 665 | 0 | 19.2 |
| 110 | 0.0257 | 1055 | 0.0026 | 765 | 0 | 19.2 |
| 125 | 0.0170 | 1181 | 0.0052 | 806 | 0.00047 | 24.3 |
| 140 | 0.0223 | 1261 | 0.0059 | 861 | 0 | 29.4 |

Table 1 Apparent rate of infection (r) and area under disease progress curve (AUDPC) of Fusarium wilt of pigeonpea

of more moisture in the vegetative stage than in the reproductive stage. The disease progress curve as a result varied slightly from that of typical monocyclic disease pattern.

When the growth curve was analysed in terms of apparent rate of infection (r), the initial rwas 0.0037 and reached the highest r of 0.0271 at 95 DAS in the susceptible ICP 2376 (Table 1 and Figure 1). In moderately susceptible genotype, C-11, the initial rate of infection was 0.0045 but reached the highest rate of 0.0136 at 95 DAS. The resistant genotype ICP 8863 had a very low infection rate of 0.0067 in the logarithmic phase till the end. It is inferred that as the amount of disease increases, the amount of susceptible host roots available for infection would not remain constant. Therefore, at the termination of the growing eason the growth curve reached a plateau. The initial low r in the logarithmic phase accelerated due to higher colonization in the next phase.

Area under the disease progress curve (AUDPC) is another way of expressing the course of disease development and has been considered a reliable and convenient

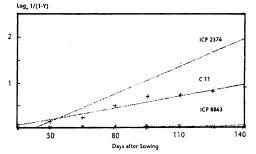


Figure 1. Progress of Fusarium wilt in vertisol

procedure for epidemic analysis. The greatest change in AUDPC value was recorded between 80 and 95 DAS in both susceptible and moderately susceptible genotypes.

Vanderplank (1963) utilizing the data of Ware and Young (1933) depicted a logarithmic growth rate followed by nonlogarithmic disease progress for cotton wilt incited by Fusarium oxysporum f.sp. vasinfectum. The rate of disease increase in the present experiment was similar to that of cotton wilt except for a short duration in each phase of the disease. Such log transformation and calculation of r has been stressed as a superior technique for epidemic analysis by Vanderplank (1963). On the other hand, Wilcoxson et al. (1975) considered estimation of AUDPC to be a more appropriate technique for epidemic analysis than the r value. In case of pigeonpea wilt, both the amount of wilt (indicated by AUDPC) and the time at which wilt occurs (indicated by r) are important. Wilting before pod set causes a total loss and wilting at later stages, results in a partial loss. Hence both r and AUDPC are important in epidemic analysis.

Inoculum density and disease (ID-D)

Wilt was not observed at 1:200 dilution of wilt sick soil (15 cfu/g soil) in susceptible ICP 2376, but it occurred at 1:100 (34 cfu/ g) dilution (Figure. 2) with incidence increasing up to 1:10 dilution (261 cfu/g soil). Wilt incidence in moderately susceptible C-11 was observed at 1:25 dilution (111 cfu/g_soil) and increased at 1:5 dilution (517 cfu/g soil). No wilt was observed at 1:200, 1:100 and 1:50 dilutions. The resistant ICP 8863 had a very low wilt incidence (5%) in undiluted sick soil. In other crops, the number of propagules of Fusarium in soil also had a direct relationship with severity of wilt (Netzer, 1976). However, Cook (1968) did not detect a plateau for Fusarium foot rot of cereals, even at 104 propagules/g of soil. He concluded that in addition to inoculum, there are other factors involved in disease severity. Douglas (1970) favoured the theory of ID-D only over an intermediate range of inoculum concentrations.

Log-log transformation suggested by Baker et al. (1967) was utilized for the study of ID-D relationship in pigeonpea

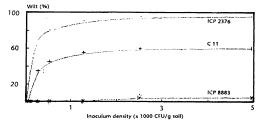


Figure 2. Inoculum density and disease relationship in Fusarium wilt of pigeonpea

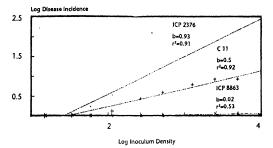


Figure C. Inoculum density and disease model for Fusarium wilt of pigeonpea

wilt (Figure 3). The ID-D indicated a slope value of 0.93 and 0.5 for susceptible and moderately susceptible genotypes, respectively with 91% correlation (Figure 3)

In this model, it is assumed that the root tip moves through the soil and activate increasing proportions of inoculum, Slope values similar to the present values have been reported for related host-pathogen systems, Mitchell and Hurwitz (1975) demonstrated a slope of near 1.0 for Pythium infection of rve. In damping-off of radish, the slope values conformed to this model (Benson and Baker 1974), However, variation in slope values, especially in vascular wilt, is common because of the influence of symptom expression by environmental factors and the absence of symptom expression in the host plant (Baker and Phillips 1962).

The infection rate and AUDPC obtained in this study confirm the resistance of ICP 8863 to will. The r value in a single figure reveals the nature of the pathogen, variety of the host-plant under cultivation and prevalent environmental conditions. The r value coupled with AUDPC for wilt is helpfulin evaluation of pigeonpeas for wilt resistance and for studying the influence of crop seed mixtures, rotations and pathotypes on wilt incidence. Similarly, the ID-D is useful in studying inoculum density in relation to crop sequence, crop density, environmental effects and for comparison of control measures.

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