## 1 First report of Fusarium acuminatum on pigeonpea in India

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Pigeonpea (*Cajanus cajan* (L.) Millsp.) is the most important protein rich grain legume crop 7 being cultivated worldwide. During surveys (2010-2012) conducted in major pigeonpea 8 growing states in southern and central India (Andhra Pradesh, Karnataka and Maharashtra), 9 rapid mortality of pigeonpea plants was observed. This occurred in all of the surveyed areas 10 with disease incidence of 20-60% irrespective of cultivars and crop growth stage. Symptoms 11 included chlorosis, drooping and rolling of the leaves followed by rapid mortality of whole 12 plant. Pinkish growth on infected stems and branches was observed and inner layer of the 13 infected stem had brown discoloration. Xylem vessels of the infected plants were healthy and 14 did not show any blackening. 15

Isolations from infected stem tissues consistently yielded cultures of *Fusarium* sp. on potato 16 dextrose agar (PDA) medium. Monoconidial isolation from three separate isolates was used 17 to establish pure cultures. The morphological characters of the fungus were consistent with 18 descriptions in Fusarium keys (1) for Fusarium acuminatum (Ellis & Everhart). The mean 19 colony growth was 86 mm after 7 days; with white aerial mycelium, developing brownish 20 pigmentation in the center on PDA. The dorsal side of the colony had rose to burgundy 21 pigmentation. Macroconidia were broadly falcate with 3-5 septa, and 3 to  $8 \times 39$  to  $64 \mu m$ . 22 Microconidia were absent and chlamydospores formed in chains, 20 to 50 µm. Koch's 23 postulates were established on seedlings of pigeonpea (cv. ICP 7119) using root dip 24 inoculation of 10-day old seedlings. The roots were immersed in a conidial suspension 25

(6×10<sup>6</sup> conidia/ml) for 2-3 minutes; the control plants roots immersed in sterilized distilled 26 water in beaker. Inoculated seedlings were transplanted into pre-irrigated pots (12 cm) 27 containing sterilized vertisol and sand (3:1). Five seedlings were used for each of 3 28 replications. Inoculated plants were kept in the greenhouse at  $28\pm2^{\circ}C$  and irrigated with 29 sterilized water. Inoculated plants developed symptoms identical to those observed in the 30 field and disease incidence reached 100% within 96 hours after inoculation. Experiment was 31 conducted twice with two independent sets of plants. No symptoms were observed in water-32 inoculated control plants. The rDNA internal transcribed spacer (ITS sequence) was 33 amplified with ITS1 and ITS4 primers (2). The amplicons of both forward and reverse (438 34 bp) were sequenced and submitted to GenBank (Accession no. JX177431). A BLASTn 35 search revealed 100% sequence similarity to the nucleotide sequence of Fusarium 36 37 acuminatum (Ellis & Everhart) (GenBank Accession no. HQ 443205). To our knowledge, 38 this is the first report with confirmed molecular identification of F. acuminatum on pigeonpea. Occurrence of F. acuminatum on various plant species have been reported by 39 Summerell et al. (3). Presence of F. acuminatum from soils of pigeonpea fields have been 40 reported, however, no information on location, symptoms, plant mortality and identification 41 of pathogen has been provided (4). 42

## 43 **References**

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