

SEED BORNE NATURE AND INFECTIVITY OF *FUSARIUM SOLANI* CAUSING CHILLI WILT: ITS EARLY DETECTION FROM SEEDS AND SOIL USING CONVENTIONAL AND MOLECULAR METHODS

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ABSTRACT

Chilli wilt is most destructive emerging disease in south India and causing huge damage to the crop production. The pathogen remains in soil and seed for long time and causes damage year after year. Seed samples of seven popularly growing chilli varieties and rhizosphere soil from hot spots were collected from south Indian states and analysed for the frequency of microflora associated with seeds and causation of wilt disease. The maximum disease incidence during was recorded at Bellary district of Karnataka (29.48% PDI) and Kurnool district of Andhra Pradesh state. Maximum population (13.2×10^5 cfu/g of soil) of *F. solani* was recorded from the rhizosphere soils of these regions. Among the microflora associated with seeds, the frequency of *F. solani* was more (10-56.50 %) which also varied with cultivars with maximum infection on Byadgi dabbi which is the highly susceptible variety. Further, a quick and early detection for the *Fusarium solani* from seed and infected soil was made by using conventional methods viz., standard blotter, rolled towel, seedling symptom and transmission test and polymerase chain reaction (PCR) based molecular method. A significant reduction in germination and vigour index of infected seeds were recorded when compared with healthy control. Translation elongation factor 1 α gene (TEF-1 α) was used for detection of pathogen from seed and rhizosphere soil and an amplified product of 658 bp was observed confirming *F. solani* infection in seed and soil. PCR based detection method was found to be the most sensitive and highly useful in early and quick detection of *F. solani* from seed and rhizosphere soil which can be used in disease management strategies.

INTRODUCTION

Chilli (*Capsicum annum* L.), despite its fiery “hotness” is one of the very popular spice and vegetable crop grown worldwide known for its medicinal and health benefiting properties. During past few years the insect and diseases are increasing in chilli cultivation and taking a heavy toll. Among the diseases, wilt caused by fungal pathogen *Fusarium solani* is emerging as a serious problem in major chilli growing areas of south India. The disease causes a loss up to 2-87% especially in some of the popular varieties viz., byadgi dabbi, byadgi kaddi and other hybrids developed by public and private sectors (Madhukar et al., 2002; Devika Rani, et al., 2007; Raghu, et al., 2013). The disease causes seedling death in nursey if the infected seeds are sown even in non-infected soil and reduces germination and seedling vigour. In main field the plants start wilting gradually and all the leaves become yellow and drooping occurs but leaves will not detach from the plants. Gradually whole field is infected and plants wilt followed by death (Fig.1 & 2). The disease is caused by *Fusarium solani* which is seed and soilborne fungi (Ascomycota: Sordariomycetes: Hypocreales: Nectriaceae). Which is a cosmopolitan soil inhabiting fungus which is known to be phylogenetically diverse. Infected (or) contaminated seeds serve as major source of inoculum for large number of plant

pathogens which may infect the seeds and survive as spore (or) resting structures on (or) within the seeds (Neergaard, 1977).

Some of the earlier works revealed the importance of seed borne detection of plant pathogens. Sajeesh et al. (2014) reported the pathogenic ability of seed borne fungi and proved through transmission studies by seedling symptom test and pot culture studies. The histopathological studies revealed the location of seed borne fungi in the infected soybean seeds. Most of the observations the fungal pathogens were occupied in the pericarp, hilum and endosperm regions. Boyer et al. (2014) detected eight different fungal species namely, *A. alternata*, *A. triticina*, *B. sorokiniana*, *C. lunata*, *D. tetramera*, *A. flavus*, *A. niger* and *F. semitectum* from twenty four wheat cultivars seeds by standard blotter paper method. Among them, incidence of *A. alternata* (41.5%) was dominant. Ellis et al. (2011) conducted a study on *F. graminearum* causes seed decay and damping-off of soybean. This study evaluated the effect of inoculum density of *F. graminearum*, temperature, and fungicide seed treatments on disease development.

There is a need to develop tools which permit rapid, sensitive and specific diagnosis in mixed contaminated samples for timely management of chilli wilt. Polymerase chain reaction (PCR) which has emerged as a major tool for the diagnosis of phytopathogenic fungi and contributed to the alleviation of

some of the problems associated with the detection and containment of plant pathogens can be used in these detection studies (Henson and French 1993; Martin *et al.*, 2000; Ghignone and Migheli, 2005). The objectives of the present study is to identify the hot spots of fusarium wilt based on disease incidence, estimation the frequency of microflora associated with chilli seed and rhizosphere soil to study the relation between pathogen population and disease severity. To develop an early and quick detection technique for the *F. solani*, from seed and soil by employing both conventional and molecular techniques.

MATERIALS AND METHODS

Source of isolates and collection of seed and soil samples

A roving survey was conducted for two seasons (2012-13 & 2013-14) in major chilli growing areas of Karnataka, Andhra Pradesh (Present Andhra Pradesh and Telangana), and Maharashtra states. Seeds from harvested/stored lots (heaps), infected plant and soil samples from the rhizosphere of naturally infected soil were collected. The disease incidence was calculated by the following formula.

$$\text{Percent disease incidence(\%)} = \frac{\text{Number of plant infected}}{\text{Total number of plants examined}} \times 100$$

All samples were brought to the laboratory and stored at -80°C. The samples were further processed for isolation and identification of different microflora by following different seed health testing methods given by ISTA. Isolation of *F. solani* from soil was carried out by serial dilution technique on specific media. The pathogen population (cfu/g of soil) was correlated with disease incidence. The pathogen pure cultures were stored at 4 °C for further use.

Detection methods

Blotter Method

The collected seed samples were analyzed for the presence of seed borne fungal microflora by blotter method given by International rules for Seed Testing (ISTA, 1999). Four hundred seeds were tested for each variety with three replications. Chilli seeds were placed on three layers of moist blotting paper in each glass petridis and incubated at 27 ± 1°C under 12 hrs light and 12 hrs darkness for 7 days. Each seed was observed under stereo binocular microscope in order to record the presence of fungal colony based on growth habit. The pathogens associated with seeds were detected by following the keys given by Mathur and Kongsdal (2003) based on the growth characters. Temporary slides were prepared for proper identification of fungi under compound microscope at 10x and 40x and identified with the help of Keys suggested (Booth, 1971). The results were presented as percent incidence of individual pathogen. The fungi from the incubated seeds were transferred to PDA for further confirmation and estimation of frequency of the microflora. The culture was incubated at 27 ± 10°C for seven days. Per cent frequency (PF) of occurrence was calculated by applying the following formula (Javaid *et al.*, 2006). PF = (No. of seeds on which fungus appear/ Total No. of Seeds) × 100

Rolled Paper Towel Method

Germination ability of the seeds was determined in the

laboratory at room temperature (27 ± 1°C) by following the method developed by Warham (1990). 400 seeds were randomly taken from each variety and 100 seeds were placed between a pair of moist paper towels by maintaining three replication. The towels were rolled and the ends were closed by threads and covered by polythene paper to prevent drying. After 10 days of incubation, the observations like germination percentage, non germinated seed (hard seed and rotten seed), post-emergence death, shoot length, root length, vigor index and disease incidence were recorded. Vigour of the seedling was determined by using the formula given by Abdul Baki and Anderson (1973).

$$\text{Vigour Index} = (\text{Mean of root length} + \text{Mean of shoot length}) \times \text{Percentage of seed germination}$$

Seedling symptom and transmission studies

Chilli seeds collected from various surveyed areas were further examined by seedling symptoms test. Test tubes (100 x 16 mm) were filled with 5-6 ml of 2 percent water agar and prepared slants. Infected seeds were placed in test tubes (one seed/tube) and incubated at 25 ± 1°C with 12 h of alternate light and dark conditions. When the germinated seedling reaches the rim of test tube, cotton plug was removed and observation was recorded based on the symptom (Khare, 1996). Both naturally infected and artificially inoculated seed were used for this study.

For transmission studies, 60 seeds (variety: Byadgi dabbi) were sown in pots (20 seeds/pot) containing sterile soil. Plants were kept at glass house with automatic temperature and humidity control system. The pots were watered and observed daily for 15 days and percent germination, seed rot and seedling death were recorded and compared with control. The pathogen from the infected seedling was re-isolated on PDA for the confirmation.

Extraction of Genomic DNA

The isolates of *Fusarium solani* were cultured on potato dextrose agar and subsequently on potato dextrose broth (PDB) at 27 ± 1°C for 7 days. After incubation, the mycelial mats were harvested by filtration and stored at -80°C for genomic DNA extraction by following the protocol of Murray and Thompson (1980) with slight modifications. Filtered mycelium (3 g) was ground to a fine powder in liquid nitrogen, transferred to 1 ml of CTAB extraction buffer (0.1M Tris, 1.5 M NaCl, 0.01M EDTA and 2 % CTAB) in a 2-ml tube and kept at 65°C for 90 minutes with occasional stirring. An equal volume of chloroform: isoamyl alcohol (24:1) was added to all and the centrifuged at 10,000 rpm for 20 min at 4°C. The upper supernatant was pipetted out and transferred to a fresh tube and mixed with 100 µl of sodium acetate (3 M) and 600 µl of ice-cold isopropanol and centrifuged at 10,000 rpm for 10 min at 4°C. The pellet was washed with 70 % ethanol, dried at room temperature and finally dissolved in T₁₀E₁ (10 mM Tris-Cl, pH 8.0; 1 mM EDTA) for further use. The DNA was quantified using spectrophotometer and quality analysis was done on 1.0 % agarose gels.

Primers and PCR conditions.

The TEF-1 α gene was amplified using specific primers Fs4-f (5'ATCGGCCACGTCGACTCT3') and Fs4 (5'GGCGTCTGTTGATTGTAGC3') reported by Abd-Elsalam *et al.*, 2006 and

Arif *et al.*, 2012. The primers were synthesized by Bangalore Genei Pvt Ltd. Bangalore and supplied as lyophilized products of desalted oligos. PCR amplification was carried out in a final volume of 20 μ l, which consisted of 2X PCR assay buffer with 15 mM $MgCl_2$ - 2 μ l, 2.5 mM dNTPs mixture 1 μ l, , Primer (5PM/ μ l) 1 μ l forward and reverse each, *Taq* DNA polymerase (6.0U μ l⁻¹) (Bangalore Genei, India) -0.5 μ l , and Template DNA (25 ng/ μ l)-1 μ l . All the reactions were carried out in a Corbett Research gradient thermocycler (96 well system) supplied by M/s JH BIO Innovation Pvt. Ltd R.T. Nagar, Bangalore. The PCR reactions were carried out with an initial denaturation at 95°C for 5 min, followed by 40 cycles of denaturation at 90°C 1 for min, annealing at 58°C for 1 min, 72°C for 2 min, and final extension at 72°C for 20 min. The amplified products were resolved by electrophoresis in 1.2 % agarose gels with 1X TBE (Tris Borate EDTA) buffer, 0.5 μ g ml⁻¹ of ethidium bromide and loading buffer (0.25% Bromophenol Blue in 40% sucrose). Four μ l of the loading dye was added to 20 μ l of PCR product and loaded to the agarose gel. Electrophoresis was carried at 75 V for 1 h. The gel was visualized under UV light, and photographed with a gel documentation unit (Syngene Inc., Cambridge, UK).

RESULTS AND DISCUSSION

Survey and identification of hot spots for the chilli wilt

Seedborne pathogens possess a serious threat to better crop health and establishment. The pathogens in close association with seeds making it niche for its long-term survival, introduction into new areas and widespread dissemination of pathogens. Survey during 2012-13 and 2013-14 for chilli wilt disease is presented in Table 1. The data indicated that, the maximum disease incidence was recorded in Bellary district of Karnataka (PDI: 29.48%) and Kurnool district of Andhra Pradesh (PDI: 34.90%) (Fig-5). A significant disease incidence and crop losses also recorded from Raichur, Koppal, Haveri and Belgaum districts of Karnataka, Guntur and Ananthpur districts of Andhra Pradesh states. These areas are identified as hot spots for disease incidence. The disease incidence also varied with variety to variety with maximum disease incidence was recorded in byadgi dabbi (range: 30-80%) followed by sitara (20-60 %). A positive correlation between disease incidence



Figure 1: Disease free field

and pathogen population was observed in all the surveyed places with maximum average population 13.2×10^{-5} /g of soil was recorded from these hot spots indicating the direct relation between disease incidence and pathogen population

Identification of microflora associated with chilli seeds and detection of seed borne *F. solani*.

Chilli fusarium wilt is becoming more and more serious in traditional chilli growing areas of south India. Its early and exact detection and diagnosis is very first step to ensure efficient disease management. Careful examination of uprooted, plants must be confirmed for the absence of external root symptoms and presence of dark-brown discoloration in xylem tissues of roots and stem will help in disease diagnosis.

Standard blotter method

A total of 40 seed samples were analysed in this study to assess the frequency of microflora associated with seeds by employing conventional seed health testing methods as described in material and methods. In all the samples the common microflora observed was *Fusarium solani*, *Colletotrichum* spp., *Aspergillus* spp., *Mucor* and *Rhizopus* spp. The frequency of *F. solani* was more with 10-56.50 percent followed by *Colletotrichum* spp. with 5-15 per cent (Table 2). *Fusarium solani* caused white mycelium growth on the seed and the microscopic observations reveal that, it produced microconidia with single cell ovoid shape, macroconidia with 3-4 septa, sickle shaped with blunt ends. Chlamydospores were formed either solitary or in chain at the middle or end of the mycelium. These morphological characters confirmed the infection of *F. solani* in seeds.

Rolled paper towel method

The germination percentage and vigour index was significantly reduced when tested against *F. solani*. The germination percentage of *F. solani* infected seeds was significantly low (10 %) when compared with hot water treated seeds (97.50%) which was taken as control (Table 3). The seeds which show low incidence of *F. solani* shown higher incidence of *Aspergillus* and *Penicillium* spp. a significant frequency of *Colletotrichum* spp. also observed resulting in poor germination and seedling vigour index (Table 3). The decline in the seed germination and seedling vigour are due to the toxins produced by these pathogens. Many workers



Figure 2: Completely infected field.

Table 1: Average disease incidence of chilli wilt disease in major growing states of south India.

State	District	Taluk	Per cent disease incidence			
			2012-13	2013-14	Pooled	
Karnataka	Bagalakot	Kaladgi	-	8.5	8.5	
		Mean	-	8.5	8.5	
	Belagavi	Saundatti	6.76	8.65	7.7	
		Bailahongal	17.07	24.82	20.94	
	Bellary	Mean	11.91	16.73	14.32	
		Hospet	12.25	13.38	12.81	
	Chickmagalur	Bellary	Siraguppa	34.42	49.93	42.18
			Bellary	28.31	46.49	37.4
		Kampli	21.73	29.38	25.55	
		Mean	24.18	34.79	29.48	
		Tarikere	7.23	-	7.23	
	Chitradurga	Kadur	Birur	5	-	5
			Mean	6	-	6
		Hosdurga	Mean	6.08	-	6.08
			Hosdurga	8.58	7.82	8.2
	Davangere	Chitradurga	Chitradurga	8.36	10.23	9.29
			Hiriyur	11.33	13.75	12.54
		Mean	Mean	9.42	10.6	10.01
			Davangere	6.02	8.5	7.26
	Dharwad	Jagaluru	Jagaluru	8.49	10.75	9.62
			Mean	7.25	9.62	8.44
		Dharwad	Dharwad	11.3	14.23	12.76
			Hubli	6.44	10.82	8.63
	Gadag	Kundgol	Kundgol	10.07	11	10.53
			Kalghatgi	5.88	13.25	9.56
		Mean	Navalagund	5.8	7.31	6.56
			Mean	7.9	11.32	9.61
	Haveri	Gadag	Gadag	7.18	8.19	7.68
			Nargund	9.67	13.09	11.38
		Mean	Mean	8.42	10.64	9.53
			Haveri	20.07	27.94	24
	Andhra Pradesh	Koppal	Ranebennur	11.98	22.63	17.3
			Byadgi	25.57	37.62	31.6
			Hirekerur	13.38	18.88	16.13
			Hanagal	8.5	10.25	9.38
			Shiggaon	9.51	16.88	13.19
		Raichur	Mean	14.83	22.36	18.6
			Koppal	20.3	27.08	23.69
			Mean	20.3	27.07	23.69
			Sindnur	9.38	19.13	14.25
Manvi			13.86	26.75	20.31	
Uttara Kannada	Raichur	Raichur	26.69	38.49	32.59	
		Mean	16.64	28.12	22.38	
	Halyala	Halyala	25.25	-	25.25	
		Joida	20	-	20	
Maharashtra	Kurnool	Mean	22.63	-	22.63	
		Mantralaya	26.01	34.88	30.44	
		Emmiganur	29.12	41.15	35.13	
		Adoni	32.52	45.72	39.12	
		Mean	29.21	40.58	34.9	
	Guntur	Sattanapalli	25.33	34.03	29.68	
		Guntur	21.97	28.85	25.41	
		Mean	23.65	31.44	27.54	
	Ananthpur	Ananthpur	19.34	27.88	23.61	
		Mean	19.34	27.88	23.61	
Solapur	Pandrapur	Pandrapur	6.13	9.57	7.85	
		Sangola	5.81	8.07	6.94	
	Mean	Mean	5.97	8.82	7.39	
		Grand mean	14.75	21.06	17.36	
Standard deviation		8.65	12.66	10.47		

carried out investigations and detected various pathogen microflora associated with chilli seeds such as, *Collectotrichum capsici*, *Curvularia lunata*, *Fusarium solani*, *Alternaria* spp. *F.*

oxysporum (Mridha and Siddique, 1989; Hasmi, 1989). Similar reports were given by the studies of Solanke (2001) and Asamul (2001) along with other pathogens such as *Aspergillus flavus*,

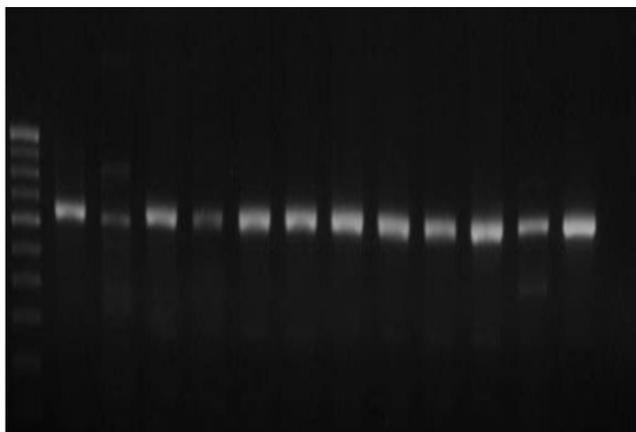


Figure 3. Amplification pattern of TEF 1a gene specific to *Fusarium solani* from seed samples of different cultivars. Lane 1-Marker 100 kb, 1-Byadgi dabbi, 2-Sitara, 3-Nayanatara, 4-Gayathri, Guntur local, 6-Byadgi kaddi, 7-Uttal

collected from surveyed area were sown in pots (20 seeds/ pot) containing sterile soil. Plants were kept at glass house with automatic temperature and humidity control system. The pots were watered and observed daily for 15 days and percent germination, seed rot and wilt incidence were recorded and compared with control. The infected seeds showed lower germination (10.25-50.25) and higher seedling mortality (5-60%). Whereas the germination (98.50%) was more and seedling death (2.50%) was very low in healthy control treatment.

Detection of *F. solani* in chilli seeds and rhizospheric soil by Specific amplification of *Fusarium solani* using TEF 1 α .

Earlier it was believed that seed borne pathogens could be only detected only by conventional methods like, plating the seeds on blotters or on PDA. But with the advent of PCR based methods, it became very easy to detect seed borne pathogens in a mixed and contaminated infections by designing the specific markers. This early and quick detection methods are

Table 2 : Incidence and frequency of *Fusarium solani* in seed samples collected from different parts of south India by Standard Blotter method

Sl No	Place of collection	Sources	Variety	Germination (%)	% incidence of <i>F. solani</i>
1	Kurgod (Bellary, KAR)	Farmer's field	Byadgi kaddi/ B. dabbi	16.5	60
2	Nelahal (Raichur, KAR)	Farmer's field	Byadgi kaddi/ B. dabbi	10.25	52.25
3	Basavapura(Koppal, KAR)	Farmer's field	Byadgi kaddi/ B. dabbi	25.75	35.25
4	Shidenur (Haveri, KAR)	Farmer's field	Byadgi kaddi, B. dabbi, Sitara, Nayanatara	40.25	30.25
5	Inchal (Belgaum, KAR)	Farmer's field	Sitara, Nayanatara	15.1	25.25
6	Amminabavi(Dharwad, KAR)	Farmer's field	Sitara, Nayanatara	26.25	4.025
7	UAS Dharwad (KAR)	UAS horticulture field	Byadgi Dabbi, Byadgi Kaddi, GPM-9	25.5	10.25
8	Betageri (Gadag, KAR)	Farmer's field	Byadgi kaddi/ B. dabbi	30.25	40.5
9	Kattavaripalem (Guntur, AP)	Farmer's field	Guntur local, Gayathri	65.5	13.25
10	Madire (TS)	Farmer's field	Dabbi, Hybrids, Guntur, Brahma	50.25	12.25
11	Emmiganur (TS)	Farmer's field	Dabbi, Hybrids, Guntur	25.52	30.52
12	Pandrapura (MH)	Farmer's field	Dabbi, Hybrids, Guntur	75.5	5.5

Table 3 : Germination percentage of chilli seeds infected by *Fusarium solani*

Treatment	Seed rot	Root rot	Healthy seedlings	Germination percentage	Seedling Vigoure index
Infected	160	20	20	10	180
Control (hot water treated)	0	0	195	97.5	9750
percentage	80	10	10		

A. niger, *Rhizopus stolonifer*. Varied frequency of microflora associated with infected chilli seeds were reported by Padaganur and Naik (1991). They reported *Colletotrichum capsici* (75.5 %), *Fusaium* spp (16.25%) and *Alternaria* spp. (5 %).

Seedling symptom and transmission test.

Seedling symptom test

Selected byadgi dabbi seed samples collected from surveyed locations subjected to seedling symptom test under two percent water agar in test tubes. The germinated seedling shown wilting symptom 30 days after inoculation. The symptoms were observed as gradual wilting and drooping of leaves. Later the plants died. The dead plants were cut into small bits and placed on PDA to confirm *F. solani* infection.

For transmission studies, seeds of variety byadgi dabbi

more and more useful in quarantine stations in order to check the entry of new pathogen thereby minimise the losses due to possible outbreaks.

Based on survey data for the disease incidence and severity, seven popularly growing genotypes and 12 rhizospheric soil samples from hot spots were collected and analysed for seed and soil borne nature of *F. solani* using *Fusarium solani* species specific primer sets Fs4-f (5'ATCGGCCACGTCGACTCT3') and Fs4 (5'GGCGTCTGTTGATTGTTAGC3'). The total genomic DNA from the pathogen isolated from infected seed and soil was used as a template. An amplification product of A 658 bp region of TEF1 α gene specific to *Fusarium solani* (Fig. 3 & 4) confirmed that the pathogen is seed as well as soil borne. This PCR based detection will help in early detection of the pathogen and avoids further losses. Detection of *Fusarium solani* based on morphological discrimination requires special

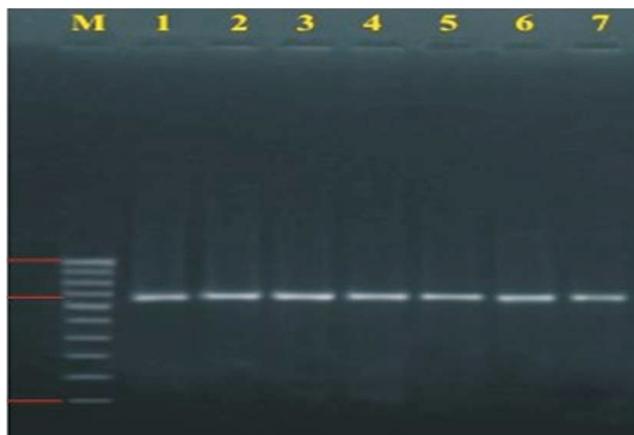


Figure 4 : Amplification pattern of TEF 1 gene specific to *Fusarium solani* from Rhizosphere soil of infected field. Lane 1: Marker 100 KB, 1 Kurgod (Bellary, KAR), 2 Nelahal (Raichur, KAR), 3 Basavapura, (Koppal, KAR), 4 Shidenur (Haveri, KAR), 5 Inchal (Belgaum, KAR), 6 Amminabavi(Dharwad, KAR), 7 UAS Dharwad (KAR), 8. Betageri (Gadag, KAR), 9.Kattavaripalem (Guntur, AP), 10 Madire (TS), 11 Emmiganur (TS), 12 Pandrapura (MH).

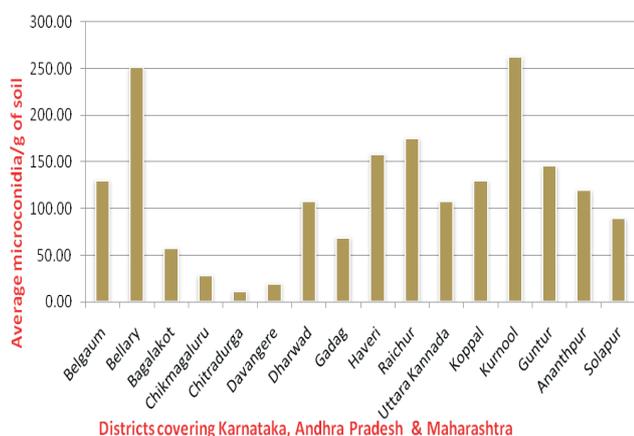


Figure 5: Average incidence of disease in major growing regions of Karnataka, Andhra Pradesh, Telangana state and Maharashtra

skill and the expertise of taxonomists or specialists. Arif *et al.* (2012) carried out the detection of *F. solani* causal agent of wilt and rots in many plant species using PCR based methods by amplification of target genes such as, internal transcribed spacer (ITS), rDNA and transcription elongation factor (TEF-1 α). As there is a saying in sanskrit "Subheeje sukshetre sampadhe jayathe" which means healthy seeds in disease free soil will germinate and establish well and ultimately gives good yield. So right selection of seeds become very important. Confirmation of disease free seeds before sowing by conventional and molecular detection will minimises further crop losses.

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REFERENCES

- Ahmad, Salman., Ansari, Mohammad Shafiq. and Hussain, Mazhar. 2014. Effect of insecticides on management of *Helicoverpa armigera* in chickpea agro ecosystem. *Annals of Plant Protection Science*. **22** (1): 107-111.
- Ali, Ghulam Abbas., Saleem, Akhtar., Zafar, Abbas. and Muhammad, Aslam. 2008. Efficacy of Neem (*Azadirachta indica* A. Juss) products against *Helicoverpa armigera* (Hubner) on chickpea. *Science International (Lahore)*. **20**(4): 281-283.
- Alves, SB. and Lopes, R. B. 2008. Controle microbiano de pragas na América Latina. avanços e desafios. Piracicaba: FEALQ. P.414.
- Bhushan, S., Singh, R. P. and Shanker, Ravi. 2011. Bioefficacy of neem and Bt against pod borer, *Helicoverpa armigera* in chickpea. *J. Biopesticides*. **4**(1): 87-89.
- Bhushan, S., Singh, R. P. and Shanker, Ravi. 2012. Biopesticidal management of yellow stem borer(*Scirpophaga incertulas* Walker) in rice. *The Bioscan*. **7**(2):317-319.
- Chandel, Rajeev., Lyall, Hemant. and Bhamba, DR. 2014. Efficacy of insecticides and neem products against *Helicoverpa armigera* on chickpea. *Ann. Pl. Protection Science*. **22**(1): 205-206.
- Gundannavar, K. P., Lingappa, S. and Giraddi, R. S. 2007. Study of dose mortality response between instars of *Helicoverpa armigera* (Hubner) and *Metarhizium anisopliae* (Metschinkoff) Sorokin. *Karnataka J. Agricultural Sciences*. **20**(1):140-141.
- Haque, J. and Ghosh, A. B. 2007. Effect of *Beauveria bassiana* (Bals.) Vuill, *Metarhizium anisopliae* (Metsch.) Sorokin and Nimbecidine on some insect pests. *Environment and Ecology*. **25**(1): 209-211.
- Kale, S. N. and Men, U. B. 2008. Efficacy of microbial insecticides and their combinations against *Helicoverpa armigera* (Hubner) on chickpea. *J. Biological Control*. **22**(1):205-208.
- Katole, S. R., Nimbalkar, S. A., Kolhe, A. V., Ghuguskar, H. T. and Yadgirwar, P. V. 2000. Performance of some IPM modules against *Helicoverpa armigera* on chickpea. *PKV research J.* **24**(1): 51-53.
- Kavitha, E., Kingsley, S., Revathi, N. and Sathivel, M. 2009. Insecticidal activity of neem derivatives against okra fruit borer *Helicoverpa armigera* Hubner. *International J. Agricultural Sciences* **5**(2): 528-530.
- Kulkarni, K. A., Kambrekar, D. N. and Gundannavar, K. P. 2005. Management of *Helicoverpa armigera* Hubner on chickpea through biopesticides. *Karnataka J. Agricultural Sciences*. **18**(4):1114-1116.
- Kumar, Vinod and Chowdhry, P. N. 2004. Virulence of entomopathogenic fungi *Beauveria bassiana* and *Metarhizium anisopliae* against tomato fruit borer, *Helicoverpa armigera*. *Indian Phytopathology*. **57**(2): 208-212.
- Moorthy, Dakhsana., Anandhi, P., Elamathi, S. and Simon, Sobita. 2011. Evaluation of bio- rational insecticides for management of *Helicoverpa armigera* in chickpea. *Annals of Plant Protection Science*. **19**(1):207-209.
- Murray, DAH., Lioyd, R. and Buddington, J. 2000. Potential in Australia for a *Helicoverpa baculovirus*. Abstract. International Congress of Entomology, Aug 2000, Igassu Falls, Brazil. PP. 21- 25
- Nahar, P., Yadav, P., Kulye, M., Hadapad A., Hassani, M., Tuor, U., Keller, S., Chandele, A. G., Thomas, B. and Deshpande, MV. 2004. Evaluation of indigenous fungal isolates, *Metarhizium anisopliae* M34412, *Beauveria bassiana* B3301 and *Nomuraea rileyi* N812 for the control of *Helicoverpa armigera* (Hubner) in pigeonpea field. *J. Biological Control*. **18**(1):1-7.
- Pawar, C. S., Bhatnagar, V. S. and Jadhav, D. R. 1986. Heliothis

species and their natural enemies with their potential for biological control. In Proceed. Indian Acad. Sci. (Animal Sciences). **95**: 697-703.

Puri, S. N., Sharma, O. P., Murthy, K. S. and Rajshe, O. 1998. Preparation of neem seed kernel extract. " Handbook on diagnosis and integrated management of cotton pests." NCIPM New Delhi, p.106.

Rao, G. V., Rao, V. R., Gottumukkala, S. J., Vidya, M. S., Srinivas, V. and Linga, O. R. 2011. Efficacy of botanical extracts and entomopathogens on control of *Helicoverpa armigera* and *Spodoptera litura*. *African J. Biotechnology*. **10(73)**:16667-16673.

Regupathy, A., Armes, N. J., Asokan, G., Jadhav, D. R., Soundarajan, R. P. and Russell, D. A. 1997. Best method for insecticide resistance management of *Helicoverpa armigera*. In:International Conf. Integrated Approach to Combating Resistance. A.L. Devonshire (ed.), IACR, Rothamsted, Harpenden, UK. p.116.

Rijal, J. P., Yubak Dhoj, G. C., Thapa, R. B. and Kafle, L. 2008. Efficacy of *Metarhizium anisopliae* and *Beauveria bassiana* against *Helicoverpa armigera* in chickpea under field condition in Nepal. *Formosan Entomology*. **28**: 249-258.

Rijal, J. P., Yubak Dhoj, G. C., Thapa, R. B. and Kafle, L. 2008 a. Virulence of native isolates of *Metarhizium anisopliae* and *Beauveria*

bassiana against *Helicoverpa armigera* in chickpea under field condition in Nepal. *Formosan Entomology*. **28**: 21-29.

Singh, B. and Yadav, R. P. 2005. Field efficacy of some microbial agents against *Helicoverpa armigera* Hub. on chickpea. *J. Applied Zoological Researches*. **16(1)**:5-6.

Singh Rajendra, Singh Sarika, Apalwal Meena and Anandhi P. 2012. Management *Helicoverpa armigera* on chickpea through biopesticides. *Annals of Plant Protection Sciences*. **20(1)**:215-216.

Sithanatham, S., Rao, V. and Ghalor, M.A. 1984. International review of crop losses caused by insects on chickpea. In: Proceed. National Semi. *Crop losses due to Insect Pests*. pp. 269-284.

Snedecor, G. W. and Cochran, W. G. 1967. Statistical Methods, Oxford and IBH Publishing Company, New Delhi. pp.1-292.

Vanladiki, H., Singh, Premjit, M. and Sarkar, P. K. 2013. Efficacy of ecofriendly insecticides on the management of diamondback moth (*Plutella xylostella*) Linn. On cabbage. *The Bioscan*. **8(4)**:1225-1230.

Wakil, Waqas., Muhammad, Ashfaq., Ghazanfar, M.U., Saleem, Akhtar. and Malhi, Z.A. 2008. Management of *Helicoverpa armigera* on chickpea. *Pakistan Entomologist*. **30(1)**: 51-54.

Wakil, W. and Ghazanfar, M. U. 2011. Effects of *Metarhizium anisopliae*, *Bacillus thuringiensis* and new chemistry insecticide on

