



Original Research Article

Study of wilt producing *Fusarium* sp. from tomato (*Lycopersicon esculentum* Mill)

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A B S T R A C T

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There are many possible causes of wilting of tomato plants. Successful treatment of the problem depends on accurate diagnosis and appropriate preventive measures. *Fusarium* wilt is caused by a fungus, *Fusarium*, that enters the plant through the roots and grows up through the vascular tissue. The fungus destroys cells of the vascular tissue, causing starvation in nearby branches of the plant. Disease development is favored by warm temperatures, dry weather, acidic soil and root-knot nematodes. A total of 36 *Fusarium* isolates were recovered from 108 samples collected from different geographic regions of Kadi. This study was based on both morphological and molecular levels that confirmed *Fusarium equiseti* is also pathogenic agent for tomato wilt.

Introduction

Tomato (*Lycopersicon esculentum* Mill.) is one of economically the most important vegetable crops in India, where it is grown both, indoors and outdoors on an area of about 20,000 ha in total. A number of economically important tomato diseases caused by fungi are transmitted by seed or transplants. Tomatoes are parasitized by a number of pathogens, including *Fusarium oxysporum* f. sp. *lycopersici* (Sacc.) (W.C. Snyder *et al.*, 2003) the causal agent of fusarium wilt, which is one of the most important species as tomato pathogen (Jones *et al.*, 1982; Agrios, 1988; Smith *et al.*, 1988). An indoor environment due to high

temperature and humidity, *F. oxysporum* f.sp. *lycopersici* can cause significant damage. The causal agent of *Fusarium* wilt is soil borne pathogen which can persist many years in the soil without a host. Most infections originate from the population associated with infected tomato debris. Healthy plants can become infected by *F. oxysporum* if the soil in which they are growing is infested with the pathogen (Farr *et al.*, 1989). However, pathogenic fungi too of the genus *Fusarium* that is the causal agents of tomato wilt cause root and basal stem deterioration and result in the wilting of vegetable plants. Browning of the vascular tissue is strong evidence of *Fusarium* wilt (Snyder and Hans, 2003).

Materials and Methods

Wilted plants of tomato were collected from different areas of Kadi Taluka at Gujarat state of India. Isolation of fungus was done from the diseased portions of tomato plants. Infected leaves and stem portions of tomato were collected from cultivated field of North Gujarat. Infected stem and leaves sections were surface-sterilized for 3 min with a 0.01% Hgcl₂ solution, rinsed twice in sterile distilled water and dried in a laminar flow cabinet. Potato Dextrose Agar (PDA) was used for fungal isolations. The plates were incubated at 25°C in incubator for 5-7 days. *Fusarium* isolates were subcultured on PDA, using a single spore technique (J. F. Leslie and B. A. Summerrell, 2006). Cultural characters were observed by eye and microscopic examination. Colony morphology was observed from PDA plates. Morphological identifications of isolates were made using the criteria of Gerlach and Nirenberg, 1982; J. F. Leslie and B. A. Summerrell, 2006.

Microscopic Observation

To confirm the exact identify of pathogen morphological studies of the mycelium and conidiospores were done by measuring the length and breadth of the spore under high power magnification using ocular micrometer. Photographs were taken by attached camera with light microscope.

DNA Extraction

Fusarium isolate was grown on PDA plates for 7 days and mycelia were harvested. Total DNA was extracted from ground mycelium of isolate (~100 mg wet weight) using a Genei fungal DNA extraction Kit (Genei,

India) according to the manufacturer's instructions done by Agarkar Research institute, Pune.

BLAST

The approach for identifying an isolate of *Fusarium* using the FUSARIUM-ID database it involves obtaining a pure culture of an isolate, extracting genomic DNA, amplification of the TEF gene region, and sequencing. BLAST is then used to identify the closest matches between the unknown sequence and those contained in the FUSARIUM-ID sequence database.

Results and Discussion

One hundred and eight (108) wilted plants were collected from different areas of Kadi. Total of 36 tomato fields in Kadi taluka were sampled during disease season. Each field was arbitrarily divided into five circular plots approximately 100 m in diameter and two to four samples were randomly taken from each plot. Samples were pooled in each field and two infected plant from each field were selected and used for pathogen isolation. A total of 36 *Fusarium* isolates were recovered from 108 samples collected from different geographic regions (Table-1).

Morphological Identification

Isolations from the leaves and stems of diseased plants yielded fungus is characterized by the development of abundant white aerial mycelium which turns pinkish by keeping in daylight on Potato Dextrose Agar (PDA) (figure-1 & 2). The colonies produced macro- and micro-conidia within 3-4 days at 25±2°C.

Table.1 Sample Collection from Different Geographic Regions

Sr. NO.	Village	Field	No. of sample
1	Nani Kadi	4	12
2	Sujatpura	4	12
3	Narsinhpura	3	09
4	Kasva	4	12
5	Vidaj	3	09
6	Balasar	4	12
7	Rangpurda	4	12
8	Borisana	4	12
9	Sadara	3	09
10	Daran	3	09
Total	10	36	108

Figure.1 Growth of *Fusarium* on PDA

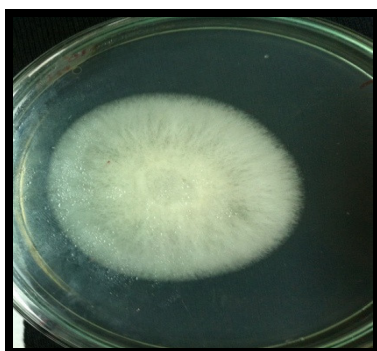


Figure.2 *Fusarium* mycelia

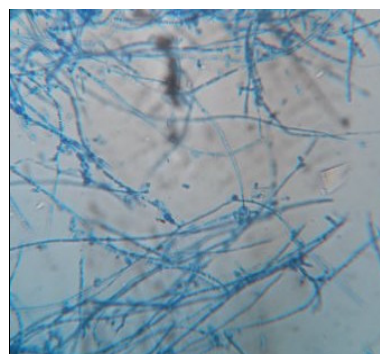


Figure.3 Observation of *Fusarium* conidia under Hemocytometer

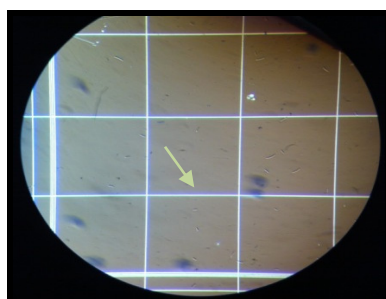


Figure.4 *Fusarium* conidia (45X)



Score	Expect	Identities	Gaps	Strand
852 bits(461)	0.0	475/482(99%)	0/482(0%)	Plus/Minus

Range 1: 11 to 492 [GenBankGraphics](#) [Next Match](#) [Previous Match](#)

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Query 1 TACGGCGTGGCCGCGACGATTACCAGTAACGAGGTGTATGATTACTACGCTATGGAAGCT 60
      |||
Sbjct 492 TACGGCGTGGCCGCGACGATTACCAGTAACGAGGTGTATGATTACTACGCTATGGAAGCT 433

Query 61 CGACGTGACCGCCAATCGATTTGGGGAACGCGGGTTACCGCGAGTCCCAACACCAAGCTG 120
      |||
Sbjct 432 CGACGTGACCGCCAATCGATTTGGGGAACGCGGGTTACCGCGAGTCCCAACACCAAGCTG 373

Query 121 AGCTTGAGGGTTGAAATGACGCTCGAACAGGCATGCCCGCCAGAATACTGGCGGGCGCAA 180
      |||
Sbjct 372 AGCTTGAGGGTTGAAATGACGCTCGAACAGGCATGCCCGCCAGAATACTGGCGGGCGCAA 313

Query 181 TGTGCGTTCAAAGATTTCGATGATTCACTGAATTTGCAATTCACATTACTTATCGCATTT 240
      |||
Sbjct 312 TGTGCGTTCAAAGATTTCGATGATTCACTGAATTTGCAATTCACATTACTTATCGCATTT 253

Query 241 TGCTGCGTTCTTCATCAGTGCCAGAACCAAGAGATCCGTTGTTGAAAGTTTTGATTTATT 300
      |||
Sbjct 252 TGCTGCGTTCTTCATCAGTGCCAGAACCAAGAGATCCGTTGTTGAAAGTTTTGATTTATT 193

Query 301 TGTTTGTACTCAGAAGTTCCACTAAAAACAGAGTTTAGGGTCCTCGGGCGGGCCGTC 360
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Sbjct 192 TGTTTGTACTCAGAAGTTCCACTAAAAACAGAGTTTAGGGTCCTCGGGCGGGCCGTC 133

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Sbjct 132 CCTTTTTACAGGGCGCGGGCTGATCCGCCGAGGCAACGTATAGGTATGTTACAGGGGTT 73

Query 421 TGGGAGTTGTAAACTCGGTAATGATCCCTCCGCTGGTTACCAACAAGGACCTTGTTACG 480
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Sbjct 72 TGGGAGTTGTAAACTCGGTAATGATCCCTCCGCTGGTTACCAACAAGGACCTTGTTACG 13

Query 481 AC 482
      ||
Sbjct 12 AC 11
    
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