

Molecular Variations among some Isolates of *Fusarium oxysporum* f.sp. *lycopersici* and Response of Different Tomato Cultivars and Seedling Age to Infection

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Survey of tomato wilt was carried out through two successive growing seasons (2013 and 2014) in farms' located at 4 sites belonging to four Egyptian Governorates (Behera, Minofiya, Ismailia and Minia). The disease incidence in these sites ranged between 14.3 in Ismailia Governorate and 37.5% in Minofiya Governorate. Isolation on Potato Agar medium from naturally infected plants collected from some Egyptian Governorates resulted in obtaining many fungal isolates that proved to be pathogenic to tomato Super-Strain B cultivar on pathogenicity test. The agent responsible for wilt of tomato in Egypt was identified as *Fusarium oxysporum*. The pathogenic capabilities of eight fungal isolates representing different localities in different Governorates, *i.e.* Behera, Minofiya, Ismailia and Minia were investigated. Pathogenicity test using eight isolates of *Fusarium oxysporum* isolated from naturally infected tomato plants showing typical symptoms of wilt revealed that they were all able to cause wilt symptoms on different tomato cultivars. In this respect, isolate of *F. oxysporum* isolated from El-Khatatba, Minofiya Governorate was the most virulent that induced the disease in tomato plants. Ten plant species were evaluated for their reactions to infection with the most virulent isolate, El-Khattba (Minofiya Governorate). The RAPD-PCR of the DNA analysis using 10-mer primer of eight isolates of *F. oxysporum* f.sp. *lycopersici* collected from different Governorates showed variations in DNA pattern. To test their response to infection with this isolate, ten tomato cultivars were evaluated to infection by the most virulent isolate (isolated from El-Khattba, Minofiya Governorate) and the results indicated that cv. Super-Strain B was the highest susceptible, whereas cv. Super-Marmand was the lowest one. Moreover, effect of tomato seedlings age of Super-Strain B (highly susceptible) and Super-Marmand (low susceptible) on infection by the most virulent isolate of the pathogen was tested.

Keywords: *Fusarium oxysporum* f.sp. *lycopersici*, host range, RAPD-PCR analysis, seedling age and tomato cultivars.

Tomato (*lycopersicon esculentum* Mill) is one of the most worldwide grown vegetables (Amini, 2009 and Hamini-Kadar *et al.*, 2014). Unfortunately, the plant is subject to many diseases caused by fungi, bacteria, viruses and nematodes that cause severe losses in yield (Nusret and Steven, 2004; Amini, 2009 and BaharMorid *et al.*, 2012). *Fusarium* wilt of tomato caused by *F. oxysporum* f.sp. *lycopersici* (Sacc.)

Snyder & Hansen is one of the most prevalent, destructive diseases worldwide that affect yield and quality (Reis *et al.*, 2005). In Egypt, Awad (1990) reported that infection by *F. oxysporum* f.sp. *lycopersici* sometimes is the main reason for restriction of expanding tomato area and also increasing production costs. The crop is highly affected during the seasons of early spring and autumn in Governorates of Upper Egypt and the newly reclaimed sandy regions of Nubaria and Ismailia.

Identification of *Fusarium* species can be achieved by their morphological characteristics on selective media (Burgess *et al.*, 1994). However, the pathogenic types or formae specialis and races cannot be identified by morphological characteristics. Three known physiological races (1, 2 and 3) of *F. oxysporum* f.sp. *lycopersici* are distinguished by their specific pathogenicity on different known cultivars carrying dominant race specific resistant genes (Masunaga *et al.*, 1998 and Sammour *et al.*, 2013). Race 1 and 2 of *F. oxysporum* f.sp. *lycopersici* have been reported in most tomato growing regions of the world (Reis *et al.*, 2005). Recently many molecular techniques are used as practical and reliable methods for identification of fungi. Molecular markers are accurate and useful tools to identify formae specialis and races of *F. oxysporum* f.sp. *lycopersici* (Livens *et al.*, 2008). Pathogenicity of different isolated fungi was also performed. Some based on polymerase chain reaction (PCR), and includes random amplified polymorphic DNA, amplified fragment length polymorphism, simple sequence repeats and sequence related amplified polymorphism (Geiser *et al.*, 2004; Al-Khatib *et al.*, 2006; Hirano and Aris, 2006 and Jacobs *et al.*, 2013). In Japan differentiation of *F. oxysporum* f.sp. *lycopersici* and *F. oxysporum* f.sp. *radicis-lycopersic* has been achieved by polymerase chain reaction PCR-based method using specific primer sets developed due to knowledge of the partial nucleotide sequences of the endopolygalacturonases genes endo (pg1) and exopolygalacturonases genes of (pg×4) of fungi (Hirano and Aris, 2006).

Tomato cultivars and hybrids reacted variably to *Fusarium* infection race1, 2 and 3 (Al-Khatib *et al.*, 2006; Hirano and Aris, 2006 and Irzykowska *et al.*, 2012). Moreover, these authors added that race1 and 2 are distributed worldwide whereas, race 3 has limited geographical distribution (Hirano and Aris, 2006).

Infection with *Fusarium* wilt fungus is affected considerably with seedlings age and the ability of the plant to tolerate or even overcome infection (Gamal El-din *et al.*, 1982, Scott and Jones, 1989 and Al-Khatib *et al.*, 2006). The present investigation was conducted to isolate and identify the causative agent of *Fusarium* wilt of tomato. Plant species were evaluated for their susceptibility to infection by the most virulent isolate. Moreover, RAPD-PCR analysis on the isolated fungi was also achieved to show variation in DNA patterns. Finally, reaction of different varieties to infection by the isolated fungus and effect of plant age were studied.

Materials and Methods

1. Occurrence of tomato wilt in four Egyptian Governorates:

Survey of tomato wilt was carried out through early spring and autumn plantations during two successive seasons, *i.e.* 2013 and 2014 in four Egyptian

Governorates, *i.e.* Behera, Minofiya, Ismailia and Minia. Percentages of disease incidence were calculated from randomly chosen fields located at the previously mentioned Governorates. Disease incidence was calculated as the following formula: Disease incidence (%) = (No. of plants showing wilt symptoms/Total No. of inspected plants) x 100.

2. Isolation and identification of fungi associated wilted tomato plants:

Isolation trials were carried out from naturally infected tomato plants by wilt collected from locations belonging to 4 Egyptian Governorates, *i.e.* Behera, Minofiya, Ismailia and Minia. Roots and stems of diseased plants were thoroughly washed under running tap water to remove all adherent soil particles. Diseased materials were cut into small pieces (0.5 cm) and surface sterilized using 1% sodium hypochlorite for 2 minutes then washed several times in sterilized distilled water. The pieces were dried between folds of sterilized filter paper then transferred onto the surface of potato sucrose agar medium containing streptomycin (25µg/ml) in Petri-dishes. Plates were incubated at 25°C for 3 days. The emerged fungi were picked-up, purified using the hyphal tip and/or single spore techniques adopted by Dhingra and Sinclair (1995). The purified fungal isolates were identified according to the keys given by Gilman (1957) and Nelson *et al.* (1983) as *Fusarium oxysporum*.

3. Pathological studies:

3.A. Pathogenicity test using eight isolates of *Fusarium oxysporum* under greenhouse conditions:

Pathogenic ability of eight isolates of *F. oxysporum*, previously isolated and identified from naturally infected tomato plants, to induce wilt in tomato plants was evaluated under greenhouse conditions. The fungal isolates were grown separately on autoclaved sorghum grain sand medium (100g, washed dried sorghum grains, 100g, washed dried coarse sand and 65ml, tap water per bottle) in 500 ml glass bottles. Inoculation was carried out using uniform agar discs (5 mm. diam.) bearing 4 days old fungal growth of any of the tested isolates. The bottles were incubated at 28°C for two weeks to obtain sufficient growth of the fungal isolates.

Fertile soil was taken from the surface layer of soil of the experimental farm; Faculty of Agric. Minia Univ. and was sterilized using formalin solution (5%). Formalin disinfested clay pots (30-cm-diam.) were filled with sterilized soil at the rate of 3kg/pot. The potted soil was then artificially infested with the desired inoculum prepared at the rate of 3% (w/w), then watered two times during one week before planting. In check treatments, equal amounts of uninoculated substrate were added in pots (Gabr *et al.*, 1998). Tomato seedlings (cv. Super-Strain B) grown for 30 days in seed boxes filled with autoclaved peat-moss vermiculite (1:1 w/w) were uprooted and transplanted in the pots at the rate of 3 seedlings/pot. Four replicate pots were used for each fungal isolate. Pots were irrigated directly after transplanting and subsequently as when necessary.

Disease assessment:

At 30 days after transplanting the following assessments were calculated:

Disease index of foliar browning:

Disease severity of foliar yellowing was determined by rating each leaf on the severity of wilt symptoms and yellowing according to 0-4 scale and computing the average grade for plant as a whole according to the following formula: % of foliar yellowing = (Sum of foliar yellowing value / (4 x Total number of leaf) x 100 (El-Zawahry, 1984; Fakhouri and Buchenaure, 2003 and Song *et al.*, 2004). In the present study, the following numerical grades were used:

0= Healthy plants.

1=1-less than 25% of plant leaflets are yellow (slight chlorosis, wilting or stunting).

2=25-less than 50% of plant leaflets are yellow (moderate chlorosis, wilting or stunting).

3=50-less than 75% of plant leaflets are yellow (severe chlorosis, wilting or stunting).

4=75- less than 100% of plant leaflets are yellow (very severe chlorosis, complete wilting or dead plant).

Disease index of vascular browning:

Disease index of vascular browning was determined by estimating the internal discoloration (browning) area of vascular bundle by making longitudinal and transverse section of root according to the scale described by (Gothoskar *et al.*, 1953).

0= no brown discoloration in vascular bundles of root and the crown and stem.

1=1-less than 25% of vascular root bundles are brown.

2=25- less than 50% of vascular root bundles are brown.

3=50- less than 75% of vascular root bundles are brown.

4=75- less than 100% of vascular root bundles are brown.

The percentage of internal discoloration was calculated following the formula: % of vascular browning = (Sum of vascular browning values/ (4xTotal number of plants) x 100.

3.B. Host range of F. oxysporum using the most virulent isolate:

This experiment was carried out under greenhouse conditions using the highly pathogenic isolate of *F. oxysporum*, the most virulent (isolated from El-Khatatba, Minofiya Governorate) causing severe tomato wilt incidence for testing its ability to induce wilt on other plant species belonging to different families (Rowe, 1980). Ten plant species belonging to families, solanaceae, cucurbitaceae, leguminosae, chenopodiaceae and gramineae were tested in addition to plant species, *i.e.* pepper (*Capsicum frutescens* L.) Elpaso hybrid, eggplant (*Solanum melongena* var. *Esculentum* Nees) Hanen hyb, chillies (*Capsicum* sp.) Cayenne Lang Slim hyb., Jimsonweed (*Datura stramonium* var. *linne*), potato (*Solanum tuberosum*) cv. Kara, petunia (*Petunia hybrida*), cucumber (*Cucumis sativus* L.) Novo hyb, soybean (*Glycine max* L. Merr) cv. Giza 22, sugar beet (*Beta vulgaris* L.) cv. Gloria and wheat (*Triticum aestivum* L.) cv. Giza 164. Again some species of the following families were used to differentiate between *F. oxysporum* f.sp. *lycopersici* and *F. oxysporum* f.sp. *radicis-lycopersici* (Jarvis and Showoemaker, 1978; Rowe, 1980; Bahar-Morid *et al.*, 2012 and Boix-Ruiz *et al.*, 2015), cucurbitaceae (cucumber), leguminosae (soybean), chenopodiaceae (sugar beet) and gramineae (wheat).

Inoculum preparation, soil infested and assessment of disease incidence were carried out 60 days after sowing as mentioned before under pathogenicity test. Seeds and grains were surface sterilized as mentioned before and sown in artificially infested soil. Control treatments were used but without inocula, plants were observed daily for development of disease symptoms.

3.C. RAPD-PCR analysis of genetic variation among isolates of F. oxysporum f.sp. lycopersici isolated from naturally infected tomato plants:

This experiment was carried out in the laboratory of Agricultural Research Centre (ARC), Ministry of Agriculture. Isolates of *F. oxysporum* f.sp. *lycopersici*, the causal of Fusarium wilt of tomato were grown in 50 ml potato broth in conical flasks for 15 days at 28°C.

Fungal growth on duplex media:

Petri dishes were used instead of liquid shacked cultures to grow the different fungal isolates for DNA isolation. Disposable polystyrene Petri dishes (4 cm) contained 1800µL solid medium (potato dextrose agar), to which a layer of liquid medium (1400 µL peptone yeast glucose) was added. The fungal isolates were cultured by inoculating a small loop from stock onto the medium prepared in Petri dishes that were subsequently incubated at 28°C for 2-3 days. Mycelium was harvested from the medium using sterilized inoculating loops then transferred into sterile 1.5-mL microcenterfuge tubes. Fungal cells were pelleted by centrifugation for 15 min at 4000 rpm a deep well swing-bucket rotor (microcentrifuge5804 R; Eppendorf). The mycelium pellet was washed with 600 mL TE (Triesedita) buffers and centrifuged again for 5 min at 4000 rpm and finally; TE (Triesedita) buffer was decanted.

DNA extraction:

A modification of the traditional sodium dodecyl sulphate (SDS) extraction procedure was adopted. Fresh fungal pellets were homogenized in 400 µL sterile salt in homogenizing buffer (200 mM Trise-HCl, pH8.5, 250 mM NaCl, 25 mM EDTA, 0.5% SDS). Next, 6 µL 20 mg/mL RNase A were added and mixed thoroughly. The samples were incubated at 65°C for 10 min, after which 130 µL 3 M sodium acetate, pH 5.2, were added to each sample. Samples were vortexed for 30 second at maximum speed, when incubated at 20°C for 10 min. The lysate was centrifuged at 13,000 rpm at 4°C for 15 min, and the supernatant was transferred into fresh tubes. An equal volume of isopropanol was added to each sample, and mixed well, and samples were incubated at -20°C for 10 min. Samples were then centrifuged at 6000 rpm for 20 min at 4°C. The DNA pellets were washed twice using 700µL washing solution (100 and 70% ethanol, respectively). The DNA pellets were subsequently air dried in an oven at 40°C for at least 10 min. The resultant DNA pellet was then re-suspended in 100µL 1XTE (10 mM Trise-HCl, 1 mM EDTA) buffer, pH 8.0 (Abd-Elsalam *et al.*, 2007).

DNA quantification and gel documentation:

Seven microliters of the isolated DNA and 3µL of 10×loading dye were loaded in a lane of 1.5% (w/v) agarose gel containing 0.05µg/ml ethidium bromide, to check the quality of the DNA. For quantitative measurements, a charge-coupled

device camera imaging system and UVIssoft analysis (Gel Documentation and Analysis Systems, Uvitec, Cambridge, UK) were used to capture the image and to calculate the band intensities.

RAPD-PCR analysis:

Analysis was performed in 25- μ l reaction volumes containing PCR buffer (Promega, Mannheim, Germany), 0.2 RAPD-PCR, analysis was undertaken using 10-mer primer (MWG, Germany). RAPD mmol/l dNTPs, 0.5 mmol/l primer (5-d AATCGGGCTG-3), 4.0 mmol/l MgCl₂, 1.25 units of Taq Polymerase (Promega, Mannheim, Germany) and 10–20 ng genomic DNA. PCR reactions were carried out in a T-Gradient thermal cycler (Biometra, Germany) using the following profile: 94 °C for 1 min, 36°C for 1 min and 72°C for 1 min for 30 cycles, and a final extension at 72 °C for 5 min. Following amplification, the samples were separated by electrophoresis in 1.4 % agarose gel, stained with 0.5 μ g/ml of ethidium bromide and viewed under ultra-violet light. A 300-to 1500-bp ladder (Promega, Mannheim, Germany) was used as a molecular mass marker.

3.D. Evaluation of ten different tomato cultivars to infection by the most pathogenic isolate of F. oxysporum f.sp. lycopersici:

Seedlings of ten tomato cultivars namely; Super-Strain B, Prichard, Peto 95, Fayrose, Super-Magic, Yara, Super-Balady, Super-Shahd, Marmand and Super-Marmand were tested for their response to infection by the most pathogenic isolate of *F. oxysporum* f.sp. *lycopersici* under greenhouse conditions. Healthy, thirty days old seedlings of the aforementioned tomato cultivars were transplanted in pots 30 cm containing soil already infested with the pathogenic isolate as previously mentioned. Four seedlings were sown in each pot and four replicate pots were used for each cultivar. Transplants were irrigated directly and subsequently when necessary and as recommended. Seedlings were observed daily and disease severity was recorded after 30 days of transplanting as previously described under pathogenicity test.

3.E. Effect of tomato seedling age on the infection by the most pathogenic isolate of F. oxysporum f.sp. lycopersici:

This experiment was carried out under greenhouse conditions to study the susceptibility of tomato seedlings of different ages to infection by the most pathogenic isolate of *F. oxysporum* f.sp. *lycopersici*. In this respect, susceptible tomato cultivar (Super-Strain B) and the resistant one (Super-Marmand) were used. Four tomato seedlings aging, 20, 25, 30 and 40 days old, representing any tested cultivar were evaluated to infection by the pathogenic isolate. Seedlings of four ages of the two aforementioned tomato cultivars were transplanted in sterilized pots (30-cm diameter) previously inoculated as mentioned under pathogenicity test. All agricultural practices and procedures were carried out as recommended. Results were taken after 30 days of transplanting as mentioned before.

Statistical analysis:

Data were subjected to statistical analysis of variance. The experimental design (S) of all studies was a completely randomized with three replications, analysis of variance (ANOVA) of the data was performed with the MSTAT-C statistical package (A) micro-computer program for the design, management, and analysis of

agronomic research experiments. Michigan State Univ., USA. Least significant difference (LSD) was used to compare treatment means (Gomez and Gomez, 1984).

Results

1. Occurrence of tomato wilt disease in four Egyptian Governorates:

Field observations of randomly chosen tomato fields located at four Egyptian Governorates showed that naturally infected plants were suffered from leaf chlorosis, epinasty, drooping of leaves and the whole plants exhibited different degrees of wilt symptoms. Moreover, when the diseased plants were split longitudinally they showed different degrees of browning in roots and stem base. In most surveyed fields, diseased plants developed chlorosis, epinasty and clear wilt symptoms. The natural infection percentages varied in different surveyed Governorates as well as in the two successive sampling years. Generally, percentage of infection ranged from 14.3-37.5% on the average (Table, 1). However, infection of tomato plants in fields of Minofiya Governorate was higher than any of the other inspected Governorates through the two successive seasons, being 37.5% on the average. However, the lowest average percentage of infection by the disease was recorded from Ismailia Governorate (14.3%).

Table 1. Occurrence of tomato Fusarium wilt at the different surveyed Governorates during seasons 2013 and 2014

Governorate	Disease incidence (%)		Mean
	Season 2013	Season 2014	
Behera	17.9	18.1	18.0
Minofiya	39.3	35.7	37.5
Ismailia	14.3	14.3	14.3
Minia	25.0	21.4	23.2
Mean	24.1	22.4	-
L.S.D at 0.05%	13.33	10.93	

2. Isolation and identification of fungi associated with wilted tomato plants:

Isolation from diseased tomato plants showing wilt symptoms showed that *Fusarium* sp. found to be associated with wilted tomato plants under investigation. Preliminary identification of the obtained isolates based on cultural, morphological characteristics and microscopic examinations according to Gilman (1957); Barnet and Hunter (1972); Booth, 1985 and Burgess *et al.* (1994) revealed that the isolated fungal isolates were *F. oxysporum*. Subsequently identification was confirmed by Assuit University Mycological Institute (AUMI). Fungal isolates isolated on PDA grew at 25°C. The culture changed in colour from white peach, salmon and purple to violet. The mycelium was striate, felted to floccose and sometimes wrinkled in older cultures (three weeks). The microconidia are oval-ellipsoid, cylindrical, straight or curved, from 5 to 12 µm by 2.2 to 3.5 µm and are produced from short simple phialides arising laterally on the hyphae or from short sparsely branched conidiophores. The macroconidia are produced sparsely and are borne on elaborately branched conidiophores. They are generally thin-walled and pointed at both ends; three-septate spore ranging in size from 27 to 46 µm by 3 to 4.5 µm.

Chlamydo spores are most commonly found, generally found and formed singly or in pairs intercalary or terminally.

Data in Table (2) indicate that fungi belonging to four genera were isolated from farms located at four Egyptian Governorates, Behera, Minofiya, Ismailia and Minia. *Fusarium oxysporum* was the most prevalent as constituted 26.7% in Minofiya, followed by *Alternaria alternata*. *Aspergillus niger* and *Trichoderma harzianum* that were also isolated but at lower frequencies, respectively. These results are in harmony with those reported from Egypt and other parts of the world (Kordali and Demirci 1998; Pushpa *et al.*, 1999; Yan *et al.*, 2004 and Mohamed, 2007).

Table 2. Occurrence and frequency (%) of fungi isolated from tomato plants collected from four different Governorates

Governorate	Isolated fungus	Frequency (%) [*]
Behera	<i>Fusarium oxysporum</i>	20.0
	<i>Alternaria alternata</i>	15.0
	<i>Aspergillus niger</i>	3.3
	<i>Trichoderma harzianum</i>	1.7
Minofiya	<i>Fusarium oxysporum</i>	26.7
	<i>Alternaria alternata</i>	20.0
	<i>Aspergillus niger</i>	20.0
	<i>Trichoderma harzianum</i>	8.3
Ismailia	<i>Fusarium oxysporum</i>	21.7
	<i>Alternaria alternata</i>	16.7
	<i>Aspergillus niger</i>	6.7
	<i>Trichoderma harzianum</i>	3.3
Minia	<i>Fusarium oxysporum</i>	23.3
	<i>Alternaria alternata</i>	18.3
	<i>Aspergillus niger</i>	10.0
	<i>Trichoderma harzianum</i>	5.0
Total		220.0

* Frequency (%) = Number of the isolated isolates x 100/Total number of the isolated fungi

3. Pathological studies:

3.a. Pathogenicity test using the eight isolates of *F. oxysporum*:

In this experiment eight fungal isolates of *F. oxysporum* isolated from different localities belonging to four different Governorates were tested for their ability to infect tomato plants. Data of the pathogenicity test (Table, 3) indicate that all isolates of *F. oxysporum* isolated from naturally wilted tomato plants produced the symptoms of Fusarium wilt expressed as leaves epinasty associated with foliar yellowing and browning of the vascular bundle on cv. Super-Strain B tomato cultivar. Isolate of *F. oxysporum* isolated from El-khatatba (Minofiya Governorate) produced the highest percentage of foliar yellowing and wilt (57.2%) which was significantly higher than any of the tested isolates followed by isolate of Behdal (44.1%) and isolate of Derwa, (27.2%). On the other hand, infection with the remainder of the isolates from different Governorates ranged from moderately to weakly. Regarding the vascular browning produced by the tested fungal isolates, the

same trend was obtained with that of foliar yellowing and wilts (Table, 3). Moreover, reisolation from artificially inoculated plants resulted in obtaining the originally isolated fungi proving Koch's postulates.

Table 3. Pathogenicity test of *F. oxysporum* isolates on tomato Super-Strain B cultivar under greenhouse conditions

Fungal source	locality	Disease severity (%)	
		Foliar yellowing and wilt (%)	Vascular browning (%)
Behera	Housh-essa	22.7	32.1
Minofiya	Sheben El-koam	15.5	31.3
	Berket El-sabaa	10.2	17.1
	El-khatatba	57.2	66.3
Ismailia	El-kassasen	19.9	29.0
Minia	Derwa	27.2	35.2
	Mallawy	24.6	36.5
	Behdal	44.1	49.6
Control		0.0	0.0
Mean		24.6	33.0
L.S.D at 0.05%		14.11	19.35

3.b. Host range of *F. oxysporum*:

Data in Table (4) show that the tested fungus infected some plant species belonging to the family solanaceae with varying degrees when these plant species were sown in artificially infested soil with the fungal inocula. Artificially infected species included pepper (*Capsicum frutescens* L.), eggplant (*Solanum melongena* var. *Esculentum* Nees), chillies (*Capsicum* sp), Jimsonweed (*Datura stramonium* var. *linne*), potato (*Solanum tuberosum*) and petunia (*Petunia hybrida*). Some plant species responded positively and produced wilt symptoms ranged from 7.0 to 40.9%. High disease severity was recorded in pepper (40.9%) followed by eggplant, potato and jimsonweed. Low disease severity was recorded on chillies and petunia, being 24.0 and 7.0%, respectively. On the other hand, other plant species belonging to cucurbitaceae, i.e. cucumber (*Cucumis sativus* L.), leguminosae, i.e. soybean (*Glycine max* L.Merr), chenopodiaceae, i.e. sugar beet (*Beta vulgaris* L.) and Graminae, i.e. wheat (*Triticum aestivum* L.) were not infected. Disease symptoms recorded on solanaceae species were yellowing, epinasty, drooping, losing turgidity of the leaves. Discoloration of vascular bundle was also observed on the previously mentioned hosts. The highest vascular browning percentage (58.0%) was also recorded from pepper plants followed by eggplant, being 49.3%. No crown rot and root rot symptoms were recorded suggesting the isolate under investigation belongs to *F. oxysporum* f.sp. *lycopersici*.

3.c. RAPD-PCR analysis of genetic variation among isolates of *F. oxysporum* f.sp. *lycopersici* isolated from naturally infected tomato plants:

The obtained results (Figs 1 and 2) obviously show dendrogram derived from RAPD-PCR. It is clear that there was similarity among DNA of the pathogenic isolates of *F. oxysporum* f.sp. *lycopersici* isolated from different localities. This similarity level was 85.43 among the tested isolates. However, dendrogram indicated

that these isolates can be divided into 5 main groups, *i.e.* group1 that had similarity level accounted 98.70% and included two isolates from El-Khatatba No.1 and Berket El-sabaa No.3, (Minofiya Governorate). The second group had similarity

Table 4. Host range of *F. oxysporum* El-khatatba isolate

Tested plant species		Disease severity (%)	
English name	Scientific name	Foliar yellowing and wilt (%)	Vascular browning (%)
Pepper	<i>Capsicum frutescens</i> L	40.9	58.0
Eggplant	<i>Solanum melongena</i> var. <i>Esculentum</i> Nees	34.4	49.3
Potato	<i>Solanum tuberosum</i>	30.1	35.8
Chillies	<i>Capsicum</i> sp	24.0	28.5
Datura	<i>Datura stramonium</i> var. <i>linne</i>	25.7	29.5
Petunia	<i>Petunia hybrida</i>	7.0	6.1
Cucumber	<i>Cucumis sativus</i> L.	0.0	0.0
Soybean	<i>Glycine max</i> L.Merr	0.0	0.0
sugar beet	<i>Beta vulgaris</i> L.	0.0	0.0
Wheat	<i>Triticum aestivum</i> L.	0.0	0.0
Control		0.00	0.00
L.S.D at 0.05%		7.55	13.31

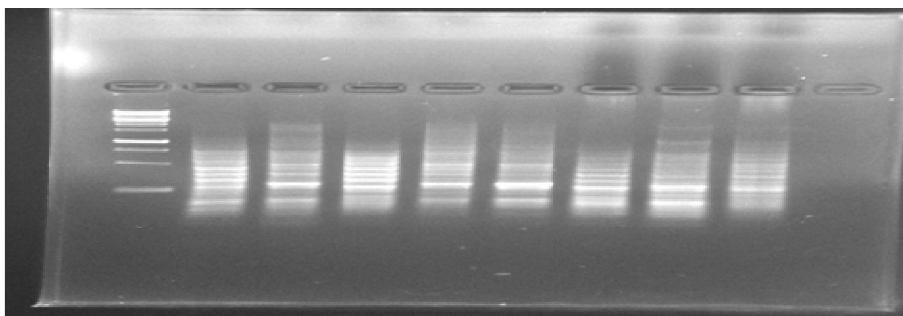


Fig. 1. RAPD analysis of DNA performed on eight isolates of *F. oxysporum* f.sp. *lycopersici*.

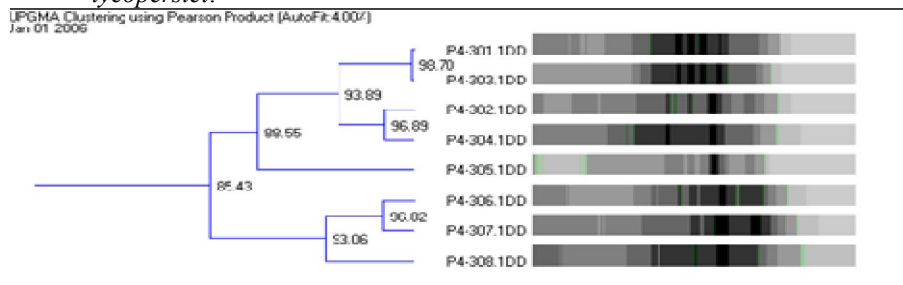


Fig. 2. Dendrogram derived from RAPD (Random amplified polymorphic DNA) profile analysis of eight isolates of *F. oxysporum* f.sp. *lycopersici*.

level 96.89% and included isolates No.2 from Sheben El-koam, (Minofiya Governorate) and isolate No.4 from Derwa, (Minia Governorate). Group 3 included isolate No.5 isolated from Mallawy, (Minia Governorate) had similarity level 88.55% with the two groups mentioned before. Group 4 included isolate No.6 isolated from Behdal, (Minia Governorate) and isolate No.7 isolated from El-kasaseen, (Ismailia Governorate) with similarity level 96.82%. Group 5 which included isolate No.8 isolated from Housh-essa, (Behera Governorate) with similarity level 93.06% with the isolates belonged to group 4.

3.d. Evaluation of ten different tomato cultivars to infection by F. oxysporum f.sp. lycopersici isolated from El-Khatatba, (Minofiya Governorate) under greenhouse conditions:

Data presented in Table (5) clearly indicate that all tomato cultivars tested were liable to infection by the highly pathogenic isolate of the causal pathogen with different degrees. The highest average percentages of foliar yellowing and wilt (43.5%) and vascular browning (55.9%) were occurred in Super-Strain B cultivar. Meanwhile, Super-Marmand recorded the lowest average percentage in this respect, being 4.1% of foliar yellowing and wilt and 3.8% of vascular browning, respectively. Results (Table, 5) also indicate that it is possible to classify tomato cultivars tested into three different categories according to their response to infection by the agent tested. The first category included the relatively resistant cultivars, *i.e.* cvs. Super-Marmand, Marmand, Super-Shahd and Super-Balady. The second category included the relatively moderately resistant cultivars, *i.e.* cvs. Yara and Super-Magic. Meanwhile, the third category included the highly susceptible cultivars to infection (cvs. Super-Strain B, Prichard, Peto 95 and Fayrose). It is worthy to mention here that the disease symptoms were developed earlier on the highly susceptible cultivars than on the moderately or relatively resistant cultivars.

Table 5. Evaluation of ten different tomato cultivars to infection by *F. oxysporum* f.sp. *lycopersici* isolated from El-khatatba, (Minofiya Governorate) under greenhouse conditions

Cultivar	Disease severity (%)	
	Foliar yellowing and wilt (%)	Vascular browning (%)
Super-Strain B	43.5	55.9
Prichard	38.1	44.5
Super-Magic	35.4	45.5
Peto 95	31.5	42.0
Yara	24.7	39.2
Fayrose	19.8	36.8
Super-Balady	18.5	29.5
Super-Shahd	11.2	22.2
Marmand	7.5	17.0
Super Marmand	4.1	3.8
Control	0.0	0.0
Mean	21.3	30.6
L.S.D at 0.05%	9.98	8.34

3.e. Response of tomato seedlings of different ages to infection with the most virulent isolate of *F. oxysporum* f.sp. *lycopersici*:

This experiment was designed to study the effect of tomato seedling age on infection using the most virulent isolate of *F. oxysporum* f.sp. *lycopersici* on the susceptible cultivar (Super-Strain B) and the resistant one (Super-Marmand). Results in Table (6) indicate that, generally, plants grown from younger tomato seedlings were more susceptible to infection with the fungus than those developed from the older ones. Young seedlings (20 days old) were the most susceptible seedlings, followed by those of 25 and 30 days old, while 40 days old seedlings were the lowest susceptible ones. This was also true for the susceptible cultivar Super-Strain B. Similar trend was obtained with the relatively resistant cultivar Super-Marmand. Again, plants responded more positively by increasing age indicating that resistant was increased progressively by increasing age. Data also showed that there were significant differences among all tested plant ages on their reaction to the two tested cultivars, Super-Strain B and Super-Marmand. Additionally, the resistant cv. Super-Marmand responded more positively regarding its resistance to infection with the tested fungus.

Table 6. Response of tomato seedlings of different ages to infection with the most virulent isolate of *F. oxysporum* f.sp. *lycopersici* (El-khatatba) on the susceptible cultivar Super-Strain B and the relatively resistant one Super- Marmand

Tomato age (Days)	Disease severity (%)			
	Foliar yellowing and wilt (%)		Vascular browning (%)	
	Susceptible (Super-Strain B)	Resistant (Super-Marmand)	Susceptible (Super-Strain B)	Resistant (Super-Marmand)
20 days	49.0	16.2	68.2	31.6
25 days	38.1	6.0	52.3	15.3
30 days	26.3	3.1	36.3	11.5
40 days	16.9	0.8	15.4	4.5
Control	0.0	0.0	0.0	0.0
Mean	26.1	5.2	34.4	12.6
L.S.D at 0.05%	12.30	6.99	18.66	7.48

Discussion

Over the last few years, tomato (*lycopersicon esculentum* Mill) wilt disease has become common and inflicted marked losses in yield in many Egyptian Governorates according to our field observations and complains of the vegetable growers. Natural infection recorded ranged from 14.3 to 37.5%. These results agree with those reported before in Egypt and other parts of the world (Awad, 1990, Mariatt *et al.*, 1996, Enespa and Dwivedi, 2014 and El-Mohamedy *et al.*, 2014). Fusarium wilt caused by race of *Fusarium oxysporum* f.sp. *lycopersici* was sever in Arkansas and threatened to eliminate commercial tomato production in the state (Goode, 1966). During the course of this investigation, eight fungal isolates were

isolated from the roots and stems of naturally infected wilted plants obtained from some Egyptian Governorates. The isolated fungi were identified and the most frequent was *F. oxysporum* according to the morphological, cultural characteristics and microscopic examination (Gilman, 1957, Booth, 1971, Barnett and Hunter, 1972, Barnett and Hunter, 1997, Palmer *et al.*, 2010, Salehzadeh, 2012 and Mc Govern, 2015). In pathogenicity tests, the isolated eight fungal isolates caused wilt in some tomato cultivars similar to that obtained under, naturally infected conditions. Generally, all fungal isolates had the potentiality to infect tomato plants although they were varied in their pathogenicity from weakly to highly pathogenic.

Fusarium oxysporum f.sp. *lycopersici* has previously been reported to be main cause of tomato wilt in Egypt (El-Zawahry, 1984, El-Shami, 1987, Farag, 2011 and Sammour *et al.*, 2013 and Radwan *et al.* 2016) and other parts of the world (Rodriguez *et al.*, 2003, Song *et al.*, 2004, Amini, 2009, Nijue *et al.*, 2012, Jacobs *et al.*, 2013, Enespa and Dwivedi, 2014 and Hamini-Kadar *et al.*, 2014). Pathogenicity tests are the primary means to distinguish different pathogenic *Fusarium* strains, but they do not indicate whether isolates of a given physiologic race or forma specialis are genetically related (Katan *et al.*, 1989). Some strains of *F. oxysporum* are responsible for vascular wilt disease of many plants of economical important and these strains show high level of host specificity and are classified on this basis into formae specialis and races (Hirano and Aris, 2006 and El-Kazzaz *et al.* 2008). Several methods have been employed in the study of the genetics of specialized forms and races of *F. oxysporum* due to techniques based on the analysis of nucleic acids including random amplified polymorphic DNA (RAPD) (Boix-Ruiz, *et al.* 2015). However, host range studies revealed that the fungal isolates are specific in infection on solanaceae species and it did not infect species from other plant families' *i.e.* Cucurbitaceae, leguminosae, Chenopodiaceae and Graminae. Furthermore, symptoms produced by these isolates were typically of vascular wilt, caused by *F. oxysporum* since they produced symptoms spread up in the vascular bundles rapidly. No crown and root rot symptoms were obtained due to artificial inoculation of the aforementioned species and families. Therefore, these findings suggest that the fungal isolates under investigation are *F. oxysporum* f.sp. *lycopersici* (Jarvis and Shoemaker, 1978, Menzies *et al.*, 1990 and Boix-Ruiz *et al.*, 2015). Traditional criteria used to differentiate species of the fungus are based on plant host, colony appearance, morphological characteristics of the conidia and teleomorph (Gordon and Martyn, 1997). Subsequently Woo *et al.*, (1996) and Gordon and Martyn (1997) reported that *F. oxysporum* has about 80 formae specialis (*i.e.* pathotypes specific to species) and several subdivided into races (specific to cultivars within a species). Furthermore, classification should take into account some consistent characters added and described by Burgess *et al.*, (1994) such as morphological characteristics, *i.e.* shape and mode of formation microconidia, presence or absence of chlamydospores, colony morphology and growth rate on PDA. Furthermore proper identifications of formae specialis and races have significant application in the diagnoses of disease in addition to its importance in quarantine measures for plant materials (El-kazzaz *et al.* 2008, Gao and Zhang, 2013 and Mc Govern, 2015).

In general, the obtained results from RAPD-PCR revealed that basically there were DNA variations among isolates of *F. oxysporum* f.sp. *lycopersici* isolated during this investigation. Regarding similarity, level, there is similarity among DNA of the pathogenic isolates isolated from different localities in the tested Governorates and affecting tomato plants. This similarity level was 85.43% among the tested isolates. However, dendrogram indicated that isolates might be divided into 5 main groups, *i.e.*, group 1 that has similarity level accounted for 98.70 and includes two isolates from El-khatatba and Berket El-sabaa, (Minofiya Governorate). The second group has similarity level 96.89 and includes 2 isolates from Sheben El-koam, (Minofiya Governorate) and isolate 4 from Derwa, (Minia Governorate). It is not surprising that isolate 1 and 3 have similarity level of 98.7, simply because they developed and revolutionized under the same environmental conditions and they are from the same native geographic origin but it is surprising that isolate two from Sheben El-koam, (Minofiya Governorate) and isolate four from Derwa, (Minia Governorate) had high similarity level of 96.89% although they were isolated from two localities belonging to two Governorates that remotely far from each other. This might be due to trade of seeds and movement of the pathogen between the two Governorates through seeds (El-Wakil *et al.*, 1998, Reis *et al.*, 2005, Sammour *et al.*, 2013 and Mc govern, 2015). Group 3 included isolate No. 5 isolated from Mallawy (Minia Governorate) has similarity level of 88.55%. This might be due to geographic origin of the isolates. Group 4 included isolate No. 6 isolated from Behdal (Minia Governorate) and isolate No. 7 isolated from El-kasaseen, (Ismailia Governorate) had similarity level 96.82% and this can be interpreted as above. Group 5 which included isolate No. 8 isolated from Housh-essa, (Beheira Governorate) had similarity level 93.06% recorded. Surprisingly, isolate 6 from Behdal and isolate 4 from Derwa, (Minia Governorate) and from Kasaseen, (Ismailia Governorate) have high similarity levels (96.89%) with isolate 1 and 2 of El-khatatba and Berket El-sabaa, (Minofiya Governorate, 98.70%) this may be due to trade of seeds and movement of pathogen through seeds.

In conclusion, the results obtained by RAPD-PCR showed variations in DNA levels (85.43) in the tested isolates of the pathogen and the possibility to separate the isolates into 5 groups. Grouping the isolates by RAPD-PCR was not related to virulence or geographic origin. However RAPD-PCR relatively was able to relate isolate 2 from El-Khatatba and isolate 3 from Berket El-sabaa, (Minofiya Governorate) to their origin to some extent. Such results obtained with RAPD-PCR are in line with results reported herein (Novack and Kohn, 1988, Saeed and Abo-Elseoud, 1990, Lima and Menzes, 2002, Sallam, 2004 and Mohamed, 2010) who worked on other different plant pathogenic fungi. Several methods have been employed in the study of the genetics of specialized forms and races of *F. oxysporum* due to techniques based on the analysis of nucleic acids including random amplified polymorphic DNA (RAPD) (Boix-Ruiz *et al.*, 2015). The authors added and concluded that RAPD marker had only limited usefulness in correlating pathogenicity among the isolates and races. In contrary, Riveros *et al.* (2001) compared RAPD-PCR with classical, morphological and pathogenicity of Fusarium and found that the obtained results were inconsistent.

In this study the tested cultivars under investigation showed variable responses to infection by the most virulent fungal isolate El-Khatatba (Minofiya Governorate) of *F. oxysporum* f.sp. *lycopersici*. Some proved highly susceptible such as Super-Strain B, Prichard, Peto 95 and Super-Magic; others were less susceptible to infection, such as Yara and Fayrose while, the remainders proved relatively resistant such as Super-Balady, Marmand, Super-Shahd and Super-Marmand. Results obtained in this investigation are in agreement with those found by many investigators in Egypt and other parts of the world (El-Zawahry, 1984; Moustafa and Khafagi, 1992; Moustafa, 1999; Rodriguez *et al.*, 2003; Reis *et al.*, 2005; Al-Khatib *et al.* (2006); Abdalla, 2007; Yousef, 2007; El-Kazzaz *et al.*, 2008; Kapoor, 2008; Amini, 2009; Dordevic *et al.*, 2012; Steinkellner *et al.*, 2012 and Radwan *et al.*, 2016). Moreover, the following Egyptian varieties and hybrids, *i.e.* Super red, Hybrid Zomoreda TH99806, V.F.N-8, Nema rock, Nema 1400, the local Hybrid Master 100 and Improved Saria were found relatively resistant to root-Knot nematode and hence may decrease infections with Fusarium wilt caused by *F. oxysporum* f.sp. *lycopersici*. Root-Knot nematode can induce susceptibility in normally resistant cultivars to Fusarium wilt fungi on host plant (Mai and Abawi, 1987 and France and Abawi, 1994). Subsequently, it has been reported that localization of the vascular infection is widely recognized as a primary resistant mechanism to Fusarium wilt in tomato plants and the distribution of the fungus is limited in resistant cultivars while extensive colonization occurs in susceptible ones (Rodriguez *et al.*, 2003). Subsequently, it has been reported that most of the common varieties of tomato and bringal in heavily infested fields with Fusarium are susceptible and fungicides should be frequently used to control the disease (Kapoor, 2008).

Generally, percentage of disease index of Fusarium wilt was decreased progressively with increase of seedlings age and the older seedlings were more relatively less susceptible to infection than younger ones by the isolate tested (isolate El-khatatba). These results are partially in agreement with those reported by Al-Khatib *et al.* (2006) and Amini (2009). It is reported that although has been an extremely useful characteristic virulence of *F. oxysporum* f.sp. *lycopersici* of differentiating between isolates of the fungus, it is still single traits since virulence has been shown to be influenced by number of factors including temperature, host range and method of inoculation (Kraft and Haglund, 1978; Williams, 1981; Hart and Endo, 1981; El-Kazzaz *et al.*, 2008 and Steinkellner *et al.*, 2012). On the other hand Steinkellner *et al.* (2012) found that infection by *F. oxysporum* f.sp. *lycopersici* on tomato varieties was not affected with plant age.

A c k n o w l e d g m e n t

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الأختلافات الجزيئية بين عزلات الفطر
Fusarium oxysporum f.sp. *lycopersici*

وحساسية أصناف الطماطم واعمار الشتلات للاصابة بالفطر

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تم عمل حصر لمرض ذبول الطماطم خلال عامي 2013 و 2014 م في حقول تابعة لمواقع منتمية لأربع محافظات مصرية هي البحيرة، المنوفية، الاسماعيلية والمنيا حيث تراوحت نسبة الاصابة بهذا المرض في هذه المواقع ما بين 14.3 و 37.5%. وقد أوضح اختبار المرضية بأستخدام 8 عزلات من فطر *F. oxysporum* سبق عزلها وتعريفها من نباتات طماطم كانت مظهرة لأعراض الذبول أن عزلة الخطاطبة المعزولة من نباتات الطماطم المنزرعة في الخطاطبة (محافظة المنوفية) كانت أشد العزلات المختبرة. كما قيمت عشرة من الأنواع النباتية والتي تتبع خمس فصائل مختلفة تحت ظروف الصوبة لمعرفة استجابتها للعدوى بأقوى العزلات قدرة مرضية المعزولة من نباتات الطماطم المنزرعة في الخطاطبة (محافظة المنوفية) وظهرت النتائج أن الفلفل كان أكثر العوائل اصابة ببتبعة الباذنجان ثم الشطة، الداتورا، البطاطس والبيبتونيا بينما الأنواع النباتية الاخرى (الخيار، فول الصويا، بنجر السكر والقمح) لم يظهر عليها أعراض الاصابة. وقد اشار تحليل RAPD-PCR وجود اختلافات جزيئية واضحة بين عزلات الفطر الممرض المعزولة من المحافظات الأربع في ال DNA أيضا قيمت عشرة من أصناف الطماطم المنزرعة تحت ظروف الصوبة لمعرفة مدى استجابتها للعدوى بأقوى العزلات قدرة مرضية وظهرت النتائج أن الصنف سوبر سترين ب كان أكثر الاصناف اصابة في حين كان الصنف سوبر مارمند أقلها اصابة. كذلك تم اختبار تأثير عمر الشتلات من صنفى سوبر سترين ب (الاكثر قابلية للاصابة) وسوبر مارمند (الاقل قابلية للاصابة) على القابلية للاصابة بأقوى عزلة من المسبب المرضي وتبين من خلال النتائج ان شتلات الصنف سوبر سترين ب عمر 20 يوم كانت هي الاكثر اصابة بالذبول مقارنة بباقي الاعمار.