IDENTIFICATION AND PATHOGENICITY OF FUSARIUM SPECIES ASSOCIATED WITH HEAD BLIGHT OF WHEAT IN IRAN

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Abstract

In order to determine Fusarium species associated with wheat heads, 95 spike samples were collected from different geographic regions and wheat growing areas in the southwest, west, northwest and north of Iran during 2006-2008. Samples were collected from plants showing head scab symptoms. A total of 280 Fusarium strains were isolated and identified. All strains belonged to 19 Fusarium species according to morphological characters. All Fusarium strains were evaluated to test their pathogenicity on wheat which stands that F. graminearum, F. culmorum, F. crookwellense, F. trichothecioides and F. chlamydosporum were the most pathogenic to wheat's head. This report is the first in-depth study to identify of Fusarium spp. from wheat in Iran.

Introduction

Wheat is one of the basic agriculture crops which provide the food requirement in Iran (Farshadfar et al., 2008). Fusarium species are responsible for many economical important plant diseases such as head blight on cereals (Niaz & Dawar, 2009). Fusarium head blight (FHB), known as scab, is a disease of different cereals can leads to the decrease quality and quantity of wheat's grains (Bottalico & Perrone, 2002). In the field, FHB is recognized by the premature bleaching of infected spikelets and during wet weather there is whitish or pinkish fluffy fungal mat on the infected heads. Several Fusarium spp. cause scab in wheat head, including F. graminearum, F. crookwellense, F. culmorum, F. equiseti, F. chlamydosporum, F. avenaceum, F. semitectum, F. verticillioides and F. camptoceras (McMullen et al., 1997). Fusarium spp. can also produce mycotoxins which have impact on animal and human health (Gamanya & Sibanda, 2001). Among these mycotoxins, zearalenone, fumonisins and trichothecenes potentially could occur in a variety of foodstuffs and feedstuffs, which are responsible for variety of toxic effects including infertility and reproductive problems, mutagenic and carcinogenic (Nelson et al., 1991; Thiel et al., 1991; Shephard et al., 2005). Wheat is one of the most significant sources of mycotoxin contamination in human and animal diets. The incidence of mycotoxin in maize has been investigated from main production areas and a region of high cancer prevalence in Iran (Ghiasian et al., 2006). So it was required to identification of Fusarium spp. from other crops such as wheat.

Materials and methods

Collection of glume samples and isolation of Fusarium spp.: Ninety five spike samples showing disease symptoms were collected from various fields at spike stage, from wheat growing areas in the southwest, west, northwest and north of Iran during 2006-2008 (Table 1). The glumes were surface sterilized with 0.05% Sodium hypochlorite solution for 3 min, rinsed twice with sterilized distilled water and placed on Pepton-Pentachloronitrobenzene agar (PPA) plates that was proposed by Nash and Snyder (1962) and water agar (WA). All the plates were incubated under specify

standard incubation conditions (Salleh & Sulaiman, 1984) for 48 h. The resulting single-spored *Fusarium* colonies were transferred onto potato dextrose agar (PDA) plates for further studies.

Identification of Fusarium spp. through morphological characterization: To study the growth rates and pigment production of Fusarium spp. all the strains were transferred onto PDA plates and incubated at ambient temperature. Ten replications were maintained for each Fusarium strain. For microscopic observations, all the strains of Fusarium were transferred to carnation leaf agar (CLA) (Fisher et al., 1982) and potassium chloride agar (KClA) (Fisher et al., 1983) mediums. The species were identified on the basis of macroscopic characteristics such as pigment production and growth rates on PDA plates, as well as their microscopic features including size of macroconidia, presence of microconidia and its production in chains or false heads, type of conidiogenous cells (monophialidies and polyplialidies conidiophores) and also absence or presence of chlamydospores (Gerlach & Nirenberg, 1982). Identification of species was based on species description of Gerlach & Nirenberg (1982) and Leslie & Summerell (2006).

Pathogenicity Test: All of the identified Fusarium species were tested for their pathogenicity on healthy wheat seedlings variety Zarrin. The healthy wheat's were inoculated with each isolate 10-14 days after heading. Prior to inoculation, 10 spikes per pot were randomly selected. For inoculation, all of the isolates were grown on PDA plates and conidial suspension of each individual strain was prepared by scrapping the mycelium with sterile distilled water onto 7 day-old cultures, shaken thoroughly, and the concentration was adjusted to 1×10⁶ conidia mL⁻¹ using a haemocytometer. Plants were inoculated at anthesis stage by injection of 5 uL of macroconidial suspension in each spiklets using point-inoculation method by hypodermic syringe (Dubin et al.,

After 30 min of the inoculation, the experiments were carried out in the polyethylene humidity chamber conditions maintained at 25°C, 100% RH for 48 h. After incubation, plants were transferred to the greenhouse. The negative control plants were injected with 5 uL sterile distilled water per spike and for positive control; plants

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were injected with 5 uL of aggressive *F. graminearum* isolate. In the greenhouse, symptoms of FHB were observed at 28 days after inoculation and rated for disease severity according to the method of Xue *et al.* (2004). Also percentage of infected spikelet's (IS) was estimated at 4 weeks after inoculation, when plants were at the soft dough stage. Disease severity was estimated visually *In*

Situ for each inoculated spike on a 0 to 9 (no visible FHB symptoms to severely diseased, spike dead) scale described by Xue *et al.* (2004). Disease severities and percentages of IS from all plants in each pot were averaged and the means per pot of percent IS were used in the statistical analysis.

Table 1. Fusarium spp., identified on wheat samples

Place of sample	Total No. of spike	No. of spike samples infected	•
collection	samples	with Fusarium spp.	Fusarium spp. identified
Ahvaz	5	5	F.ve, F.se, F.pr, F.ny, F.an, F.sc, F.eq, F.gr, F.ve,
			F.ch
Ahar	3	3	F.eq, F.ox, F.pr, F.an, F.sc, F.ve, F.ny, F.gr, F.ch,
			F.se
Andimeshk	4	4	F.ch, F.an, F.sc, F.pr, F.ny, F.gr, F.ve, F.so
Amol	3	3	F.gr, F.pr, F.ve, F.ny
Babol	2	2	F.gr, F.ve
Bahar	3	3	F.gr, F.an, F.so, F.ve, F.pr, F.ny, F.eq, F.cr
Babolsar	2	2	F.gr, F.pr, F.ve, F.ny
Behshahr	2	2	F.gr, F.pr
Bijar	3	3	F.eq, F.so, F.ny, F.se, F.ox, F.av, F.gr, F.ve, F.cr
Bisotun	5	5	F.sc, F.ve, F.pr, F.ny, F.gr, F.se, F.av, F.eq, F.so,
			F.ox, F.co, F.an, F.sa, F.tr, F.ch, F.cu, F.sub, F.la
Eilam	4	4	F.ve, F.ny, F.gr, F.se, F.so, F.eq, F.av
Hamadan	4	4	F.gr, F.an, F.so, F.ox, F.sc, F.ve, F.pr, F.ny, F.eq,
			F.cr
Kermanshah	3	3	F.eq, F.so, F.sc, F.ve, F.pr, F.ny, F.gr, F.se, F.ox,
			F.av,
Khoy	2	2	F.sc, F.gr, F.se, F.eq, F.ve, F.pr, F.ox, F.av
Lahijan	4	4	F.gr, F.sc, F.pr, F.ny, F.se
Langaroud	3	3	F.sc, F.ve, F.pr, F.ny, F.gr
Mahabad	3	3	F.av, F.cu, F.gr, F.eq, F.so, F.sa, F.ve, F.se, F.ox,
			F.cr
Myaneh	3	3	F.av, F.cu, F.gr, F.eq, F.ve, F.se, F.ox, F.cr
Nour	2	2	F.pr, F.ny
Sanandaj	4	4	F.gr, F.sc, F.pr, F.ny, F.se, F.eq, F.ve, F.av
Ramsar	3	3	F.gr, F.pr, F.ve
Rasht	5	5	F.sc, F.ve, F.pr, F.ch, F.sub, F.se, F.av, F.eq, F.ny,
			F.gr, F.so, F.ox, F.co, F.an, F.sa, F.la
Salmas	3	3	F.pr, F.ny, F.gr, F.se, F.co, F.an, F.sa, F.tr
Sari	5	5	F.pr, F.ny, F.gr, F.se, F.co, F.an, F.sa, F.tr, F.ch,
			F.sub, F.la
Takab	5	5	F.eq, F.ox, F.se, F.av, F.gr, F.ve, F.sa, F.cu, F.sp,
	_	_	F.cr
Tabriz	5	5	F.ve, F.gr, F.sa, F.av, F.se, F.eq., F.cu
Urmia	5	5 E	F.ve, F.gr, F.co, F.se, F.pr, F.eq, F.su

F.av=F. avenaceum, F.cu=F. culmorum, F.an=F. anthophilum, F.co=F. compactum, F.ch=F. chlamydosporum, F.eq=F. equiseti, F.gr=F. graminearum, F.pr=F. proliferatum, F.ox=F. oxysporum, F.sa=F. sambucinum, F.cr=F. crookwellense, F.sc=F. scripi, F.ny=F. nygamai, F.sp=F. sporotrichioides, F.so=F. solani, F.la=F. lateritium, F.tr=F. trichothecioides, F.se=F. semitectum, F.ve=F. verticillioides.

Results

Ninety five spike samples were collected and analyzed for the occurrence of Fusarium spp.. All the spike samples were found positive for Fusarium species. A total of 280 Fusarium strains were isolated and identified through morphological characters. According to morphological characters all these strains belonged to 19 Fusarium spp., were F. avenaceum, F. compactum, F. culmorum, F. equiseti, F. graminearum, F. nygamai, F. anthophillum, F. oxysporum, F. proliferatum, F. trichothecioides, F. crookwellense, F. chlamydosporum, F. sambucinum, F. scripi, F. solani, F. sporotrichioides, F. semitectum, F. lateritium and F. verticillioides (Table 1 & 2). Of the Fusarium isolates collected from the wheat growing areas in Iran, F. graminearum was the most prevalent with frequency of 28%, followed by F. verticillioides and F. proliferatum with frequency of 14% and 10% respectively

(Table 3). Four days after inoculation, disease symptoms were observed on the heads. Initially, FHB symptoms appeared on water-soaked then on lose their chlorophyll and mycelium developed abundantly in the infected spikelets and the infection spreads to adjacent spikelets or through the entire head. The results in pathogenicity tests indicated that F. graminearum had the greatest IS (75%), followed by F. culmorum (65%) F. crookwellense (50%), F. trichothecioides (50%), F. chlamydosporum (50%), F. avenaceum (45%), F. verticillioides (45%) and F. sporotrichioides (45%) are the most pathogenic on the heads and caused severe blighting of wheat heads following inoculation in the greenhouse (Table 3). Therefore, these eight species were considered highly to moderately pathogenic. F. proliferatum, F. equiseti and F. nygamai did not show blight heads but sometimes caused damage to inoculated spikelets (Table 3).

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		Table 2. Mor	phological chara	Table 2. Morphological characters of Fusarium spp. associated with wheat head blight.	n spp. associated	with wheat head	i blight.		
Name of species	Chlamydospo	Pigmentation	Number of	Microconidia	Types of conic	Types of conidigenious cells	General morphology	rphology	Macroconidia
	res	on PDA	septa		Poly	Mono	Apical cell	Basal cell	size (µm)
F. avenaceum	1	Yellow	3-5	+		+	Tapered to	Nfs	$40-69 \times 3.2-4.1$
							pomea		
F. anthophilum		Violet	3-4	+	+	+	curved	Pdfs	$33-66 \times 2.5-4.5$
F. crookwellense	+	Red	5	,	,	+	Tapered to	Fs	$32-72 \times 4.1-6.7$
							pointed		
F. culmorum	+	Red	3-4	,	,	+	Rounded	Notch	$32-54 \times 4.0-7.0$
F. chlamydosporum	+	Red	3-5	+	+	+	Curved and	Nfs	$34-44 \times 3.0-4.5$
							pointed		
F. compactum	+	Red	5			+	Tapered, elongate	ş.	24-65 × 3.0-6.3
F. equiseti	+	Brown	5-7	,	,	+	Tapered	Fs	$45-85 \times 3.2-5.6$
							,elongate		
F. graminearum	+	Red	5-7		,	+	Tapered	Fs	$35-75 \times 4.0-6.5$
F. lateritium	+	Beige	4-7	+		+	Hook or break	Nfs	$35-72 \times 3.6-6.0$
F. nygamai	+	Violet	3-5	+	+	+	Short and	Nfs	$27-57 \times 2.1-5.0$
							tapered		
F. oxysporum	+	Violet	3	+	,	+	Curved	Fs	$30-58 \times 3.0-5.8$
F. proliferatum	,	Violet	3-5	+	+	+	Curved	Pdfs	$23-60 \times 3.0-5.0$
F. solani	+	Ream	3-5	+	,	+	Rounded	Nfs	$34-66 \times 3.6-6.0$
F. semitectum	+	Brown	3-5		+	+	Curved and	Fs	$35-58 \times 3.0-5.0$
							Tapered		
F. sambucinum	+	Red	3-5	,	,	+	pointed	Fs	$35-56 \times 3.0-5.1$
F. sporotrichioides	+	Red	3-5	+	+	+	Curved and	Nfs	$26-53 \times 3.0-5.5$
							Tapered		
F. scirpi	+	Violet	2-9	+	+	+	Tapered and elongate	Fs	46-80 × 3.9-6.0
F. trichothecioides	+	Red	3-5		,	+	Tapered to	Fs	$36-56 \times 3.3-5.5$
							pointed		
F. verticillioides		Violet	3-5	+		+	Curved	Nfs	$36-62 \times 2.5-4.5$
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+= Presence, -= Absence, Poly= Polyphialidic, Mono= Monophialidic, Pdfs= Poorly developed foot shape, Nfs= Notch or foot shape, Fs= Foot shape

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Name of Species	samples from the largest area of wheat Infected spikelets (%)	Frequency (%)
•	•	1 1 7
F. avenaceum	45	8
F. anthophilum	0	2
F. crookwellense	50	6
F. culmorum	65	6
F. chlamydosporum	50	3
F. compactum	0	1
F. equiseti	10	3
F. graminearum	75	28
F. lateritium	0	2
F. nygamai	6	3
F. oxysporum	0	2
F. proliferatum	12	10
F. solani	0	1
F. semitectum	0	2
F. sambucinum	0	2
F. sporotrichioides	45	3
F. scirpi	0	1
F. trichothecioides	50	3
F. verticillioides	45	14

Discussion

In our study identification of *Fusarium* spp., and their pathogenicity on wheat was investigated in the largest area of wheat plantation in Iran. The fungal isolation assays made on spike samples collected throughout the Kermanshah, Kurdistan, Hamadan, Khuzestan, Western Azarbaijan, Estern Azarbaijan and Eilam provinces clearly indicate that the members of section *Discolor* could be pathogenic to the wheat and suggested that *F. graminearum* and *F. culmorum* could be the main causal agent of head blight. Several studies have shown that *Fusarium* spp., in section *Discolor* can be readily isolated from cereal grains and according to this, results obtained in this study are in agreement with the previous findings of Nicholson *et al.*, (2003) and Dawson *et al.*, (2004).

Investigations on the pathogenicity of these species indicated that F. graminearum and F. culmorum are the most important pathogens and have a main role to cause FHB on wheat in mentioned provinces. The results of this study are in agreement with the previous literatures from north and south of Iran (Zare & Ershad, 1997; Moosawi-Jorf et al., 2007). In this survey all of the Fusarium species isolated from spike samples indicates that Fusarium species are capable contaminants of grains of wheat and may produce mycotoxins. On the other hands the occurrence of mycotoxins in wheat by Fusarium species is of great concern worldwide, and their presence in processed feeds and foods seems unavoidable (Bottalico & Perrone, 2002). Of the Fusarium isolates collected from mentioned provinces, F. graminearum was the most prevalent with a frequency of 28%, followed by F. verticillioides and F. proliferatum with a frequency of 22% and 18% respectively. Similar results were found by other researchers in north and south of Iran (Zare & Ershad, 1997; Moosawi-Jorf et al., 2007; Kachuei et al., 2009).

This study concludes that *Fusarium* spp., can cause FHB on wheat crop and may produce mycotoxins which have impact on human and animal health. This report is the first in-depth study to identity, pathogenicity and

distribution of *Fusarium* spp. from wheat in Iran. We believe that this study will serve as a basis for further identification of *Fusarium* species using molecular techniques. Mycotoxin profiles produced by these species are under progress.

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