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Fusarium graminearum

(Fusarium Head Blight)

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Background Information

Common Names:

Fusarium head blight; FHB, head blight of maize

Scientific Name:

Fusarium graminearum (anamorph = asexual stage), Gibberella zeae (teleomorph = sexual stage)

Synonyms:

Botryosphaeria saubinetii, Dichomera saubinetii, Dothidea zeae, Fusarium roseum, Gibbera saubinetii, Gibberella roseum, Gibberella saubinetii, Sphaeria saubinetii, Sphaeria zeae

Taxonomy:

Kingdom: Animalia; Phylum: Ascomycota; Class: Sordariomycetes; Order: Hypocreales; Family: Nectriaceae

Crop Hosts:

Wheat (*Triticum* spp. L.), barley (*Hordeum vulgare* L.), oat (*Avena sativa*), maize (*Zea mays*), triticale (x *Triticosecale*)



Figure 1. Typical signs of *Fusarium graminearum* on wheat. Photograph from USDA ARS, http://www.ars.usda.gov/Main/docs.htm?docid=9756.

Introduction

Fusarium graminearum Schwabe [teleomorph Gibberella zeae (Schweinitz) Petch], is of world-wide importance on small grain cereals and corn, occurring under a wide range of soil and environmental conditions (CAB International 2003; Gilchrist and Dubin 2002; Parry et al. 1995; Stack 2003). Since the early 1990s, fusarium head blight (FHB) caused primarily by F. graminearum has become one of the most significant cereal diseases faced by producers in central Canada and the prairie region, and the midwestern United States (e.g., Gilbert and Tekauz 2000; McMullen et al. 1997b; Tekauz et al. 2000). Fusarium graminearum was identified by CIMMYT to be a major limiting factor to wheat production in many parts of the world (Stack 1999). The fungus can produce several mycotoxins, including deoxynivalenol (DON) and zearalenone. In non-ruminants, feed contaminated with DON can reduce growth rates, while zearalenone can cause reproductive problems (Charmley et al. 1996; D'Mello et al. 1999). Barley (Hordeum vulgare L. emend. graminearum infected with F. Bowden) contaminated with mycotoxins can also cause quality problems for the malting and brewing industries (Schwarz 2003). Fusarium graminearum has also been linked to human illnesses (Goswami and Kistler 2004).

Fusarium graminearum is a genetically diverse species, with eleven distinct lineages currently known as the FG complex (Qu et al. 2008). There is little gene flow within these lineages, and all are well suited to infect their hosts in warm and wet climates (O'Donnell et al. 2000; Hope et al. 2005). According to Qu et al. (2008) one lineage in China can begin infection below 15 °C, and all lineages can overwinter on crop debris in any climate wheat is grown. This leaves high risk of continual infection, especially with rotations of less than two years between host crops.

Environmental conditions, especially temperature and moisture, are the key factors influencing the distribution and severity of fusarium head blight caused by *F. graminearum* (Shaner 2003; Stack 1999; Sutton 1982; Xu 2003). Moisture appears to be the most important environmental factor influencing the severity of infection caused by *F. graminearum* in small grain cereals, given

that fusarium head blight development can occur at temperatures that range from approximately 9 °C to 30 °C (Anon. 2011; de Wolf et al. 2003; McMullen et al. 1997a; Shaner 2003; Stack 1999).

Known Distribution

Fusarium graminearum has been reported wherever wheat is grown (Sutton 1982; CAB International 2003; Goswami and Kistler 2004), and infection has reached epidemic proportions in the United States over the last decade (O'Donnell et al. 2000). In South America, F. graminearum persists in southern Brazil and northern Argentina, while in Africa it plagues eastern South Africa and countries along the south coast of the Mediterranean (Wang et al. 2011). Fusarium graminearum persists within all of central Europe and southwestern Russia, and infects wheat fields grown along the eastern coast of China, and further inland where irrigation is used (Hope 2005; Ou et. al, 2008). In North America, increased levels of Fusarium head blight and percentage seed infection with F. graminearum have been associated with wheat under irrigation compared with dryland production (Clear and Patrick 2010; Strausbaugh and Maloy 1986; Turkington et al. 2005).

Description and Biology

Airborne ascospores produced by *G. zeae*, the sexual stage of *F. graminearum*, fall on flowering spikelets of wheat, germinate and enter the plant through natural openings, such as degrading anther tissue or stomates (Bushnell et al. 2003; Trail 2009). The fungus then grows, spreading through the xylem and pith of the wheat. As colonization continues, tissue becomes bleached and necrosis occurs (Figure 1).

Following infection, the fungus expresses genes for mycotoxin production, including DON, which causes shriveled, undersized grains known as tombstones (Trail 2009).

Fusarium graminearum has both sexual and asexual lifecycles (Figure 2) (Mathre 1997; Parry et al. 1995). Sexual reproduction takes about two weeks, and since F. graminearum is homothallic, it does not need two parents. Meiosis produces ascospores, which are forcibly discharged into the air through flask-shaped perithecia. Asexual conidia are produced during especially wet periods, and are moved via rain-splash dispersal. Conidia also overwinter on crop residues.

Host Crops and Other Plants

Fusarium graminearum affects many cereal crops. The main host crops are wheat (Triticum spp.), and barley (Hordeum spp.); however, rye (Secale cereale) and triticale (X Triticosecale) can also be affected (Parry et al. 1995). Fusarium graminearum causes head blight or 'scab' on rice (Oryza spp.), oats (Avena spp.) and Gibberella stalk and ear rot disease on maize (Zea spp.), and can also infect other plant species without causing

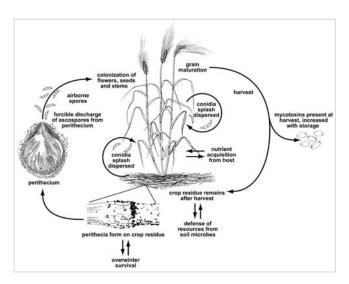


Figure 2. Fusarium graminearum life cycle (Trail 2009).

disease symptoms. Other host genera include *Agropyron*, *Agrostis*, *Bromus*, *Calamagrostis*, *Cenchrus*, *Cortaderia*, *Cucumis*, *Echinochloa*, *Glycine*, *Hierochloe*, *Lolium*, *Lycopersicon*, *Medicago*, *Phleum*, *Poa*, *Schizachyrium*, *Secale*, *Setaria*, *Sorghum*, *Spartina*, and *Trifolium* (Farr et al. 1989; Goswami and Kistler 2004).

Potential Distribution

CLIMEX (Sutherst et al. 2007) was used to infer potential distribution and abundance of *F. graminearum* based on knowledge regarding prevailing conditions where the species exists. CLIMEX calculates an Ecoclimatic Index (EI) that describes the suitability of locations for growth and survival. The EI is defined by variables that reflect conditions during the growing season (Growth Index) combined with variables that describe the effect of stress (Stress Index).

Temperature parameters were initially based on published information related to FHB biology (Anderson 1948; Anon. 2011; Clear and Patrick 2010; Doohan et al. 2003; McMullen et al. 1997; Parry et al. 1995; Shaner 2003; Sutton 1982; Xu 2003). Adjustments were made to reflect the known distribution and severity of FHB in the eastern prairies of Canada, especially Manitoba, and Prince Edward Island on the east coast of Canada (Clear and Patrick 2010; Gilbert and Tekauz 2000; Martin and Johnston 1997; Martin and MacLeod 1991; Tekauz et al. 2000).

Moist conditions resulting from high relative humidity, dew, rainfall or irrigation during flowering are critical for head infection (Mathre 1997; Wiese 1987). Much research indicates that high levels of moisture are associated with infections (e.g., Anderson 1948; McMullen et al. 1997a; De Wolf et al. 2003). Overall, adequate periods of either high humidity or surface wetness over successive days are required for FHB epidemics to occur. Soil moisture parameters were fitted iteratively to distribution data on FHB at Manitoba and Prince Edward Island where FHB is an established problem. As suggested pre-

Table 1. CLIMEX Parameter Values for Fusarium graminearum

Parameter	Description	Value
Moisture		
SMO	lower soil moisture threshold	0.2
SM1	lower optimum soil moisture	0.45
SM2	upper optimum soil moisture	1.5
SM3	upper soil moisture threshold	2.5
Temperatur	re	
DV0	lower threshold	9 °C
DV1	lower optimum temperature	20 °C
DV2	upper optimum temperature	25 °C
DV3	upper threshold	35 °C
Heat Stress		
TTHS	heat stress temperature threshold	35 °C
THHS	temperature threshold stress accumulation rate	0.005 week ⁻¹
Dry Stress		
SMDS	soil moisture dry stress threshold	0.2
HDS	stress accumulation rate	-0.01 week ⁻¹
Wet Stress		
SMWS	soil moisture wet stress threshold	2.5
HWS	stress accumulation rate	0.01 week ⁻¹
Threshold A	nnual Heat Sum	
PDD	number of degree-days above DV0 needed to complete one generation	300 °C days
Irrigation So	cenario	
	2.5 mm day-1 as top-up throughout the year	

viously by Yonow et al. (2004) for *Pyrenophora semeniperda* (Brittlebank and Adam) Shoemaker, the lower soil moisture parameter (SM0) was set to 0.2.

Cold Stress was not used in the model, as F. graminearum overwinters in the coldest grain-growing areas. Wet Stress parameters have minimal impact worldwide, although some Wet Stress does accumulate in small areas of Colombia, Sierra Leone, and in parts of Asia. Heat Stress precludes persistence across the globe where temperatures regularly exceed 35 °C. There are areas of limiting Dry Stress on all continents. A PDD value of 300 was set to limit the distribution in Norway, whilst retaining as suitable the areas shown in Lanseth and Elen (1997) and Koziak et al. (2004). The PDD value has minimal impact in the southern hemisphere, only precluding F. graminearum from persisting in the coldest areas of the Andes and in Tierra del Fuego. Its impact is greatest in the northern hemisphere, where most of the Arctic Circle and some surrounding areas are made unsuitable.

To verify the model (Table 1) for temperate northern climates, results were compared with independent data for the observed distribution of FHB for northern European countries (Baumgardt et al. 2008; Börjesson 2010; Bottalico 1998; Bottalico and Perrone 2002; CAB International 2003; Elen et al. 1997; Fredlund et al. 2008; Gagkaeva et al. 2004; Henriksen 1999; Hofgaard et al. 2010; Kosiak et al. 1997, 2003, 2004; Langseth and Elen 1996; 1997; Waalwijk et al. 2003; Yli-Mattila et al. 2002, 2008, 2010). The EI values are consistent with the known northern European occurrences of this disease and the projected range is also consistent with distribution data available elsewhere (Fernandes et al. 2004) (Figure 3).

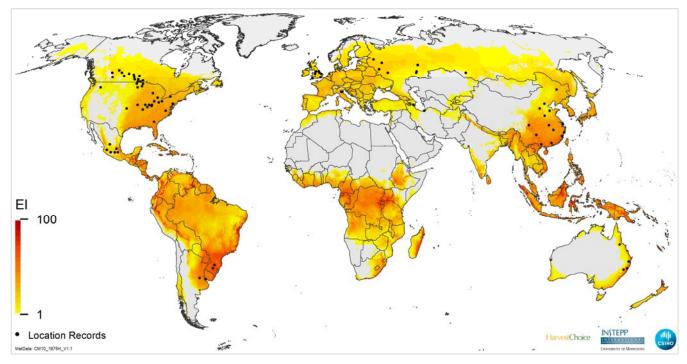


Figure 3. Modelled global climate suitability for *Fusarium graminearum* under dryland production. Location records taken from the Global Biodiversity Information Facility (GBIF).

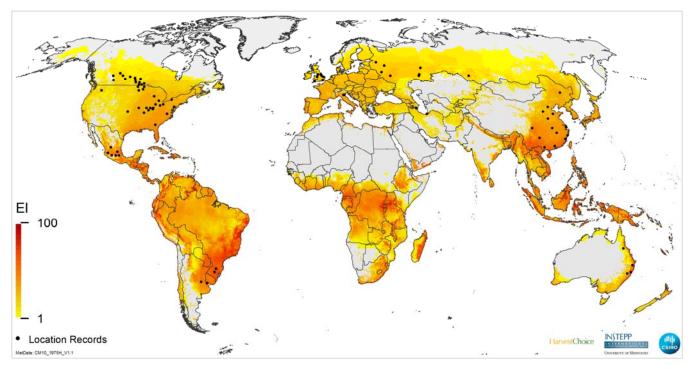


Figure 4. Modelled global climate suitability for *Fusarium graminearum* as a composite of natural rainfall and irrigation based on the irrigation areas identified in Siebert et al. (2005). Location records taken from GBIF.

An irrigation scenario of 2.5 mm day-1 applied as top-up increases both the potential range of *F. graminearum* as well as the suitability of areas already within the modelled range (Figure 4).

Potential Impact in Africa

The projected potential distribution in Africa (Figure 5) matches the known distribution of *F. graminearum*, and indicates areas at climatic risk.

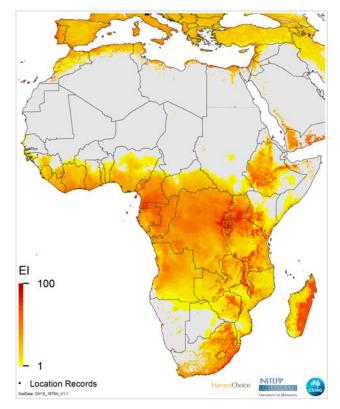


Figure 5. Modelled climate suitability of Africa for *Fusarium graminearum* as a composite of natural rainfall and irrigation based on the irrigation areas identified in Siebert et al. (2005). Location records taken from GBIF.

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Revision History

Figure 4 was a duplicate of Figure 3 in the original September 2014 publication. The revised March 2016 version contains the correct figure.