Root Rot of Dry Edible Bean Caused by *Fusarium graminearum*

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**Abstract**

*Fusarium graminearum* was identified as a root pathogen of a diverse array of dry edible bean genotypes under both field and greenhouse conditions in North Dakota. In comparisons under controlled conditions, root rot caused by *F. graminearum* was equal or greater than that caused by *F. solani* f. sp. *phaseoli*. Out of eleven dry bean genotypes evaluated in controlled conditions, Eclipse, VAX 3, and T-39 had the lowest root rot severity values for both *F. graminearum* and *F. solani* f. sp. *phaseoli*. A significant and positive correlation between genotype response to *F. graminearum* and *F. solani* f. sp. *phaseoli* indicates that genetic resistance to both pathogens may be related.

**Introduction**

Fusarium root rot is one of the major yield-limiting diseases of dry edible bean (*Phaseolus vulgaris* L.) in North Dakota and Minnesota production regions (13). More dry edible beans are produced in North Dakota and Minnesota than in any other region in the United States. Dry edible bean market classes grown in North Dakota and Minnesota include black, great northern, red kidney, navy, pinto, and a few other specialty classes. Previously, most Fusarium root rot of dry edible bean was considered to be caused by *Fusarium solani* (Mart.) Sacc. f. sp. *phaseoli* W.C. Snyder & H.N. Hansen (Fsp) (6). Recently, several reports of *F. graminearum* Schwabe (Fg) causing disease on below-ground tissues of dicotyledonous plants have been published (1,4,5,7,8,12,17). In North Dakota and Minnesota, Fg is very common and causes Fusarium head blight on barley (*Hordeum vulgare* L.) and wheat (*Triticum aestivum* L.) (16). The objectives of our research were to: (i) determine if a pathogenic association between Fg and dry edible bean occurs under natural field conditions; (ii) test Fg isolates from dry edible bean to compare aggressiveness with an Fsp isolate; and (iii) evaluate dry edible bean genotypes for resistance to root rot caused by Fg and Fsp.

**Dry Edible Bean Genotype Field Study**

**General methods.** Dry edible bean genotypes Eclipse (black bean), Maverick (pinto bean), Norstar (navy bean), Red Hawk (dark red kidney bean), and Rojo Chiquito (small red bean) were planted into a field in Fargo, ND, at the North Dakota State University Agricultural Experiment Station on 1 June 2006 into a Fargo clay soil (fine, smectitic, frigid Typic Epiaquerts). Planting depth was approximately 4 cm. Each plot was 4 rows wide (38 cm centers) and 4.6 m long. The field had been planted to spring wheat and used for Fusarium head blight fungicide research trials in 2005. The experimental design was a randomized complete block (RCB) with 4 replications. The field study originally was designed to be a dry bean cultivar yield and performance test, but served as a source of *Fusarium* isolates for further studies and as a preliminary trial in which dry bean genotypes could be compared for root rot severity.
Fusarium graminearum from roots. Fusarium pathogens causing root rot were identified through isolations as follows. On 29 June, twenty plants per plot were arbitrarily chosen and dug, collected, and taken to the laboratory. Roots were removed from the plants, washed to remove soil, and cut into sections approximately 1 cm in length. Root pieces from all plants were combined, and 50 root pieces were selected arbitrarily, surface sterilized in a 0.5% solution of NaOCl for 2 min, rinsed in sterile distilled water three times, and placed into petri dishes containing Komada’s medium (14).

All Fusarium isolates that grew from the plated roots appeared to be Fg according to morphological characters (15). To confirm the identification, six isolates were sent to the Fusarium Research Center (Penn State University, University Park, PA). All six isolates were confirmed as Fg through morphological characters by the Fusarium Research Center, and for five of the isolates, partial translation elongation factor 1-alpha sequences were generated for identification as described by Geiser et al. (11). The sequences from the five isolates were 99% to 100% identical to Fg. The mean number of plated root pieces (out of 50) in which Fg was isolated from each genotype is shown in Table 1. These data were analyzed using the general linear models procedure (PROC GLM) of SAS (Version 9.2, SAS Institute Inc., Cary, NC), and means were compared using Fisher’s protected least significant difference (LSD), where $\alpha = 0.05$. Genotypes did not significantly differ for the number of roots in which Fg was recovered.

<table>
<thead>
<tr>
<th>Genotype</th>
<th>Root rot severity (1 to 7)$^x$</th>
<th>Root pieces with $F. graminearum$ (no.)$^y$</th>
</tr>
</thead>
<tbody>
<tr>
<td>Red Hawk</td>
<td>4.3 a</td>
<td>6.3 a</td>
</tr>
<tr>
<td>Rojo Chiquito</td>
<td>4.3 a</td>
<td>3.0 a</td>
</tr>
<tr>
<td>Maverick</td>
<td>3.5 b</td>
<td>6.8 a</td>
</tr>
<tr>
<td>Norstar</td>
<td>2.4 c</td>
<td>2.3 a</td>
</tr>
<tr>
<td>Eclipse</td>
<td>2.0 c</td>
<td>2.0 a</td>
</tr>
</tbody>
</table>

$x$ Means within a column followed by the same letter are not significantly different according to Fisher’s protected least significant difference ($\alpha = 0.05$).

$y$ Represents the mean number of root pieces that yielded $F. graminearum$ colonies. A total of 50 root pieces collected from each plot were plated onto Komada’s medium.

Root rot severities. Dry edible bean genotypes were assessed for root rot severity in the field. On 12 August, ten plants from each plot were dug, roots were washed to remove soil, and roots were rated for root rot severity using a 1 to 7 scale described by Schneider and Kelly (18), where: 1 = healthy roots with no discoloration of root or hypocotyl and no reduction in root mass; 2 = 0.1 to 0.2 cm small reddish brown lesions at the base of the hypocotyl, with normal root mass and size; 3 = increase in intensity and size and coalescing of localized root/hypocotyl lesions approximately 180° around the stem, with lesions from 0.5 to 1 cm and 10 to 20% root discoloration but no reduction in root mass size; 4 = increase in intensity of discoloration and size of hypocotyl lesions, with lesions extending and completely encircling the stem, 5 to 10% root mass reduction, and 95% of the roots discolored; 5 = increasingly discolored and extended hypocotyl lesions, with 100% of the roots intensely reddish-brown and 20 to 50% root mass reduction; 6 = hypocotyl lesions encircling the stem extending up to 2 cm, intense root mass discoloration, and 50 to 80% root mass reduction; and 7 = pithy or hollow hypocotyl with very extended lesions, 80 to 100% root mass reduction, and functionally dead. Data were analyzed using PROC GLM in SAS, and means were compared using Fisher’s protected LSD, where $\alpha = 0.05$. 

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The dry edible bean genotypes in the field differed significantly for root rot severity. The genotype Eclipse had the lowest root rot severity rating, which was significantly lower than the root rot ratings for all other genotypes except Norstar (Table 1). The genotypes Red Hawk and Rojo Chiquito had the same root rot rating, which was significantly greater than the ratings for all other genotypes.

**Pathogenicity and Aggressiveness of* Fusarium graminearum* Isolates**

**General methods.** Three Fg isolates (denoted as R-10062, R-10063, and R-10064) collected from the dry edible bean genotype field study in Fargo, ND, and one Fsp isolate originally collected from a dry bean field near Staples, MN, were used to inoculate the roots of the dry bean genotype Montcalm (dark red kidney bean) to determine their pathogenicity and aggressiveness. A potted plant assay with a modified inoculum layer technique (3) was used in the greenhouse. Inoculum was prepared by growing each isolate on autoclaved sand-cornmeal for 1 week at 22 to 25°C under a 12-h light/dark cycle. Two pre-germinated seeds of ‘Montcalm’ were planted in 9-cm plastic pots containing steam-pasteurized potting mix (Sunshine Mix no. 1; Sun Gro Horticulture Ltd., Seba Beach, AB, Canada). The seeds were covered with a thin layer of potting mix, followed by 10 g of Fsg or Fsp sand-cornmeal inoculum and more potting mix (approximately 5 g). As negative controls, a non-infested sand-cornmeal mixture and a no sand-cornmeal mixture treatments also were included. Pots were watered to saturation after planting in the greenhouse and lightly watered on alternate days. The greenhouse temperature ranged from 20 to 22°C. Natural sunlight was supplemented with fluorescent lights that were set for a 12-h light/dark cycle. Each individual pot represented an experimental unit, and each experiment consisted of five replications per treatment. Eighteen days after planting, plants were removed from the soil and the roots were washed. Root rot severity was evaluated using a 1 to 7 scale (18). Pieces of symptomatic and healthy root tissue (negative controls) were surface sterilized in a 0.5% solution of NaOCl for 2 min, rinsed in sterile distilled water three times, and placed into petri dishes containing potato dextrose agar (PDA; Becton Dickinson and Company, Sparks, MD) to recover *Fusarium* species, and the identities of *Fusarium* species were confirmed by morphology characteristics (15). The experiment was repeated. The statistical design was a RCB. Data were analyzed using PROC GLM in SAS. No significant ($P \leq 0.05$) experiment × isolate interaction was observed, therefore, the data from the two experiments were combined for analysis. Means were compared using Fisher’s protected LSD, where $\alpha = 0.05$.

**Pathogenicity and aggressiveness of isolates.** All Fg isolates and the Fsp isolate caused significantly greater root rot severity on the dry bean genotype Montcalm compared to the negative control treatments (Table 2). Both Fg and Fsp cultures were recovered from roots that were inoculated with the respective isolates, no *Fusarium* cultures were recovered from negative control treatments. Fg isolates differed in the level of root rot severity they caused, but the range of root rot severity values was not wide (range of 2.5 to 3.0). Roots inoculated with the Fg isolates had significantly greater root rot severity than those inoculated with the Fsp isolate. Root rot symptoms caused by Fg and Fsp were similar (Fig. 1).
Table 2. Comparison of three *Fusarium graminearum* (Fg) isolates and a *F. solani* f. sp. *phaseoli* (Fsp) isolate for their aggressiveness in causing root rot severity on ‘Montcalm’ dry bean using an inoculum layer inoculation technique in the greenhouse.

<table>
<thead>
<tr>
<th>Isolate</th>
<th>Root rot severity (1-7)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Fsp</td>
<td>1.7 c</td>
</tr>
<tr>
<td>Fg R-10062</td>
<td>2.5 b</td>
</tr>
<tr>
<td>Fg R-10063</td>
<td>2.8 ab</td>
</tr>
<tr>
<td>Fg R-10064</td>
<td>3.0 a</td>
</tr>
<tr>
<td>Non-infested sand-cornmeal control</td>
<td>1.0 d</td>
</tr>
<tr>
<td>No sand-cornmeal control</td>
<td>1.0 d</td>
</tr>
</tbody>
</table>

x Results reported were averaged over two trials.

y Values followed by the same letter are not significantly different according to Fisher’s protected least significant difference ($\alpha = 0.05$).

**Evaluation of Dry Bean Genotypes for Resistance to *Fusarium graminearum* and *Fusarium solani* f. sp. *phaseoli***

**General methods.** The dry bean genotypes Eclipse (black bean), Matterhorn (great northern bean), Maverick (pinto bean), Montcalm (dark red kidney bean), Norstar (navy bean), Othello (pinto bean), Red Hawk (dark red kidney bean), Rojo Chiquito (small red bean), T-39 (black bean), VAX 3 (small red bean), and Vista (navy bean) were evaluated for their resistance to Fg and Fsp using a “paper towel inoculation method” described by Bilgi et al. (3). This method had been used previously to screen dry bean genotypes for resistance to Fsp (3). To determine if this method would work with Fg, a preliminary trial was conducted with ‘Montcalm.’ The results of the preliminary trial indicated that Fg caused significantly greater root rot severity on Montcalm compared to the negative control treatments using this method (Mathew, Bradley, and Rasmussen, unpublished). The paper towel inoculation methods used to evaluate dry bean genotypes for resistance to Fg, briefly described, were: seeds were planted into 266-cm$^3$ drinking cups filled with vermiculite and placed into a growth chamber maintained at 14 h of light and 10 h of darkness with day and night temperatures of 21 and 18°C, respectively. After plants grew in the vermiculite for 10 days, they were removed from the cups, washed, and placed onto layers of bleached paper towels. Roots within the paper towels were inoculated with sterilized wheat grains infested with either an Fg isolate originally collected from a wheat field in Ransom County, ND (designated as isolate R4), or an Fsp isolate originally collected from a dry bean field near Staples, MN. After inoculating, plants were wrapped inside paper towels,
moistened with sterilized distilled water, and placed in a plastic bag inside the
growth chamber. Plastic bags were placed in an aluminum stand that allowed
the plants to be upright. Plants were removed 10 days after inoculation and the
roots were rated using a 1 to 7 root rot severity scale (18). Two roots were inside
each plastic bag, and each plastic bag was considered an experimental unit. The
experimental design was a RCB with 4 replications, and the experiment was
repeated. Data were analyzed using PROC GLM in SAS. No significant
\( P \leq 0.05 \) experiment \( \times \) treatment interaction was observed, therefore, the data
from the two experiments were combined for analysis. Least-square means
(\textit{LSMEANS}) were calculated and \textit{t}-test comparisons among dry bean genotypes
and between \textit{Fusarium} isolates were made using the \textit{PDIFF} option in SAS.

Spearman’s rank correlation (PROC \textit{CORR SPEARMAN}) was used in SAS to
determine the relationship between genotype responses to \textit{Fg} and \textit{Fsp}.

\textbf{Reaction of dry bean genotypes.} A significant \( P < 0.0001 \) dry bean
genotype \( \times \textit{Fusarium} \) isolate interaction was detected; therefore, data are
presented by \textit{Fusarium} isolate (Table 3). The dry bean genotype Rojo Chiquito
had the greatest root rot severity caused by \textit{Fg} compared to all other genotypes.
The dry bean genotype T-39 had the least root rot severity caused by \textit{Fg}
compared to all other genotypes, but was not significantly different than VAX 3.
Similar to the \textit{Fg} root rot reaction, genotype Rojo Chiquito had the greatest root
rot severity caused by \textit{Fsp} compared to all other genotypes, but was not
significantly different than Montcalm. The genotypes Eclipse, VAX 3, and T-39
had the least root rot severity caused by \textit{Fsp} compared to all other genotypes.
For all genotypes except Matterhorn and Othello, the \textit{Fg} isolate caused
significantly \( P \leq 0.05 \) more severe root rot compared to the \textit{Fsp} isolate.
Spearman’s rank correlation analysis indicated that a positive (Spearman’s rank
correlation coefficient = 0.89) and significant \( P < 0.0001 \) correlation was
present between genotype response to \textit{Fg} and \textit{Fsp}.

\begin{table}
\centering
\caption{Root rot severity reactions of dry edible bean genotypes
inoculated with \textit{Fusarium graminearum} or \textit{F. solani} f. sp. \textit{phaseoli}
using a paper towel method in a growth chamber\textsuperscript{x}.
}
\begin{tabular}{llll}
\hline
Genotype & \textit{Fg}\textsuperscript{y} & \textit{Fsp}\textsuperscript{y} & \textit{P > F}\textsuperscript{z} \\
\hline
Rojo Chiquito & 6.2 a & 5.0 a & 0.0001 \\
Red Hawk & 5.4 b & 4.5 bc & 0.0001 \\
Montcalm & 5.3 b & 4.9 ab & 0.0153 \\
Maverick & 5.1 bc & 4.2 cd & 0.0001 \\
Matterhorn & 4.6 cd & 4.3 cd & 0.0510 \\
Othello & 4.5 d & 4.1 d & 0.0510 \\
Vista & 4.3 de & 3.2 e & 0.0001 \\
Norstar & 4.0 e & 3.5 e & 0.0008 \\
Eclipse & 3.5 f & 1.9 f & 0.0001 \\
VAX 3 & 3.2 fg & 1.9 f & 0.0001 \\
T-39 & 3.1 g & 2.1 f & 0.0001 \\
Mean & 4.5 & 3.6 & 0.0001 \\
\hline
\end{tabular}
\textsuperscript{x} Results reported were averaged over two trials.
\textsuperscript{y} Root rot severity (1 to 7) caused by \textit{Fusarium graminearum} (\textit{Fg})
or \textit{F. solani} f. sp. \textit{phaseoli} (\textit{Fsp}). Values followed by the same
letter within a column are not significantly different according to
least-square mean \textit{t}-tests (\( \alpha = 0.05 \)).
\textsuperscript{z} \textit{P}-value for the comparison between \textit{Fg} and \textit{Fsp} within a row.
\end{table}
Discussion

From our research results, we report here the first confirmation of a pathogenic association between Fg and dry edible bean based on field infection. Chongo et al. (8) reported that when inoculated under controlled conditions with an Fg isolate collected from a wheat head, dry bean had severe root rot and reduced seedling emergence. Under field infection conditions in our study, Fg caused root rot on five different dry bean genotypes, each representing a different market class. As a root rotting pathogen, Fg was equal or had significantly greater aggressiveness on dry bean roots compared to Fsp under controlled conditions in our research. In comparisons of Fg and Fsp in our research trials, only one Fsp isolate was used. Comparisons of additional Fsg and Fsp isolates from diverse locations are needed to verify if Fg generally causes greater root rot severity on dry bean compared to Fsp.

Dry bean genotypes differed in their response to Fg based primarily on differences in disease severity. This indicates that a type of partial resistance to Fg root rot may be present in some genotypes. Three genotypes (Eclipse, VAX 3, and T-39) had Fg root rot severity values lower than all other genotypes tested. All three of these genotypes represent market classes that have been developed from the Mesoamerican gene pool. This is interesting in that genotypes developed from the small-seeded Mesoamerican gene pool generally have been less susceptible to Fsp root rot compared to genotypes developed from the large-seeded Andean gene pool (2). In addition, these same three genotypes (Eclipse, VAX 3, and T-39) had the lowest Fsp root rot severity values compared to other genotypes. Bilgi et al. (3) had the same finding in both field and controlled conditions. Considering that the same three genotypes had the lowest root rot severity ratings for both Fusarium species and that a strong and significant correlation existed between genotype reactions to Fg and Fsp, it is likely that genotypes that have resistance to Fg will have resistance to Fsp, and vice versa. Additional research is needed to better understand this relationship and the number of genes involved in root rot resistance.

In light of our findings, growers that plant dry edible bean following wheat or barley fields that were affected by Fusarium head blight may need to consider root rot management tactics such as planting a cultivar with a high level of Fusarium root rot resistance or applying a fungicide seed treatment. Ellis et al. (10) reported that fungicide seed treatments could reduce disease on soybean seedlings caused by Fg. Our findings may have implications beyond dry bean production. Rotation with nongramineous crops has been a management practice used to help control Fusarium head blight (caused by Fg) of wheat and barley (9,19). More research is needed to understand how the pathogenic relationship between Fg and dry bean will affect Fg diseases in other crops. In addition, the prevalence of Fg root rot on dry bean is not known, and focused surveys are needed to determine this.

Acknowledgements

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Literature Cited


