

Diseases Caused by Fungi and Fungus-Like Organisms

First Report of Crown Rot Caused by *Fusarium redolens* on Wheat in Kazakhstan

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Plant Dis. 105:3302, 2021; published online as <https://doi.org/10.1094/PDIS-10-21-0015-PDN>. Accepted for publication 24 March 2021.

Fusarium crown rot, caused by several species within the genus, is a major constraint that results in significant losses in wheat production worldwide. In June 2019, diseased wheat plants with typical symptoms of crown rot including discoloration on the first two or three internodes of the stem just above the soil line and stunted, dry rotted, and discolored roots were collected in several bread wheat fields during the maturity stage in Almaty, East Kazakhstan, and Karaganda Regions of Kazakhstan. For each field, approximately 20 tillers were randomly sampled. Symptomatic tissues were surface sterilized in 1% NaClO for 2 min, rinsed with sterile distilled water three times, air dried in a laminar flow hood, and then transferred to Petri dishes containing one-fifth-strength potato dextrose agar (PDA). After incubating in the dark at 23°C for 5 days, 79 single-spore isolates showing cultural and microscopic characteristics of *Fusarium* were obtained on PDA and Spezieller-Nährstoffarmer agar (SNA). Colonies were initially white but later produced a beige to pink diffusible pigment in PDA. Microconidia that formed on aerial monophialides were hyaline, zero to one septum, oval- to kidney-shaped, and measured 4.3 to 10.3 × 1.9 to 3.4 μm (average 7.8 × 2.6 μm), and macroconidia were straight to slightly curved, three to five septate, and measured 18.7 to 38.8 × 2.9 to 6.6 μm (average 29.9 × 4.7 μm), with foot-shaped basal cells on SNA. Chlamydospores were present on PDA. Sequence analysis based on portions of translation elongation factor 1α (*TEF1*) and the nuclear ribosomal internal transcribed spacer region (ITS rDNA) loci with primers EF1/EF2 (O'Donnell et al. 1998) and ITS1/ITS4 (White et al. 1990) identified 29 of the 79 isolates as *Fusarium redolens* Wollenw. The sequences of

the five representative isolates had 99.85% similarity to those of *F. redolens* strains available in GenBank (e.g., ITS, MT435063; *TEF1*, GU250584). The *TEF1* (MW403914 to MW403918) and ITS rDNA (MW397138 to MW397142) sequences of the isolates were deposited in GenBank. The morphological features are consistent with the described features of *F. redolens* (Leslie and Summerell 2006). To confirm pathogenicity of the five isolates, five pregerminated seeds of wheat cultivar Seri 82 were placed in a 9-cm-diameter pot filled with a sterile potting mix containing equal volumes of peat, vermiculite, and soil. An approximately 1-cm-diameter 7-day-old mycelial plug of each isolate was individually placed in contact with the seeds. Seeds were covered with the same potting mix, and then the pots were maintained for 4 weeks in a growth chamber at 23°C with a 12-h photoperiod. The experiment was conducted twice with three replicate 15-cm-diameter pots with five plants per pot. Controls were inoculated with sterile agar plugs using the same procedure. After 4 weeks, all the inoculated plants showed stunted growth with brown discoloration in most parts of the crown and roots, whereas no symptoms were observed in the control plants. The mean severity of the disease for each isolate was between 2.1 and 2.7 according to the scale of 1 to 5 described by Gebremariam et al. (2015). The pathogen was reisolated from crowns of diseased plants but not from asymptomatic control tissues and was identified morphologically based on the methods described above, fulfilling Koch's postulates. Although several morphological features are shared by *F. oxysporum* and *F. redolens*, Baayen et al. (2001) showed that these species could be easily distinguished using molecular data. The pathogen was previously reported as *F. redolens* associated with crown rot of wheat in Turkey (Gebremariam et al. 2015) and Saskatchewan, Canada (Taheri et al. 2011). The presence of *F. redolens* causing crown rot is confirmed in the six wheat fields surveyed in Kazakhstan, for the first time. This pathogen may pose a risk for wheat production, and further studies needed to determine the impact on the crop in Kazakhstan.

References:

- Baayen, R. P., et al. 2001. *Phytopathology* 91:1037.
Gebremariam, E. S., et al. 2015. *Plant Dis.* 99:1280.
Leslie, J. F., and Summerell, B. A., eds. 2006. *The Fusarium Laboratory Manual*. Blackwell, Oxford, U.K.
O'Donnell, K., et al. 1998. *Proc. Natl. Acad. Sci. U.S.A.* 95:2044.
Taheri, E. A., et al. 2011. *Can. J. Plant Pathol.* 33:559.
White, T. J., et al. 1990. Page 315 in: *PCR Protocols: A Guide to Methods and Applications*. Academic Press, San Diego, CA.

The author(s) declare no conflict of interest.

Keywords: *Fusarium* crown rot, *F. redolens*, surveillance, wheat

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