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Pathogen Identification of Fusarium Spp. And Chemotypes Associated With Wheat Crown Rot in Hebei Province of China

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Abstract : Fusarium crown rot (FCR) diseased wheat seedlings were collected from different wheat-growing counties in seven different regions (Baoding, Cangzhou, Handan, Hengshui, Langfang, Shijiazhuang, and Xingtai) in Hebei province, China from 2019 to 2020. One-hundred twenty-two Fusarium isolates were isolated from crown rot diseased wheat seedlings and identified morphologically, confirmation was undertaken molecularly, and species-specific PCR was utilized to verify the morphological identification of F. psuedograminearum, F. graminearum, F. asiaticum, and F. culmorum. The predominant Fusarium species associated with wheat crown rot in the Hebei province were F. psuedograminearum, F. graminearum, F. asiaticum, and F. culmorum with isolation frequency of 85.25%, 12.30%, 1.64%, and 0.81%, respectively. All the Fusarium strains isolated from the different wheat-growing fields were qualitatively tested for toxigenic chemotypes using toxin-specific primers and chemotaxonomically classified into DON, 3-ADON, 15-ADON, and NIV. Among F. psuedograminearum identified, 84.62% were classified as DON chemotypes, 6.73% as 15-ADON chemotypes, 3.84% as 3-ADON chemotypes, and 4.81% of them had NIV as detected by the toxin-specific PCR results. Most of the F. graminearum isolates produced 15-ADON, and only two isolates had NIV chemotypes. F. asiaticum and F. culmorum produce chemotype of 15-ADON and 3-ADON, respectively. Pathogenicity test results showed that F. pseudograminearum and F. graminearum had strong pathogenicity, and F. asiaticum and F. culmorum had moderate pathogenicity to wheat in Hebei province.

Keywords: crown rot, pathogen, wheat, Fusarium species, mycotoxin

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