Diseases Caused by Fungi and Fungus-Like Organisms

First Report of *Nigrospora sphaerica* Associated with Leaf Spot Disease of *Crossandra infundibuliformis* in India

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Crossandra (Crossandra infundibuliformis (L.) Nees.) is one of the main floriculture crops in Karnataka. In March to June 2020, a characteristic leaf spot disease of unknown etiology with an incidence of 10 to 12% (~30 ha area evaluated) was observed in Southern Karnataka (Mysore, Mandya). Initially, the symptoms developed as small specks (3 to 8 mm), characterized by circular to irregular shapes in the beginning and coalesced to form larger lesions. Ten samples were collected and the associated fungal pathogen isolated on potato dextrose agar (PDA) medium amended with chloramphenicol (60 mg/liter). Briefly, infected leaves were cut into small pieces and surface sterilized with 2% sodium hypochlorite solution, rinsed three times with sterile distilled water (SDW), blot dried, inoculated onto PDA medium, and incubated at room temperature (27 ± 2°C) for 3 to 5 days. Fungal colonies developed from the segments were subcultured through hyphal tipping to fresh PDA plates to get pure cultures. Twelve pure cultures were obtained. Mycelia were initially white and eventually turned gray. The conidia were black, single celled, smooth, spherical to subspherical, 9 to 18 μ m in diameter (n = 50), and borne singly on a hyaline vesicle at the tip of each conidiophore. The identity was initially established as Nigrospora sp. based on culture and morphology (Deepika et al. 2021). To confirm the identity of fungal isolates, molecular sequence analysis was performed for two representative isolates (CIT1 and CIT2). ITS-rDNA, tub2, and EF-1 α gene were amplified using primers ITS1/ITS4, T1/T22, and EF1-728F/986R (Glass and Donaldson 1995; O'Donnel and Cigelnik 1997; White et al. 1990), then purified and sequenced. The BLASTn analysis of ITS, tub2, and EF-1 α showed 99 to 100% similarity with reference sequences from GenBank of Nigrospora sphaerica (ITS: 520 bp, KX985935 - LC7312; MH854878 - CBS:166.26; tub2: 357 bp, MZ032030 - WYR007, 350 bp, KY019606 – LC7298, KY019522 – LC4278, KY019520 – LC4274; *EF-1α*: 472 bp, KY019397 - LC7294, KY019331 - LC4241; MN864137 - HN-BH-3) and the sequences were deposited in GenBank (ITS: OL672271 and OL672272; tub2: OL782120 and OL782121; EF-1a: ON051604 and ON051605; Wang et al. 2017). The associated fungal pathogen was identified as N. sphaerica (Sacc.) Mason (Chen et al. 2018; Deepika et al. 2021) based on the cultural, morphological, microscopic, and molecular characteristics. Pathogenicity tests were conducted on healthy plants (crossandra cv. Arka; n = 30) grown under greenhouse conditions (28 ± 2°C; 80% RH). Inoculations were made with conidial suspension (18-day-old N. sphaerica isolate CIT1, 10⁶ conidia/ml) prepared in SDW, and healthy plants sprayed with SDW (n = 10) served as controls. All the plants were covered with polyethylene bags for 24 to 48 h and observations were made at regular intervals. Typical necrotic lesions developed on 16 plants 12 days after inoculation, but no symptoms were observed on control plants. The associated pathogen was reisolated from diseased leaves and its identity confirmed based on morphology and cultural characteristics. Earlier, N. sphaerica was associated with various tree species as an endophyte, and recently several reports have described it to cause disease on various crop plants (Deepika et al. 2021). However, there are no previous reports of N. sphaerica causing leaf spot disease on C. infundibuliformis from India. Early diagnosis of this leaf spot disease will help the floriculturist adopt suitable management practices to avoid significant economic loss.

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