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Parasporendocladia W.P. Wu & Y.Z. Diao

Parasporendocladia was introduced by Wu & Diao (2022) to accommodate Sporendocladia bactrospora. The type species, Parasporendocladia bactrospora W.P. Wu & Y.Z. Diao was first introduced as *Phialocephala bactrospora* (Kendrick 1961), then later reclassified as Sporendocladia bactrospora based only on conidial development, but without molecular data (Wingfield et al. 1987). Later, the phylogenetic analysis of ITS, SSU, and LSU sequences showed that S. bactrospora belongs to Microascales (Hypocreomycetidae, Sordariomycetes), while the type species of Phialocephala, P. dimorphospora belongs to Helotiales (Leotiomycetidae, Leotiomycetes) (Grünig et al. 2002, Jacobs et al. 2003). On the other hand, the type species of Sporendocladia, S. fomusa, belongs to Chaetosphaeriaceae (Chaetosphaeriales, Sordariomycetes) (Wu & Diao 2022). Based on molecular data, S. bactrospora was distinct from species of Phialocephala and Sporendocladia, thus the genus Parasporendocladia was introduced (Wu & Diao 2022). It is difficult to distinguish Parasporendocladia morphologically from Phialocephala and Sporendocladia (Wu & Diao 2022). Parasporendocladia has conidiophores that are solitary to aggregated, erect or flexuous, branched at the apex, septate, brown, and smooth (Wu & Diao 2022). The conidiogenesis cells are holoblastic and the conidia are cylindrical, hyaline, aseptate, smooth, truncate at both ends and extruded in long chains (Wu & Diao 2022). Parasporendocladia bactrospora (≡ Phialocephala bactrospora) was first isolated as a saprobe from Populus trichocarpa (Kendrick 1961). Although the species has been reported as a saprobe on dead material of Clushia melchiorii in Brazil and Fagus sylvatica in Czechoslovakia (Barbosa et al. 2007, Kubatova 1992), some studies have reported it as a plant pathogen associated with wounds on native broadleaved trees in Norway (Roux et al. 2014) and canker and dieback of Juglans regia in Iran (Sohrabi & Mohammadi 2023).

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Published online 5 April 2024