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Salinomyces Czachura & Piatek

Salinomyces was introduced by Czachura et al. (2021) with S. polonicus Czachura & Piatek as the type species. The name is derived from the environment inhabited by the isolates collected from brine samples (Czachura et al. 2021). As described by Czachura et al. (2021), members of the genus typically have greenish grey or blackish colonies. The mycelia consist of hyphae which are hyaline, pale brown to brown, septate, branched verrucose or smooth (Czachura et al. 2021). Brown chlamydospores and multicellular bulbil-like structures are sometimes present (Czachura et al. 2021). The conidiogenous cells are intercalary on hyphae which may be simple or with inconspicuous collarettes (Czachura et al. 2021). The conidia are aseptate to 3-septate, pale brown to brown, smooth or rugose (Czachura et al. 2021). The genus is closely related to *Hortaea*; however, analysis of combined ITS, LSU and rpb2 loci revealed that this genus formed a well-supported clade separate from other species of *Hortaea* (Czachura et al. 2021). Moreover, they also differ morphologically from the members of *Hortaea* based on the appearance of the conidiomata and conidia (Czachura et al. 2021). Salinomyces polonicus was isolated from brine samples collected from Southern Poland; thus, it is moderately halophilic (Czachura et al. 2021). Another species, previously known as Hortaea thailandica (Crous et al. 2009), was reassigned to this genus after analysis of combined ITS, LSU, and rpb2 genes showed that it is a sister species of S. polonicus. This was subsequently renamed Salinomyces thailandicus (Crous & K.D. Hyde) Czachura & Piątek. This species was previously isolated from corals (Mitchison-Field et al. 2019) and monumental sites close to the sea (Isola et al. 2016). Thus, it is believed that this species is also halotolerant. The sexual morph is unknown.

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