

**Outlineoffungi.org - Note 199, [Haplohelminthosporium](#)**

**[Haplohelminthosporium](#)** Konta & K.D. Hyde

Based on morphological characteristics and phylogenetic analyses of combined ITS, LSU, SSU, and tef1- $\alpha$  sequence data, the monotypic genus [Haplohelminthosporium](#) was established by [Konta et al. \(2021b\)](#) with *H. calami* as the type species. [Haplohelminthosporium](#) is distinguished by its unbranched conidiophores arising solitarily or fasciculate from the stroma-like bulbous basal cells that are hyaline in the middle, brown to red-brown at 1–2-cells above the base, pale brown to red-brown and curved at the apical cell with well-defined non-cicatrized small pores and with a single olive-brown conidium arising from each conidiophore ([Konta et al. 2021b](#)) (M. Erdoğan).

**Reference**

Konta S, Hyde KD, Eungwanichayapant PD, Karunarathna SC et al. 2021b – Multigene phylogeny reveals *Haploanthostomella elaeidis* gen. et sp. nov. and familial replacement of *Endocalyx* (Xylariales, Sordariomycetes, Ascomycota). Life 11(6), 486. [Doi 10.3390/life11060486](https://doi.org/10.3390/life11060486)