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Haploanthostomella Konta & K.D. Hyde

Based on phylogenetic analyses of a combined dataset of ITS, LSU, *rpb2*, and *tub2* nucleotide sequence data as well as unique morphological characteristics, [Konta et al. \(2021a\)](#) introduced [Haploanthostomella](#) to accommodate *H. elaeidis* (the type species) collected from dead leaves and rachis of *Elaeis guineensis* (M. Erdoğdu).

Reference

Konta S, Hyde KD, Karunarathna SC, Mapook A et al. 2021a – Multi-Gene phylogeny and morphology reveal *Haplohelminthosporium* gen. nov. and *Helminthosporiella* gen. nov. associated with palms in Thailand and a checklist for *Helminthosporium* reported worldwide. Life 11(5), 1–53. [Doi 10.3390/life11050454](https://doi.org/10.3390/life11050454)