

Outlineoffungi.org - Note 154, [Engleromyces](#)

[Engleromyces](#) Henn.

[Zhou et al. \(2021\)](#) have recently been the first to generate DNA sequence data of a species of [Engleromyces](#) and finally confirmed the placement of this genus, whose species form massive stromatal on bamboo that are used in folk medicine, in the *Xylariaceae*. Previously, it had been only retained in the family based on chemotaxonomy and ascospore morphology (cf. [Wendt et al. 2018](#)) (M. Stadler).

Reference

- Zhou H, Wang QT, Tong X, Hou CL. 2021 – Phylogenetic analysis of *Engleromyces sinensis* and identification of cytochalasin D from culture. *Mycological Progress* 20, 1343–1352 (2021)
[Doi 10.1007/s11557-021-01739-z](#)
- Wendt L, Sir EB, Kuhnert E, Heitkämper S et al. 2018 – Resurrection and emendation of the *Hypoxylaceae*, recognised from a multi-gene genealogy of the *Xylariales*. *Mycological Progress* 17:115–154. [Doi: 10.1007/s11557-017-1311-3](#)