## Outlineoffungi.org - Note 987 Allotrechispora

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Allotrechispora L.W. Zhou & S.L. Liu

<u>Allotrechispora</u> was introduced by Liu et al. (2022) to accommodate <u>Allotrechispora</u> <u>gatesiae</u> L.W. Zhou, S.L. Liu & T.W, as the type species based on the morphological characteristics and phylogenetic analysis of ITS, LSU, tef1-α, and rpb2 sequence data. The type species was isolated from fallen trunk of <u>Atherosperma moschatum</u> in Australia. This genus has three species including <u>A. daweishanensis</u>, <u>A. gatesiae</u>, and <u>A. xantha</u>. This genus generally occurs as annual, resupinate, cream to cinnamon-buff, corticioid, thin and soft basidiomes, and loosely attached to the substrates. Hymenophore is white to cinnamon-buff when fresh, smooth to tuberculate, and cracked with age. Hyphal system is monomitic, generative hyphae with clamp connections, absence of isodiametric subhymenial hyphae, hyaline and thin-walled. Basidiospores are ellipsoid to ellipsoid, or oblong shape, hyaline with thin-walled, smooth, inamyloid, indextrinoid, and acyanophilous. Basidia are subcylindrical to subclavate, with four sterigmata, hyaline, thin-walled, and occurred clamp connection. Cystidia are absent. Phylogenetically *Fibrodontia* is closest genus to <u>Allotrechispora</u>. The taxonomic placement of <u>Allotrechispora</u> is in <u>Allotrechisporaceae</u>, <u>Allotrechisporales</u>, <u>Agaricomycetes</u>, <u>Agaricomycotina</u>, <u>Basidiomycota</u>.

## Reference

Liu SL, He SH, Wang XW, May TW, et al. 2022 – *Trechisporales* emended with *segregation* of *Sistotremastrales* ord. nov. (*Basidiomycota*). Mycosphere 13(1), 862–954. https://doi.org/10.5943/mycosphere/13/1/11

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