

## Outlineoffungi.org - Note 962 *Brownlielloideae*

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***Brownlielloideae*** S.Y. Kondr., Kärnefelt, Elix & A. Thell

The subfamily *Brownlielloideae* was established to accommodate eight genera namely *Brownliella*, *Lazarenkoella*, *Marchantiana*, *Raesaeneniana*, *Streimanniella*, *Tarasginia*, *Tayloriella* and *Thelliana* based on the phylogenetic analyses ([Kondratyuk et al. 2015](#)). The subfamily was named after the type genus *Brownliella* S.Y. Kondr., Kärnefelt, Elix, A. Thell & Hur and the members were mainly reported from the Southern and Northern hemispheres ([Kondratyuk et al. 2015](#)). The family formed a clade outside *Teloschistaceae* based on the recent phylogenetic analyses conducted by Wilk et al. ([2021](#)) based on a subset of DNA. The taxa of *Brownlielloideae* dispersed among all the subfamilies mostly within *Teloschistoideae* but also within *Caloplacoideae* and *Xanthorioideae* in the phylogenetic analyses and the generic type *Brownliella* itself nested within *Teloschistoideae* ([Wilk et al. 2021](#)). The DNA sequences data used in the establishment of the subfamily was found to have “chimeric” DNA data set whereas concatenated dataset contained ITS and LSU loci from the lichen mycobiont, but SSU from a fungal contaminant ([Vondrák et al. 2018](#), [Llewellyn 2019](#), [Wilk et al. 2021](#)). Thus, *Brownlielloideae* was synonymized under *Teloschistoideae* ([Wilk et al. 2021](#)). The taxonomic placement of this subfamily *Teloschistoideae* is in *Teloschistaceae*, *Teloschistales*, *Lecanoromycetidae*, [Lecanoromycetes](#), [Pezizomycotina](#) and *Ascomycota*.

### References

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