



SHORT COMMUNICATION

Validation of two fungal names in *Marasmius* Fr. (Marasmiaceae)Frances E. Guard^a, Matthew D. Barrett^b, Arian Farid^c, Matthew Smith^d & Teresa Lebel^e

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When publishing *Marasmius vagus* F.E.Guard, M.D.Barrett & Farid (Crous *et al.* 2020) and *Marasmius lebeliae* F.E.Guard (Crous *et al.* 2019), the authors omitted the appropriate herbarium code when designating the types, which were lodged in the Queensland Herbarium (BRI). They only cited the type specimen accession numbers as “AQ1008080” and “AQ799986”, respectively. Although the acronym AQ means ‘Accession Queensland’, and all collections with an AQ number are curated at the Queensland Herbarium, no direct reference was made in either publication to BRI. The names are therefore invalid as per Art. 40.7 of the *International Code of Nomenclature* (Turnland *et al.* 2018). We here validate both names, by presenting the corrected type citations along with references to the original descriptions. As required under Art F.5.7 (May *et al.* 2019), new identifiers for the names have been obtained from MycoBank.

Marasmius vagus* F.E.Guard, M.D.Barrett & Farid, sp. nov.*Mycobank number: MB837473.**

Marasmius vagus F.E.Guard, M.D.Barrett & Farid, *Persoonia* 44: 415 (2020), *nom. inval.* [Mycobank number: MB833552]; see here for validating description.

Holotypus: Australia, Queensland, Mt Carbine, S16°34'44.1" E145°11'13.7", in savanna grassland leaf litter, 7 Mar. 2018, *F. Guard & S. McMullanFisher SMF3041* (BRI AQ1008080; ITS and LSU sequences: GenBank MT117839 and MT110674)

Marasmius lebeliae* F.E.Guard, sp. nov.*Mycobank number: MB837474.**

Marasmius lebeliae F.E.Guard, *Persoonia* 42: 427 (2019), *nom. inval.* [Mycobank number: MB828485]; see here for validating description.

Holotypus: Australia, Queensland, Dilkusha Nature Refuge, Maleny, Site 1, in leaf litter and *Casuarina* needles under *Elaeocarpus grandis* and *Allocasuarina cunninghamiana*, in regenerating subtropical rainforest, 3 Feb. 2018, *F. Guard F2018011* (BRI AQ799986; ITS sequence GenBank: MK211200).

Acknowledgements

We would like to thank Tom May (Royal Botanic Gardens Victoria) for pointing out the nomenclatural issue and providing suggestions regarding validation of names.

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