Unearthing high species- and sectional-level diversity of *Agaricus* (Agaricaceae) in Australia and New Zealand

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The genus Agaricus contains over 550 species worldwide and includes both edible and poisonous species. However, Australian and New Zealand Agaricus are underrepresented in recent phylogenetic studies and their diversity remains incompletely understood. Thirty-two species of Agaricus have been described from Australian and New Zealand type material and the majority of these are unplaced in the current subgeneric classification. Field and herbarium-based observations of Australian and New Zealand Agaricus suggest that current species do not accurately delimit the morphological and genetic diversity. We sought to improve our understanding of Agaricus species diversity in Australia and New Zealand, to estimate their relationships, and to place species within a global Agaricus phylogeny. Sequence data were generated for three genetic markers (ITS1&2, tef1-a, LSU) for 371 Agaricus individuals. The evolutionary history of ~460 Agaricus individuals was inferred using maximum likelihood and Bayesian inference analyses. Agaricus subgenera and sections were resolved, and many Australian and New Zealand species were phylogenetically placed for the first time. The resulting phylogeny indicates that there is considerable undescribed Agaricus biodiversity in this region, with many misidentified taxa and cryptic species groups. A strongly supported clade containing only Australian and New Zealand species has been identified, which would represent the first Australasian-only lineage in the genus. The vast and undescribed diversity of Agaricus in Australia and New Zealand uncovered during this study has implications for determinations of rarity and exotic introductions, foragers and poisons on-call services, and potentially to the global classification of the genus.