

## Author Correction: Genome expansion and lineage-specific genetic innovations in the forest pathogenic fungi *Armillaria*

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In the version of this Article originally published, it was incorrectly stated that “16,687 protein-coding genes were inferred for the most recent common ancestor (MRCAs) of *Armillaria*”; the value was incorrect and it should have read “15,787”. This has now been corrected.

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