

## Supplementary Information

Review title: Progress and Prospects of Mycorrhizal Fungi Diversity in Orchids

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The following Supplementary Information is available for this article:

### **Table S1 | Recommended primers for high-throughput sequencing of orchid mycorrhizal partners.**

| Primer name       | Sequence (5'-3' direction) | Reference                     |
|-------------------|----------------------------|-------------------------------|
| Modified ITS1ngs  | TCCGTAGGTGAACCTGC          | Oja <i>et al.</i> (2015)      |
| Modified ITS1Fngs | GGTCATTTAGAGGAAGTAA        | Oja <i>et al.</i> (2015)      |
| ITS1              | TCCGTAGGTGAACCTGCGG        | White <i>et al.</i> (1990)    |
| Modified ITS4ngs  | TCCTSCGCTTATTGATATGC       | Oja <i>et al.</i> (2015)      |
| ITS4Tul2          | TTCTTTTCCTCCGCTGAWTA       | Oja <i>et al.</i> (2015)      |
| ITS1F             | CTTGGTCATTTAGAGGAAGTAA     | Gardes and Bruns (1993)       |
| ITS86F            | GTGAATCATCGAATCTTTGAA      | Turenne <i>et al.</i> (1999)  |
| ITS4              | TCCTCCGCTTATTGATATGC       | White <i>et al.</i> (1990)    |
| ITS4Tul           | CCGCCAGATTCACACATTGA       | Taylor (1997)                 |
| TW14ngs           | CTATCCTGRGRGAAAYTTC        | Tedersoo <i>et al.</i> (2014) |

**Oja, J., Kohout, P., Tedersoo, L., Kull, T., and Kõljalg, U. (2015).** Temporal patterns of orchid mycorrhizal fungi in meadows and forests as revealed by 454 pyrosequencing. *New Phytol.*

**205**, 1608-1618. doi: 10.1111/nph.13223

**White, T. J., Bruns, T., Lee, S., and Taylor, J. (1990).** “Amplification and direct sequencing of fungal ribosomal RNA genes for phylogenetics,” in PCR Protocols: A Guide to Methods and Applications, eds. **M. A. Innis, D. H. Gelfand, J. J. Sninsky, and T. J. White** (San Diego, CA, USA: Academic Press), 315-322.

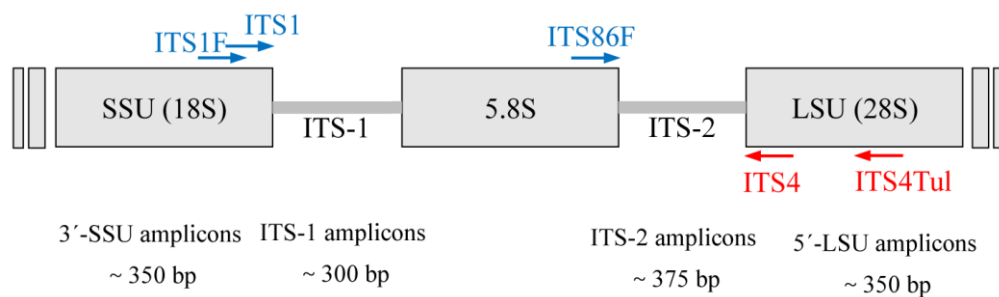
**Gardes, M., and Bruns, T. D. (1993).** ITS primers with enhanced specificity for basidiomycetes-application to the identification of mycorrhizae and rusts. *Mol. Ecol.* **2**, 113-118. doi:

10.1111/j.1365-294X.1993.tb00005.x

**Turenne, C. Y., Sanche, S. E., Hoban, D. J., Karlowsky, J. A., Kabani, A. M. (1999).** Rapid identification of fungi by using the ITS2 genetic region and an automated fluorescent capillary electrophoresis system. *J. Clin. Microbiol.* **37**, 1846–1851. doi: 10.1128/JCM

**Taylor, D. L. (1997).** The evolution of myco-heterotrophy and specificity in some North American orchids. dissertation. Berkeley: University of California.

**Tedersoo, L., Bahram, M., Pölme, S., Kõljalg, U., Yorou, N. S., Wijesundera, R., et al. (2014).** Global diversity and geography of soil fungi. *Science* **346**:1256688. doi: 10.1126/science.1256688



**Figure S1 | Schematic representation of primers recommended for MiSeq PE300 and HiSeq PE250 amplicon sequencing. Blue for forward primer, red for reverse primer.**