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| **Table S3** All EM fungal lineages and relative abundances in this study. |
| Lineage | GH (%) | HGL (%) | SHWL (%) | Total (%) |
| /tricholoma | 6.39  | 53.10  | 50.69  | 36.72  |
| /tomentella-thelephora | 40.21  | 3.55  | 0.39  | 14.72  |
| /suillus-rhizopogon | 18.18  | 2.05  | 22.90  | 14.38  |
| /piloderma | 2.38  | 33.59  | 0.26  | 12.08  |
| /wilcoxina | 0.73  | 2.55  | 15.06  | 6.11  |
| /russula-lactarius | 12.03  | \_ | 0.01  | 4.01  |
| /sebacina | 2.86  | 0.11  | 8.59  | 3.85  |
| /cortinarius | 10.82  | 0.04  | \_ | 3.62  |
| /geopora | 0.06  | 0.85  | 1.79  | 0.90  |
| /pseudotomentella | 0.37  | 2.20  | \_ | 0.86  |
| /tomentellopsis | 2.27  | \_ | \_ | 0.76  |
| /tuber-helvella | 0.83  | 0.78  | 0.28  | 0.63  |
| /inocybe | 1.78  | 0.01  | 0.01  | 0.60  |
| /hebeloma-alnicola | 0.04  | 0.91  | \_ | 0.32  |
| /cenococcum | 0.80  | 0.01  | \_ | 0.27  |
| /amphinema-tylospora | 0.15  | 0.01  | 0.03  | 0.06  |
| /hygrophorus | 0.03  | 0.11  | \_ | 0.05  |
| /Helotiales1 | \_ | 0.10  | \_ | 0.03  |
| /ramaria-gautieria | 0.07  | \_ | \_ | 0.02  |
| /genea-humaria | \_ | 0.01  | \_ | <0.00  |
| /otidea | 0.01  | \_ | \_ | <0.00  |