Supplementary Table S7. *Amylostereum areolatum* genetic diversity based on allelic diversity.

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| Population | Na | Alleleb | 1-Dc | Hexpd | Evennesse |
| Australia | 84 | 2.000 | 0.459 | 0.462 | 0.946 |
| Austria | 3 | 1.820 | 0.330 | 0.390 | 0.940 |
| Czech Republic | 4 | 1.640 | 0.260 | 0.300 | 0.930 |
| New Zealand | 14 | 2.450 | 0.510 | 0.530 | 0.930 |
| North America | 6 | 3.180 | 0.520 | 0.560 | 0.770 |
| South Africa | 55 | 2.550 | 0.420 | 0.430 | 0.880 |
| Spain | 8 | 2.450 | 0.520 | 0.550 | 0.980 |
| Mass-rearing strains | 5 | 2.550 | 0.540 | 0.600 | 0.910 |
| Total | 179 | 5.640 | 0.580 | 0.580 | 0.820 |

a N: Number of fungal isolates per collection site

b Allele: Mean number of observed alleles per loci

c 1-D: Mean number of Simpson index

d Hexp: Mean number of Nei's 1978 gene diversity

e Evenness: distribution of alleles abundance per population