Supplementary Table S5. *Deladenus siricidicola* genetic diversity based on allelic diversity.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
|  | Population | Na | Alleleb | 1-Dc | Hexpd | Evennesse |
| All populations | Australia | 117 | 4.33 | 0.50 | 0.50 | 0.79 |
| New Zealand | 16 | 2.17 | 0.28 | 0.29 | 0.65 |
| South Africa | 21 | 1.58 | 0.07 | 0.07 | 0.50 |
| South America | 6 | 2.08 | 0.37 | 0.40 | 0.81 |
| Spain | 12 | 1.92 | 0.18 | 0.19 | 0.74 |
| North America | 16 | 2.00 | 0.16 | 0.16 | 0.62 |
|  | Total | 188 | 6.25 | 0.60 | 0.60 | 0.79 |
| Australian sub-populations | New South Wales | 34 | 3.33 | 0.35 | 0.36 | 0.61 |
| South Australia | 12 | 2.17 | 0.36 | 0.37 | 0.76 |
| Victoria | 66 | 3.67 | 0.51 | 0.52 | 0.87 |
| Tasmania | 3 | 2.00 | 0.43 | 0.55 | 0.93 |
|  | Commercial | 2 | 1.42 | 0.20 | 0.26 | 0.96 |

a N: Number of strains 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