**Supplementary table 4.** Diversity statistics of STRUCTURE inferred populations

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| **Group** | **N** | **Na** | **Ne** | **I** | **He** | **PP (%)** |
| A | 16 | 1.47±0.036 | 1.29±0.014 | 0.29±0.011 | 0.18±0.008 | 69.24 |
| B | 11 | 1.04±0.041 | 1.21±0.014 | 0.21±0.011 | 0.13±0.007 | 48.07 |
| C | 14 | 1.462±0.037 | 1.30±0.014 | 0.30±0.011 | 0.19±0.008 | 69.98. |
| Mean | 13.7 | 1.32±0.051 | 1.27±0.008 | 0.26±0.006 | 0.17±0.004 | 62.43 |

Na: number of different alleles; Ne: number of effective alleles; I: Shannon's information index; He: Expected Heterozygosity; PP: percentage of polymorphic loci.