Table S4. The final SNP (single nucleotide polymorphisms) datasets, including number of SNPs, number of individuals, mean depth of coverage, and the standard deviation of depth of coverage. Estimation of effective migration surfaces is abbreviated to EEMS.

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| Dataset | Number of SNPs | Number of Individuals | Mean Depth | Standard Deviation of Depth |
| Phylogenetic | 5,529 | 75 | 56.98180744 | 92.39357691 |
| ADMIXTURE A | 4,653 | 73 | 57.12188336 | 92.18153906 |
| ADMIXTURE B | 4,338 | 70 | 57.16327472 | 92.68282951 |
| EEMS | 4,497 | 59 | 55.24484496 | 93.79999071 |