**Supporting Information Table S4**. Mean estimated Ln likelihood, standard deviation, and ΔK (Evanno et al. 2005) calculated across five replicates of STRUCTURE analysis for all eighteen populations for K=1-5, using a burn-in of 500,000 followed by 1,000,000 MCMC repetitions under a correlated allele frequencies with admixture model.

|  |  |  |  |
| --- | --- | --- | --- |
| *K* | *Ln Lik* | *σ* | Δ*K* |
| 1 | -44971.9 | 0.13 | N/A |
| 2 | -39169.1 | 0.86 | 6250.02 |
| 3 | -38753.8 | 17.09 | 0.34 |
| 4 | -38332.5 | 3.93 | 36.97 |
| 5 | -38056.6 | 173.04 | 1.59 |