**Table S3.** Summary of results for effective population size calculated from the individual inbreeding rate and through the individual coancestry rate and the number of equivalent subpopulations.

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| Parameter  Breed | Population set | Effective population size calculated | | Number of equivalent subpopulations |
| through individual inbreeding rate (NeFi) | through individual coancestry rate (NeCi) |
| PRá | Historic | 49.02 | 83.33 | 1.70 |
| Current | 43.86 | 69.44 | 1.58 |
| PRE | Historic | 49.02 | 8.98 | 0.18 |
| Current | 48.54 | 8.90 | 0.18 |
| Há | Historic | 52.63 | 24.27 | 0.46 |
| Current | 53.19 | 23.47 | 0.44 |