**Table 3. Pairwise number of diagnostic allozyme loci between the primary genetic groups identified by PCoA.** Lower = number of fixed differences (10% tolerance for all shared alleles combined); upper = unbiased Nei’s Distance. Also shown are the observed heterozygosity counts (HO) and standard errors (S.E.) for each group.

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
| Lineage | KN  (20) | KE  (96) | KEm  (16) | KSxKE  (9) | KS  (39) | KWm  (27) | KW  (26) |
| KN | - | 0.16 | 0.14 | 0.15 | 0.29 | 0.31 | 0.41 |
| KE | 4 | - | 0.04 | 0.07 | 0.28 | 0.30 | 0.41 |
| KE­m | 4 | 0 | - | 0.04 | 0.23 | 0.25 | 0.38 |
| KSxKE | 5 | 0 | 0 | - | 0.08 | 0.10 | 0.18 |
| KS | 12 | 11 | 5 | 0 | - | 0.05 | 0.09 |
| KWm | 12 | 9 | 5 | 1 | 1 | - | 0.07 |
| KW | 18 | 17 | 16 | 4 | 3 | 2 | - |
| HO | 0.060 | 0.059 | 0.134 | 0.182 | 0.090 | 0.074 | 0.017 |
| ±S.E. | 0.016 | 0.010 | 0.026 | 0.033 | 0.021 | 0.018 | 0.008 |