**Table S1**. Details of 10 microsatellite loci (Skirnisdottir et al. 2013) used in this study, including allele sizes found across study populations.

|  |  |  |  |
| --- | --- | --- | --- |
| Locus | Primer sequence (5' - 3') | Dye | Allele size range (bp) |
| Multiplex 1 |  |  |  |
| *Clu29* | F: CGCGCGGTCAGCTCATCCTTAG | PET | 126-142 |
| R: TCGCGTGACGGACAGGTTTCG |
| *Clu34* | F: TCTGCGATAGTAGCGTCAGGGTTC | NED | 179-219 |
| R: AGGCCGGCTGATCAAGAGCAC |
| *Clu36* | F: CACGGCGAGTCAGACGAGGC | 6-FAM | 183-203 |
| R: GCTGCCGCTACTCCGCACAG |
| *Clu45* | F: GCGCAGGAATGCGCCTGAAG | PET | 268-298 |
| R: ACCGCAGCTTGTTGGGCAGG |
|  |  |  |  |
| Multiplex 2 |  |  |  |
| *Clu12* | F: CCACAACCGGTGGGTCCCG | 6-FAM | 188-202 |
| R: ACGCTCCTTCTGATCTTCGCCC |
| *Clu26* | F: CGAGAGAGGAGAACGCACGGC | 6-FAM | 97–121 |
| R: GGCACAAGTGCATGGGCACG |
| *Clu33* | F: TCATGCAAGCATTTGAGCGCCG | VIC | 173–193 |
| R: TGTTGCCTTGTAACTGCGCTTGAG |
| *Clu37* | F: CTTCACAGGTCGGGCGACGG | PET | 200-226 |
| R: GCACAGCGATGACGCTTGCAG |
| *Clu40* | F: TGGGCATACAGGTCTGAACACGC | NED | 246-266 |
| R: GCCACCTGCTGCAGCCTCTC |
| *Clu44* | F: CCGGCCCAGCCTGCCTTATG | 6-FAM | 269-287 |
| R: TGCCTGGAAACAGTGTATGGCAC |