|  |  |  |  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Locus | No. of alleles | No. of adult individuals | HO | HE | PIC | NE-1P | NE-2P | NE-PP | NE-I | NE-SI | HW | F(Null) |
| Nl3 | 9 | 33 | 0.79 | 0.80 | 0.77 | 0.57 | 0.39 | 0.20 | 0.07 | 0.37 | NS | -0.001 |
| G6 | 11 | 33 | 0.85 | 0.85 | 0.81 | 0.50 | 0.33 | 0.15 | 0.05 | 0.35 | NS | -0.014 |
| Nl6 | 8 | 33 | 0.52 | 0.75 | 0.69 | 0.67 | 0.50 | 0.31 | 0.11 | 0.41 | NS | 0.184 |
| P11 | 10 | 33 | 0.88 | 0.86 | 0.83 | 0.46 | 0.30 | 0.13 | 0.04 | 0.34 | NS | -0.015 |
| P223 | 14 | 33 | 0.85 | 0.84 | 0.81 | 0.50 | 0.33 | 0.15 | 0.05 | 0.35 | NS | -0.015 |
| P217 | 16 | 33 | 1.00 | 0.91 | 0.89 | 0.35 | 0.21 | 0.07 | 0.02 | 0.31 | NS | -0.057 |
| H29 | 15 | 33 | 0.88 | 0.90 | 0.88 | 0.37 | 0.23 | 0.08 | 0.02 | 0.31 | NS | 0.000 |
| H23 | 21 | 33 | 0.94 | 0.93 | 0.92 | 0.28 | 0.16 | 0.04 | 0.01 | 0.29 | ND | -0.014 |
| P219 | 6 | 33 | 0.85 | 0.76 | 0.71 | 0.65 | 0.47 | 0.29 | 0.10 | 0.40 | NS | -0.060 |

**Table S1** Results from allele frequency calculations with CERVUS v. 3.0 (Kalinovski et al. 2007).

For allele frequency calculations, only individuals were caught as adults (n=33) were used. HO: observed heterozygosity, HE: expected heterozygosity, PIC: mean polymorphic information content, NE-1P: non-exclusion probability for first parent, NE-2P: non-exclusion probability for second parent, NE-PP: non-exclusion probability for parent pair, NE-I: non-exclusion probability for identity, NE-SI: non-exclusion probability for sib identity, HW: Hardy-Weinberg  equilibrium test, NS: not significant; ND: not done; F (Null): null allele frequency.

Number of individuals: 33

Number of loci: 9

Mean number of alleles per locus: 12.222

Mean proportion of loci typed: 0.9935

Mean expected heterozygosity: 0.8450

Mean polymorphic information content (PIC): 0.8115

Combined non-exclusion probability (first parent): 0.00100702

Combined non-exclusion probability (second parent): 0.00002239

Combined non-exclusion probability (parent pair): 0.00000001

Combined non-exclusion probability (identity): 3.890E-0013

Combined non-exclusion probability (sib identity): 0.00006890