

S3 Table. Summary statistics for DNA sequence data.

Locus	N	Alignment Length	Parsimony Informative Sites	Unique Seqs	Segregating sites, S	Average nucleotide differences, k	Nucleotide diversity, π
mtDNA	225	2097	72	78	106	8.72	0.0043
PRLR	208	541	9	12	9	0.95	0.0018
R35	195	658	5	12	8	0.71	0.0019
scar298	213	341	13	14	13	1.92	0.0057
scar875	212	287	12	20	13	1.27	0.0044