

Supplementary table 2. Microsatellite loci developed for *M. auretenra*: locus name, forward and reverse primer sequences, number of alleles and GenBank accession number.

LOCUS	PRIMER SEQUENCES (5' - 3')	GENBANK ACCESSION NO	UNIVERSAL TAIL	FLUORESCENT DYES
MAUR_1	F: AACGATCAAAGTCGATGGAGACGTAAGGC R: TCCACTCAAGAACAATGTTGATACACAGGC	MT799827	Blacket A	6-FAM
MAUR_6	F: CGACTGGATTGAAGTTACTAGAGTGCGG R: CTCAAGTCATCCAAGGTCAGCAGCG	MT799828	Blacket A	6-FAM
MAUR_7	F: AATAGGCGCGTTCAGAGAACAAGGG R: CATGTCTAATCCCCATGATTGCGG	MT799829	Blacket A	6-FAM
MAUR_8	F: TTCTTATTCTGTGCTTCCTTTGCTGGG R: ATGCCAGGTTGCCTCTGATTGGC	MT799830	Blacket A	6-FAM
MAUR_12	F: GTGAATAAATGAATAACTGAGTACCTCCCG R: GAAGAAACGTCAACAGTGAGAGGGG	MT832345	Blacket A	6-FAM
MAUR_14	F: TTTAGGAAGGGAAGTGCCTGTTCCG R: TCACGGCTAATAAATTTACGTGCG	MT832346	Blacket A	6-FAM
MAUR_16	F: CTTCGGCAGGATCATTTGTAATCGG R: TTTAGCAGCTTGGTGCCAAACCC	MT832347	Blacket A	6-FAM
MAUR_17	F: GAGTTCCAAGGAGTTTTGAATTGCCC R: GATGTAGAATCGCAGGTTCCCTGGG	MT832348	Blacket A	6-FAM
MAUR_20	F: GGATAATGTTGTAGTCCATGCCTTGCC R: GCAATTAATTATCCCATGAAGCTCTGGC	MT832349	M13-mod B	ROX
MAUR_21	F: AAGTAGTTGCTTGACTCTTTTCTGAAGCG R: ACCTCCACCTTGACAGCTCTTTCCC	MT832350	M13-mod B	ROX
MAUR_23	F: TATGCGCACATTCAACTAGCATCCC R: TGTAAAGGCTTGGTTTCTTCATGCG	MT832351	M13-mod B	ROX
MAUR_24	F: GAAATCTTTGTTTCAAGGAGGAAGGGG	MT832352	M13-mod B	ROX

MAUR_26	R: TCTACCTTGGCTTGTTTCATCAGGGG			
	F: GTGGATGAGGAAGAGAGCAGTGGC	MT832353	M13-mod B	ROX
MAUR_30	R: CTGTCATGAGATCCAAACCTCCGC			
	F: TCCATAGTCTCACGATTGAGCG	MT832354	M13-mod B	ROX
MAUR_34	R: GAACAGTATGTTTACCCAGATCCC			
	F: CCTGGCTCCAAGTTGAAAGTAGGC	MT832355	M13-mod B	ROX
MAUR_35	R: TCAAGTTTGAAGACTGCAAGTAATCCC			
	F: GGGTATACACCCTGAAGTTTCACATAGCC	MT832356	M13-mod B	ROX
MAUR_36	R: CTCCGTAGGAATCCGAGCATTACCC			
	F: TCATGGTTGAGAGGTCATATTTTAGCCC	MT832357	M13-mod B	ROX
	R: ATAAAACACAAAGGCTGGTCACGGC			