Table 2. Primers used in the present study for microsatellite markers amplification and sequencing.

Locus	Primer	Sequence (5'– 3')	Tm(°C)
Proq01	Proq01_Fwd	AAGCGTCGCAGATCTAATGC	64.3
	Proq01_Rev	GACAGACCCTCGATGTTTGC	64.7
Proq01_2	Proq01_2_Fwd	ACTGTGAAACGAGCCTCCTG	64.4
	Proq01_2_Rev	ACACCATTGCCATCCATACC	64.4
Proq01_3	Proq01_3_Fwd	TGTGTACTCCACAGCGGCTA	64.5
	Proq01_3_Rev	TTGTCTTTCGGGTGTCCAAT	64.3
Proq02	Proq02_Fwd	GCCGAAGAAGGAGCTGATCT	64.8
	Proq02_Rev	GAGGACGGAACGATTCTCAA	64.1
Proq02_2	Proq02_2_Fwd	CCACTGTTAGAATCGCTGGG	64.5
	Proq02_2_Rev	CGTGAACGTGGAGTTGACTG	64.5
Proq03	Proq03_Fwd	AACCAGTCGATCTGTTCCCA	64.5
	Proq03_Rev	ATTTGCAATATGCTGGGTCG	64.6
Proq03_2	Proq03_2_Fwd	TAGAACACAAGGCATTGGCA	64.2
	Proq03_2_Rev	TCCAAATGAAGCGGGAAGTA	64.3
Proq03_3	Proq03_3_Fwd	GGGACTTCCTTGGCGTATCT	64.2
	Proq03_3_Rev	ATGGATGATTCTACGCCTCG	63.9
Proq04	Proq04_Fwd	TGAAGGTTATTGAAGAAAGACCG	63.0
	Proq04_Rev	CAAATCTCGCCCACCAAAC	65.4
Proq04_2	Proq04_2_Fwd	CGTTGGATAACCACTACGCA	63.5
	Proq04_2_Rev	CGATCGAATCCCATTTCACT	63.7
Proq04_3	Proq04_3_Fwd	ATGGTGGGTGCAGGGATT	65.4
	Proq04_3_Rev	CACCGTCAGCACTACCATTG	64.2
Proq05	Proq05_Fwd	TCCCTGCCGTCTGATAGTTC	64.2
	Proq05_Rev	AAGGTGCTGTGGACTGGTTC	64.2
Proq07	Proq07_Fwd	AAAGTCTGGATGTGAGGGCA	64.7
	Proq07_Rev	GATCTCTTGGTTGGAATGCG	64.5
Proq07_2	Proq07_2_Fwd	CCATGAACTGCCTTACGCTT	64.0
	Proq07_2_Rev	ATCGCGGTTGCTCTATTTGA	64.5
Proq07_3	Proq07_3_Fwd	CCATGAACTGCCTTACGCTT	64.0
	Proq07_3_Rev	ATCGCGGTTGCTCTATTTGA	64.5
Proq09	Proq09_Fwd	TCCGTTCAGGAACTGTCGAT	64.7
	Proq09_Rev	TCCATGGCAGTTGCTTCTTT	64.6
Proq10	Proq10_Fwd	GCCTTGAGTTGTAACCAATCCTTT	65.1
	Proq10_Rev	TCCTAGATGTTCCCGATTGGT	64.4
Proq10_2	Proq10_2_Fwd	GCCTCCCAGTTCATGACAAC	64.6
	Proq10_2_Rev	CTGCCGAAACTGCTTGCTAT	64.3
Proq11	Proq11_Fwd	ACACCCAATCACTACGACGG	64.8
	Proq11_Rev	TGAAGTGAGGACCTTTGGGA	64.6
Proq14	Proq14_Fwd	TCTTCGCATAGGGAGTTGGA	64.6
	Proq14_Rev	TGGTAGAATACCGTTCCCGA	64.1
Proq16	Proq16_Fwd	TTGAGGATTTCCGGAGACAA	64.5
	Proq16_Rev	ATGCGCAATAAGACCCAAGA	64.4
Proq17	Proq17_Fwd	TATCGTCCGCACTAAGGGAA	64.3
	Proq17_Rev	TGCTTCATTTCCGAAGGTGT	64.5
Proq17_2	Proq17_2_Fwd	GATCGGAAACCCAGGAATTT	63.8
	Proq17_2_Rev	GGGCCATATCCCATTCTTGA	65.7
Proq18	Proq18_Fwd	TCAGCACAATCAGTTCACGC	65.2
	Proq18_Rev	TCAGCATTTGCTGCTGTTGT	64.7

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