

Supplementary Table S3: SSR loci employed, primer sequences (in 5' to 3' order), and their main features.

Locus	Forward	Reverse	Core motif	Ta (°C)	Allelic size range (bp)	Reference
AMPA095	GCCAAAGTTAGGAAGGAGAATG	TTCATATTCAAACCCAATGAC	(AC)	56	79-87	[28]
AMPA112	CAAAGTGTCATTGTTCTTGCTAG	AAGGCCAAAACCTCCAGTAGAAG	(AG)	55	206-236	[28]
AMPA111	ACTGACAGCAGCAGAATGTGTC	TATCATCAGGGCTTTTGTGTG	(CTT)	55	171-207	[28]
AMPA113	CAACTCTGGTTTTCTGTCTCTC	CCTTCAGACTAGCTGAGCTTCC	(CT)	55	116-152	[28]
AMPA124	TCAACCTGTAATAACCTAGCTTGC	AGCCGATGACTTGATTTTCCTC	(GA)	55	152-174	[28]
UDAp-401	AAACCCTAGCCGCCATAACT	GCTAAAGGCCTCCGATACC	(TC)	56	152-176	[29]
UDAp-410	TTGTTGACAAGAAGAAAACAAAGC	CAACGGGTTGGTTTCAGAAG	(AG)	56	168-184	[29]
UDAp-414	CAAGCACAAGCGAACAAAAT	GGTGGTTTCTTATCCGATGC	(AG)	56	130-164	[29]
UDAp-415	AACTGATGAGAAGGGGCTTG	ACTCCCGACATTTGTGCTTC	(GA)	56	297-300	[29]
UDAP-419	TTCTTTTGGGATTGGTCTCG	GATTTTAAATAACCAACCAGCTTC	(GA)	56	135-149	[29]
UDAp-420	TTCCTTGCTTCCCTTCATTG	CCCAGAACTTGATTCTGACCA	(CT)	56	96-140	[29]
UDAp-446	CCTCCCCTAGATTTTCAGC	CGTGCTTGGGACATAGATCA	(GA)	54	150-180	[19]

Ta: temperature of annealing