Table S1 Primers and Probes Sequence(5'-3') of each SNP for PCR and LDR

Polymorphism	Primers and probes sequence(5'-3')	Length
rs494379 (A/G)	Forward: GGTCAAAGAGTACTCCATGGTCTT	90bp
	Reverse: CCTGGGAGCAAGGTGTGC	
Common probe	P-CTTTTTCAAAAAGACCATGGAGTACTTTTTTTTTTTTTT	
Discriminating probes	rs494379_A: TTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTT	98bp
	rs494379_G: TTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTATGACCATCAGAACCAGC	100bp
rs5749511 (C/T)	Forward: CTGCTGACGTTCAGGGAAAT	157bp
	Reverse: TGGGTAGGAAAAGCAAGCAC	
Common probe	P-TCACCCCTTTCCACTGCTCCTTTCCCTTTTTTTTTTTTT	
Discriminating probes	rs5749511_C:	102bp
	TTTTTTTTTTTTTTTTTTTTTTTTGATTGGTGGCCAGCCA	
	rs5749511_T :	144bp
	TTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTT	
rs3025058 (5A/6A)	Forward: FAM—TGGCCAAATATTTTCCCTGT	123bp
	Reverse: ACATGGGTCACGGCACCT	124bp
rs3918242 (C/T)	Forward: CAGCCTGGTCAACGTAGTGA	118bp
	Reverse: AAGCAATTCTCCTGCCTCAG	
Common probe	P-TGCGCCACCACGCCTGGCTAAATTTTTTTTTTTTTTTTT	
Discriminating probes	rs3918242_C: TTTTTTTTTTTTTTTCCCGAGTAGCTGGTATTATAGGCG	82bp
	rs3918242_T: TTTTTTTTTTTTTTTTTCCCGAGTAGCTGGTATTATAGGCA	84bp
rs17576 (A/G)	Forward: CTCGCCCCAGGACTCTA CAC	175bp

Reverse: GTGGAGGTACCTCGGGTCGGG

Common probe	P-CCC(AG)CAAGCCCTCCGGGAACTGTGGTTTTTTTTTTTTTTTT	
Discriminating probes	Rs17576_A:TTTTTTTTTTTTTTTTTGCACTCACCTGCCCCAAGGCCAGCT	168bp
	rs17576_GT:TTTTTTTTTTTTTTTTTGCACTCACCTGCCCCAAGGCCAGC	173bp