

S4 Table. Primers for Mutation Analysis of *SLCO2A1* Gene

Amplicon	Forward (5'–3')	Reverse (5'–3')	Size (bp)	RFLP*
Exon 1	GCCTGGAGTTGTCCGAGTAAG	TTTGACACCCGAGGAAAAGAG	620	
Exon 2	ACAGAATTTGAAGGCGGACAG	TGATGACGTTTCATCCTGGAG	655	
Exon 3	AATGCTTTGATCCTCCTCCTG	CAGAGGGAAAACCAAATTGC	676	
Exon 4	GCTGTTGGTTTAGGGTCCATC	CCCACATTCCACCTCTCTTTC	786	
Exon 5	AAGGGCAGAAAGTGAGTTTGC	GCAGGTCTCTTTGGAAGTTGG	570	
Exon 6	AATTCAGCCACCCTCAACAC	CAATAGCTGGGAGAGGGAATG	742	
Exon 7-8	GATTGCCAACCAGGAAAACCTG	AGGTGCTGTTTGCTTCAGGAG	778	<i>HpyCH4IV</i>
Exon 9	AGTTGCCACGTGAGAGATCAG	GTGTAGGCAAGGCAATCCTG	860	
Exon 10	TGCTTTGACTGGTTTCTGCTC	TGGTTGTCTTGACGCCTACTG	747	<i>HapII, BsrI</i>
Exon 11	GCAAAGAACCTTGCACATTG	GCCTCCCTCTGCAATAAACAC	710	
Exon 12	AAACCGTCCACATGGATTTTG	CAGCATCCTTCTCTCCACTCC	643	
Exon 13	GTGGCCCTTCATGTTCTCTTC	GCCCGTGTATCTCCACTCTG	564	
Exon 14	CAAATGAGGACTGGGGTTTTTC	GAAGACCACAAAAGGGACAGG	617	

**HpyCH4IV*, *HapII*, and *BsrI* were used for genotyping of c.940+1G>A, c.1372G>T, and c.1461+1G>C mutations, respectively. RFLP, restriction fragment length polymorphism.