

Table S2. Primer sequences for 14 polymerase chain reaction (PCR) fragments, and the sizes of 13 *SCARB1* exons.

Fragment	Forward Primer Sequence	Reverse Primer Sequence	Size of PCR Fragment (bp)	Size of Exon (bp)
Exon 1	5' -CCCATAGACGTTTTGGCTCA-3'	5' -AGCAACCTGTCCACACTCCT-3'	896	379
Exon 2	5' -TCCCTGCTGTGTTTTCTCTG-3'	5' -CTGGTGATTCGCACCTGTAA-3'	844	158
Exon 3	5' -TGGCTTGGAGAGATGAGAGA-3'	5' -TGGGAAACTCAGAACCCACT-3'	752	142
Exon 4	5' -CTGTCTTGTGAGGGCTGAG-3'	5' -CAGGTTGCCAAGAGTCAAGA-3'	848	204
Exon 5	5' -CTCCTAGAAAGCTCCCAAGC-3'	5' -TTCACCTCCTGTGTTCAAGC-3'	791	96
Exon 6	5' -CTCACCTGCTCACCACACTT-3'	5' -TGATGTCTCAGCACCCAGAT-3'	770	116
Exon 7	5' -ACAGGTGTGAGCCATAATGC-3'	5' -GGTATCCAGGAGAGCACGAT-3'	989	167
Exon 8	5' -AGGGACACACTCCTGTGGAT-3'	5' -ACAGAACTTCACACGGGACA-3'	1042	119
Exon 9	5' -TCAGGAGAGGAGATCCCAGT-3'	5' -CGTGTAGGAAACAGCTTGGA-3'	906	74
Exon 10	5' -TGAGGGTAAGAAATGGCAGA-3'	5' -ACAAGCTAGGCCAGAAGGAA-3'	862	52
Exon 11	5' -TCCTTCTGGCCTAGCTTGTT-3'	5' -GGTGCTGACTTGATGAATGG-3'	990	147
Exon 12	5' -ACATGCGGGTAAACTCAACA-3'	5' -AGCCTTGCTCCTGTCTTTCT-3'	1033	129
Exon 13 (1)	5' -GCTCTGCCCTCACTGTATT-3'	5' -TGTCAGTTTAGGCTGGAGGA-3'	978	
Exon 13 (2)	5' -ATGGAGTGAGCACAAGATGC-3'	5' -TAAGGGGAAAAGGGCTAACA-3'	1038	959

RefSeq of *SCARB1*: hg19, NM_005505 (CHIP Bioinformatics, University of Florida, <http://snpper.chip.org/>).