

Supplementary Table 5. Sequencing primers for *CD209*, *CLEC4M (CD209L)* and *CLEC4G (LSECtin)*

Region amplified ^a	PCR primers		Sequencing Primers ^b		Product size
	Forward (5'-3')	Reverse (5'-3')	Forward (5'-3')	Reverse (5'-3')	
<i>CD209</i>					
Promoter_1	ACTGTGTTACACCCCCTCCA	CACACATGTGGTAAATGCACA			789
Promoter_2, exon 1	CTCCCCCTTCCCAGAATC	CAAAAATGAGGACAGCAGCA			695
exon 1, exon 2	AGGTACCTGCTTCCACCTCC	TGGATGACAGATCCCTACCC	CCCAAACCTCAACCTCCTCAG	TTGGGAGGAGAGACGAGAGA	903
exon 3, exon 4	TCCAGATACTCACCCACTGC	CAGGGACCTGGGAATGTAGA	ACTTATGGAGCTGGGGACCT	GATTCCCAGGTGTTGGAGTG	1164
exon 4	CACCACACAACGACCATCTC	AGGTCCCCAGCTCCATAAGT			616
exon 5	CGGAGCATTACACATGTTC	GCTGGGCAGGTACAGGAATA			414
exon 6	CTGCTGGTGGAAATTCTAGGC	AGCAAGAGGGTGTGAGGAGA			534
exon 7_1	TCTACACCAGGGGAAATTGG	AGTCCAGCATAACACACGCAG			470
exon 7_2	TTGGTTGACTGGGGGAATTA	GCTGGAACGACGACAAATG			587
exon 7_3	TCCCAAAGTCGTGGTAGGAG	TTTGATCGGGGACTTACTGG			595
exon 7_4	TGCAACTTGGAACTCACCT	TGGGATTTGGGACTCTGAAC			592
exon 7_5	TCCCAACATCCAGTCCCTACC	TTGGCATTCTTCTCCGACT	AAGCAAGCCTGGAGTACAGC	CAGGTGAGGTTCCAAGTTGC	605
exon 7_6	CGAAAAGGTGACATTTCCAGA	GCAGCACATCCTCACAGAAA			699
exon 7_7	GCATCTTTGGGGTCAACATC	TGGCGTTTCTTTCGTTTCTC			922
<i>CLEC4M (CD209L)^c</i>					
Promoter_1	TGCCCTTGATGTTATTAGACCC	TTCCACCCTCTGATCCTCAC			662
Promoter_2, exon 1, exon 2_1	CGATCTCAGGCTCTGACTCC	GGCCATGTATCTGCTGGAAT			363
exon 2_2	AAGATCCAACAACCAAGTGGC	CTTCCCCAGGACCCAGTAAT			517
exon 3	GGGAGGTGGACGCAGGTA	TCCTGCTCTAGAGTGGGGAG			520
exon 4_1	ATGAACAGAGCCTGGAGTGG	AGCTTGGATTTCTCTGGCAA			499
exon 4_2, exon 5	GTTGCCAGAGAAATCCAAGC	GTACCTGGGACCCCTTCATT			884
exon 5, exon 6	CTGTCCAAGGACTGGACAT	CCAGGGAAAGCTCAAGTCAG			1031
exon 6, exon 7_1	CTGACTTGAGCTTCCCTGG	ACTGCCACTAAATCCGCAC			1228

exon 7_2	CAGCAAGAGAGGCAGGAACT	GCTCCTTCAACCAGCAAGTC			762
exon 7_3	GACCTGAGTGGGATGCATT	AGCCTCACAGTTCTCATGCC			1256
<i>CLEC4G (LSEctin)</i>					
Promoter_1	ATGCCTGAAATGGTGCTAGG	CCAAATTGTAACCCCTGCTG			532
Promoter_2, exon1	TGGATCAAGGTCAGGTTCAA	TTGCTGTACCTGGTGGTGTC			520
exon 1	AAGGCCAAGAAAATGGGAAG	AGGACTGTGGTGACCAGGAC			506
exon 2	GAGGGAAGAGTCGAGACAGG	GGCCCAAAGATGTACACACA			465
exon 3, exon 4, exon 5	TGCACATCGTGTGAGCTGTA	GGATTATGGCTGGGGTCAG	GGAGATGGAGTCTCGAGTGC	GTCACCTGGGAGGACTCAGG	986
exon 6, exon 7	CAGGGGTCCTGACCTTTAGC	AAAGGAGGGGACATCTGAGC			582
exon 8	TGTGCCAAAGACGACGTG	AGCCTGCCCATCCCTAAGTA			406
exon 9	ACCCCGAGTTCATCCTCTGT	GGCCAAGAGGTCACTGAGAG			701

Notes:

^a In regions encompassing introns, the corresponding introns were also sequenced.

^b These primers were used as sequencing primers in addition to PCR primers to cover the whole region.

^c Numbering of *CLEC4M (CD209L)* exons is according to NM014257