

Table S2. Primers for q-PCR amplification. Annealing temperature for all primers is 60°C.

Gene name	Forward primer (5'-3')	Reverse primer (5'-3')	Primer concentration (nM)	Amplification product (bp)
<i>DPRX</i>	GGACCTGAACCCAGGCGCAC	TGCATCTGGTCCTTGCCTTTACGA	200/200	78
<i>NALP12</i>	GCACCTTTGAGCGGATAAAC	AAGGCATGTTGACTGGTTCC	200/200	116
<i>MYADM</i>	CCACCAGATCTTTCTCCGTG	AAGATGACGTCGTGGTGGTT	200/200	96
<i>PRKCG</i>	GCCACGAATTTGTGACCTTC	TAGCTATGCAGGCGGAACTT	200/200	94
<i>CACNG7</i>	TACTATCGGGCCTCTCCTTG	GACCACCCGTAGCGATAATG	200/200	118
<i>CACNG8</i>	GAGGATCTGCTGCCTGGA	TAAGGATGGGGAAGATGCTG	200/200	140
<i>CACNG6</i>	AGTTGGAGCCGTCTGCTTT	TACTCGTAGGTGAGGCGTGG	200/200	135
<i>VSTM1</i>	TGGCAGAAGGGACGCTATGA	GCTTGGGCGGTTTCTCATT	200/200	101
<i>OLT-2</i>	CGGTGGTCCCTGCCAACAGC	GAAATTCGGCCGCGCCCTCT	200/200	138
<i>OSCAR</i>	TGATCCTCCAGCTGCTGAC	CCATGGCTTAGGGTGGTATG	200/200	101
<i>NDUFA3</i>	ACAAAGATGGCTGCGAGAGT	CTCAATGGGGGACAGAATTAC	200/200	113
<i>TFPT</i>	TCGCAGAAAGTACCAGGCACT	CTGCAGCCTCCGAGTTATCC	200/100	109
<i>PRPF31</i>	CAGCAAGCAAGCCAAAGCT	CCGGATGAACTTATGGATGATG	200/200	133
<i>LOC100288135</i>	CTGCCGACTTGCCAGGTG	CTGGTGGAGGGGGTGCCTCG	200/200	142
<i>CNOT3</i>	CGCAAATGGAACGGTTCAAAG	GTATTCGTGAGCCACTGGCCA	200/200	136
<i>LENG1</i>	CTCAGCAAGAGGCCCGTACA	AGCAGCTCCCGAAACAGGTC	200/200	127
<i>TMC4</i>	CTCATCTGCCTCCTGTCA	GCTGTAGCTGGTCAGAGCCT	200/200	90
<i>LENG4</i>	AACATGACGGTGCAGTGGT	GATGGTCAGGAAGCTCAGGT	200/100	147
<i>LENG5</i>	ATGAGGTGACTCGCTGGTTC	CACCTCCACCACCAGCAT	200/200	106
<i>RPS9</i>	CTGCTGACGCTTGATGAGAA	CAGTTTCATCTTGCCCTCAT	200/100	105
<i>LILRB3</i>	TGCTATGGTGCACACAACCT	ACAGGGAGACGGTGCATAGA	200/200	97
<i>LILRB6</i>	TCACGGTCTGATCTGTCTC	CCAGCTGATCACAGAGCCT	100/200	110
<i>LILRB5</i>	CCGCTGCACAGCTGAGTCC	GGTTGGGGAGGGTGCCTGC	200/50	122
<i>LILRB2</i>	GCAGCTGATGCCCACTCCG	GGCCTGCAGGTGTGGAGATG	200/200	236
<i>LILRA3</i>	CACTCAGCTCCAACCCTAC	GACTTGTTTTGTGGTGGCT	200/200	100
<i>LIR9</i>	TGATCTGAGTCTGCTGTGG	CAGCTGTGCAGATGGATGAG	200/200	105
<i>ILT7</i>	AACCCCTACCTGTGTCTCA	GGGCAGTCTTGAATCTGAC	200/100	106
<i>LAIR1</i>	AGCCCCATAAATGGTCTGAG	AGGTGCATGCTCATTGTGAC	200/200	98
<i>TTYH1</i>	CACCAGTTGGTGGCACTG	GAGTAGCAGGAAGAGCAGGC	200/200	108
<i>LENG8</i>	CAACTACCACCGCTTTTTC	AGGGCGGAAGGTTTGTAT	200/200	124
<i>LENG9</i>	ACAGCCCACACAGGAAAGAC	ACTGAGCACCATTGGAAGC	100/200	94
<i>LOC100288231</i>	GCAGCGTGTGCCGGAAGTC	GCGCTGACCCCTCCCGTGC	200/200	130
<i>CDC42EP5</i>	GGCTAGAGCTGGAGTCGTGA	ATCAGGCCGCTTCTTGG	200/200	105
<i>LAIR2</i>	TGTCTCCACACCTCACTGCT	GAGATGGAGGGTCTGGGAA	200/200	94
<i>LILRA2</i>	CACTCAGCTCCAACCCTAC	CTGTGTAATCCTGGGGGTGT	200/100	147
<i>LILRA1</i>	CACTCAGCTCCAACCCTAC	GACTTGTTTTGTGGTGGCT	200/200	100
<i>LILRB1</i>	GTCTCAGGACCGTCTGGGGGC	TCTGGGATCCGACCCGGTG	200/200	106
<i>LILRB4</i>	CTGCCGTGAAGAACACACAG	CACCTTGGCATACTGACTG	100/200	98
<i>KIR3DL3</i>	ACAGATGCTTCGGCTCTTTC	ACGTGCAGGTGTCTGGAGTT	200/200	100
<i>KIR2DL3</i>	AACTCGAGTGACCCACTGCT	AACATGCAGGTGTATGGGGT	200/200	108
<i>KIR2DL1</i>	CCCCTGCTTGTCTGTCA	CCAATCAGAATGTGCAGGTG	200/200	105
<i>KIR2DL4</i>	ACCCACTGCCTGTTCTGTG	ATCACAGCATGCAGGTGTCT	200/200	103
<i>KIR3DL1</i>	ATGTTGCTCATGGTCGTCAG	AGACAGGAAGGGCTTGCC	200/200	96
<i>KIR2DS4</i>	AAATAGTTGGCCTTACCCA	GGAGGATGGTGAAAGGGATT	200/200	107
<i>KIR3DL2</i>	AGGTCCAATCAGAACATGC	AGGTCCAATCAGAACATGC	200/200	95
<i>FCAR</i>	AAACAGACCACCTCCTGTG	ATCACAGGACTCGATTTGGC	200/200	107
<i>NCR1</i>	CCACAGCCCACCGAGGGACA	GTGTCTGCAGGAAAGTGGGGTC	200/200	148
<i>NALP7</i>	GGAAGCAAGACCTGACCTGA	CTCGTTCAGCTGCTCCAGA	200/200	121
<i>NALP2</i>	ATGCACCGAATGGATCTGTC	CGTTCCTCCGTGTTATCCC	200/200	110
<i>GP6</i>	AGCTTGTGGTCACAGGAACC	ATGAGACGGTCAGTTCAGCG	200/200	105