

Table S2. Sequences of oligonucleotide primers used for conventional and quantitative RT-PCR studies.

Gene name	GenBank Accession N ^o	Sequence 5'-3'	Product Size (bp)	Position (Exons)
Conventional PCR				
<i>ACTB</i> *	NM_001101	Forward : GCGGGAAATCGTGCGTGCGTGACATT Reverse : GATGGAGTTGAAGGTAGTTTCGTG	232	615-846
<i>KIR3DL2</i>	NM_006737	Forward : CGGTCCCTTGATGCCTGT Reverse : GACCACACGCAGGGCAG	369	546-914 (4-5)
		Forward : GAAGACAGAAGCCACGTTCCCATC T Reverse : GTTTGACCACACGCAGGGCAG	732	187-918 (3-5)
		Forward : CAACTTCTCCATCGGTCCCTTGATG Reverse : GTCTCCCTAGAAAACCCCTCAAGA	841	534-1374 (4-9)
		Forward : CGGGGGACAGAACAGTGAATAGGC Reverse : GTCTCCCTAGAAAACCCCTCAAGA	242	1133-1374 (8-9)
		Forward : GAAGACAGAAGCCACGTTCCCATCT Reverse : GTCTCCCTAGAAAACCCCTCAAGA	1188	187-1374 (3-9)
<i>KIR3DL1</i>	NM_013289	Forward : ACATCGTGGTCACAGGTCC Reverse : TCGTATGTCACCTCCTC	557	641-1197
Quantitative PCR				
<i>SF3A1</i>	NM_005877	Forward : TGCAGGATAAGACGGAATGGAAACTGA Reverse : GTAGTAAGCCAGTGAGTTGGAATCTTTG	179	2141-2319
<i>CD3D</i> †	NM_000732	Forward : GGTACTGGCTACCCTTCTCTCGCA Reverse : CTTGTAATGTCTGAGAGCAGTGTGCCAC	141	30-270
<i>KIR3DL2</i>	NM_006737	Forward : CAACTTCTCCATCGGTCCCTTGATG Reverse : GTTTGACCACACGCAGGGCAG	385	534-918
<i>KIR3DL1</i>	NM_013289	Forward : GGACATCGTGGTCACAGGTCC Reverse : GCCTGGAATGTTCTGTTGACCTTGC	195	639-833
<i>PLS3</i> ‡	NM_005032	Forward : CCAGTTTGGCAGTTGTGGATTTAATTGAT Reverse : ACAAGGTCTTCAGGGAGAGCATAAC	172	1652-1823

Nucleotide positions of amplified fragments were calculated from the start codon according to the sequences previously reported.

bp, base pair.

*The *ACTB* gene encodes for β -actin

†The *CD3D* gene encodes for the delta chain of CD3

‡The *PLS3* gene encodes for T-plastin