

Supplementary Table S1. Sequence of primers used for RT-PCR studies.

Primer name	Accession Nr.	Sequence_Fw	Sequence_Rv
<i>A20</i>	NM_009397	AAGCTCGTGGCTCTGAAAAC	TTCCTCAGGACCAGGTCAGT
<i>Ahr</i>	NM_013464	CTGGTTGTCACAGCAGATGCCT	CGGTCTTCTGTATGGATGAGCTC
<i>Areg</i>	NM_009704	GCAGATACATCGAGAACCTGGAG	CCTTGTCATCCTCGCTGTGAGT
<i>Armcx3</i>	NM_027870	GTGAATGCCGAAAATCAGCGCAG	GGTAAGCAATCTGAGTCCAGCC
<i>Ccl2</i>	NM_011333	GCTACAAGAGGATCACCAGCA	GTCTGGACCCATTCTTCTTG
<i>Ccl5</i>	NM_013653	CCACTTCTTCTCTGGGTTGG	GTGCCACGTCAAGGAGTAT
<i>Cdkn1a</i>	NM_007669	GCAGAATAAAAGTGCCACAGG	AAAGTTCCACGTTCTCGGG
<i>Cdkn2a</i>	NM_001040654	TGTTGAGGCTAGAGAGGATCTTG	CGAATCTGCACCGTAGTTGAGC
<i>Ctnnb1</i>	NM_001165902	GTTTCGCTTCATTATGGACTGCC	ATAGCACCTGTTCCCGCAAAG
<i>Cxcl1</i>	NM_008176	ACCCAAACCGAAGTCATAGCC	CTCCGTTACTTGGGGACACC
<i>Cyld</i>	NM_001128170	CACCTACTGGGAAGAACGGA	ACGGTCTTGATGTACTGTCCTATA
<i>Dok1</i>	NM_001291799	TTTTCTGCCTTGAGATGCT	GCTCCAGGATTTGACTCTGC
<i>Dok2</i>	NM_010071	ATGGTCAGGATGGAGGAGC	ATATAACACGGCTGCGAACC
<i>Dsp</i>	NM_023842	TACACCTCAGGGCTGGAACTC	GTAGTCTCCAGACCTCGTAAGC
<i>Duba</i>	NM_138604	AGCGGGCTACAACAGTGAAG	AAGGCCTTTTCAAACCAAGT
<i>Erbin</i>	NM_001289473	GCCCTGAGACACCCTGAGA	CAACCGCACAAACAACTTC
<i>Errfi1</i>	NM_133753	GCCTACAATCTGAACCTCCCTG	CCTTGGAGATGGACCACACTCT
<i>Gdf15</i>	NM_011819	AGCCGAGAGGACTCGAACTCA	GGTTGACGCGGAGTAGCAGCT
<i>Glb1</i>	NM_009752	CAAGACAGTGGCTGAAGCTCTG	GAGGAAGCGTTGTTCCGTACAG
<i>Il6</i>	NM_031168	TGATGCACTTGACAGAAAACA	ACCAGAGGAAATTTCAATAGGC
<i>Il10</i>	NM_010548	ACAGCCGGGAAGACAATAACT	CCTGCATTAAGGAGTCGGTTA
<i>iNos</i>	NM_010927	TTCTGTGCTGTCCAGTGAG	TGAAGAAAACCCCTTGTGCT
<i>IrakM</i>	NM_028679	CACTGCTGGGAGAGCTTTG	CCAGCCAGCTGTTTGAAAGT
<i>Irf4</i>	NM_013674	TGCAAGCTCTTTGACACACA	CAAAGCACAGAGTCACCTGG
<i>Lancl1</i>	NM_021295	CTCCTGAGTTCTCACATCGCTTG	GGTGGAGGTAAAGCACAGCGAT
<i>Ly6d</i>	NM_010742	CATGCTTTAGCCATGATGGAGGC	ATGTCATCAGCTCTTGGTGCCC
<i>Mcpip1</i>	NM_153159	CCTGTGGTCATCGACGGAAG	GAAGGATGTGCTGGTCTGTGATA
<i>Muc1</i>	NM_013605	AGTGCTCTGACGTGAAGTCAC	GGGAGGGAACTGCATCTCATTC
<i>Nox4</i>	NM_015760	CGGGATTGCTACTGCCTCCAT	GTGACTCCTCAAATGGGCTTCC
<i>Otud7b</i>	NM_001025613	GGAGGTGAAGTTACATCTGCTGC	TCAGGAGTGGACCTGGGTTTCAT
<i>Pias4</i>	NM_021501	CCAACCGCATTACTGTCACCTG	CGTCTTCAACCTCTGTAGCAGG
<i>Prodh</i>	NM_011172	ATCGCATCCAGGGCTGAGATTG	GTCAATGAGGCTGTCCAGTCC
<i>S100a9</i>	NM_001281852	TGGTGGAAGCACAGTTGGCAAC	CAGCATCATACACTCTCAAAGC
<i>Ship</i>	NM_010566	GCTGTTCCGGAATTGTGTTT	GTGAAGAACCTCATGGGGAC
<i>Sigirr</i>	NM_023059	AGAGGTCCAGAAAGAGCCAT	ACACTGGGGCTAGAGAACA
<i>Sike1</i>	NM_025679	TTCAGGTGGACGATAACCAA	GAGATTCACTGCTGATGGACAG
<i>Slc9a3r1</i>	NM_012030	ATCTGCCTCCAGCGATACCAGT	AGCCAAGGAGATGTTGAGGTCC
<i>Slc22a6</i>	NM_008766	GAAAGGCTGTCTGGCTTCTCT	ATCAGTGGGCTCACTATGCTGC
<i>Slc22a8</i>	NM_031194	AGGCTCTGAAGACACTCCAACG	ACCTTGGCTGAGGTGATGTCCT
<i>Smurf2</i>	NM_025481	CCAATGCCATCAACCGCCTCAA	GTGCCTATTCCGGTCTCTGGACT
<i>Snx14</i>	NM_172926	GTGGCTGAAGGCGAAGATGACT	CGAGGAGTGTTAGGTGTGCTCA
<i>Socs1</i>	NM_001271603	ACTTCTGGCTGGAGACCTCA	ACAAGCTGCTACAACCAG
<i>Socs3</i>	NM_007707	AAGGCCGGAGATTTCGCT	AACTTGCTGTGGGTGACCAT
<i>Stx4a</i>	NM_009294	GTCCAGCAATTTGTCGAGCTC	TCCTCGTCAGACACCATTCAG
<i>Tank</i>	NM_001164071	GCTTCCAGAATGGGTACGTG	TGGTAGGAATGCCAGCTCTC
<i>Tgfb</i>	NM_011577	GGAGAGCCCTGGATACCAAC	CAACCCAGGTCCTTCTCTAAA
<i>Tnfa</i>	NM_013693	AGGGTCTGGGCCATAGAAGT	CCACCAGCTCTTCTGTCTAC
<i>Tollip</i>	NM_001347562	GCGGGTCTCTGTGCAGTT	TGTGGGTGTTATACGGAGGAA
<i>Trim32</i>	NM_001161782	GAAACAGCGGAGTTCTGAGGCA	AACCAGCACCTCACTCTGACTG

<i>Tslp</i>	NM_021367	GCAAATCGAGGACTGTGAGAGC	TGAGGGCTTCTCTTGTCTCCG
<i>Twist1</i>	NM_011658	GATTCAGACCCTCAAAGTGGCG	AGACGGAGAAGGCGTAGCTGAG
<i>Vamp3</i>	NM_009498	AGACCAGAAGCTCTCGGAGCTA	ACCAGGACACTGATCCCTATCG
<i>Zranb1</i>	NM_207302	GGCTATTCCTCAATGCCTGTGTG	AGAAGGACGGTTCAGCAAGCGT
<i>hA20</i>	NM_001270507	GGACTTTGCGAAAGGATCG	TCACAGCTTTCGCATATTG
<i>hCCL2</i>	NM_002982	AGAATCACCAGCAGCAAGTGTCC	TCCTGAACCACTTCTGCTTGG
<i>hCCL5</i>	NM_002985	CCTGCTGCTTTGCCTACATTGC	ACACACTTGGCGGTTCTTTCGG
<i>hCXCL1</i>	NM_001511	AGCTTGCCTCAATCCTGCATCC	TCCTTCAGGAACAGCCACCAGT
<i>hIL-6</i>	NM_000600	AGTGAGGAACAAGCCAGAGC	GTCAGGGGTGGTTATTGCAT
<i>hINOS</i>	NM_000625	GCTCTACACCTCCAATGTGACC	CTGCCGAGATTGAGCCTCATG
<i>hMCPIP1</i>	NM_025079	AAGAGGAAAAGGAGGGGCAGC	ATGTGCTGGTCTGTGATGGG
<i>hTNFa</i>	NM_000594	CTCTTCTGCCTGCTGCACTTTG	ATGGGCTACAGGCTTGCTACTC