

**Supplementary Table 1**

<b>Gene</b>	<b>Forward Primer (5' to 3')</b>	<b>Reverse Primer (5' to 3')</b>
<i>Spi1</i>	ATTGCGCTGTACCAGTTCCTGC	TGGACGAGAACTGGAAGGTACC
<i>Spib</i>	CAAGCGCATGACGTATCAGAA	GCTGTCAAACCTGGTAGGTGAG
<i>Junb</i>	CTGAAACCCACCTTGCGCG	CCGAAAAGTAGCTGCCTGCC
<i>Il7r</i>	GCGGACGATCACTCCTTCTG	AGCCCCACATATTTGAAATTCCA
<i>Jchain</i>	GAACTTTGTATACCATTTGTGACAGACG	CTGGGTGGCAGTAACAACCT
<i>H2-ob</i>	AGGCGGACTGTTACTTCACC	ATCCAGGCGTTTGTTCCTACTG
<i>Cd37</i>	GTCCTTTGTGGGTTTGTCTT	GAGACAGCGCAGCTCCTTTAG
<i>Tlr2</i>	TCTGGGCAGTCTTGAACATTT	AGAGTCAGGTGATGGATGTCG
<i>Cxcr4</i>	AGGAAACTGCTGGCTGAAAAGG	GGAATTGAAACACCACCATCCA
<i>Ccl5</i>	TGCCACGTCAAGGAGTATTTT	AACCCACTTCTTCTCTGGGTTG
<i>Cdkn2a</i>	GCCCAACGCCCCGAACTCTTTC	GCGACGTTCCAGCGGTACACA
<i>Batf</i>	CCAGAAGAGCCGACAGAGAC	GAGCTGCGTTCTGTTTCTCC
<i>Ces1d</i>	GGAGAGTCAGCAGGAGGTTTC	GAGGGACACCCACTCTCAG
<i>Abcd3</i>	GGCCTGCACGGTAAGAAAAGT	CCGCAATAAGTAACAAGTAGCCT
<i>Acsm2</i>	ACTAATACCCATACAGTGGGGC	CTGAAGATCTCTTGCCAGCCTTC
<i>Ehhadh</i>	TGGCTCTAACCGTATGGTCC	CTATGATCCGCCTCTGCAA
<i>Me1</i>	AGTATCCATGACAAAGGGCAC	ATCCATTACAGCCAAGGTC
<i>Acot12</i>	CCACCACCTTGGAGAAGATAAA	GTGAGGTCAGACAAGAGATGATAAG
<i>Ugt2b38</i>	TGCGCCACAAAAGGGCTAA	ACACAAGAGAGTAGGAAGCCG
<i>Fmo2</i>	ACTCAGAGCAACGGAAAGGA	CCTGGGAATGACTTGAGTGG
<i>Nr1h4</i>	GCAGGGAGAAAACGGAAC	TCTGTACATGACTGGTTGCC
<i>Acsm5</i>	CAGGTTCCAATAACCACCT	CCGTCAAACAGTGCCTCAG
<i>Crot</i>	GAACGGACATTTCAGTACCAGG	CTTCATTTGCGAATGGTTTCACT
<i>Coq10b</i>	GAGGACTATCAGCATTTTCGTTCC	AGTGCGGTTTTACCAAGGTTAC
<i>c-Jun</i>	GCCTACGGCTACAGTAACCC	GAGAAGGTCCGAGTTCTTGG
<i>c-fos</i>	GGGGACAGCCTTTCCTA	CTGTCACCGTGGGGATAAAG
<i>Jund</i>	GAAACGCCCTTCTATGGCGA	CAGCGCGTCTTTCTTCAGC
<i>Il-6</i>	GTTCTCTGGGAAATCGTGGA	GGTACTCCAGAAGACCAGAGGA
<i>Gapdh</i>	CTTTGTCAAGCTCATTTCTCTGG	TCTTGCTCAGTGTCTTGC