

SUPPLEMENTAL MATERIAL

Stock et al., <http://www.jem.org/cgi/content/full/jem.20151853/DC1>

Table S1. Primer sequences for real-time qPCR

Gene	Forward primer (5'-3')	Reverse primer (5'-3')	Amplicon
Gapdh	CCAGGTTGTCTCCTGCGACTT	CCTGTTGCTGTAGCCGTATTCA	133
IL-1 $\alpha$	AACGTCAAGCAACGGGAAG	AAGGTGCTGATCTGGGTGG	125
IL-1 $\beta$	GCTACCTGTGCTTTCCCGT	ATCTCGGAGCCTGTAGTGC	163
IL-3	GGTCTTGCCAGCTCTACCA	GTATCCCGCCACTGATTGA	99
IL-4	ATGGATGTGCCAAACGCTCT	AAGCACCTTGAAGCCCTAC	78
IL-5	AGCAATGAGACGATGAGGCT	AGCATTTCACAGTACCCCC	125
IL-6	AGAGTTGTGCAATGGCAATT	TCATGTACTCCAGGTAGCTATGG	182
IL-10	TAAGGTGGCCACACTTGAG	GTTTTAGGGATGAAGCGG	209
IL-17 $\alpha$	TCATCCCTCAAAGCTCAGCG	TTCATTGCGGTGGAGATCC	113
IL-17f	GAAGTGACCCCGTAAACAG	TTTGGGGTTCTTCCGAGCTG	97
IL-18	GGCTGCCATGTCAGAAGACT	CAGTCTGGTCTGGGGTTCAC	241
IL-22	TTGACACTTGTGCGATCTCT	CCTTAGCACTGACTCCTCGG	280
IL-33	AACTTCTCTGCTATCCACG	TCATTGTATGTACTCAGGGAGCC	143
TNF $\alpha$	ACTGAACTTCGGGGTGATCG	TGATCTGAGTGTGAGGGTCTGG	97
Cox2	CTTCGGGAGCACAACAGAGT	AAGTGGTAACCGCTCAGGTG	155
5L0	CTACGATGTCACCGTGGATG	GTGCTGCTTGAAGATGTGAA	235
FLAP	TGAAGCAAGCATGGATCAAG	AAGTGGGTACGCATCTACG	209
LT4AH	CCTTGGGGCACATAAAGCGA	TGAATGCAGAGCCGTAACCA	94
LTC4S	GCTCTTCTGGCTACCGTCAC	CAGGTAGAACAGTCCGCACA	246
VEGF $\alpha$	ACGATGAAGCCCTGGAGTGC	TGTGCTGTAGGAAGCTCATCTC	114
MMP2	ACAAGTATGAGAGTGCACCAG	GTGTAGATCGGGGCCATCAG	205
MMP9	TAAGGACGGCAAATTTGGTT	CTTTAGTGGTGCAGGCAGAG	128
IFN- $\alpha$	ACCTCCACCAGCAGCTCAATGACC	AGCACATTGGCAGAGGAAGACAG	214
IFN- $\beta$	CGTGGGAGATGTCCTCAACT	AGATCTCTGCTCGGACCACC	117
IFN- $\gamma$	TCTTGATATCTGGAGAACTGG	AGCTCATTGAATGCTTGGC	253
M-CSF	GATGTGGTGACCAAGCCTGA	AATCATCCAAGCCAAGCCA	133
G-CSF	AGTGAGGAAGATCCAGGCCAGC	AGAGCAGCCACTCAGGGAAGC	136
GM-CSF	CCAGCTCTGAATCCAGTCTCTC	TCTCTCGTTTGTCTTCCGCTGT	172
Ccl1	TGCTGCTGGCTGCCGTGTGG	TTGAGGCGCAGCTTTCTCTACC	198
Ccl2	TCTTTCTCCACCACCATTGC	AGCTTCTTTGGGACACCTGCTG	206
Ccl3	TCTCCACCACATGCCCTTGC	AGGCAGTCGGGGTGTGAGC	96
Ccl4	ACACCTCCCGGAGCTTCCAC	AGAAGCATCAGGGCTGAGGAGG	231
Ccl5	CCTCGTGCCACGTCGAAGG	TGGCACACACTTGGCGGTTCC	97
Ccl7	TCTGCCACGCTTCTGTGCC	AACAGCTTCCAGGGACACCG	178
Ccl17	TGCTGCCTGGATTACTTCAAAGG	AGCCCTGGCTTCTCAAATGC	220
Ccl22	ACTACATCCGTACCCTCTGC	TAGCTTCTTACCAGACCTGC	146
Cxcl1	ATCCAGAGCTTGAAGGTGTTGC	AAGGCAAGCCTCGCGACC	90
Cxcl2	AGTTTGCCTTGACCTGAAGC	AGGCTCCTCTTCCAGG	94
Cxcl3	TCACCACAGCCCTTCGCAC	TCACCGTCAAGCTCTGGATGG	229
Cxcl4	GTCGCTGGGTGTTTCGAG	ATAGGGGTGCTTCCCGTCC	277
Cxcl5	TTGGAGGTGATCCCTGCAGGTC	ATCACAGGAGCTTCTGGATCCAG	101
Cxcl9	TGAAGTCCGCTGTTCTTTTCC	AGTGATCGTGCCTCGGCTG	123
Cxcl10	GTCCTAATTGCCCTTGGT	TCTTGCTTCGGCAGTTAC	119
Cxcl12	ATCGCCAGAGCCAACGTCAAGC	TGCCCTTGCATCTCCACG	226
Ccr1	TCAACTCCATCCAACCCCTCC	TGCTCACCAAGTGGCCAATTCC	145
Ccr2	TCAGTGCCTGCAAAGACCAG	ACGGTGTGGTGGCCCTTC	201
Ccr3	ATGAGTGGGACCACCCTGTG	CATGTTGCCAGGAGGCCGA	107
Ccr4	TACCCGAGCGGACGATTTC	AAGGAGTAGAGAGGAGGAGG	168
Ccr5	TGAGACATCCGTTCCCTTAC	TGCTGACATACCATAATCGATG	103
Cxcr2	TACCTGGCCATTGTACATGC	TCAAACGGGATGTATTGTTACC	190
Cxcr3	AGCCGGAGCACCAGCCAAGC	TGTAGAGGGCTGGCAGGAAGG	199
Cxcr4	TGAGGCGTTTGGTGCTCC	ATGAAGTAGATGTTGGGACGG	197
human Gapdh	TGCACCACCAACTGCTTAGC	GGCATGGACTGTGGTCATGAG	87
human GM-CSF	GCCAGCCACTACAAGCAGCAC	CAAAGGGGATGACAAGCAGAAG	118