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| ***Table S2.******Nucleotide sequences, base pair length, and accession number used to design primers for quantitative real-time RT-PCR.*** |
| Primer | Sequence | Length | Accession number |
| mβ-actin F | 5’-CCCAGGCATTGCTGACAGG-3’ | 141 | X03672 |
| mβ-actin R | 5’-TGGAAGGTGGACAGTGAGGC-3’ |  |  |
| mIL-1β F | 5'-TATCACTCATTGTGGCTG-3' | 76 | NM\_008361.3 |
| mIL-1β R | 5'-ATGTCCTCATCCTGGAAG-3' |  |  |
| mIL-17a F | 5'-CCAGACGGCCCTCAGATTAC-3' | 103 | NM\_010552.3 |
| mIL-17a R | 5'-CACTTGGCCTCCCAGATCAC-3' |  |  |
| mIL-6 F | 5'-TTTCCTCTGGTCTTCTGGAG-3' | 112 | NM\_031168.1 |
| mIL-6 R | 5'-CTGAAGGACTCTGGCTTTGT-3' |  |  |
| mMCP-1 F | 5'-CTGCCTAATCCACAGACTG-3' | 142 | AJ238892 |
| mMCP-1 R | 5'-GCCTGAACAGCACCACTA-3' |  |  |
| mTNF-α F | 5’-CGTGCTCCTCACCCACAC-3’ | 133 | M13049 |
| mTNF-α R | 5’-GGGTTCATACCAGGGTTTGA-3’ |  |  |
| mTGF-β F | 5′CAACTTCTGTCTGGGACCCT3 | 150 | NM\_011577 |
| mTGF-β R | 5′TAGTAGACGATGGGCAGTGG3′ |  |  |
| aap F | CTT GGG TAT CAG CCT GAA TG | 310 | FN554767.1 |
| aap R | AAC CCA TTC GGT TAG AGC AC |  |  |
| mIL-12p35 F | CAGATAGGAAACAAAGAAAGAT | 129 | NM\_001159424 |
| mIL-12p35 R | GGAGATGAGATGTGATGG |  |  |
| mIL-4 F | GCTATTGATGGGTCTCAACC | 112 | NM\_021283.1 |
| mIL-4 R | GCCGATGATCTCTCTCAAGT |  |  |
| mIL-10 F | GGGCTTCTTTCTAAATAGTTC | 135 | NM\_010548.1 |
| mIL-10 R | GATCTTAGCTAACGGAAACA |  |  |
| CCL20 F | CAGCACTGAGCACATCTACT | 128 | NM\_016960.2 |
| CCL20 R | CAGCAGCACCGACATCAA |  |  |
| S100A8 F | CGACACCTTCCATCAATACTCT | 92 | NM\_009114 |
| S100A8 R | TTGCCAACTGTGCTTCCA |  |  |
| S100A9 F | GACATCAATAGTGACAAT | 143 | NM\_013650 |
| S100A9 R | TATTCTGTAGACATATCCA |  |  |
| CXCL1 F | GTCATAGCCACACTCAAG | 93 | NM\_008176 |
| CLCL1 R | GACACCTTTTAGCATCTTTTG |  |  |