**Supplementary Table 1.** Primers used in the qRT-PCR analyses.

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| --- | --- | --- | --- |
| **Target Gene** | **Primer Sequences** **(5’-3’)** | **GenBank Number** | **Product Length (bp)** |
| **Human** |   |   |   |
| *INOS* | F- ATGCCCGATGGCACCATCAGAR- TCTCCAGGCCCATCCTCCTGC | NM\_000625.4 | 394 |
| *IL-12* | F- GCCACAAAAATCCTCCCTTGR- CTCCCTAGTTCTTAATCCAC | NM\_001354583.2 | 217 |
| *CCR7* | F- ACAGCCTTCCTGTGTGGTTTR- ATGATGGAGTACATGATAGG | NM\_001301714.2 | 191 |
| *CXCL10* | F- GAAATTATTCCTGCAAGCCAATTTR- TCACCCTTCTTTTTCATGTAGCA | NM\_001565.4 | 67 |
| *TNFα* | F- GCCCCAATCCCTTTATTACCR- CACATTCCTGAATCCCAGGT | NM\_000594.4 | 145 |
| *IL-1β* | F- CAGTGGCAATGAGGATGACTTGR- AGTGGTGGTCGGAGATTCGT | NM\_000576.3 | 125 |
| *ARG1* | F- GTTTCTCAAGCAGACCAGCCR- GCTCAAGTGCAGCAAAGAGA | NM\_000045.4 | 149 |
| *IL-10* | F- GAGGCTACGGCGCTGTCAR- TCCACGGCCTTGCTCTTG | NM\_001382624.1 | 57 |
| *SPHK1* | F- CATCCAGAAGCCCCTGTGTAGR- GTCTTCATTGGTGACCTGCTCAT | NM\_001142601.2 | 100 |
| *YM-1* | F- CCAGTGCTGCTCTGCATACAR- ATGCCGTAGAGCGTCACATC | NM\_001276.4 | 189 |
| *RELMα* | F- CGTCCTCTTGCCTCCTTCTCR- ACAAGCACAGCCAGTGACAG | NM\_032579.3 | 227 |
| *MMP9* | F- GTGGCGGCGCATGAGTR- AGGTGCCGGATGCCATT | NM\_004994.3 | 134 |
| *ZO-1* | F- AGAAGGATGTTTATCGTCGCATTR- CCAAGAGCCCAGTTTTCCAT | NM\_001330239.4 | 164 |
| *IL-4* | F- CCGTAACAGACATCTTTGCTGCCR- GAGTGTCCTTCTCATGGTGGCT | NM\_001354990.2 | 209 |
| *TSG-6* | F- TCATGTCTGTGCTGCTGGATGR- GGGCCCTGGCTTCACAA | NM\_007115.4 | 67 |
| *GAPDH* | F- ACAGTCAGCCGCATCTTCTTR- GACAAGCTTCCCGTTCTCAG | NM\_001357943.2 | 205 |
| **Mouse** |   |   |   |
| *INOS* | F- CCAAGCCCTCACCTACTTCCR- CTCTGAGGGCTGACACAAGG | NM\_001313922.1 | 127 |
| *IL-12* | F- AGTGACATGTGGAATGGCGTR- CAGTTCAATGGGCAGGGTCT | NM\_001303244.1 | 133 |
| *CCR7* | F- AAAGCACAGCCTTCCTGTGTR- AGTCCACCGTGGTATTCTCG | NM\_001301713.1 | 115 |
| *IL-6* | F- CCTTCCTACCCCAATTTCCAAR- AGATGAATTGGATGGTCTTGGTC | NM\_031168.2 | 89 |
| *TNFα* | F- ACGGCATGGATCTCAAAGACR- AGATAGCAAATCGGCTGACG | NM\_001278601.1 | 138 |
| *IL-1β* | F- AGTTGACGGACCCCAAAAGR- AGCTGGATGCTCTCATCAGG | NM\_008361.4 | 75 |
| *ARG1* | F- CAGAAGAATGGAAGAGTCAGR- CAGATATGCAGGGAGTCACC | NM\_007482.3 | 250 |
| *IL-10* | F- CCAGTTTTACCTGGTAGAAGR- TGTCTAGGTCCTGGAGTCCA | NM\_010548.2 | 324 |
| *CD163* | F- CCTGGATCATCTGTGACAACAR- TCCACACGTCCAGAACAGTC | NM\_001170395.1 | 67 |
| *YM-1* | F- AGAAGGGAGTTTCAAACCTGGTR- GTCTTGCTCATGTGTGTAAGTGA | NM\_009892.3 | 109 |
| *GAPDH* | F- ACAGTCAGCCGCATCTTCTTR- GACAAGCTTCCCGTTCTCAG | NM\_008084.3 | 248 |