

Supplemental Table 1. Primer sequences

| Gene | Refseq ID | Forward Primer | Reverse Primer | Probe (UPL) | Probe | Control (PCR fragment) |
|--------------|-----------|------------------------------|-----------------------------|-------------|----------|------------------------|
| IL-1 β | NM_008361 | 142/tgacggaccctcaaaagat | 216/gaagctggatgctctcatctg | # 26 | ctgggctg | 98-598 (501 bp) |
| IL-6 | NM_031168 | 327/gctaccaactggatataatcagga | 404/ccaggtagctatggactccagaa | # 6 | ttctctg | 65-542 (478 bp) |
| TNF α | NM_013693 | 428/ctgtagcccacgtcgtagc | 524/ttgatccatgccgttg | # 25 | tggaggag | 281-699 (419 bp) |
| IL-10 | NM_010548 | 184/cagagccacatgctcctaga | 262/tgtccagctgtcctttgtt | # 41 | cttcagcc | 92-497 (406 bp) |
| CCL2 | NM_011333 | 142/catccacgtgttgctca | 217/gatcatctgctggatgagt | # 62 | acctgctg | 36-451 (416 bp) |
| TGF β | NM_011577 | 1358/tggagcaacatgtggaactc | 1428/cagcagccggttaccag | # 72 | ttctgctg | 1054-1599 (546 bp) |
| CD11c | NM_021334 | 1725/atggagcctcaagacaggac | 1787/ggatctgggatgctgaaatc | # 20 | ccagccag | 1554-2145 (592 bp) |
| LYVE-1 | NM_053247 | 758/gaagcagctgggtttgga | 847/gtagcaaacagccagcacag | # 91 | ctctctc | 623-1092 (470 bp) |
| ApoE. | NM_009696 | 370/cccaagtacacaagaactgac | 465/ccactggaccagctgtt | # 17 | aggagctg | 186-617 (432 bp) |