**Supplemental Table S1.** Primers used for qPCR. Nucleotide positions in transcripts are shown in parentheses for each primer. mRNA sequences (gene symbols are given with accession numbers in parentheses) were from the Rat Genome Database viewed at <http://www.ncbi.nlm.nih.gov/entrez>.

|  |  |  |  |
| --- | --- | --- | --- |
| Gene symbol(accession no.) | Size (bp) | Forward primer | Reverse primer |
| Csf1 (NM\_023981.4) | 150 | ACCAGGAACAGCTGGACGAT (369-388) | TGTTCAGTTTCATAGAGAGTTCCTGTAG (491-518) |
| Csf2 (XM\_340799.3) | 90 | GCGTGCTCTGGAGAACGAA (138-156) | AGGCGGGTCTGCACACAT (210-227) |
| Csf3 (NM\_017104.1) | 82 | AGCACCATGCCAATCTTCACTT (513-534) | CCAGGAAGCTCTGCAGGTATG (574-594) |
| Ptgs2 (NM\_017232.3) | 90 | GAAGAACTTACAGGAGAGAAAGAAATGG (1393-1420) | CAGCAGGGCGGGATACAGT (1464-1482) |
| Il1a (NM\_017019.1) | 95 | AGCCCATGATTTAGAAGAGACCAT (370-393) | CCTGCTTGACGATCCTTATCAAT (442-464) |
| Il1b (NM\_031512.2) | 88 | TCGTGCTGTCTGACCCATGT (457-476) | ACAAAGCTCATGGAGAATACCACTT (520-544) |
| Gapdh (NM\_017008.3) | 93 | CCAAGGTCATCCATGACAACTT(476-497) | AGGGGCCATCCACAGTCTT(550-568) |