Supplemental Table

Supplemental Table S1: PCR primer sequence

Gene Name	GenBank Accession Number	Primer Sequence	Amplicon Size
Actb	NM_007393	(s) AGGCCAACCGTGAAAAGATG (a) ACCAGAGGCATACAGGGACA	447-547 (101 bp)
Caveolin1	NM_007616	(s) CGTAGACTCCGAGGGACATC (a) CGTCATACACTTGCTTCTCAGTC	13-130 (113 bp)
Caveolin2	NM_016900	(s) GGATCCTCACCAGCTCAACTC (a) GGCTGCAGATCCACACTTTG	120-220 (101 bp)
P2rx1	NM_008771	(s) GAAAGCCCAAGGTATTCGCAC (a) TCCACCTCTACAGGACACCAG	587-679 (93 bp)
P2rx2	NM_153400	(s) ACCATGTCGGAACACAAAGTG (a) GCTGTGAACCCTCATGCTCTC	339-482 (144 bp)
P2rx3	NM_145526	(s) ACCTCACCGACAAGGACATAA (a) GTCCCCTACCCTCAAGATGG	1116-1195 (80 bp)
P2rx4	NM_011026	(s) AGTGGAGAATGACGCTGGTG (a) CTTGAGGTAGGAGGTGGTAATGTTG	608-755 (148 bp)
P2rx5	NM_033321	(s) ACCAACCTGATCGTGACTCCT (a) ACAGTCGGTGTCCTCTGAACA	522-614 (93 bp)
P2rx6	NM_011028	(s) GTAGTCTACGTGATAGGGTGGG (a) CACGAGGAAGTTGGTTACCAG	283-489 (207 bp)
P2rx7	NM_011027	(s) AGCAGAGGTGACGGAGAATG (a) GCACTTGGCCTTCTGACTTG	367-509 (143 bp)
Nlrp3	NM_145827	(s) ATTACCCGCCCGAGAAAGG (a) TCGCAGCAAAGATCCACACAG	312-452 (141 bp)
Casp1	NM_009807	(s) ACAAGGCACGGGACCTATG (a) TCCCAGTCAGTCCTGGAAATG	355-591 (237 bp)
Il1b	NM_008361	(s) GCAGCAGCACATCAACAAGAG (a) CCACGGGAAAGACACAGGTAG	243-313 (71 bp)
I16	NM_031168	(s) GGATGTCTGTAGCTCATTCTGCTC (a) CATCAGTCCCAAGAAGGCAAC	15-132 (118 bp)
Tnf	NM_013693	(s) AAGCCTGTAGCCCACGTCGTA (a) GGCACCACTAGTTGGTTGTCTTT	435-556 (122 bp)
Gsdmd	NM_026960	(s) ATCCTGGCATTCCGAGTGG (a) CTCTGGCCCACTGCTTTTCT	787-908 (122 bp)