

Table S1. Primer sequences for real-time quantitative PCR.

Gene	Gene ID	FORWARD	REVERSE
<i>Actb</i>	NM_031144.3	GGTCAGGTCATCACTATCGGCAATG	CAGCACTGTGTTGGCATAGAGGTC
<i>Adh1</i>	NM_019286.4	TGTTGGTCTGTCTGTCGTCATTGG	CCTGGATGGGTTTGGTGTAGTCTTG
<i>Aldh2</i>	NM_022407.3	CCTCCTGGCGTGGTGAACATTG	CAGCGTGACCCTCTTCAGATTGC
<i>Cyp2e1</i>	NM_031543.2	CAATCTTGTCCTTCCAACCTACCC	CTGGATCTGGAAACTCATGGCTGTC
<i>Cat</i>	NM_012520.2	AGAAACCCACAACTCACCTGAAGG	GAATCCCTCGGTCGCTGAACAAG
<i>Nrf2</i>	NM_001399226.1	TGCCACCGCCAGGACTACAG	GCGTGCTCAGAAACCTCCTTCC
<i>Ho-1</i>	NM_012580.2	CAGGTGTCCAGGGAAGGCTTTAAG	TGGGTTCTGCTTGTTCGCTCTATC
<i>IL-1β</i>	NM_031512.2	AATCTCACAGCAGCATCTCGACAAG	TCCACGGGCAAGACATAGGTAGC
<i>TNF-α</i>	NM_012675.3	ATGGGCTCCCTCTCATCAGTTCC	CCTCCGCTTGGTGGTTTGCTAC
<i>IL-10</i>	NM_012854.2	GGCAGTGGAGCAGGTGAAGAATG	TGTCACGTAGGCTTCTATGCAGTTG
<i>Fabp2</i>	NM_013068.1	TGAAACTGACGATCACACAGGAAGG	GACGCCGAGTTCAAACACAACATC
<i>Acac</i>	NM_022193.1	CGTTCGCCATAACCAAGTAGAGTCC	ATATACCTCCAGAGCCGCCATCC
<i>Cpt1a</i>	NM_031559.2	CAGGAGAGTGCCAGGAGGTCATAG	TGCCGAAAGAGTCAAATGGGAAGG
<i>Colla1</i>	NM_053304.1	TGTTGGTCCTGCTGGCAAGAATG	GTCACCTTGTTTCGCCTGTCTCAC
<i>Acta2</i>	NM_031004.2	TAGAACACGGCATCATCACCAACTG	ACATACATGGCAGGGACATTGAAGG

Table S2. Primer sequences for real-time semi-quantitative PCR.

Gene	Gene ID	FORWARD	REVERSE	ProdSize
<i>Actb</i>	NM_031144.3	GGTCAGGTCATCACTATCGGCAATG	CAGCACTGTGTTGGCATAGAGGTC	165
<i>Adh1</i>	NM_019286.4	CTTCATCAGCACCAGCACCTTCTC	CAGCCAATGACGACAGACAGACC	216
<i>Aldh2</i>	NM_022407.3	GTTTCCTCCTGGCGTGGTGAAC	AGCAATGTCCAAGTCGGCATCTG	223
<i>Cyp2e1</i>	NM_031543.2	GGAAGTGAAGACCAGCACAAC	GGGTAGGTTGGAAGGGACAAGATTG	210
<i>Cat</i>	NM_012520.2	CCAGCGACCAGATGAAGCAGTG	CTGTGGAGAATCGGACGGCAATAG	324
<i>Nrf2</i>	NM_001399226.1	ACGAGGCGGTACAAGTTTTGGAAG	AGGGACTGGAGAGATGGCTAAACC	385
<i>Ho-1</i>	NM_012580.2	GGAAGTTTCAGAAGGGTCAGGTGTC	GCTGTGTGGCTGGTGTGTAAGG	234
<i>IL-1β</i>	NM_031512.2	CGACCTGCTAGTGTGTGATGTTCC	GGTGGGTGTGCCGTCTTTCATC	256
<i>TNF-α</i>	NM_012675.3	CACCACGCTCTTCTGTCTACTGAAC	GGTATGAAATGGCAAATCGGCTGAC	365
<i>IL-10</i>	NM_012854.2	TGACAATAACTGCACCCACTTCCC	CTTCACCTGCTCCACTGCCTTG	352
<i>Fabp2</i>	NM_013068.1	TGAAAGTGAAGATCACACAGGAAGG	ACCAGAAATCTCTCGGACAGCAATC	224
<i>Acac</i>	NM_022193.1	ACAACGCCTTCACACCACCTTG	GCCTGTCATCCTCAATATCGCCATC	243
<i>Cpt1a</i>	NM_031559.2	TGGCTATGGTCAAGGTCTCTCAG	CAGCAGTATGGCGTGGATGGTG	392
<i>Colla1</i>	NM_053304.1	TCACCAGACGCAGAAGTCATAGGAG	GGAGACCACGAGGACCAGAAGG	262
<i>Acta2</i>	NM_031004.2	TAGAACACGGCATCATCACCAACTG	CATCTCCAGAGTCCAGCACAAATACC	261