

Supplementary Table S6: Primer sequences used in real-time qRT-PCR analyses

Probeset ID	Gene symbol	Forward (5'-3')	Reverse (3'-5')
	18s rRNA_v1	ccgtcggcatgtattagctc	accgtctcctacctggttga
Ssc.30843.1.A1_at	ADAMDEC1	ttctgcttgctgactcgta	aggtgagccatgcagacttc
Ssc.18933.1.A1_at	ADH4	tggatgactggtgtctcat	tcatggctcattgttcatca
Ssc.4891.1.A1_at	AGPAT9	cgactcaagcgggagtaaga	gtcgccttacgctacggagtc
Ssc.6943.1.A1_at	ANGPT1	tgatattttggatgcagtttca	catgtgtattgttttaagggttg
Ssc.8980.1.A1_at	ANGPTL4	actcttcacctgctgagga	gcaccacatacaacagcac
Ssc.1039.1.S1_at	APOC3	gacacctcccttctggacaa	ggactccttcacgctggtta
Ssc.833.1.S1_at	C1QA	cttccaggtggtgtccaagt	tggatccagacctgtctcc
Ssc.17026.1.Sa_at	C4A	gcctctttgagacgagcatc	ctaagtgaaggcggcaagag
Ssc.10460.1.S1_at	CFB	aaggaaagctgctgtgaga	agtgttggggtcagcatagg
Ssc.864.1.S1_at	CFH	cagtctatcctccgggaaca	gcattttggtggttctgacc
Ssc.16157.1.S1_at	CRP	taggaacctgaggggtgagg	ggggtcaaatcgggaactaca
Ssc.2033.1.S1_at	CRY1	ttaggacaaaattttcttgaatc	aaatttgggtcaaaacaagca
Ssc.19692.1.S1_at	CXCL2	cagggaggaggagttctctcaa	atgtccaagggaacgtcaa
Ssc.208.1.S1_at	CYP1A1	ctccccctacatgccact	agagcctggtccaaagaatg
Ssc.203.1.S1_s_at	CYP3A22 / CYP3A39	cgccctgaaaggttcagtaa	ggggcgctctgtgtttctta
Ssc.203.1.S1_at	CYP3A22 / CYP3A39	cctgaaactaagctctcaagga	caaggtctctgatgctctgg
SscAffx.21.1.S1_at	CYP7A1	caattggcaaacgttttctt	gcgtgatgtgtccttcaaaa
Ssc.21145.1.S1_at	DEFB1	gctgactgtctgcctctct	acttggccttgcactgtaa
Ssc.16236.1.S1_at	DPP4	acatgggtcttccaactcca	aatccactccagcatccac
Ssc.30467.1.A1_at	ELOVL2	tgatgagcagaccagcactt	ccgccccaatcatgatacta
Ssc.13778.1.S1_at	FCGR1A (IgG)	caagtgcaaggtaacaacg	cgttgctcttccactcaaca
Ssc.16377.1.A1_s_at	GSTA1	tgttgaagagccatggacaa	ttcagcagaggggaagttgg
Ssc.390.1.A1_at	HIF1A	tggggcatggtaaaagaaag	gtaatgctccccatccaa
Ssc.47.1.S1_at	IGFBP2	catggcctgtacaacctcaa	tggctgtggttactgcatc
Ssc.15588.1.S2_at	IGFBP3	agcccctgtaccgtaactct	ggcagaacgtctgaagatcc
Ssc.286.1.S1_s_at	IRG6 (RSAD2)	caccatgcgggaattctact	tggaaaaatgctgacaatga
Ssc.14340.1.S1_at	LITAF	ccggtggtatggttgaaag	aaaccaactctggcttgtgg
Ssc.670.1.S1_at	LYZ	aaggagcagaatgccagatg	cttcatggtcagcaaatgc
Ssc.548.1.S1_a_at	MMP7	tggcattctttaggtctgg	tgaatgtgtgtctcctgaa
Ssc.600.1.S1_s_at	MT1A	actgctctgccccacag	acagcagctgcacttctcc
Ssc.9372.1.A1_at	NFKBIZ	ggaaacatgctttgctggtt	ccaaatacattccttgggaca
Ssc.13870.1.A1_at	NR1D2	tcaagtatccttttcttcca	tgcttcggatcaaatcaaga
Ssc.15928.1.A1_at	NR3C1	ccaggtcacttccatcact	gtctcagccagactccaagc
Ssc.6323.1.S1_at	PLIN2	tctgatggcctcctcagttc	tctcaggttctgggtctct
Ssc.8177.1.A2_at	PLP1	agaagcttccaggtgcata	taggaggtctccaggggaagg
Ssc.14490.1.S1_at	PTGR1	catgaacacatcacggaagg	tggactctcgtgtcgtcttg
Ssc.2381.1.A1_at	S100A9	caaatgtcgcagatggaatg	caggtcttccaggtatggtt
Ssc.7493.1.A1_at	SIRT1	cgcagagaagcatttatttcc	ctcttctccggcagtaatgg