

Table S1. Primer sequences used for RT-qPCR

Gene symbol	Forward Primer (5' to 3')	Reverse Primer (5' to 3')	Fragment size (bp)	Tm (°C)	PCR efficiency
<i>TSC22D2</i>	AGACTCAGACCCAGCCTTTG	CAACAGGAGGCTTCACAACA	120	60	1.99
<i>ACTB</i>	CAACTGGGACGACATGGAG	GAGTCCATCACGATGCCAG	234	60	1.92
<i>GAPDH</i>	ATTCCACCCACGGCAAGTTC	AAGGGGCAGAGATGATGACC	225	60	1.96
<i>SDHA</i>	CTACAAGGGGCAGGTTCTGA	AAGACAACGAGGTCCAGGAG	141	60	1.94
<i>NR3C1</i>	CACCTGGATGACCAAATGACC	AGGGTAAAGCCATTCTCTGCTC	151	60	1.98
<i>NR3C1α</i>	TCAGAACTGGCAACGCTTTTATCAAC	AATGTCTGGAAGCAATAGTTAAGGAG	93	60	1.98
<i>FKBP5</i>	TGCCATTTACTGTGCAAACCAG	TTCTTCGGATAATGCCTCCATC	148	63	1.97
<i>TSC22D3</i>	GCAGGCCATGGATCTGGTG	GGGCTCGCCAGGGTCTTC	144	60	1.99
<i>OVGP1</i>	GGGGCACTTTCTGTGGCACT	AGCCAGGCTTTCAGGGCAAG	149	60	1.96
<i>PGR</i>	TGAGAGCACTAGATGCCGTTGCT	AGAACTCGAAGTGTCGGGTTTGGT	197	60	1.98
<i>ESR1</i>	AGGGAAGCTCCTGTTTGCTCC	CGGTGGATATGGTCCTTCTCT	234	60	1.90
<i>HSD11B1</i>	TTCTCAACCACATCACCCAC	AGTTGACCTCCATGCTTCTG	81	60	1.98
<i>HSD11B2</i>	GGTCAAGGTCAGCGTCATC	CAAGTGCTCGATGTAGTCCTC	148	60	1.96
<i>IL6</i>	ATAAGGGAAATGTCGAGGCTG	GTGGCTTTGTCTGGATTCTTTC	88	60	1.98
<i>CXCL8</i>	GCTCTCTGTGAGGCTGCAGTT	TTTATGCACTGGCATCGAAGTT	62	60	1.96
<i>PTGS2</i>	AGAGCTTCCCGATTCAAAGG	CCTCGCTTCTGATCTGTCTTG	144	60	1.94
<i>TP53</i>	GGAACAGCTTTGAGGTGCGTGTTT	ATACTCGCCATCCAGTGGCTTCTT	182	60	1.96
<i>DDB2</i>	GATTCGGGTTTACTCTGCCTC	AAATTAGGATCTGGGTATCGGC	150	60	1.97
<i>CASP3</i>	ATTGGA CTGTGGATTGAGAC	ACCAGGTGCTGTAGAATATGC	106	60	1.95
<i>BAX</i>	GCTGACGGCAACTTCAACTG	GCGTCCCAAAGTAGGAGAGG	202	60	1.96
<i>NFKBIA</i>	TGTCTTTGGGTGCTGATGTC	CTGGTAGGTGACTCTGTTGAC	140	60	1.95
<i>GADD45G</i>	ACTCTGGAAGAAGTTCGCGG	TTGTCGGGGTCCACATTGAC	164	60	1.95