

<i>Gene</i>	<i>Forward primer</i>	<i>Reverse primer</i>	<i>Probe</i>	<i>Tm probe</i>	<i>Tm fwd primer</i>	<i>Tm rev primer</i>
<b>Ppia</b>	AGGGTTCCTCCTTTCACAGAATT	TGCCATTATGGCGTGTAAAGTC	TCCAGGATTCATGTGCCAGGGTGG	73.1	61.5	62.1
<b>Gnas</b>	AGAACATCCGCCGTGTCTTC	CCTTCTTAGAGCAGCTCGTATTGG	CGTGACATCATCCAGCGCATGCAT	73.2	62.6	62.7
<b>PGK1</b>	ATTCTGCTTGACAAATGGAGC	AGGCATGGGAACACCATCA	AACTCCGTTGTCTTATGAGCCACCTGG	72.1	61.5	62.4
<b>HPRT1</b>	GCAAACCTTTGCTTTCCCTGG	ACTTCGAGAGGTCCTTTTCACC	CAGCCCCAAAATGGTTAAGGTTGCAAG	71.2	62.0	61.0
Cx40	ACTCCAGGGAGGAGGAAAGG	AGGAAGCTCCAGTCACCCAT	GGCCTCGGTCTCCTACTCTTGGCAA	71.3	61.9	61.1
Cx43	TGCCGGCTTCACTTTCATTA	GACCTTGTCCAGCAGCTTCC	AGACATGGGTGACTGGAGCGCCTTG	73.3	61.7	62.3
Cx45	TGGGTAACAGGAGTTCTGGTGA	CCTCTAGCAGGCGAGTCAGG	GGCAAACCAATTCCACCACCATGAG	71.1	61.9	62.6
Cx30.2	GGTCGTCTCTGGCTGGTGAT	ACGAACTCCTCCTGCTCGTC	TGATCTCCGCATCCTGGTGCTGG	73.3	62.6	61.9
Scn1b	ACCAGCGTCGTCAAGAAGAT	GGTCAACACCACAATGAGCA	AGGTGGTGGACAAGGCCAACAGAGA	71.0	59.9	60.6
Scn5a	GGAAGAGTCCAGCAAACAGC	GGCTTCAGAGGATGTGGTCT	AATCCCAAGTTGTGTCTGGTGGCCA	71.5	60.0	59.3
Scn10a	GCCTCATCTTCTGGCTCATC	AATGGGTTGCTTCTGGTGTC	ATCATGGGTGTGAACCTTTCGCCG	72.8	59.9	60.0
Usp16	CCTTCGGTTTGGCTCTGTCTT	GCATGCTGCTCCTGAGAATCT	GCCACAGCCCTGATGGCCACATT	73.2	62.9	62.0
Cct8	GGCTAAACAATCACATCATATGGA	CAAACGCTTCAGCAAACCTTCTTAA	GCATACTGTTCAAGCCCAGGACACGTCT	71.7	61.2	62.0
Bach1	CTGAGACGGACACGGAAGGT	GGGCATTGAATGGCAGCTT	ACCTCACAGTCTGCTCCCTGGCA	72.4	62.6	63.4
Dyrk1a	TAACAGTGGGAGAGCCAGGT	ACCTGGGACTGTGTGTCTC	AGCATCGACACAGCGGTGGACACTT	72.0	59.7	60.0
Sh3bgr	CCCTTCCAATGGAGATGTA	TGTCCTCTTTCTGTGCCTCA	CAGAGGGTGTGCAGAGGGAACAGA	71.9	59.7	59.6