

**Table:** Primers used in the Real-Time quantitative PCR analyzes.

lncRNAs	Primer sequences (5'-3')	Ta (°C)	Efficiency (%)	Amplicon (bp)	Transcript ID (Ensembl) or Accession number (NCBI)
<i>lncRNA14696</i>	F: GATAGCGGCAGTTGGGTAAA R: CAAAGAGAGGGAGAGCATCG	55	92.2	119	ENSSAUT00010014696
<i>lncRNA16861</i>	F: TGCAGTATGACTTGAGTGAAAAA R: AGTTGCTGTTTTGCCTCCTT	54	102.4	79	ENSSAUT00010016861
<i>lncRNA20194</i>	F: GGACACGTAGACGCTGTGAA R: CCTCCAAACAGTCGCTCAA	55	98.0	120	ENSSAUT00010020194
<i>lncRNA21817</i>	F: TGACAAAAACCTCAGGGATTG R: CTTGCGTGCTCTGTGGTAA	55	90.3	129	ENSSAUT00010021817
<i>lncRNA40141</i>	F: TTCCCTGTTTCTGGAGGATG R: GGAGCATTTCTGTTGTGT	55	115.3	200	ENSSAUT00010040141
<i>lncRNA54283</i>	F: CTGCTGTGATCAGCGGTTT R: AAAACGAGGTGGAAATGTGC	55	110.1	110	ENSSAUT00010054283
<i>lncRNA60660</i>	F: GCCTGACTCTCCATTGAAGC R: ACAGCTCCCATCAGCTCACT	55	91.4	147	ENSSAUT00010060660
<i>lncRNA02328</i>	F: GAGGAAGGCTGGACTGTGAG R: GTGGAGCTCTGCCCATAGAG	55	95.1	197	ENSSAUT00010002328
<i>lncRNA62925</i>	F: CGCATCCAATCAAAAGAGGT R: CAGGATCTCACCCACAACACT	55	92.2	116	ENSSAUT00010062925
<i>lncRNA31317</i>	F: GTTGTTCAAGTTTCCCTCA R: GTCGGCGTTTCTTTCCATA	55	112.5	153	ENSSAUT00010031317
<i>lncRNA43061</i>	F: TACGTTGGGAGGATGAGGAG R: TGTACGCAGCCATAAAGCAG	55	200.2	199	ENSSAUT00010043061
<i>lncRNA05337</i>	F: ATGACGCCTCCATTCACTCCTCT R: AGCTTTCCTGTGACAACGGTTGC	62	100.1	81	ENSSAUT00010005337
<i>myf5</i>	F: CCATCCAGTACATCGAGAGCC R: ATCGCCCAAAGTGTCGTTCT	56	98.9	209	JN034420.1
<i>myod1</i>	F: TTTGAGGACCTGGACCC R: CTTCTGCGTGGTGATGGA	60	96.4	139	AF478568
<i>myod2</i>	F: CACTACAGCGGGGATTCAGAC R: CGTTTGCTTCTCCTGGACTC	60	101.2	149	AF478569
<i>myog</i>	F: CAGAGGCTGCCCAAGGTCGAG R: CAGGTGCTGCCCGAACTGGGCTCG	68	96.7	182	EF462191.1
<i>mef2c</i>	F: ACAGCTTGTTGGTGCTGTTG R: AGATAGCGCGAATCATGGAC	55	96.4	144	XM_030435338.1

<i>myf6</i>	F: CATCCACAGCTTTAAAGGCA R: GAGGACGCCGAAGATTCACT	60	100.6	151	JN034421
<i>cdh15</i>	F: GGCCTTCGTGGGTAGATAGG R: TTAAACGTGCACCTGCTGTG	55	96.7	149	XM_030414763.1
<i>cav3</i>	F: CTGAGGGTGTGGACAAGGTCCG R: GTCAGGCCACGATAAACCCA	55	98.5	133	XM_030420899.1
<i>mymx</i>	F: TGCTGCGGTCCCTGGTTATC R: ACTCCTGGGATCGAATGCGG	60	92.4	144	ENSSAUG00010011859
<i>mymk</i>	F: TTCACTGCGGTTTACCACGC R: CCCACATAGAGAGAGCTGTGCTG	59	98.9	112	ENSSAUG00010019449
<i>dock5</i>	F: TCAACAGGCCCAGTAAATCC R: GGGAAGCAGTTCCATCATTC	60	95.2	115	XM_030416547.1
<i>crk-a</i>	F: AACGCCCAGAACTCTGAAGG R: ACCATGTGCCCCACCTCTAA	57	96.4	314	ENSSAUG00010004157
<i>crk-b</i>	F: ATGCCCAGCCATTCAGAAG R: TACCATGTCTCCACCTCCAGA	59	95.8	77	ENSSAUG00010012872
<i>crkl</i>	F: TCCGCCTGGTATTTTGGACC R: TGTGCTGCTTGGGTACCTGG	59	90.9	294	ENSSAUG00010019265
<i>ef1a</i>	F: CTTCAACGCTCAGGTCATCAT R: GCACAGCGAAACGACCAAGGGGA	60	96.6	263	AF184170.1
<i>rps18</i>	F: GGGTGTTGGCAGACGTTAC R: CTTCTGCCTGTTGAGGAACCA	60	97.7	163	AM490061.1
<i>rpl27a</i>	F: AAGAGGAACACAACCTCACTGCCCCAC R: GCTTGCCTTTGCCCAGAACTTTGTAG	60	97.7	160	AY188520.1

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Ta: annealing temperature; F: forward; R: reverse; Amplicon: product size (base pairs).