

Additional file 3: Table S2. List of PCR primer sequences

Application	Gene symbol	Sequence (5'→3')	fragment size (pb)	Efficiency
Terra PCR	<i>MMUT</i> NC_040271.1	F: GCAGATTTTGGCAAAGAAAG R: ACCTTCAAGGCAGCATCATA	402	-
PACE PCR	<i>MMUT</i> NC_040271.1	F1: GAAGGTGACCAAGTTCATGCTAATCCCAGATTCTTCTTGAATAATGATTTG F2: GAAGGTGCGAGTCAACGGATTGAATCCCAGATTCTTCTTGAATAATGATTTA R: GTGAAAAGTGCTCGAATTGCCAGGAA	81/82	-
Quantitative PCR	<i>MMUT</i> <i>XM_004018875</i>	F: GCTCATCAAGGAACTCAA R: TTCAATATCATCAAGCACTTG	169	1.80
	<i>GAPDH</i> NM_001190390	F: CGACTTCAACAGCGCACTC R: TGCTGTAGCCGAATTCATTG	113	1.90
	<i>YWHAZ</i> NM_001267887	F: ATTAAGTGAAGAGTCATACAA R: GTATCCGATGTCCACAAT	81	2.00
	<i>RPL19</i> <i>XM_004012836</i>	F: AATGCCAATGCCAACTC R: CCCTTTCGCTACCTATACC	149	2.00
	<i>SDHA</i> <i>XM_027980212</i>	F: GAATGGTCTGGAACACTG R: AGTAATCGTACTCGTCAAC	156	2.00

F: Forward primer, **R:** Reverse primer.