

RT-qPCR primer information

Contig id	Oligo Sequence (Forward)	Oligo Sequence (Reverse)	Amplicon Length (bp)	Primer Efficiency %
Stage 0-1				
comp14102_c0_seq1	CTTCCTACCACAGCTCAGAAAC	ACCATTGACCGATTGCACTAT	111	102.4%
comp13917_c0_seq1	TCCTGATGTTGTGCCGTATTC	CAAGGTCACAGGAGTTGTAAGT	97	103.2%
comp17500_c0_seq1	CGCATCAATCAGTCGGGATAA	GACTTCTTGTCTCTCCCAAAG	99	98.7%
comp13662_c1_seq1	AACGAGGAACTTGAGACCAATC	GGCACTTCTGTCCTGTCATATC	83	103.8%
Stage 1-2				
comp2693_c0_seq1	GGGAAGATGGTGACCGTTATT	AGGATGGTTGCCATGTGTAG	86	96.3%
comp17994_c0_seq1	CCGAAAGTGTTGTGGCAATATC	CCAAAGGTGTAGCTCTGGTTAT	99	99.8%
comp2742_c0_seq1	CCTACGTAGCATTGCCTGTATC	GTGCGACCTTCATATCCGTATC	143	95.2%
comp14332_c0_seq1	CAGTGACGGTGATGGAGTATG	GCGAATGAACCAGGAGTAAGA	120	96.3%
comp16666_c0_seq1	GACCTAGATGTCTCGTTTCGTTAG	GATCTGTCTGACTGTGGTGAAG	84	94.9%
Stage 3-5				
comp13053_c0_seq1	GTACTCTTCTCGCGGAATTGAT	ACATACCTTCACCCTTGATTG	99	101.2%
comp10800_c0_seq1	GTCAATCCCATCTCCTGAGTTT	CATGGTTGGAGAAGCTCTTGTA	97	104.4%
comp19935_c0_seq1	CATACGCGACTCGTGGTAATA	CAGAGATGGAATGTGGTTGATTG	107	96.1%
comp17465_c0_seq1	GGTTGAAGGCAGCAATGTAATC	GGTAGGTCCTCTTCACTTCTTG	101	94.7%