

Supplementary table 2: Primers used for Sanger sequencing.

Gene name	Sequence
NODE_3507_length_3801_cov_287.490107_g1332_i0_4_Fr	GTTGTACCAGCCCGTGATCT
NODE_3507_length_3801_cov_287.490107_g1332_i0_4_Re	GCACCAAGACCATCAAGGAT
NODE_10194_length_2189_cov_334.651786_g3124_i2_4_Fr	CCATTCTTTTCGATGGCTGT
NODE_10194_length_2189_cov_334.651786_g3124_i2_4_Re	CTTTGGCTTCTCCGACTCTG
NODE_10326_length_2171_cov_27.087678_g3450_i0_4_Fr	CGGTATCCGTTGTCTCTGGT
NODE_10326_length_2171_cov_27.087678_g3450_i0_4_Re	TCCATATGCCTAGCCGACTC
NODE_29952_length_617_cov_2.156475_g11635_i0_4_Fr	CCAAGCAGAACTTCCACCTC
NODE_29952_length_617_cov_2.156475_g11635_i0_4_Re	GCAGGAGCGACGACTTAGAA
UnnamedSample_HQ_transcript_125_1_Fr	CCTCGACGAAGACGAAAAAG
UnnamedSample_HQ_transcript_125_1_Re	GAGAACCGGGTGTACTGGAA
UnnamedSample_HQ_transcript_15376_3_Fr	AGGGATTTCTCGCCCTCTAC
UnnamedSample_HQ_transcript_15376_3_Re	GACCGACTTGATGGTGTCTCT
UnnamedSample_HQ_transcript_16148_2_Fr	CCACGGTCTACAACGACCTT
UnnamedSample_HQ_transcript_16148_2_Re	CGGCAGGTAGAATTCGTAGG
UnnamedSample_HQ_transcript_16178_1_Fr	CGCAGAGAGGAAGAAGATGG
UnnamedSample_HQ_transcript_16178_1_Re	TTCGTGCCTCTGTTGTTGAG
UnnamedSample_HQ_transcript_3435_3_Fr	CTCGAGCAGAACATCCCTTC
UnnamedSample_HQ_transcript_3435_3_Re	AGTGGTGTGAGCTCGTCCTT
UnnamedSample_HQ_transcript_4172_3_Fr	TGAGGAGCGAAACAACCTCT
UnnamedSample_HQ_transcript_4172_3_Re	TGCTTCTGCTGCTTCTCGTA