

Table S1 Primers used for qPCR in this study

Gene	sequence (primer forward 5'→3')	sequence (primer reverse 5'→3')	primer efficiency (%)	Tm (°C)
<i>ve11</i>	AGCCTTATGTGCCTCACT	GCAGGAGCAGAGTAGTTG	100	57.2
<i>lae1</i>	ACTGGAGATTGACTGGATGC	TTCTGCGTCTGGTAGCCTC	100	62.9
Trire2:23171	GGTGGCTGTAGATGTGTT	TGAGAGTCGTCGTTGAAG	100	57.2
Trire2:112083	GGTGAAGAGATACGGTAT	TGAGTTGGAGTAGATGTC	100	52.6
Trire2:109101	GAATCCAGTGGCATTCTATC	CGACCCTCAATAATATCATAACC	100	57.1
Trire2:69187	GCCTCTACCAATGAACCTC	CCGAGACAGGACATAATC	97.2	56.1
Trire2:107071	CGATTGCCGTCTACGATAT	GCTCAGTTGTCCAGTCAT	100	56.1
Trire2:111494	ATCAACCAGACAGACAAC	AGATATAGACTTGCTCACATAG	95	52.6
Trire2:122824	CGTATGGCACCATCATTGG	CTGGGAAGACTCGTCGGAG	100	64
Trire2:39587	CTTGACAACCCGCCAAACAG	CGATGGAGTAAAAGACACCG	87.5	62
Trire2:82041	CTCATCGTCTTCCCCATTC	CATGCTTGTGTTGTGCCCTG	100	62
Trire2:69904	CGGGAGACATTTGCCTTTATC	CAGAAGCAGGTCGCAGAGTAG	86	64
Trire2:5647	CTACAACCCCGATAACCTC	CGTGGAAGATGCGGTAGTAC	98.5	60
Trire2:110744	CCTCTACGCCTCGTCAATC	CAATGGCGGAGAAGAAGAAGC	85	62
Trire2:110339	GATGCCGCTGATTGCCAC	CGAGATCGGTGATGATGTCG	91	65
Trire2:62462	CACGAACCCAACGCTCCAAC	CGAAGTATGCCCACTCCAG	95.5	63
Trire2:121990	CTCAAGGTAGTTGCGGTG	CGTCATCTGGTGGCTGTTG	90	60
Trire2:76763	CGGCAGAGGTTGAGCAATAC	CGGTAACACGGAGGCGTTTC	89	64
Trire2:105224	CACCGCTTGCCATCATCTTC	CAAGAGGAATCCCAAGACAC	100	61
<i>sar1</i>	TGGATCGTCAACTGTTCTACGA	GCATGTGTAGCAACGTGGTCTTT	64	64