Table S1. Primer sequences used for RT-qPCR analysis

Gene ID	Gene Product Name	Primer sequence (5′-3′)	Amplicon
TERG_02940	bZIP ap1 transcription factor	F: GCCACTTTGATCCAGTTCTG R: TCAGGTAGAGGGAAAGCATC	103 bp
TERG_01117	Stress response transcription factor SrrA/Skn7	F: TCTACTCTTTCTCCTGCGTC R: CCTGTATCTTGTTCACCGC	117 bp
TERG_06759	C2H2 transcription factor (Seb1)	F: CGAACTGCTCAGAGTAATGTG R: GGTGCTTTGTGGATTGGTC	120 bp
TERG_07131	Phosphotransmitter protein Ypd1	F: TGTCCCAACTAGGTCACTTC R: TGCCAGACTCGTCCTTTT	114 bp
TERG_07855	Response regulator ssk1	F: GGATGGCGGTTTGTCAAC R: TGGAGGCTACAGGTAATCTTC	95 bp
TERG_07326	Glutathione S- transferase	F: GAGCACAAGAAGAAGCTGGA R: AGCCAAGAAGGGAAGGTAATAG	127 bp
TERG_01685	Glutathione S- transferase	F: GACTTCAGACAGTGCAAGGA R: ACCAGATTCCCAGAGGTAGA	105 bp
TERG_04073	Glutathione synthetase	F: CGTTCACGACGCATATTCAATG R: GACCGATTCTCGCCTTACTTATC	99 bp
TERG_01349	Glutathione peroxidase	F: TCAACTACGGCGTCACTTTC R: AGACCAGGCATCTCCTTCT	100 bp
TERG_08069	Glutathione reductase	F: CCCGTTCTACTCCTCCTCTATT R: TAGTCGCCAGTGTTCTGTTG	101 bp
TERG_02005	Catalase	F: CAGGAGTCCGCTCATCAAAT R: CTTGTATGTGTGGCCAGAGTAG	96 bp
TERG_01252	Catalase A	F: ATTGGGATCTCGTGGGAAAC R: GGCCTTGAGGGACTTCATTAT	114 bp
TERG_03174	Siderochrome iron transporter	F: TGTCGACGCAATGGAAG R: AGGTTGATTCCCAGTAAAGCC	85 bp
TERG_00697	Nonribosomal siderophore peptide synthase	F: GACTGTGGTTCCCACTGAAA R: TCGGTTGGCTGCTGTAAAT	105 bp
TERG_04949	Siderophore iron transporter mirC	F: CCATTCTGGGAGACATCCAAA R: GAAGAAAGCCAATGCACATCC	102 bp
rpb2	DNA-directed RNA polymerase II core subunit RPB2	F: TGCAGGAGGTTTGATGAAGA R: GCTGGGAGGTACTGTTTGATCAA	59 bp
gapdh	Glyceraldehyde-3-phosphate dehydrogenase	F: GCGTGACCCAGCGATGTAGT R: CGGTGGACTCGACGATGTAGT	62 bp