

**Table S2. Primer sequences used for qRT-PCR analysis of drought-related genes in *PtrbHLH66* homolog silencing lemon**

Name	Accession number	Primer sequences (5'-3')	
		Forward	Reverse
<i>CISOD</i>	AF318938.1	CCGTTTACTGGTCGTTT	TGACACCCTCAGTTCCAC
<i>CIPOX</i>	KT374296.1	CCGTAGACGAGATGGTGA	CGTAAAGACGCTTGGAGA
<i>CICAT</i>	KF640697.1	CGGTATTGTGGTTCCTGG	CCCTCGTGGTGATTGTTG
<i>CIP5CS</i>	XM_025093600.1	AGGAAATGCTGCCACCCA	CGGAAGAAGAAAGCGAGT
<i>CINCED3</i>	AB219179.1	CAGCAACCCAAGGAAA	TGCGTCCAAGGCTAAA
<i>CINCED9</i>	AB219172.1	AACAGCCTTACCCGAATC	AGCGAAGTTACCGCCAAT
<i>CIZEP</i>	AB114670.1	GGTCAACGCTATGAAGGT	CACTACGGCCAAACAAAT
<i>CLABA2</i>	NM_001288867.1	TTGGGATTCTAAGATAAAGG	GACAGCAGGCGATGAG
<i>CLAAO3</i>	XM_006487735.3	AAGTTGGCACAATCCTAT	ATTCCTGTCTCATCCCTC
<i>CIDREB1A</i>	XM_006490258.3	GGTCATCGCCAGAGTCACGT	GACTTCTTGTTTCGGCTCCCT
<i>CIDREB2A</i>	XM_006478914.3	GTCTGTCGGCAATTCATCTC	CACCCATCATCAAATCACTC
<i>CIDREB3</i>	XM_006421783.2	GGGTCTCCGAAATCCGTGAG	TGCTGAGTTGCCCTTGATGC
<i>CIRD20</i>	XM_025102476.1	GACTTATTCAGGGTTGCG	CTCGGTGTCGTATGTTCC
<i>CIRD29A</i>	XM_006421853.2	AAAGGTGCTGTAGGTTTCGTG	GTTGTTATTGTTGCGTCCTG
<i>AtERD</i>	XM_025097036.1	TCCGAATCCTCCACTT	AACAGCACAGCCGTAA
<i>CIACTIN</i>	XM_006464503.3	CGTATGAGCAAGGAAATCACAG	ATTGATCCTCCAATCCAAACAC