

Table S1: Primers used for QPCR analysis

Gene	Arabidopsis Genome Initiative ID	Forward Primer (5'-3')	Reverse Primer (5'-3')
<i>ACT2</i>	At1g49240	CCGAGCAGCATGAAGATTAAG	CATACTCTGCCTTAGAGATCCACA
<i>CER1</i>	AT1G02205	AAGGATGGGAAATGCATGAG	TGATGTGGAAGGAGGAGAGG
<i>CER2</i>	AT4G24510	GGCGAAACAACACGAATG	TCGGATATCCAACCATTTTC
<i>CER3/WAX2</i>	At5g57800	GGGAAGCAGCCATGAAGTAT	TTCTGCTTGCATCTCCTTT
<i>CER4</i>	At4g33790	ACCGTGGACCAACAAGAAG	GCAATCAAGTAGCGTATGGTCA
<i>CER5/WBC12</i>	At1g51500	TTTCTTAATTTGTTTTGCACATACG	TTCTCCAAAATAACAATGCTGAA
<i>WBC11</i>	At1g17840	TAGCACGCAGAAGGATGAAG	CCATCTGTTTCGTGTAGCAATG
<i>CER7</i>	At3g60500	CCCGTTACTAAGTCATCGAGCA	CCCTAATTCTCCGGTTACTTCC
<i>KCS6/CER6</i>	At1g68530	TCTAGCTCGGTGAAGCTCAAG	AAGCTCAACGGCGACAATAG
<i>KCR</i>	At1g67730	CTACTTCTCCGACCATCCA	CTGGGTTACGAGCAACGAGT
<i>ECR/CER10</i>	At3g55360	CTGGAGCTTTGGTGCTTACA	CTGGGGTCCCTCAGATTCTT
<i>LACS1</i>	At2g47240	GCTCAAGAATCGGAGGAACA	CACACACATCATCGCAACAC
<i>LACS2</i>	At1g49430	GCAAAGAGGTCCATGGCATA	GATGAGAGGATCCGAACACAA
<i>LACS3</i>	At1g64400	TGAAAAGACCTCAGCTTCTCAA	AAACCTGAGAGATTTGTCCAACA
<i>ATT1</i>	At4g00360	GAGAAACGACGTCGTTGTTCTTAACG	AACCCGAGAAGCTGCATTACACGG
<i>GPAT4</i>	At1g01610	TCCTGAAGGCACAACGTGTA	ACATCCCTTGCTTGCAGTTC
<i>GPAT8</i>	At4g00400	AATGACTGTCAACGTTGGTGGCAAG	GCGAGTAAGTTTCGGTGCATTCTGAAG
<i>LCR</i>	At2g45970	TGAAATCGACTTGGGGAGAG	GCCAGATCCTTACCCAGACA
<i>HTH</i>	At1g72970	TTCTTCGGGAGAGACTTGA	CCAACACAATTCACAAACAGA
<i>BDG</i>	At1g64670	CACATTAGCCGCACAGTCAG	TCAAGTACGTTCCGCATCCTG
<i>SHN2</i>	At5g11190	CGCTAAGACCAATTTCCCTGT	GCGTTCAAAAAGCTCAGATAATG
<i>SHN3</i>	At5g25390	GAAGATGATCAAATGGCGATG	TACCCTCGATTATTATTAACCTCA
<i>MYB30</i>	At3g28910	CAACGTCTTCTCTGCTGAA	GTCTTCGGCGAGTTTTTCAC
<i>MYB41</i>	At4g28110	CTTAGACTCGCTCATGTCAACG	CTTCCGGAATTCCAAAGTCG
<i>ACR4</i>	At3g59420	TCGAGATCCTAAGCGGAAGA	GGGTCCAAGAGTGCCTTTAT