

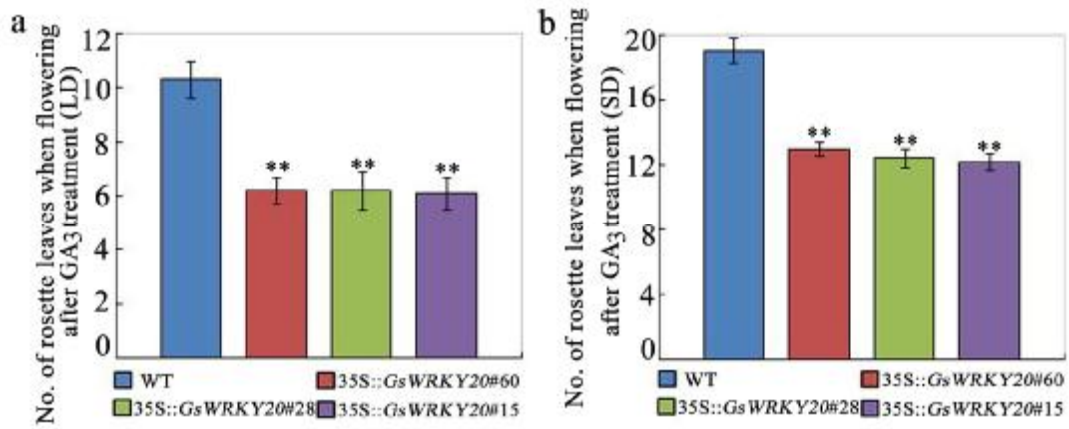
Locus or Accession number of the genes

Gene symbol	Gene locus or Accession number	Gene symbol	Gene locus or Accession number
<i>GAPDH</i>	DQ355800	<i>AtWRKY12</i>	AT2G44745
<i>ACTIN2</i>	AT3G18780	<i>AtWRKY53</i>	AT4G23810
<i>FLC</i>	AT5G10140	<i>AtWRKY63</i>	AT1G66600
<i>CO</i>	AT5G15840	<i>AtWRKY70</i>	AT3G56400
<i>SOC1</i>	AT2G45660	<i>AtWRKY75</i>	AT5G13080
<i>FT</i>	AT1G65480	<i>TaWRKY5</i>	EU665434.1
<i>API</i>	AT1G69120	<i>TaWRKY11</i>	EU665440
<i>SEP3</i>	AT1G24260	<i>OsWRKY45</i>	GQ331932
<i>AP3</i>	AT3G54340	<i>OsWRKY61</i>	BK005064
<i>AG</i>	AT4G18960	<i>OsWRKY71</i>	AB190817.1
<i>PI</i>	AT5G20240	<i>OsWRKY78</i>	BK005212
<i>SPL4</i>	AT1G53160	<i>OsWRKY89</i>	AY781112.1
<i>PRE1</i>	AT5G39860	<i>GmWRKY60</i>	EU375357
<i>EXL4</i>	AT1G75910	<i>GhWRKY15</i>	HQ651070.1
<i>EXL6</i>	AT1G75930	<i>TcWRKY53</i>	EF053036
<i>CYP77A6</i>	AT3G10570	<i>ScWRKY1</i>	AY366389
<i>ACS2</i>	AT1G01480	<i>SUSIBA2</i>	AY323206.1
<i>AtWRKY4</i>	AT1G80840	<i>MINISEED3</i>	NM_104436.2
<i>AtWRKY6</i>	AT1G62300	<i>MtSTP</i>	HM622067
<i>AtWRKY11</i>	AT4G31550	<i>TTG2</i>	AF516172

Gene-specific primers used for RT-PCR assays

Gene name	Primer Sequence (5' to 3')
<i>GsWRKY20</i>	Forward:GGCCATCAAGGATCACCATAAT-3' Reverse: GGCTCCAAGGATGTCTTTCTGA-3'
<i>GAPDH</i>	Forward: GACTGGTATGGCATTCCGTGT Reverse: GCCCTCTGATTCCCTCCTTGA
<i>ACTIN2</i>	Forward:TTACCCGATGGGCAAGTC Reverse: GTCATACGGTCAGCGATAC
<i>FLC</i>	Forward: GCTTTCTGTTCTCTGTGACGCA Reverse: CTCCAGTTGAACAAGAGCATCG
<i>CO</i>	Forward: GCTCCACACCATCAAACCTTACT Reverse: CGATTGGCAGAGTGAACCTTGAG
<i>SOC1</i>	Forward: ACACAAATAGATGAAACGAGGAAAG Reverse: GATAAAAACCTAACCAGGAGGAAGC
<i>FT</i>	Forward: ACAATCAACACAGAGAAACCACCT Reverse: CCTGAGGTCTTCTCCACCAATC
<i>AP1</i>	Forward: GCAAGCAATGAGCCCTAAAGAG Reverse: AGTGCGGATGTGCTTAAGAGC
<i>SEP3</i>	Forward: GCATGCTTCGGACACTGGA Reverse: GGCCTCTCTTGAAGGCACATT
<i>AP3</i>	Forward: GAGTGTTTGGACGAGCTTGACA Reverse CGCGAACGAGTTTGAAGTGTT
<i>AG</i>	Forward: CAACCGTTTGATTCACGGAA Reverse: GGCGGATGAGTAATGGTGATTG
<i>PI</i>	Forward: AAAATCTGATGGCTGTTCGAGC Reverse: CCATCTGGTGGTCTCGGACT
<i>SPL4</i>	Forward: GTAGCATCAATCGTGGTGGC Reverse: CTTCGCTCATTGTGTCCAGC
<i>PRE1</i>	Forward: CTTGGTCGTTTCCTTTTCAGCA Reverse: TTGATTATCGGAGATCCTTGGAG
<i>EXL4</i>	Forward: ACAGCAGGTCCTTCAAGAGCA Reverse: CCCAACGGTATCACTCCCATTA
<i>EXL6</i>	Forward: CGCTTTTGGCTTTTGGTGATTC Reverse: AACCTTCGGCAACTATATCGGT
<i>CYP77A6</i>	Forward: CGCTGCGTTTTGTATTCTTTTG Reverse: AACTTGTTTACAACGAACCTCG
<i>ACS2</i>	Forward: ACATCGCTAATTTCCAAGACTACCA Reverse: CTCTATCAAATGCGGCATAGTACG

Average rosette leaf numbers of WT and *GsWRKY20ox* plants at the time of flowering under GA3 treatment.



All values are means (\pm S.E.) from three independent experiments (at least 30 seedlings per experiment). Data were analyzed statistically using the t-test, double asterisks indicate significant differences from the corresponding WT at $P < 0.01$.