

<b>Cloning</b>			
<b>Fragment:</b>		<b>Forward</b>	<b>Reverse</b>
<i>IAA19</i> promoter		CACCGCGTCCGCTAAACACTCGTGC	CCTCTCACAATTTGATACAATTATGAAATGAGAAC
NotI- <i>GUS</i> -SacI		TTTTGCGGCCGCATGGTCCGTCTGTAGAAAC	CCCCGAGCTCTCATTGTTTGCCTCCCTG
<b>qPCR</b>			
<b>Gene:</b>	<b>AtCode</b>	<b>Forward</b>	<b>Reverse</b>
<i>AT1613320</i>	AT1613320	GTAGGACCGGAGCCAAGTCTAG	ACAGGGAAGAATGTGCTGGA
<i>SAG29</i>	AT5G13170	GCTGCATGCTATGGCTCTAC	GAGGGGAGGAGTTTTAACCA
<i>PRE1</i>	AT5G39860	CAAGGCAATCTTCAAGTGCT	CGCTCAGATTGTCAACTTCTC
<i>SAUR16</i>	AT4G38860	CGGTTTACGTAGGCGAAAAG	CACAAGGGATGGTGAGTCCT
<i>PIF4</i>	AT2G43010	CCAACCTCAGAGACGGTTAAG	GTTTCCCTAATCCCCGATG
<i>PIF5</i>	AT3G59060	CAACGGTCAGGATCTACTCG	TGGAGTTGCATTTGAAGTGA
<i>BSK5</i>	AT5G59010	CGAGCTGCTACTTGTGGATT	AGCCAAGCGGATCTATTGA

**Methods S1: Primers used in this study, related to Figures 2, S2, 3, S3, 4 and S4, and STAR Methods.**