

Supplement table 4.

Estimated period length of transgenic lines over-expressing bZIP transcription factors										
genes	At number	line	n	Period	± SE	comments	control plants			
							n	Period	±	SE
<i>bZIP48</i>	At2g04038	<i>C</i>	6	23.62	± 0.49	high RAE	17	25.01	±	0.66
		<i>G</i>	6	25.00	± 0.78		22	24.49	±	0.31
		<i>J</i>	8	24.74	± 1.19		7	24.54	±	0.58
<i>bZIP50</i>	At1g77920	<i>B</i>	11	24.79	± 0.37		12	23.87	±	0.16
		<i>D</i>	39	24.04	± 0.19		30	23.95	±	0.30
		<i>E</i>	6	24.63	± 0.24		11	24.66	±	0.20
<i>bZIP51/VIP1</i>	At1g43700	<i>E</i>	17	24.63	± 0.31		10	23.89	±	0.15
<i>bZIP58</i>	At1g13600	<i>C</i>	13	24.61	± 0.37		10	23.96	±	0.30