

Table 1 Suppl. The absorbance (A_{260}/A_{280}) ratios of RNA isolated from three biological replicates.

Treatments	Plants	Sample 1	Sample 2	Sample 3
Control	WT	2.13	2.10	2.12
	ERF96ox2-1	2.16	2.14	2.16
	ERF96ox11-5	2.13	2.09	2.08
NaCl	WT	2.09	2.10	2.12
	ERF96ox2-1	2.11	2.12	2.14
	ERF96ox11-5	2.14	2.08	2.13

Table 2 Suppl. Genes used to real time qPCR analysis, primers and different parameters.

Gene name	Primer sequences (forward and reverse)	Tm [°C]	Amplicon length [bp]	Amplification efficiency [%]
<i>ACT2</i>	GCAGGAGATGGAAACCTCAA CTTCTGGGCATCTGAATCTCTC	80.67	107	95.31
<i>ERF95</i>	TGATGAAGGACGGTCCAAATG CTCAATAACTTCCCTCCCATCATC	83.8	104	94.89
<i>ERF96</i>	GAGATACGAGATTCGAGGAAGC CTTGGCCTCTCATGGAGTAAG	84.85	112	96.77
<i>ERF97</i>	CTCAACTTCCCTCACGAGTATAAC CGCTATCGTCCAAGTACTCAAA	82.75	109	95.17
<i>ERF98</i>	TTTGGCTCGGGACATTGA GGGAAGTTGAGTATAGCGAGATG	83.8	97	94.49
<i>RD29A</i>	GATGACGAGCTAGAACCTGAAG CCTTTGTCCCTGGTGAATAA	85.17	106	92.38
<i>P5CS</i>	AGCAGCCTGTAATGCGATGG AAGTGACGCCTTTGGTTTGC	79.55	94	92.09
<i>COR15A</i>	GCCAGAAAACCTCAGTTCGTCG ATACGCCGCAGCTTCTCAGC	84.87	239	93.19
<i>KIN1</i>	TGGAGCTGGAGCACAACA GACCCGAATCGCTACTTGTTTC	85.32	105	94.24
<i>SKOR</i>	GGAGCACACCTTTACTAGAAGC GAAAGTCCCTGCGTTCTCTATG	81.44	94	90.18
<i>AKT2/3</i>	GGCCTCACTGCTTACCTTATT GACGCTGCTTCAATGCTATTC	81.89	95	93.58

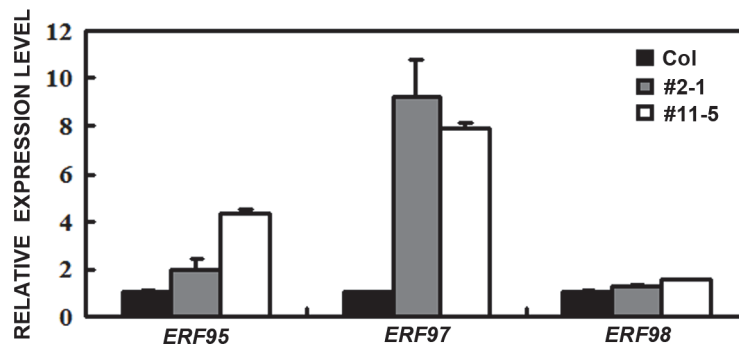


Fig. 1 Suppl. Expression of *ERF95*, *ERF97*, and *ERF98* in the *35S:HA-ERF96* transgenic plants. Total RNA was isolated from 10-d-old *35S:HA-ERF96* transgenic seedlings (#2-1 and #11-5), and real time qPCR was used to examine the expression of *ERF95*, *ERF97*, and *ERF98*. The expression of *ACT2* was used as an internal control, and the expression level of corresponding small *ERF* genes in the Col wild type seedlings was set at 1. Means $\pm$ SDs of three replicates.