

Supplemental Table S2. Primers used in this study

Name	Use	Primer sequence (5' to 3')
<i>MsGSTU17</i>	Full-length cDNA cloning	ATGAGTCAATATATGGCTG CTATTCAAGTCCATGCTTC
<i>MsGSTU17</i>	qRT-PCR	GGACAGGAAAGGGTGAAGAA GCCACATCAACATACCCAAAG
<i>AtACTIN</i>	qRT-PCR	TGTGCCAATCTACGAGGGTTT TTTCCCGCTCTGCTGTTGT
<i>AtSOS1</i>	qRT-PCR	TTGGCTATGGTTTGGATTGGA ATGCGAAGAAGGCGTAGAACA
<i>AtSOS2</i>	qRT-PCR	GCCACATCAACATACCCAAAG GTGCCACCTCGTAAATCTCTATC
<i>AtSOS3</i>	qRT-PCR	CATCCGTCACGCCATTAC TCACTCCATTTGCTTCACATC
<i>AtRD29A</i>	qRT-PCR	GTTACTGATCCCACCAAAGAAGA GGAGACTCATCAGTCACTTCCA
<i>AtKIN2</i>	qRT-PCR	TATATCGGATGCGGCAGTGG ACAACAAGTACGATGAGTACGAGA
<i>AtHKT1</i>	qRT-PCR	AAACGGCGAGAGATGTTCTTAGT GGAGCCAGATGAGACCAGAGTT
<i>MsACTIN</i>	qRT-PCR	CAAAAGATGGCAGATGCTGAGGAT CATGCACCAGTATGACGAGGTCG