

## Differentially Expressed Genes between Drought Tolerant and Sensitive Barley Genotypes in Response to Drought Stress during the Reproductive Stage.

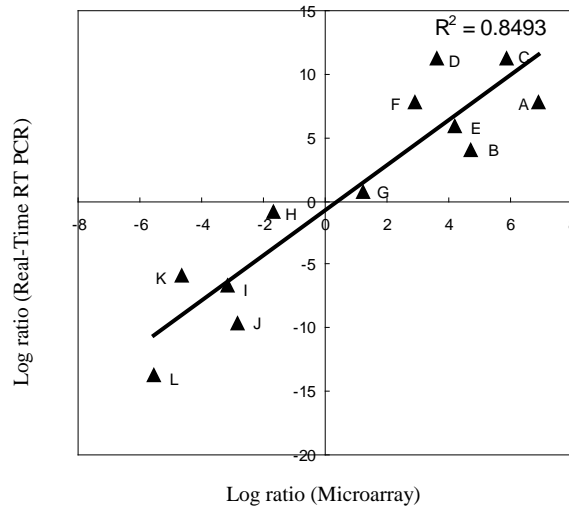
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### Supplementary Table S3 Oligonucleotide Sequences Used for Real Time RT-PCR in this Study.

Primers were designed by using Primer Express<sup>TM</sup> 1.5 software (Applied Biosystems, Foster, CA) as the manual instruction. The primer combinations which unambiguously amplified only one desired band in an agarose gel were selected as candidates for real time RT-PCR analysis. The dissociation curve analysis was carried out for each primer pair. The primer combination was used for real-time RT-PCR analysis only when its amplification product had one melting temperature.

Probe set	Annotation	Primer sequence (5'→3')
Contig10522_at	gamma-aminobutyric acid (GABA-A) receptor	CTACGGCGAAGGTCATGGATAT TGAAGCTCGAGACAACCATACTTTA
Contig11696_at	putative peptide chain release factor subunit 1	GAGAACAAGGAGGTGGAGAGAAGA TTCCAACCTCCACATTAAGCTTCAA
Contig1394_s_at	actin	TCTGTTGACTGTATTGCCGGATTA TTCAGCACCGATCAAATATAGCA
Contig16113_at	putative protein	TAGGATAGAAGAAGGTGTCGGAAAAT AAAACCTCTACCTGCAGCAGCTACA
Contig1713_s_at	dehydrin; DHN4	CACCGTTGCAGAATCAATAATAAGA CATTCGCCCTCCAACAG
Contig19029_at	bacterial-induced peroxidase precursor	GAGATCAGGAAAACTGCCATGTC GCACGCCCTGGGTATCG
Contig24415_at	<none>	ATTTTTGCTCTCGGTCCCAT GCAATAGGGAGCGTATAACACAAG
Contig26204_at	kaurene synthase	GTGGCAAGAATTGCTACGGAATAT TGTCACAACCGCATTTTCTATGTA
Contig3814_at	delta 1 pyrroline-5-carboxylate synthetase	TTCTAACCCTCGGTGCATTCTAC CTACCCTCATTGCAAAGGAAGATC
Contig422_at	chlorophyll a/b-binding protein WCAB precursor	CTGAGTTGAGAGCTGGCAGGAT GAGAGTGAATGTTGGCAAATCAAA
Contig6539_s_at	Beta-fructofuranosidase	GGGAAGATCTCCTTGAGAAGCTT GCCATGGACGGGTACACTCT
Contig7359_at	cell death suppressor protein	GTGTGGAGACATCACACGAGATC AGTATAAACATGCTGGGACAATCTCA
HVSMEi0006K11r2_s_at	NADP dependent malic enzyme	GCTCGGTTGAACATAGTGTAATTTTG GTCGGTCTGCCATTGCAAT

Supplementary Figure S1



**Supplementary Figure S1** Confirmation of Microarray Data by Real-Time RT-PCR. Correlation analysis of selected data between microarray and real-time RT-PCR experiments. The fold changes in gene expression in response to drought stress were  $\log_2$  transformed. The microarray data  $\log_2$ -values (X-axis) were plotted against the real-time RT-PCR  $\log_2$  values (Y-axis). Letter A to L in the figure represent the contig name of Contig3814\_at, Contig11696\_at, Contig1713\_s\_at, Contig10522\_at, Contig24415\_at, HVSMEi0006K11r2\_s\_at, Contig7359\_at, Contig6539\_s\_at, Contig26204\_at, Contig16113\_at, Contig19029\_at, Contig422\_at, respectively, in Affymetrix barley 1 GeneChip.