

Figure S1. Phenotypic differences between IR64 and DK151 under control and drought conditions.

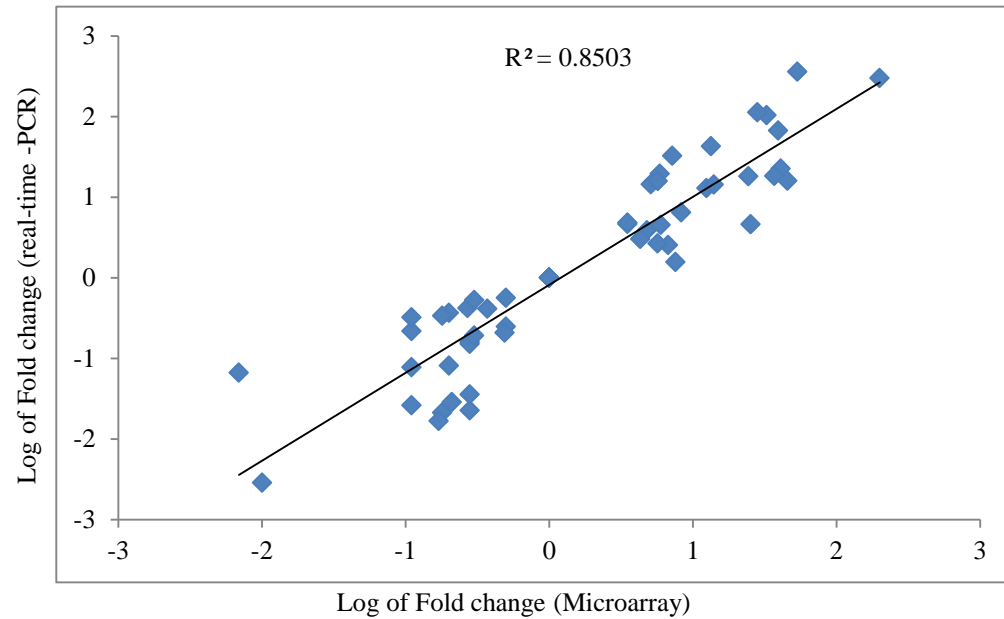


Figure S3. Validation of the expression of 49 random selected genes by qRT-PCR. Correlation analysis shows a good agreement between microarray and qRT-PCR experiments. The fold change values were transformed to log scale.