**Figure S3.** Transcript abundance of the genes of interests and the reference gene *UBI3* without normalization.

Transcript abundance of the five selected genes and the reference gene *UBI3* was determined by quantitative RT-PCR. Equal amounts of RNA were added to the RT reaction. Three independent experiments were used to determine the standard deviations.

D:\sandra_doktorarbeit\missbach_publications\missbach et al 2012a\plant physiology\submission\Fig S3.tif