S8 Fig. Expression of 23 GRF9 early responding genes in different GRF9-modified lines. Gene expression as determined by Affymetrix ATH1 microarray hybridizations (first two columns) or qRT-PCR (other columns). RNA for expression analysis was obtained from 2-week-old GRF9-IOE seedlings grown on MS medium and induced with 10 µM estradiol for the indicated time points (0.15% [v/v] ethanol as control), or from 2-week-old GRF9ox and grf9-1 seedlings grown on MS medium (WT as control). Values represent the means of replicates obtained from three sets of seedlings (except for the microarray data where each value represents one replicate).