Table S3. *GRF* expression in 35S:*miR396b* plants compared to that in wild type, as estimated by Affymetrix microarray

Description	Relative expression*
GRF1	0.81
GRF2	0.58
GRF3	0.73
GRF4	WA
GRF5	0.89
GRF6	A
GRF7	0.57
GRF8	A
GRF9	NP

*Fold change relative to wild type, normalized with gcRMA. The average of two biological replicates for each genotype is shown.
A, a gene termed 'absent' by MAS 5.0 software (Affymetrix); NP, not present in ATH1 arrays; WA, wrongly annotated in ATH1 arrays.