



Figure S4. SCAP1 regulates *AtMYB60* expression

(a) *AtMYB60* accumulation determined by quantitative PCR in manually dissected first two-leaf primordia of wild-type or *scap1* seedlings at different time points. *ACTIN* (*ACT2*) was used for normalization. Values represent the mean of three biological replicates (30 leaves / replica). Error bars = standard deviation.

(b) *SCAP1* transcript accumulations determined by RT-PCR in *pro35S:SCAP1-GR* T1 lines. Total RNA was isolated from 2-week-old seedlings and PCR was conducted for 30 cycles. Actin was used as a positive control and amplified for 25 cycles.

(c) GUS staining of double (*proAtMYB60:GUS pro35S:SCAP1-GR*) hemizygous lines. Shown are 2 weeks-old seedlings (inset, higher magnification). Bar = 1 mm (inset, 500 μ m).