



Figure S1. Characterisation of the *scap1* and *Atdof4.1* mutant alleles

(a) Schematic representation of the *SCAP1* and *AtDof4.7* loci. Grey boxes represent amiRNAs target regions and triangles represent transposon or T-DNA genomic insertion points for *scap1-2* and *dof4.7*, respectively.
 (b) Reverse Transcriptase-PCR analysis of *SCAP1* in wild type (Ler), *scap1-2*, and *AtDof4.7* in wild type (Col) and *dof4.7* plants. Total RNA was isolated from 2-week-old seedlings and PCR was conducted for 35 cycles. Actin was used as a positive control and amplified for 25 cycles.