



Supplemental Figure S2. Quantitative RT-PCR analysis of expression levels for IAA transgenes in Arabidopsis seedlings.

- (A) Relative *35S:IAA17m/mlII-1* (b) transgene expression compared to *IAA17* gene expression in nontransformed seedlings (a).
- (B) Relative *35S:IAA3m/mlII-1* transgene expression (b) compared to *IAA3* gene expression in nontransformed seedlings (a).
- (C) Relative *35S:IAA6m/mlII-1* transgene expression (b) compared to *IAA6* gene expression in nontransformed seedlings (a).
- (D) Relative *35S:IAA19m/mlII-1* (b), *IAA19:IAA19m/mlII-1* (c) and *IAA19:VP16-IAA19m/mlII-1* (d) transgene expression compared to *IAA19* gene expression in nontransformed seedlings (a).
- (E) Relative *35S:IAA12m/mlII-2* (b) and *35S:IAA12m/mlII-3* (c) transgene expression compared to *IAA12* gene expression in nontransformed seedlings (a).