

The tomato (*Solanum lycopersicum* cv Micro-Tom) natural genetic variation *Rg1* and the DELLA mutant *procera* control the competence necessary to form adventitious roots and shoots.

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Supplementary Material

Table S1. Primers used in the RT-PCR analysis

Primer	Sequence	SGN Accession*
GAPDH F	TCCATCACAGCCACTCAGAA	Solyc05g014470
GAPDH R	TCAACCACGGACACATCAAC	
GAMyb1 F	AACCACATCATCAGGTCAAGC	Solyc01g009070
GAMyb1 R	GCACACAGGAAAATCCAGAAG	
GOB F	CTCCTCGATTCCCTCTCCGTA	Solyc07g062840
GOB R	AGTGCATGCAAGGTGTTGA	
miR164 F	CCTGAGTGGAGAACGCAGGGCA	
Universal R	GTGCAGGGTCCGAGGT	At2g47585
miR164 RT	GTCGTATCCAGTGCAGGGTCCGAGGT ATTCGCACTGGATACGACTGCACG	
TKn1 F	GTGAAGGCCCTTTCGTTTG	Solyc04g077210
TKn1 R	GCTTAGGCCTATAAGGCTTGCT	
TKn2 F	GAGGCATTGAAACCATCAG	Solyc02g081120
TKn2 R	CATGCACACAAGTAATATGCAA	
PTS F	TCTAATTGATGGATGAAGG	Solyc06g072480
PTS R	TACGAGTTGTGGCAATCTAGC	
SIPIN4 F	TGCGGTTATTTTGGGATGT	Solyc05g008060
SIPIN4 R	TTCTTGTCTTAATGCTCCAAA	
AXL F	CGAGACAATTCCATTCCATGT	Solyc11g072460
AXL R	GCAACTTGCTTTCACTTGC	
IAA11 F	GTTTAACGGACGTGAAGCTG	Solyc03g120380
IAA11 R	TTATCTGCATCCTCCAATGCT	

*As annotated in the Sol Genomics Network - SGN (<http://solgenomics.net/search/loci>)

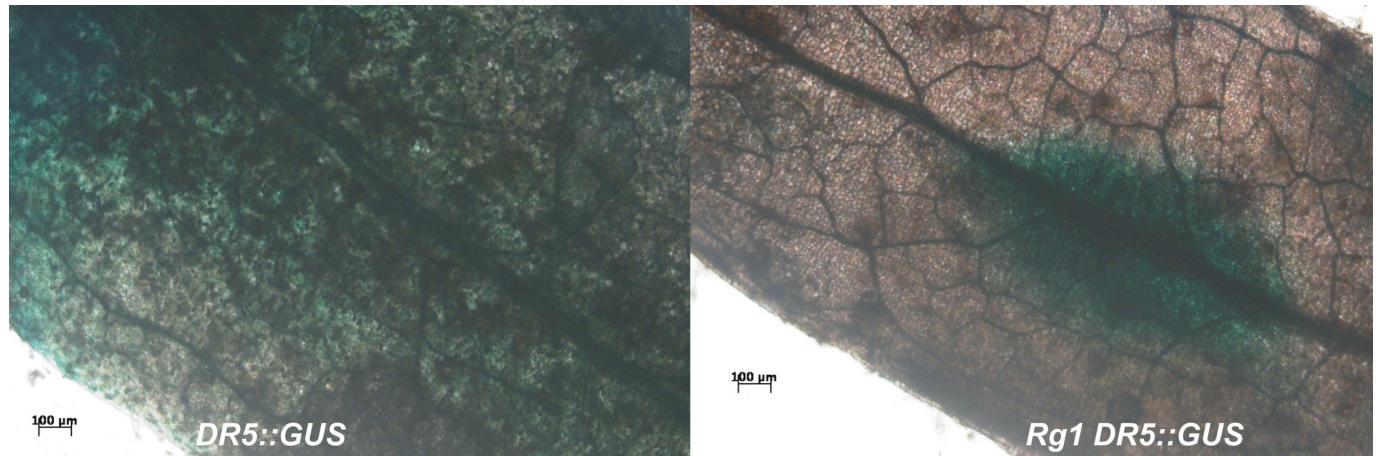


Fig. S1. GUS staining of *DR5::GUS* cotyledons in MT (left) and *Rg1* (right) genetic background. Eight-d-old cotyledons were cultivated on RIM (0.4 μ M NAA) during 2 d.