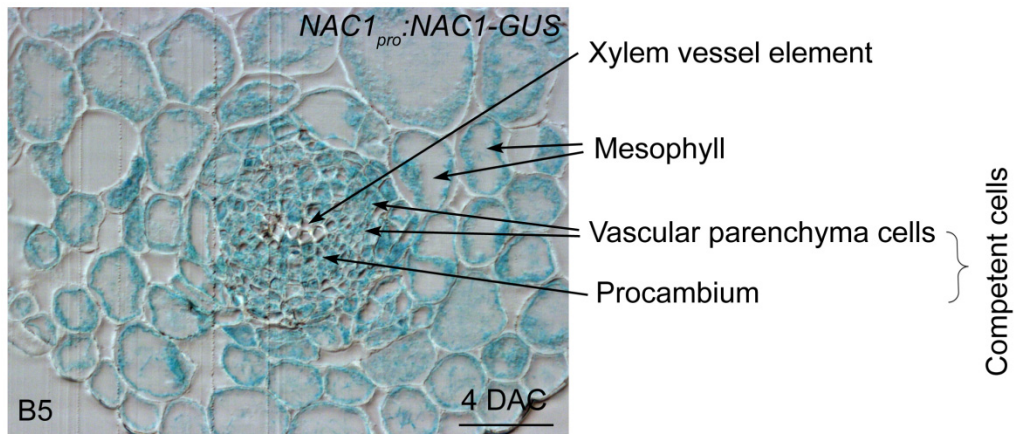


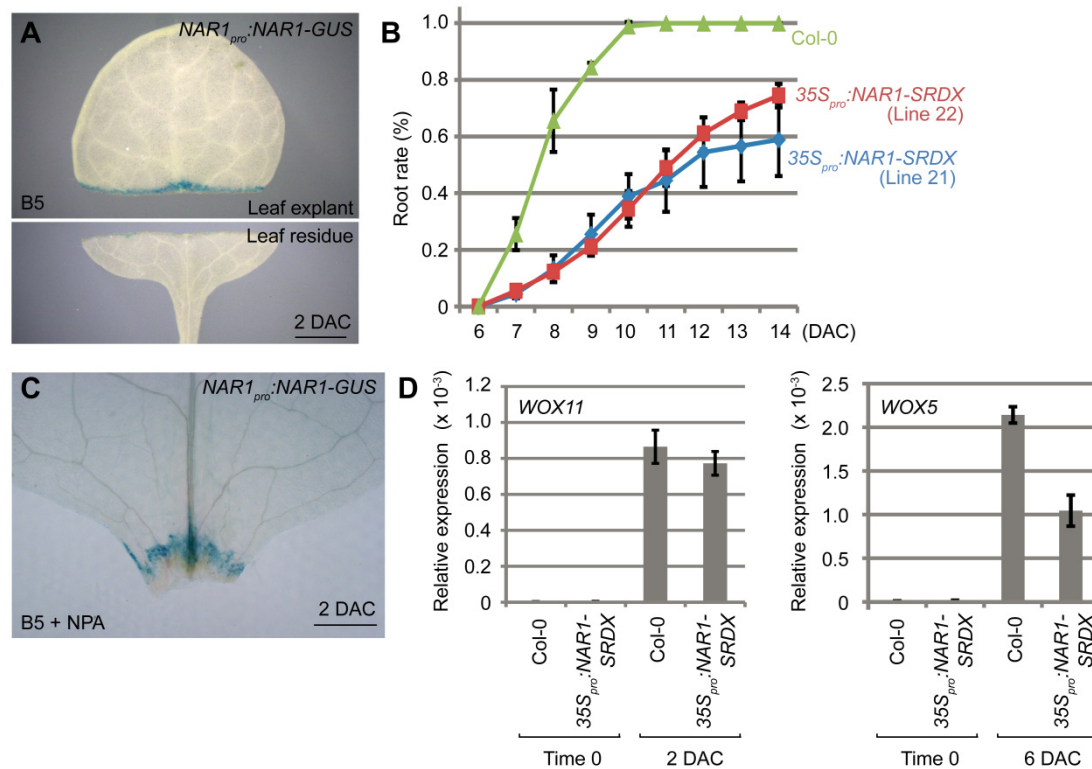
Supplementary data



Supplemental Figure S1. *NAC1* expression in the leaf explant.

Thin sectioning of 4-DAC leaf explants from *NAC1_{pro}:NAC1-GUS* cultured on B5 medium at the wounded region, showing the midrib of vasculature.

Scale bars, 50 μm .



Supplemental Figure S2. *NAR1* in *de novo* root organogenesis.

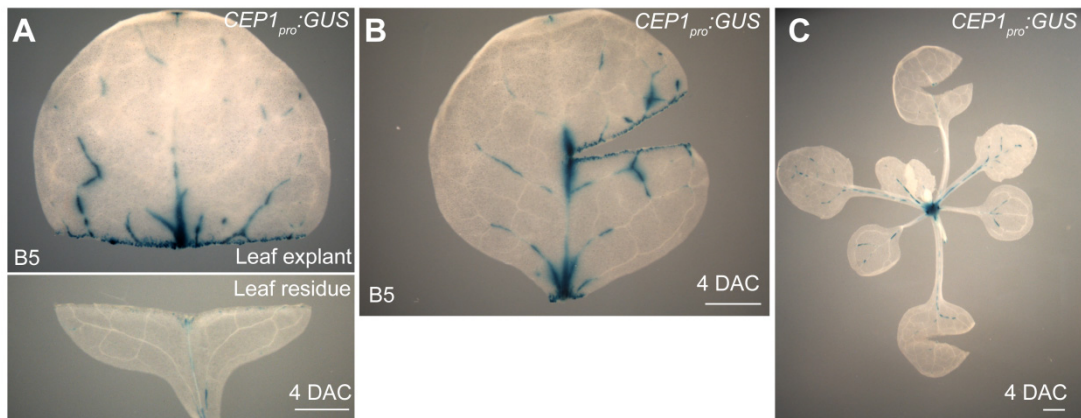
(A) GUS staining in leaf explant (upper panel) and wounded leaf residue (lower panel) of *NAR1_{pro}:NAR1-GUS* at 2 DAC.

(B) Rooting rate analyses of leaf explants from Col-0 and *35S_{pro}:NAR1-SRDX*. Bars show SD from three biological repeats, $n = 30$ in each repeat.

(C) GUS staining of *NAR1_{pro}:NAR1-GUS* in leaf explant at 2 DAC cultured on B5 medium containing NPA.

(D) qRT-PCR analysis of *WOX11* and *WOX5* transcript levels in leaf explants from Col-0 and *35S_{pro}:NAR1-SRDX*. *WOX5* was upregulated in *35S_{pro}:NAR1-SRDX* leaf explants at 6 DAC, although the upregulation level is lower than that in the wild-type leaf explants. This might be caused by the arrest of the process from root apical meristem formation to root tip emergence in *35S_{pro}:NAR1-SRDX*. Bars show SD from three PCR experiments. Results were confirmed in two biological repeats. Data from one biological repeat are shown.

Scale bars, 1 mm in (A) and 500 μm in (C).



Supplemental Figure S3. *CEPI* expression responds to wounding of leaf explant.

(A) GUS staining of *CEPI_{pro}::GUS* leaf explant (upper panel) and wounded leaf residue (lower panel) at 4 DAC

(B) Large wound on leaf explant from *CEPI_{pro}::GUS* cultured on B5 medium, showing GUS signal in wounded region.

(C) Large wound on seedling of *CEPI_{pro}::GUS*, showing no GUS signal in wounded region.

Scale bars, 1 mm in (A-C).

Supplemental Table S1. RNA-seq analysis of genes expressed in response to wounding.

Supplemental Table S2. RNA-seq analysis of genes upregulated in *pER8:3×FLAG-NAC1*.

Supplemental Table S3. List of primers used in this study.

Experiments	Primers	Sequence (5' →3')
Molecular cloning		
<i>NAC1_{pro}:NAC1-GUS</i>	NAC1 _{pro} :NAC1-F1	ccgctcgagGAGTCTGATCATAGGCAC ACG
	NAC1 _{pro} :NAC1-R1	gaagatctGCAATTCCAAACAGTGCTT G
<i>NAR1_{pro}:NAR1-GUS</i>	NAR1 _{pro} :NAC1-F1	acgcgtcgacGGAGAGAAGTGGCAATG AGAG
	NAR1 _{pro} :NAC1-R1	gctctagaGTAATTCCATGCATGGCTTG
<i>CEP1_{pro}:GUS</i>	CEP1 _{pro} :GUS-F1	ccgctcgagTGTTATAATCTGATTAGAC TAA
	CEP1 _{pro} :GUS-R1	tccccgggATTTTCTATGGTGATTTT TGTATTTACTTGTG
<i>EXT1_{pro}:GUS</i>	EXT1 _{pro} :GUS-F1	acgcgtcgacGTAGGATTTTGATGGCAG TGTTG
	EXT1 _{pro} :GUS-R1	cgggatccTGTGTTATGTGTTTTGCTTT GTG
<i>NAC1_{pro}:NAC1-SRDX</i>	NAC1 _{pro} :NAC1-SRDX-F1	ccgctcgagGAGTCTGATCATAGGCAC ACG
	NAC1 _{pro} :NAC1-SRDX-R1	ccaatgcattggttctgcagTCAAGCGAAACCC AAACGGAGTTCTAGATCCAGATCGAG GCAATTCCAAACAGTGCTTG
<i>35S_{pro}:NAC1-SRDX</i>	35S _{pro} :NAC1-SRDX-F1	gaagatctATGGAGACGGAAGAAGAG
	35S _{pro} :NAC1-SRDX-R1	acgcgtcgacTCAAGCGAAACCCAAACG GAGTTCTAGATCCAGATCGAGGCAAT TCCAAACAGTGCTTG
<i>35S_{pro}:NAR1-SRDX</i>	35S _{pro} :NAR1-SRDX-F1	tccccgggATGGAGGAGACAGAAAAG AATAAG

	35S _{pro} :NAR1-SRDX-R1	acgcgtcgacGTAATTCCATGCATGGCTTG
<i>pER8:CEP1</i>	pER8:CEP1-F1	ccgctcgagATGAAGCGATTTATTGTCTTGC
	pER8:CEP1-R1	tccgggcccTTAGAGTTCATCCTTAAGCGAGTCC
<i>pER8:CEP2</i>	pER8:CEP2-F1	ccgctcgagATGAAGAACTTCTTTTGA TATTTC
	pER8:CEP2-R1	tccgggcccCTAGAGCTCATCTTTGACATCAC
<i>pER8:3×Flag-NAC1</i>	pER8:3×Flag-NAC1-F1	acgcgtcgacATGGACTACAAAGACCATGACGGTGATTATAAAGATCATGATATCGATTACAAGGATGACGATGACAAGggggcccATGGAGACGGAAGAAGAGATGAAGG
	pER8:3×Flag-NAC1-R1	gactagtTCAGCAATTCCAAACAGTGC TTG
qRT-PCR		
<i>ACTIN</i>	ACT-F	TGGCATCA(T/C)ACTTTCTACAA
	ACT-R	CCACCACT(G/A/T)AGCACAATGTT
<i>NAC1</i>	NAC1-qRT-F1	CATCCTCCCAATCATTCTCTG
	NAC1-qRT-R1	GAATGAGTCGAGGCCTGTGA
<i>NAR1</i>	NAR1-qRT-F2	ATACGCAACAGGCTACAGAACA
	NAR1-qRT-R2	TGTCGGTCAATTGGCTCTCT
<i>WOX11</i>	WOX11-F3	CGCAACCACCAACACTTGTGACC
	WOX11-R6	AAGACATCTGTTGCATCACC
<i>WOX5</i>	WOX5-RT-F	GTGAAAGGTCTGAAGCTTACG
	WOX5-RT-R	GTACTGGTTATTGCCTCTAGC
<i>CEP1</i>	CEP1-qRT-F1	GACTCACAAGTGAGCTAGTG
	CEP1-qRT-R1	CTACTACCGCAACTCCATGG

<i>CEP2</i>	CEP2-qRT-F1	CGTGCTTGTGACGATAGATG
	CEP2-qRT-R1	CACACCATGGTTCAGCTCTG

Note that lower case letters represent additional nucleotides to introduce restriction sites, and the italic letters indicate SRDX or 3×FLAG.