

Article Title:

Plasmodesmata callose binding protein 2 contributes to the regulation of cambium/phloem formation and auxin response during the tissue reunion process in incised Arabidopsis stem

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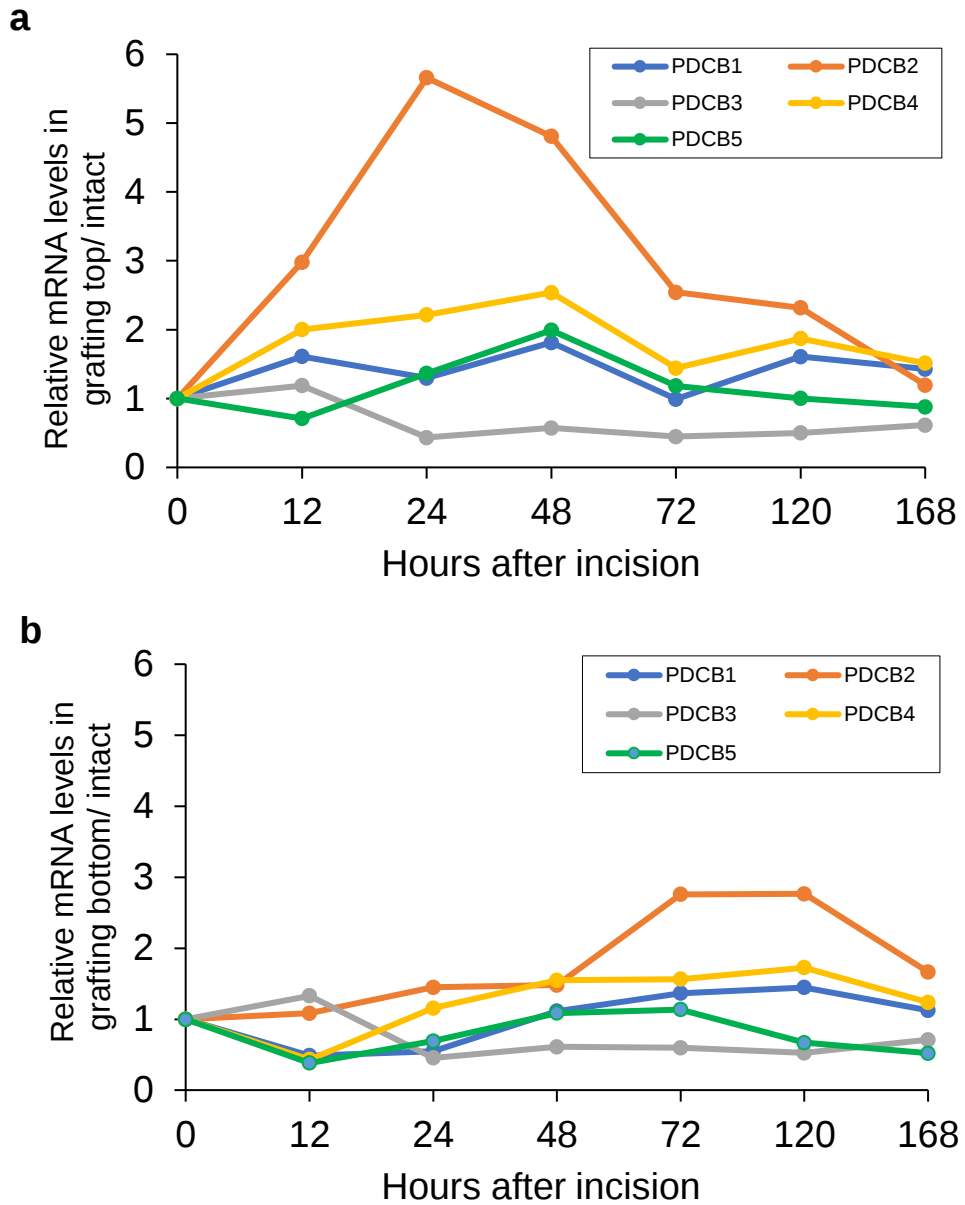


Fig. S1. *PDCB* mRNA levels at the top and bottom of grafted intact stem (Melnyk *et al.*, 2018; GSE107203).

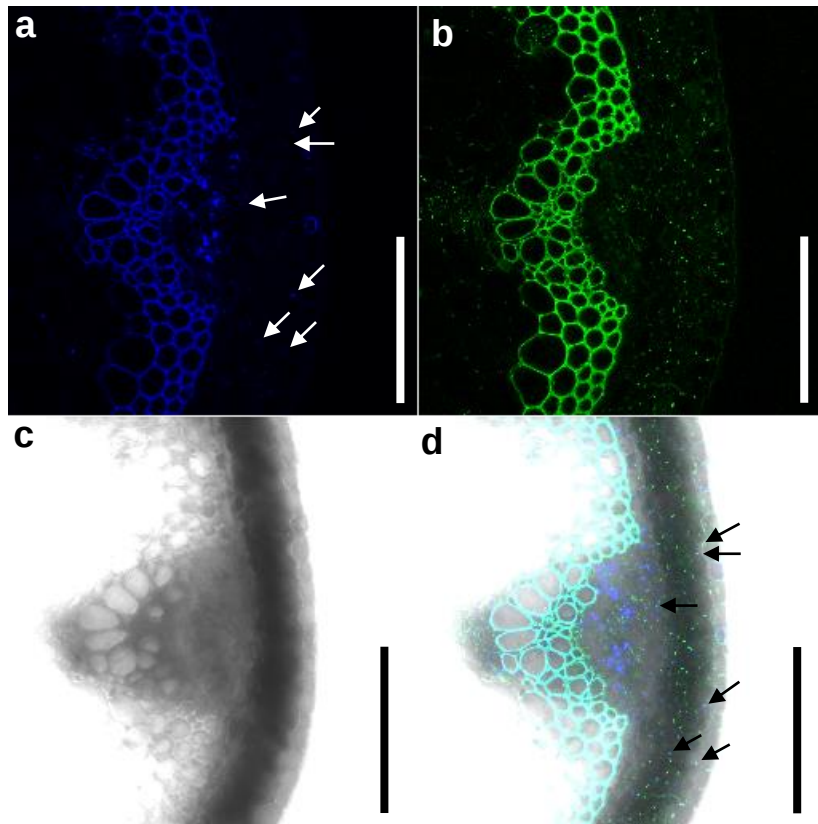


Fig. S2. Aniline blue and 35S::MP17-GFP fluorescence were coincident.

(a) Aniline blue staining, (b) 35S::MP17-GFP, (c) Bright field, (d) Merge. Bars = 100 μm .

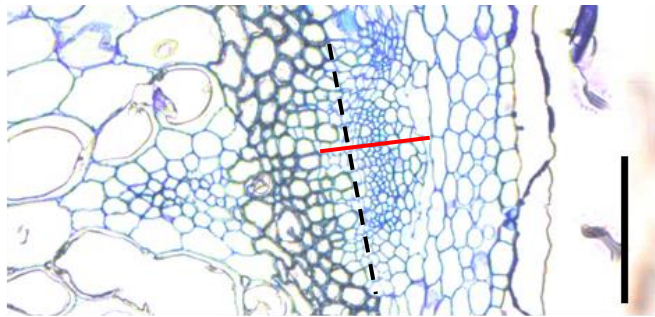


Fig. S3. Analysis of the number of cells and length of the cambium/phloem region

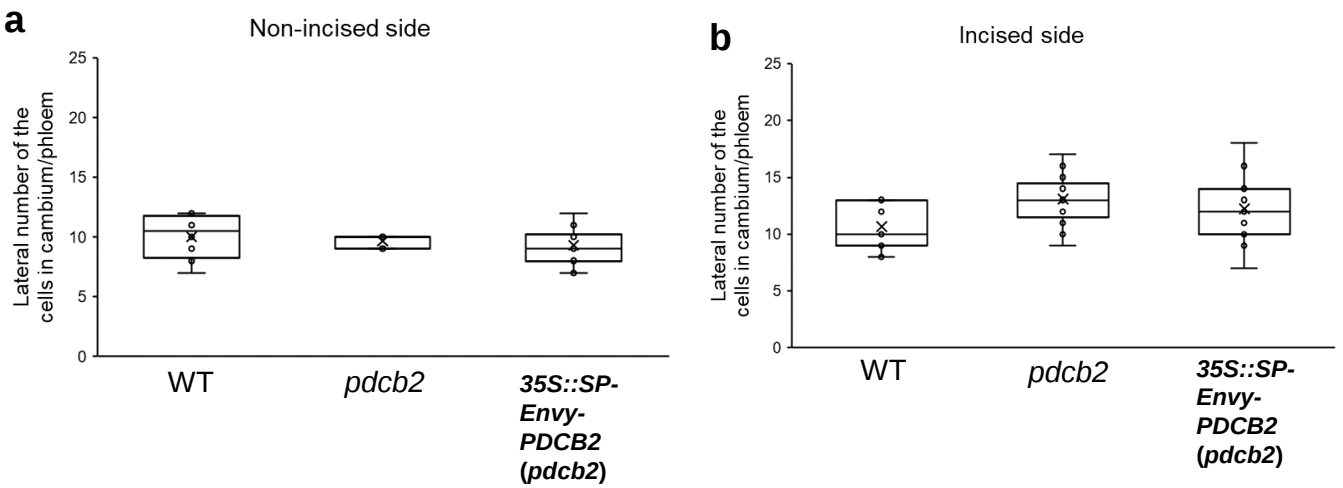


Fig. S4. Measurement of the lateral number of the cells in non-incised and incised side 7 days after incision. Values are means boxplot (WT: n=15 from six plants, *pdcb2*: n=13 from five plants, *35S::SP-Envy-PDCB2 (pdcb2)* : n=21 from seven plants). Statistical analysis was performed by Welch's t-test.

Table S1. Primer used in this study

For qRT-PCR			
AGI	Gene	Forward primer	Reverse primer
AT3G18780	ACT2	5'-CTGGATCGGTGGTTCCATTC-3'	5'-CCTGGACCTGCCTCATCATAC-3'
AT5G08000	PDCB2	5'-GTCC AAAAGGAAGCAACAGC-3'	5'-CCGGTTCCAGTAGCATCAGT-3'
AT5G61480	TDR/PXY	5'-ACATCGGACATTGCGAGAAG-3'	5'-GATCTACGTCGGCATTGAC-3'
AT1G46480	WOX4	5'-AGGTGGAACCCGACTCAAG-3'	5'-CGTACTTACCGAGTTGCAATGTG-3'
AT1G70940	PIN3	5'-TCTTTGATTAGGTTCGGGTAACTC-3'	5'-GCTCATGTGAACTTGGAAACAAG-3'
For genotyping			
AGI	Gene	Forward primer	Reverse primer
AT5G08000	PDCB2	5'-CCTGCAACTCTGCAACACAT-3'	5'-CCAGAGCTGAATGTGAAAGG-3'
	LBb1,3	5-ATTTTGCCGATTTTCGGAAC-3'	-
For cloning			
Primer name		Forward primer	Reverse primer
AtX8-GPI-Pro		5'-CATGTGTGTAGTCAACGCTCTAG-3'	5'-GGTTGGAGTTCAGAGATACAAATG-3'
Envy		5'- GTGACCACCCTGTGCTACGGCGTGCAGTGCTTCG CCCGCTACCCC-3'	5'-GTTGTGGCGGGTCTTGAAGTTG-3'
PDCB2-SP		5'- GCTCGATCCACCTAGGCTATGGCTCCTCTGGTTC TCTACCTTCTCACTCTCCTCATGGCTGGCCATAC CAGTGCCGGCCGGCCTGGA-3'	5'- TCCAGGCCGGCCGGCCACTGGTATGGCCAGCCAT GAGGAGAGTGAGAAGGTAGAGAACCAGAGGAGC CATAGCCTAGGTGGATCGAGC-3'
Envy-FP3		5'- GCCGGCCGGCCTGGAGTGAGCAAGGGCGAGGAG C-3'	-
GFP-R-linker		-	5'- AGAACCACCACCACCAGAACCACCACCACCCTT GTACAGCTCGTCCATGCCGAG-3'
PDCB2-Fj		5'- GGTGGTGGTGGTTCTGGTGGTGGTGGTTCTGCCTC ATGGTGTGTGTGCAA-3'	-
PDCB2-RP4		-	5'- CGTAGCGAGACCACAGGATCAGAGCATCATCAC GAGA-3'
BPr		5'- GGGGACAAGTTTGTACAAAAAAGCAGGCTGCTCG ATCCACCTAGGCT-3'	5'- GGGGACCACTTTGTACAAGAAAGCTGGGTCTGTAG CGAGACCACAGGA-3'