

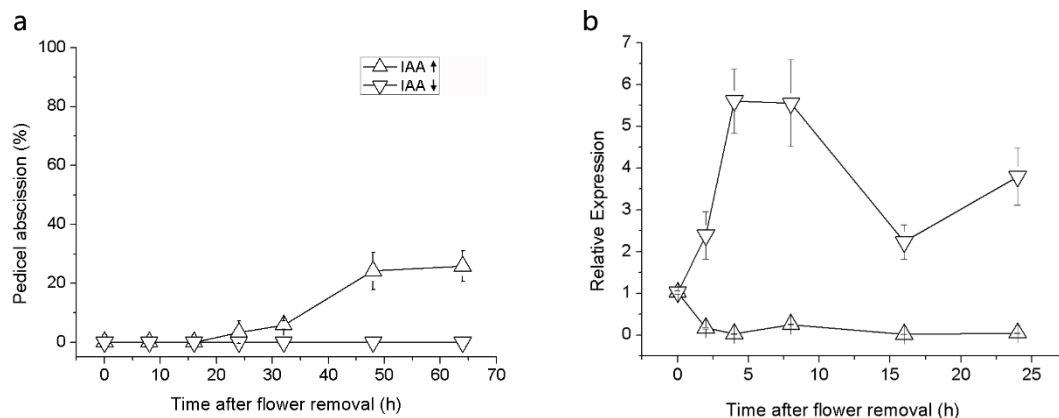
SIPIN1 regulates auxin efflux to affect flower abscission process

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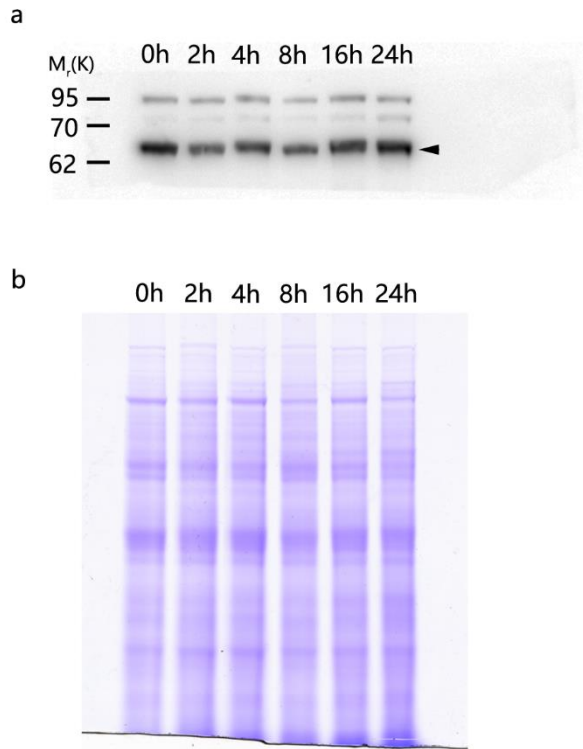
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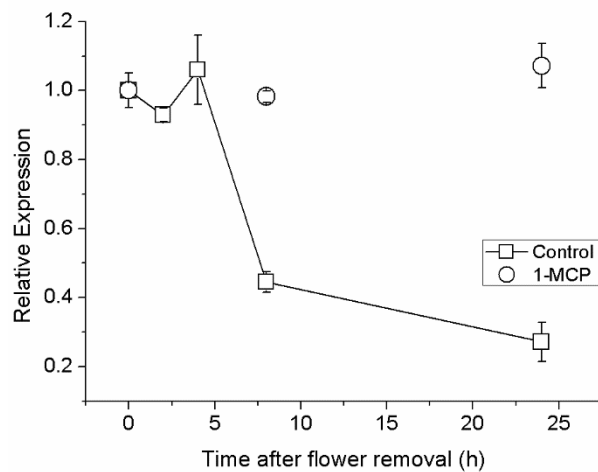
Supplementary information



Supplementary Figure S1. Abscission process and *SIPIN1* expression profiles following the IAA treatments at different ends. The arrows indicate the opposite directions of incubation. The data are presented as the mean (\pm SD) values corresponding to three independent tests. For each test of abscission rate, 40 to 50 pedicels were investigated.



Supplementary Figure S2. Full-length images in SIPIN1 western blotting experiments. a, SIPIN1 western blotting of the AZ protein samples as described in Fig. 3 a. A specific signal among the 62-70 KD (arrowhead) was detected. b, the Coomassie-staining of the AZ protein samples.



Supplementary Figure S3. *FEI2* expression profiles in the AZ during abscission. *FEI2* expression decreased at 8-24 h after flower removal, while 1-MCP treatment prevented the down-regulation of *FEI2* during abscission. The data are presented as the mean (\pm S D) values corresponding to three independent tests.

Supplementary Table S1. Primers used in this study

Primer name	Sequence (5' -3')
Forward primer of <i>SIPIN1</i> for RT-PCR	CAGGCAGCTCTACCACAAGG
Reverse primer of <i>SIPIN1</i> for RT-PCR	TGTAATCGGCAACGCAATC
Forward primer of <i>FEI2</i> for RT-PCR	TCTGGTCTCCACTGCTCAAGA
Reverse primer of <i>FEI2</i> for RT-PCR	CCAAGGGAACCGTAGCACAG
Forward primer for vector pSPT19 construction (Eco RI)	GG <u>GAAATTC</u> ACTTATTGTTCTTGGGGTTCTTGCT
Reverse primer for vector pSPT19 construction (Hind III)	GG <u>AAGCTT</u> TTTTCCATCTTCTTTAACTCTGCT
Forward primer of <i>SIPIN1</i> for VIGS construction (Eco RI)	CCG <u>GAAATTC</u> TGGTACTGCTGGGAGAACTCAAAC
Reverse primer of <i>SIPIN1</i> for VIGS construction (BamHI)	CGC <u>GGATCC</u> GCTTTGTTGCCATTGTTACTAGGTG
Forward primer of <i>SIPIN1</i> for bait construction of Y2H (Eco RI)	CCG <u>GAAATTC</u> AGAGGTGCAAGACTGCTAATTTCT
Reverse primer of <i>SIPIN1</i> for bait construction of Y2H (BamHI)	CGC <u>GGATCC</u> GGACTGTGCAATTATAGCAGGC
Forward primer of <i>SIPIN1</i> for vector construction of BiFC (KpnI)	CGG <u>GGTACC</u> AGAGGTGCAAGACTGCTAATTTCT
Reverse primer of <i>SIPIN1</i> for vector construction of BiFC (SpeI)	CGG <u>ACTAGT</u> GGACTGTGCAATTATAGCAGGC
Forward primer of <i>PP2C</i> for vector construction of BiFC (KpnI)	CGG <u>GGTACC</u> ATGACTGGAGGCAAAGAAATCATCG
Reverse primer of <i>PP2C</i> for vector construction of BiFC (Eco RI)	CCG <u>GAAATTC</u> TTGGAACCTTACAACACTACGCAGG

Supplementary Table S2. Phosphopeptides of SIPIN1

Phosphorylated peptide and site (in lowercase)	pRS	Mascot	Relative quantity	
	Score	Score	12h/0h	24h/0h
DDFsFANR	108	35	0.582	0.378
FNNYHGGAAPQsNsNtHyPAPNPGMFsPsNNGNk	57	11	0.700	0.761

