

Supporting Information

Article title: CARK1 phosphorylates ABA receptors subfamily III members

Authors: Xiaoyi Li[¶], Xiangge Kong[¶], Qi Huang[¶], Hu Ge, Qian Zhang, Liang Zhang, Gaoming Li , Lu Peng, Zhibin Liu, Jianmei Wang, Xufeng Li, Ying Yang*

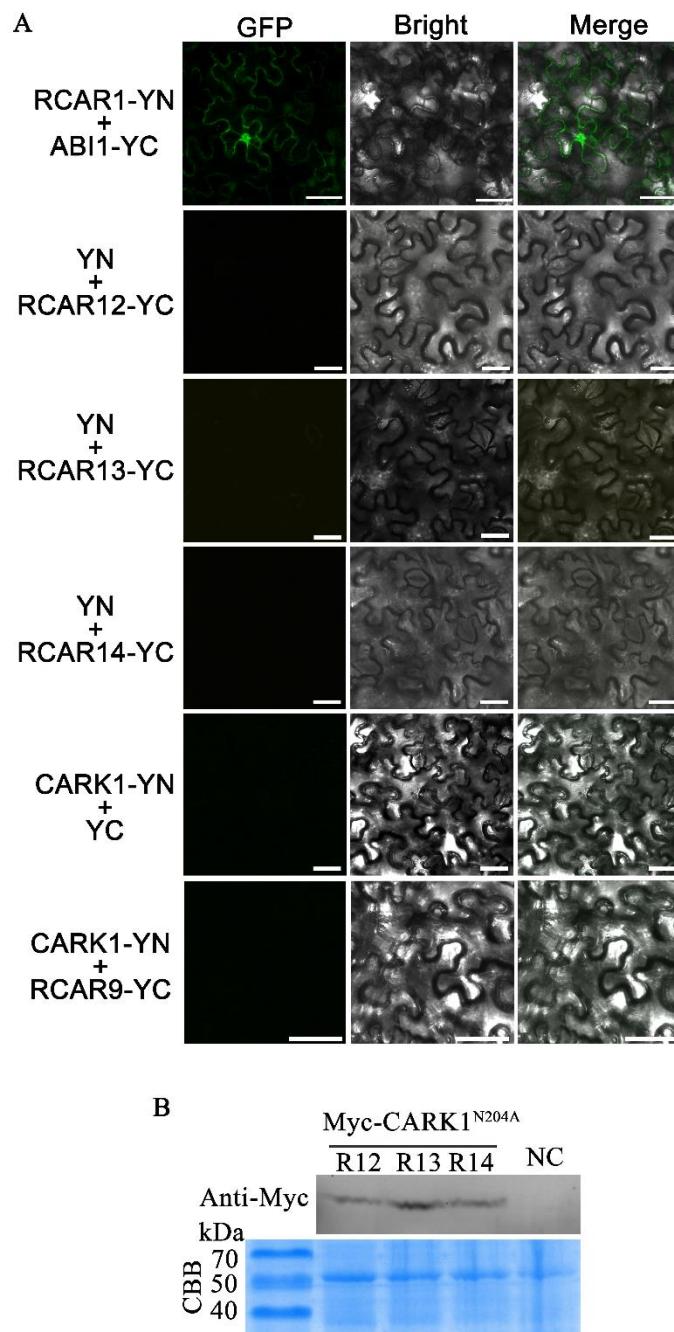
*Author for correspondence: Yi Yang (yangyi528@scu.edu.cn)

[¶]These authors contributed equally to this work

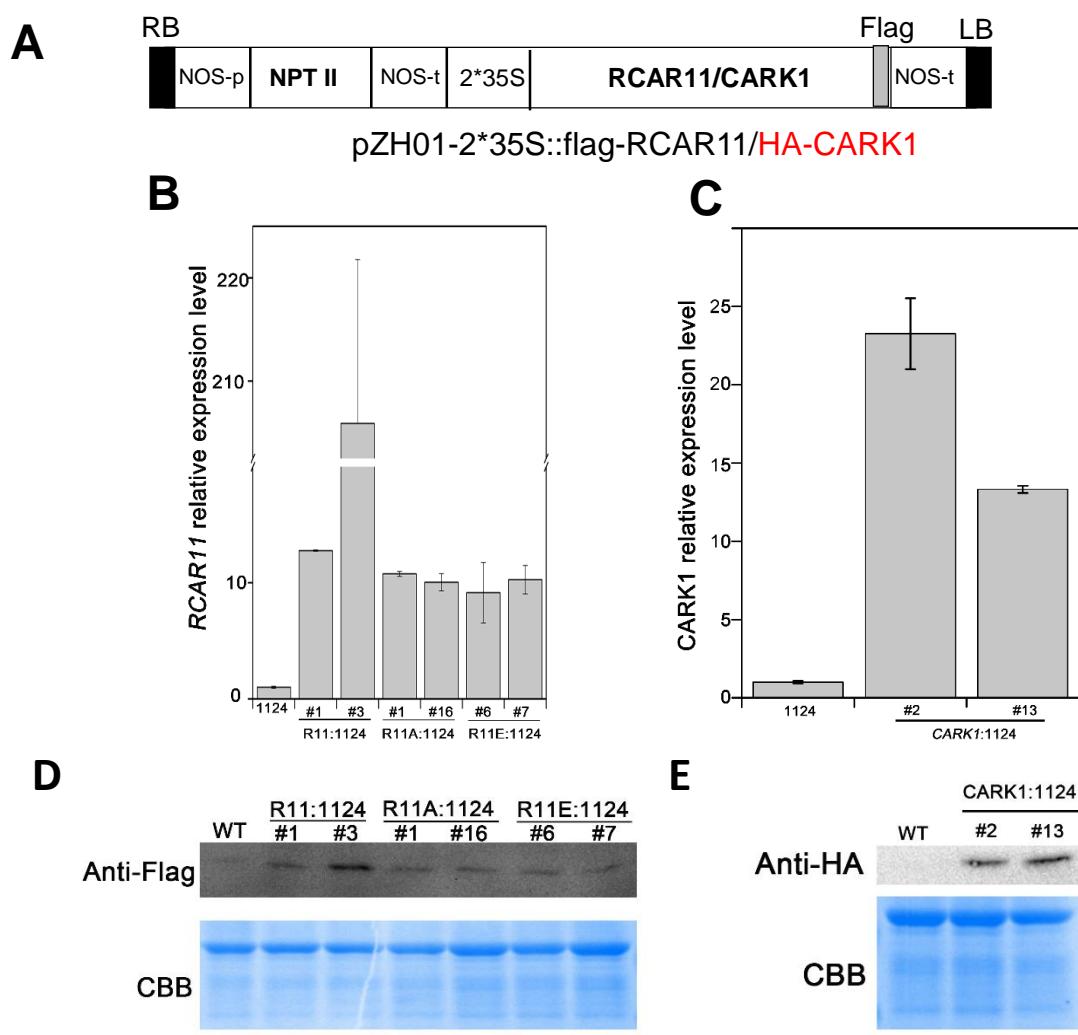
The following Supporting Information is available for this article:

Supporting Online Figures (Fig. S1—S2)

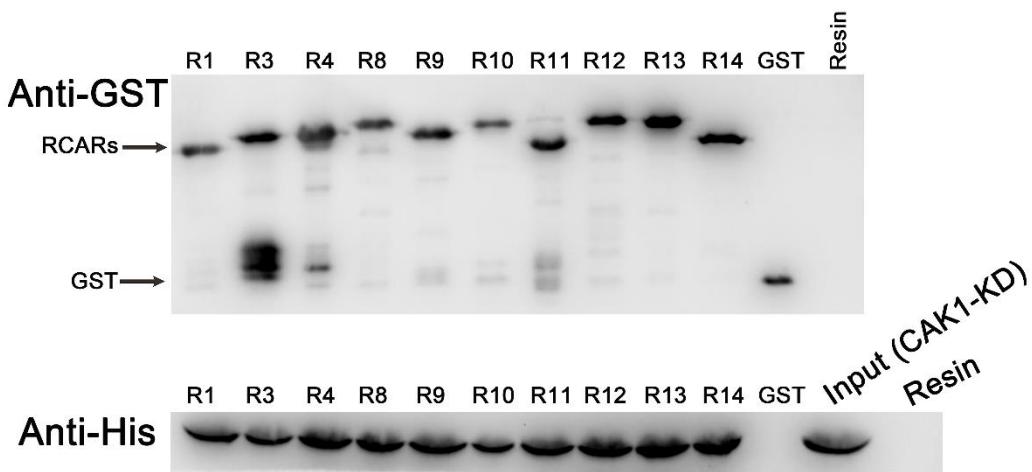
Supporting Online Table (Table S1)



Supplementary Fig. S1 Controls of BiFC assay. A, Co-expression of RCAR1-YN and ABI1-YC (positive control) in leaves of *N. benthamiana*, yielded fluorescence signals both in the cytosol and in the nucleus. No signal of YFP fluorescence was detected after co-expression of YN and with RCAR12-YC, RCAR13-YC, RCAR14-YC, YC and **RCAR9-YC** with CARK1-YN. Scale bar, 50 μ m. **B,** Western blot analyzing CARK1^{N204A} expression levels in tobacco epidermal cells, co-infiltration CARK1^{N204A} with RCAR12, RCAR13 and RCAR14, respectively. NC, Crude extracts without plasmids transfection were used as the negative control.



Supplementary Fig. S2 Construct used for plant and identification of transgenic lines by qRT-PCR and protein levels. **A**, Construct used for transgenic plant, RB, right T-DNA border, LB, left T-DNA border; Nos-p, NOS promoter; NOS-t, NOS terminator, 35S, cauliflower mosaic virus (CaMV) 35S promoter; NPTII, neomycin phosphotransferase II. **B and C**, Real time-PCR analysis of the 1124, R11:1124, R11A:1124, R11E:1124, CARK1:1124. Expression levels of *RCAR11* or *CARK1* transcripts were determined by qRT-PCR using gene-specific primers. ACTIN2/8 mRNA level were used as the internal control. **D and E**, western blot analyzing protein levels of transgenic plants. Anti-Flag was used to detect RCAR11:1124, RCAR11A:1124 and RCAR11E:1124, and anti-HA was used to detect CARK1:1124. CBB (Coomassie Brilliant Blue) stain is shown as loading control.



Supplementary Fig. S3 *In vitro* GST pull-down assay of CARK1-KD interaction with RCARs.

6xHis fused CARK1-KD was incubated with, GST-tagged RCARs as indicated, respectively. The RCARs proteins, GST or the negative control (Resin) were analyzed by anti-GST (up) and immunoblotting with antibodies against the His-tag (bottom).

Table S1 Primers used in this paper

Assays	Genes	Primer sequences (5`-3`)
Primers for GST-pull down	RCAR1	F (<i>EcoRI</i>): GGAATT CATGGACGGCGTTGAAGGCG R (<i>Xhol</i>): CCGCTCGAGTCACTGAGTAATGTCCTGAGAAGCC
		F (<i>EcoRI</i>): GGAATT CATGGAAGCTAACGGGATTG R (<i>Xhol</i>): CCGCTCGAGTCAAGCGTAATCTGGAACATC
	RCAR4	F (<i>EcoRI</i>): GGAATT CATGAACGGTGACGAAACAAAGAAG R (<i>Xhol</i>): CCGCTCGAGTATCTTCTTCTCCATAGATTCT
		F (<i>BamHI</i>): GGAATT CATGGAAGCTAACGGGATTG R (<i>EcoRI</i>): CCGCTCGAGTCAGCGTAATCTGGAACATC
	RCAR8	F (<i>EcoRI</i>): GGAATT CATGCCAACGTCGATACAGTT R (<i>Xhol</i>): CCGCTCGAGTTACGAGAATTAGAAGTGTCTCG
		F (<i>EcoRI</i>): GGAATT CATGCCAACGTCGATACAGTT R (<i>Xhol</i>): CCGCTCGAGCAGAGACATCTTCTTCTGCTC
	RCAR11	F (<i>BamHI</i>): CGCGGATCCATGGCGAATTCAAGAGTCCTC R (<i>SmaI</i>): TCCCCCGGGCCTAACCTGAGAAGAGTTGT
		F (<i>BamHI</i>): CGCGGATCCATGGCGAATTCAAGAGTCCTC R (<i>SmaI</i>): TCCCCCGGGCCTAACCTGAGAAGAGTTGT
	RCAR12	F (<i>EcoRI</i>): GGAATT CATGAATCTTGCTCCAATCCA R (<i>Xhol</i>): CCGCTCGAGTCAGGTCGGAGAACCGCTGG
		F (<i>EcoRI</i>): GGAATT CATGAGCTCATCCCCGG R (<i>Xhol</i>): CCGCTCGAGTTATTCATCATCATGCATAGGTG
	RCAR9	F (<i>BamHI</i>): CGGGATCCATGCCAACGTCGATACAGTT R (<i>SalI</i>): TCCGTCGACCGAGAATTAGAAGTGTCTCG
		F (<i>BamHI</i>): CGCGGATCCATGGCGAATTCAAGAGTCCTC R (<i>SalI</i>): TCCGTCGACAACCTGAGAAGAGTTGT
	RCAR13	F (<i>BamHI</i>): CGCGGATCCATGAATCTTGCTCCAATCCA R (<i>SalI</i>): TCCGTCGACGGTCGGAGAACCGCTGG
		F (<i>BamHI</i>): CGCGGATCCATGAGCTCATCCCCGG R (<i>SalI</i>): TCCGTCGACTTCATCATCATGCATAGGTG
Primers for BiFC	RCAR12	F (<i>BamHI</i>): CGCGGATCCATGGCGAATTCAAGAGTCCTC R (<i>SalI</i>): TCCGTCGACAACCTGAGAAGAGTTGT
		F (<i>BamHI</i>): CGCGGATCCATGAATCTTGCTCCAATCCA R (<i>SalI</i>): TCCGTCGACGGTCGGAGAACCGCTGG
	RCAR13	F (<i>BamHI</i>): CGCGGATCCATGAATCTTGCTCCAATCCA R (<i>SalI</i>): TCCGTCGACGGTCGGAGAACCGCTGG
		F (<i>BamHI</i>): CGCGGATCCATGAGCTCATCCCCGG R (<i>SalI</i>): TCCGTCGACTTCATCATCATGCATAGGTG
Primers for CoIP	RCAR12	F (<i>BamHI</i>): CGCGGATCCATGGCGAATTCAAGAGTCCTC R (<i>SalI</i>): TCCGTCGACAACCTGAGAAGAGTTGT
		F (<i>BamHI</i>): CGCGGATCCATGAATCTTGCTCCAATCCA R (<i>SalI</i>): TCCGTCGACGGTCGGAGAACCGCTGG
	RCAR13	F (<i>SalI</i>): TCCGTCGACATGGGCTGCTTGGTTGTGT R (<i>XbaI</i>): GCTCTAGAATACTGGGTTCTGTGTGGAGTC
		F (<i>SalI</i>): TCCGTCGACATGGGCTGCTTGGTTGTGT R (<i>XbaI</i>): GCTCTAGAATACTGGGTTCTGTGTGGAGTC
Primers for site-directed mutagenesis	RCAR12 ^{T105A}	CGAGTGGATGCGCGCGACGTGAAC CGCGCATCCCACTCGCATCTCGAAATC
		GATTAAGTTGGGGCGATAAGGGAAG CGCCCCAACCTTAATCTCTTGTGC
	RCAR13 ^{T101A}	GGTGACGTGGAGCGGTAGAGAAG CGCTCCGACGTACCCATCACCGGAG
		CGCTCCGACGTACCCATCACCGGAG