CORRECTION

Correction: FIN219/JAR1 and cryptochrome1 antagonize each other to modulate photomorphogenesis under blue light in *Arabidopsis*

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There is an error in panel A of Fig 2. Specifically, the sample order should read 'Col-0, *FIN219-OE, cry1, FIN219-OE/cry1, cry2, FIN219-OE/cry2, cry1cry2, FIN219-OE/cry1, cry2*, not 'Col-0, *FIN219-OE, cry1, cry2, FIN219-OE/cry1, cry1cry2, FIN219-OE/cry2, FIN219-OE/cry1-cry2*'. The authors have provided a corrected version here.



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Fig 2. FIN219 and CRY1 antagonize each other under blue light. (A) CRY1 and CRY2 negatively regulate FIN219 protein level under blue light. Western blot analysis of FIN219 protein level in wild-type Col-0, *cry1*, *cry2* mutants and transgenic seedlings grown under blue light. The signal was detected by FIN219 monoclonal antibody. Blue light: 2.2 μmol•m⁻²•s⁻¹. The number below each blot represents the level of the indicated protein. The level of wild-type Col-0 was arbitrarily set to 1. The asterisk (*) indicates nonspecific bands. (B) *FIN219* overexpression in *GUS-CCT1* seedlings abolishes GUS-CCT1 fusion protein in blue light. Western blot analysis of protein levels in Col-0, *fin219-2*, *FIN219* overexpression line (*FIN219-OE*), *FIN219-OE/GUS-CCT1*, *GUS-CCT1* and *cry1* seedlings grown in blue light for 3 days. The blots were detected by antibodies against FIN219 and GUS-CCT1 and HY5. Blue light: 2.2 μmol•m⁻²•s⁻¹. RPN8 was a loading control. The asterisk (*) indicates nonspecific bands. The number below each blot represents the level of the indicated protein. The level of wild-type Col-0 was arbitrarily set to 1. (*C) GUS-CCT1*, *GUS-CCT1* and *cry1* seedlings grown in blue light for 3 days. The blots were detected by antibodies against FIN219 and GUS-CCT1 and HY5. Blue light: 2.2 μmol•m⁻²•s⁻¹. RPN8 was a loading control. The asterisk (*) indicates nonspecific bands. The number below each blot represents the level of the indicated protein. The level of wild-type Col-0 was arbitrarily set to 1. (*C) GUS-CCT1* transcripts detected in transgenic seedlings of *FIN219-OE/GUS-CCT1* under blue light. Quantitative Real-time PCR (qPCR) analysis of transgenic seedlings shown in B. Total RNAs were extracted from transgenic seedlings shown in the figure and subjected for qPCR analysis. *Ubiquitin 10* (*UBQ10*) was an internal control. (D) GUS-CCT1 fusion proteins were stable in *fin219-2/GUS-CCT1* and *PGR219/GUS-CCT1* seedlings grown in blue light for 4 days. Total proteins extracted from seedlings

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Reference

 Chen H-J, Fu T-Y, Yang S-L, Hsieh H-L (2018) FIN219/JAR1 and cryptochrome1 antagonize each other to modulate photomorphogenesis under blue light in *Arabidopsis*. PLoS Genet 14(3): e1007248. https://doi.org/10.1371/journal.pgen.1007248 PMID: 29561841