

B1-type cyclins control microtubule organization during cell division in Arabidopsis

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Editor: Deniz Senyilmaz Tiebe

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(Note: With the exception of the correction of typographical or spelling errors that could be a source of ambiguity, letters and reports are not edited. Depending on transfer agreements, referee reports obtained elsewhere may or may not be included in this compilation. Referee reports are anonymous unless the Referee chooses to sign their reports.)

1st Editorial Decision 6th Oct 2021

Dear Arp,

Thank you for transferring your manuscript to EMBO Reports, which was previously reviewed at The EMBO Journal. Having looked at everything, I would like to invite you to submit a manuscript with minor revisions as outlined below.

I would like to apologize once again for this unusual delay. A member of the data editing team is unexpectedly out of the office due to emergency surgery.

- Please reduce the number of figures to max 7 (by combining panels) to increase the accessibility of the findings.
- The 'Data Availability' section should be moved after the 'Materials and Methods' section.
- Please fill out and include an author checklist as listed in out online guidelines (https://www.embopress.org/page/journal/14693178/authorguide)
- Please enter the funding information into the manuscript submission system, too.
- We note that all figures are submitted as a single file. Please submit them separately in high resolution.
- We realize that Fig 1A, 2B, 8C, EV2A, EV3 & EV4 panel callouts are missing. Figure 10 is not called out in the text, either.
- I note that Table EV1 is the datapoints based on which the calculations are performed in Figures 6 and 7. Therefore, please upload this table as 'source data' in two separate excel files for each figure.
- Please upload the Appendix table S1 as Table EV1. Please remember to update the text callouts.
- The tables (1-4) are currently merged with the figures. Please include the Tables and their legends in the manuscript text as the tables need to be editable.
- Papers published in EMBO Reports include a 'synopsis' and 'bullet points' to further enhance discoverability. Both are displayed on the html version of the paper and are freely accessible to all readers. The synopsis includes a short standfirst summarizing the study in 1 or 2 sentences that summarize the paper (max 35 words) and are provided by the authors and streamlined by the handling editor. I would therefore ask you to include your synopsis blurb and 3-5 bullet points listing the key experimental findings.
- In addition, please provide an image for the synopsis. This image should provide a rapid overview of the question addressed in the study but still needs to be kept fairly modest since the image size cannot exceed 550x400 pixels. For example, you can adapt Figure 10 for this.
- Our production/data editors have asked you to clarify several points in the figure legends (see attached document). Please incorporate these changes in the attached word document and return it with track changes activated.

Thank you again for giving us to consider your manuscript for EMBO Reports, I look forward to your minor revision.

Kind regards,

Deniz

Donis

Deniz Senyilmaz Tiebe, PhD Editor EMBO Reports The authors have addressed all minor editorial requests.

Dear Arp,

Thank you for submitting your revised manuscript. I have now looked at everything and all is fine. Therefore, I am very pleased to accept your manuscript for publication in EMBO Reports.

Congratulations on a nice work!

Kind regards,

Deniz

DCIII

Deniz Senyilmaz Tiebe, PhD Editor

EMBO Reports

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Corresponding Author Name: Arp Schnittger

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porting Checklist For Life Sciences Articles (Rev. June 2017)

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A- Figures

1. Data

The data shown in figures should satisfy the following conditions:

- the data were obtained and processed according to the field's best practice and are presented to reflect the results of the experiments in an accurate and unbiased manner.
 figure panels include only data points, measurements or observations that can be compared to each other in a scientifically
- meaningful way.

 graphs include clearly labeled error bars for independent experiments and sample sizes. Unless justified, error bars should not be shown for technical replicates.
- → if n< 5, the individual data points from each experiment should be plotted and any statistical test employed should be
- in the control of the guidelines on Data Presentation.

2. Captions

Each figure caption should contain the following information, for each panel where they are relevant:

- a specification of the experimental system investigated (eg cell line, species name).
 the assay(s) and method(s) used to carry out the reported observations and measurements
 an explicit mention of the biological and chemical entity(leg) that are being measured.
 an explicit mention of the biological and chemical entity(ies) that are altered/varied/perturbed in a controlled manner.

- → the exact sample size (n) for each experimental group/condition, given as a number, not a range;
 → a description of the sample collection allowing the reader to understand whether the samples represent technical or biological replicates (including how many animals, litters, cultures, etc.).
 → a statement of how many times the experiment shown was independently replicated in the laboratory.
 → definitions of statistical methods and measures:
 common tests, such as t-test (please specify whether paired vs. unpaired), simple χ2 tests, Wilcoxon and Mann-Whitney tests, can be unambiguously identified by name only, but more complex techniques should be described in the methods
 - · are tests one-sided or two-sided?

 - are tests one states on two states?

 are there adjustments for multiple comparisons?

 exact statistical test results, e.g., P values = x but not P values < x;
 - definition of 'center values' as median or average
 - · definition of error bars as s.d. or s.e.m

Any descriptions too long for the figure legend should be included in the methods section and/or with the source data.

n the pink boxes below, please ensure that the answers to the following questions are reported in the manuscript itse

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	all samples were treated the same way, each genotype was analyzed at least with three independent biological replicates
1.b. For animal studies, include a statement about sample size estimate even if no statistical methods were used.	not applicable
2. Describe inclusion/exclusion criteria if samples or animals were excluded from the analysis. Were the criteria pre-established?	not applicable
3. Were any steps taken to minimize the effects of subjective bias when allocating animals/samples to treatment (e.g. randomization procedure)? If yes, please describe.	no
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4.a. Were any steps taken to minimize the effects of subjective bias during group allocation or/and when assessing results (e.g. blinding of the investigator)? If yes please describe.	no
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S. For every figure, are statistical tests justified as appropriate?	yes
Do the data meet the assumptions of the tests (e.g., normal distribution)? Describe any methods used to assess it.	yes
Is there an estimate of variation within each group of data?	yes

Is the variance similar between the groups that are being statistically compared?	yes

C- Reagents

6. To show that antibodies were profiled for use in the system under study (assay and species), provide a citation, catalog number and/or clone number, supplementary information or reference to an antibody validation profile. e.g., Antibodypedia (see link list at top right), 1DegreeBio (see link list at top right).	only previously published ab were used, their use is fully referenced in our manuscript
Identify the source of cell lines and report if they were recently authenticated (e.g., by STR profiling) and tested for mycoplasma contamination.	not applicable

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D- Animal Models

8. Report species, strain, gender, age of animals and genetic modification status where applicable. Please detail housing	not applicable
and husbandry conditions and the source of animals.	
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compliance.	

E- Human Subjects

11. Identify the committee(s) approving the study protocol.	not applicable
12. Include a statement confirming that informed consent was obtained from all subjects and that the experiments conformed to the principles set out in the WMA Declaration of Helsinki and the Department of Health and Human Services Belmont Report.	not applicable
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F- Data Accessibility

18: Provide a "Data Availability" section at the end of the Materials & Methods, listing the accession codes for data generated in this study and deposited in a public database (e.g. RNA-Seq data: Gene Expression Omnibus GSE39462, Proteomics data: PRIDE PXD000208 etc.) Please refer to our author guidelines for 'Data Deposition'.	not applicable
Data deposition in a public repository is mandatory for: a. Protein, DNA and RNA sequences b. Macromolecular structures c. Crystallographic data for small molecules d. Functional genomics data e. Proteomics and molecular interactions	
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G- Dual use research of concern

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