

Protein complexes in cells by AI-assisted structural proteomics

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Transaction Report:

(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.)

The reviewers' comments and authors' responses are not available with this article, as the initial review process took place with another journal.

Thank you once again for submitting your manuscript "Protein complexes in cells by AI-assisted structural proteomics" to Molecular Systems Biology. I have now discussed with the team the manuscript and the reviewers' comments from the other journal. We think that the topic fits well to the scope of MSB and we find the integrative approach combining in-cell cross-linking mass spectrometry, co-fractionation mass spectrometry and protein structure modelling with AlphaFold-Multimer interesting. We would be happy to publish the study in Molecular Systems Biology, pending some editorial issues that need to be addressed.

Overall, we think that the concerns of the reviewers from the initial submission at the other journal have been satisfactorily addressed. Reviewer #1 pointed out that the work seems generally technical sound and was mainly concerned about the advance provided by the study. Reviewer #2 raised some questions regarding how much the experimental approaches contribute to the findings and had some concerns about the validations of the Yugi/YabR and PdhI/YneR interactions. We think that the clarifications and additional explanations regarding how the different approaches complement each other in generating insights into protein-protein interactions as well as emphasizing the contribution of the study is analysing these interactions in cells (and not in lysates as in previous studies) address the concerns raised by reviewer #1 and #2 regarding the advance provided by the study. Moreover, we think that the additional experiments have strengthened the conclusions about the selected validated interactions. Overall, we think that the main concerns of the reviewers from the other journal have been satisfactorily addressed and the study is well suited for publication in Molecular Systems Biology.

Before we can formally accept the study for publication, we would ask you to address the following points:

Thank you again for sending us your revised manuscript. We are now satisfied with the modifications made and I am pleased to inform you that your paper has been accepted for publication.

EMBO Press Author Checklist

Corresponding Author Name: Juri Rappsilber
Journal Submitted to: Molecular Systems Biology
Manuscript Number: MSB-2023-11544

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Reporting Checklist for Life Science Articles (updated January)

This checklist is adapted from Materials Design Analysis Reporting (MDAR) Checklist for Authors. MDAR establishes a minimum set of requirements in transparent reporting in the life sciences (see Statement of Task: [10.31222/osf.io/9sm4x](https://doi.org/10.31222/osf.io/9sm4x)). Please follow the journal's guidelines in preparing your manuscript. **Please note that a copy of this checklist will be published alongside your article.**

Abridged guidelines for 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.
- ideally, figure panels should include only measurements that are directly comparable to each other and obtained with the same assay.
- plots 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. Any statistical test employed should be justified.
- Source Data should be included to report the data underlying figures according to the guidelines set out in the authorship guidelines on Data

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(ies) 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 χ^2 tests, Wilcoxon and Mann-Whitney tests, can be unambiguously identified by name only, but more complex techniques should be described in the methods section;
 - are tests one-sided or two-sided?
 - 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.

Please complete ALL of the questions below.

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Newly Created Materials	Information included in the manuscript?	In which section is the information available? (Reagents and Tools Table, Materials and Methods, Figures, Data Availability Section)
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For every figure, are statistical tests justified as appropriate? Do the data meet the assumptions of the tests (e.g., normal distribution)? Describe any methods used to assess it. Is there an estimate of variation within each group of data? Is the variance similar between the groups that are being statistically compared?	Yes	Material and methods, Figures, supplementary figures

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In the figure legends: state number of times the experiment was replicated in laboratory.	Yes	Figures
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