

The fitness cost and benefit of phase separated protein deposits

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

This checklist is used to ensure good reporting standards and to improve the reproducibility of published results. These guidelines are consistent with the Principles and Guidelines for Reporting Preclinical Research issued by the NIH in 2014. Please follow the journal's authorship guidelines in preparing your manuscript.

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
- justified Source Data should be included to report the data underlying graphs. Please follow the guidelines set out in the author ship guidelines on Data Presentation.

2. Cantions

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 χ2 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

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

the pink boxes below, please ensure that the answers to the following questions are reported in the ma very question should be answered. If the question is not relevant to your research, please write NA (non applicable).

B- Statistics and general methods

1.a. How was the sample size chosen to ensure adequate power to detect a pre-specified effect size? ot applicable 1.b. For animal studies, include a statement about sample size estimate even if no statistical methods were used. ot applicable 2. Describe inclusion/exclusion criteria if samples or animals were excluded from the analysis. Were the criteria pre escribed in detail in the materials and methods section of the established 3. Were any steps taken to minimize the effects of subjective bias when allocating animals/samples to treatment (e.g. ot applicable ndomization procedure)? If yes, please descri For animal studies, include a statement about randomization even if no randomization was used. 4.a. Were any steps taken to minimize the effects of subjective bias during group allocation or/and when assessing result (e.g. blinding of the investigator)? If yes please describe 4.b. For animal studies, include a statement about blinding even if no blinding was done ot applicable 5. 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. es and these are described in the materials and methods section of the pape Is there an estimate of variation within each group of data?

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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	PGK1 mouse monoclonal antibody (22C5D8). Novex catalog 459250, GFP rabbit polyclonal
	antibody, Santa Cruz Biotechnology catalog sc-8334 AC. Anti-Protein G HRP conjugated, Millipore
	catalog 18-161. Rabbit anti-oligomer A11, Invitrogen catalog AH80052. Goat anti-Rabbit IgG H&L tagged with an Alexa Fluor [®] 555, abcam catalog ab150078.
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

* for all hyperlinks, please see the table at the top right of the document

D- Animal Models

 Report species, strain, gender, age of animals and genetic modification status where applicable. Please detail housing and husbandry conditions and the source of animals. 	not applicable
 For experiments involving live vertebrates, include a statement of compliance with ethical regulations and identify the committee(s) approving the experiments. 	not applicable
10. We recommend consulting the ARRIVE guidelines (see link list at top right) (PLoS Biol. 8(6), e1000412, 2010) to ensure that other relevant aspects of animal studies are adequately reported. See author guidelines, under 'Reporting Guidelines'. See also: NIH (see link list at top right) and MRC (see link list at top right) recommendations. Please confirm compliance.	not applicable

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14. Report any restrictions on the availability (and/or on the use) of human data or samples.	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	DNA sequences are provided at the Appendix section.
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'.	
Data deposition in a public repository is mandatory for:	
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b. Macromolecular structures	
c. Crystallographic data for small molecules	
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19. Deposition is strongly recommended for any datasets that are central and integral to the study; please consider the	We provide all relevant datasets at the Appendix section.
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