***eLife’s* transparent reporting form**

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**Sample-size estimation**

* You should state whether an appropriate sample size was computed when the study was being designed
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* If no explicit power analysis was used, you should describe how you decided what sample (replicate) size (number) to use

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Figure legends for Figures 1 and 3, Figure 2 – figure supplement 1. Method for qRT-PCR.

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* You should report how often each experiment was performed
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* Statistical analysis methods should be described and justified
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(For large datasets, or papers with a very large number of statistical tests, you may upload a single table file with tests, Ns, etc., with reference to sections in the manuscript.)

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* Indicate how samples were allocated into experimental groups (in the case of clinical studies, please specify allocation to treatment method); if randomization was used, please also state if restricted randomization was applied
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Please indicate the figures or tables for which source data files have been provided:

The global proteome analysis relating to Figure 2 – figure supplement 1 has been provided as Figure 2 – figure supplement 1 source data. The additional data files for Figures 1 and 3 have been provided as .xls files.