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

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* 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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* You should report how often each experiment was performed
* You should include a definition of biological versus technical replication
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* High-throughput sequence data should be uploaded before submission, with a private link for reviewers provided (these are available from both GEO and ArrayExpress)

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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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Numerical data are provided as follows:

Figure 1-source data1 for Figure 1

Figure 2- source data1 for Figure 2

Figure 3- source data1 for Figure 3

Figure 4- source data1 for Figure 4

Figure 5- source data1 for Figure 5

Figure 6- source data1 for Figure 6

Figure 1-figure supplement 1- source data1 for Figure 1-figure supplement 1

Figure 1-figure supplement 2- source data1 for Figure 1-figure supplement 2

Figure 1-figure supplement 3- source data1 for Figure 1-figure supplement 3

Figure 2-figure supplement 1- source data1 for Figure 2-figure supplement 1

Figure 2-figure supplement 2- source data1 for Figure 2-figure supplement 2

Figure 4-figure supplement 1- source data1 for Figure 4-figure supplement 1

Figure 4-figure supplement 2- source data1 for Figure 4-figure supplement 2

Figure 5-figure supplement 1- source data1 for Figure 5-figure supplement 1