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* You should state whether an appropriate sample size was computed when the study was being designed
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study.

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* You should report how often each experiment was performed
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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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Information for the replicates is described in figure legends and methods. Expression threshholds are described in the figure legends and methods. Any outliers that were not plotted are mentioned in the figure legends. High-throughput sequencing data and associated processed data have been deposited in the GEO database (GSE166544).

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* Statistical analysis methods should be described and justified
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* For each experiment, you should identify the statistical tests used, exact values of N, definitions of center, methods of multiple test correction, and dispersion and precision measures (e.g., mean, median, SD, SEM, confidence intervals; and, for the major substantive results, a measure of effect size (e.g., Pearson's r, Cohen's d)
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