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* You should state whether an appropriate sample size was computed when the study was being designed
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No experiments were performed as part of this study.

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* You should include a definition of biological versus technical replication
* The data obtained should be provided and sufficient information should be provided to indicate the number of independent biological and/or technical replicates
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* Statistical analysis methods should be described and justified
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* Report exact p-values wherever possible alongside the summary statistics and 95% confidence intervals. These should be reported for all key questions and not only when the p-value is less than 0.05.

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Details of the derivations for analytical results, simulation methods and fitting procedures are described in Appendices.

(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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Least squares fitting was used as a fitting procedure. Details of the fitting procedure are described in Appendices. The full list of parameters extracted from the fit and used for illustrations is given in Appendices.