# natureresearch

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## **Reporting Summary**

Nature Research wishes to improve the reproducibility of the work that we publish. This form provides structure for consistency and transparency in reporting. For further information on Nature Research policies, see <u>Authors & Referees</u> and the <u>Editorial Policy Checklist</u>.

#### Statistics

For	all st	atistical analyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.
n/a	Cor	nfirmed
	$\boxtimes$	The exact sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement
	$\boxtimes$	A statement on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly
		The statistical test(s) used AND whether they are one- or two-sided Only common tests should be described solely by name; describe more complex techniques in the Methods section.
$\boxtimes$		A description of all covariates tested
$\boxtimes$		A description of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons
		A full description of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient) AND variation (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)
		For null hypothesis testing, the test statistic (e.g. <i>F</i> , <i>t</i> , <i>r</i> ) with confidence intervals, effect sizes, degrees of freedom and <i>P</i> value noted <i>Give P values as exact values whenever suitable</i> .
$\boxtimes$		For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings
$\boxtimes$		For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes
$\boxtimes$		Estimates of effect sizes (e.g. Cohen's <i>d</i> , Pearson's <i>r</i> ), indicating how they were calculated

Our web collection on statistics for biologists contains articles on many of the points above.

#### Software and code

Data collection	This paper makes use of restricted access data from the National Institutes of Health, protected by the Privacy Act of 1974 as amended (5 U.S.C. 552a). De-identified data necessary to reproduce all plots and statistical analyses will be made freely available. Those wishing to access the raw data may apply for access following the procedures outlined in the NIH Data Access Policy document available at http:// report.nih.gov/pdf/DataAccessPolicy.pdf. The VentureXpert database is available via Thomson Reuters. The Global Terrorism Database is publicly available at https://www.start.umd.edu/gtd/.
Data analysis	Data analyses were conducted using Python 3.4. Regression analysis were conducted using Stata 14.

For manuscripts utilizing custom algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors/reviewers. We strongly encourage code deposition in a community repository (e.g. GitHub). See the Nature Research guidelines for submitting code & software for further information.

#### Data

Policy information about availability of data

All manuscripts must include a data availability statement. This statement should provide the following information, where applicable:

- Accession codes, unique identifiers, or web links for publicly available datasets

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- A list of figures that have associated raw data
- A description of any restrictions on data availability

De-identified data necessary to replicate results of this study will be made freely available.

### Field-specific reporting

Please select the one below that is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.

Life sciences 🛛 Behavioural & social sciences 🗍 Ecological, evolutionary & environmental sciences

For a reference copy of the document with all sections, see <u>nature.com/documents/nr-reporting-summary-flat.pdf</u>

## Behavioural & social sciences study design

All studies must disclose on these points even when the disclosure is negative.						
Study description	A quantitative study of dynamics of failure based on pre-existing datasets.					
Research sample	We collected three large-scale datasets from three domains: (1) R01 grant applications ever submitted to the National Institutes of Health (NIH), (776,721 applications by 139,091 investigators from 1985 to 2015); (2) Start-up investment records from VentureXpert database (58,111 startup companies involving 253,579 innovators); and (3) Terrorist attack data from Global Terrorism Database (70,350 terrorist attacks by 3,178 terrorist organizations from 1970 to 2017).					
Sampling strategy	No statistical methods were used to predetermine sample size.					
Data collection	This study is based on pre-existing datasets.					
Timing	The NIH dataset was collected in 2016. The VentureXpert and GTD datasets were collected in 2017.					
Data exclusions	The analysis has no data exclusions. Selection criteria within a dataset are described in the supplementary information.					
Non-participation	There are no participants in this study.					
Randomization	This is a data driven study, not a randomized experiment.					

### Reporting for specific materials, systems and methods

We require information from authors about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, system or method listed is relevant to your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.

#### Materials & experimental systems

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n/a	Involved in the study	
$\boxtimes$	Antibodies	
$\boxtimes$	Eukaryotic cell lines	
$\boxtimes$	Palaeontology	
$\boxtimes$	Animals and other organisms	
$\boxtimes$	Human research participants	
$\boxtimes$	Clinical data	

Wiethous					
n/a	Involved in the study				
$\boxtimes$	ChIP-seq				

]	ChIP-seq
1	Flow cytometry

MRI-based neuroimaging