nature portfolio

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Reporting Summary

Nature Portfolio 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 Portfolio policies, see our Editorial Policies and the Editorial Policy Checklist.

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For all statistical ar	nalyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.			
n/a Confirmed				
☐ ☐ The exact	sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement			
A stateme	ent on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly			
The statis Only comm	tical test(s) used AND whether they are one- or two-sided non tests should be described solely by name; describe more complex techniques in the Methods section.			
A descript	A description of all covariates tested			
A descript	tion of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons			
A full desc	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 h	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>			
For Bayes	For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings			
For hierar	For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes			
Estimates	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 an	d code			
Policy information	about <u>availability of computer code</u>			
Data collection	N/A			
Data analysis	All data were analyzed with the ASCETIC R package, available for download from GitHub (https://github.com/danro9685/ASCETIC).			

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 and reviewers. We strongly encourage code deposition in a community repository (e.g. GitHub). See the Nature Portfolio guidelines for submitting code & software for further information.

Data

Data analysis

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
- A description of any restrictions on data availability
- For clinical datasets or third party data, please ensure that the statement adheres to our policy

All cancer data are publicly available from the relative original publication or from the cBioPortal repository (https://www.cbioportal.org/). In particular, the Acute Myeloid Leukemia dataset Tyner et al. can be downloaded at https://www.cbioportal.org/study/summary?id=aml_ohsu_2018. Moreover, the Pan-Cancer Atlas datasets and the MSK-MET datasets can be downloaded from the cBioPortal repository at https://www.cbioportal.org/datasetshttps://www.cbioportal.org/ datasets.

ist of the Pan-Cancer Atlas datasets: Acute Myelot leukemia [https://www.cbioportal.org/study/summary?id=laml_trga_pan_can_atlas_2018] Adrenocortical Carcinoma (https://www.cbioportal.org/study/summary?id=laml_trga_pan_can_atlas_2018) Bladder Urothelial Carcinoma (https://www.cbioportal.org/study/summary?id=bloa_trga_pan_can_atlas_2018) Breast Invasive Carcinoma (https://www.cbioportal.org/study/summary?id=bloa_trga_pan_can_atlas_2018) Breast Invasive Carcinoma (https://www.cbioportal.org/study/summary?id=broa_trga_pan_can_atlas_2018) Cervical Squamous Cell Carcinoma (https://www.cbioportal.org/study/summary?id=cesc_trga_pan_can_atlas_2018) Cholangiocarcinoma (https://www.cbioportal.org/study/summary?id=coadread_trga_pan_can_atlas_2018) Colorectal Adenocarcinoma (https://www.cbioportal.org/study/summary?id=coadread_trga_pan_can_atlas_2018) Esophageal Adenocarcinoma (https://www.cbioportal.org/study/summary?id=coadread_trga_pan_can_atlas_2018) Esophageal Adenocarcinoma (https://www.cbioportal.org/study/summary?id=esca_trga_pan_can_atlas_2018) Esophageal Adenocarcinoma (https://www.cbioportal.org/study/summary?id=broa_trga_pan_can_atlas_2018) Elead and Neck Squamous Cell Carcinoma (https://www.cbioportal.org/study/summary?id=broa_trga_pan_can_atlas_2018) Kidney Chromophobe (https://www.cbioportal.org/study/summary?id=kirc_trga_pan_can_atlas_2018) Kidney Renal Clear Cell Carcinoma (https://www.cbioportal.org/study/summary?id=kirc_trga_pan_can_atlas_2018) Kidney Renal Papillary Cell Carcinoma (https://www.cbioportal.org/study/summary?id=kirc_trga_pan_can_atlas_2018) Liurg Adenocarcinoma (https://www.cbioportal.org/study/summary?id=luc_trga_pan_can_atlas_2018) Liurg Squamous Cell Carcinoma (https://www.cbioportal.org/study/summary?id=luc_trga_pan_can_atlas_2018) Mesothelioma (https://www.cbioportal.org/study/summary?id=luc_trga_pan_can_atlas_2018) Prostate Adenocarcinoma (https://www.cbioportal.org/study/summary?id=san_trga_pan_can_atlas_2018) Prostate Adenocarcinoma (https://www.cbioportal.org/s		
Research involving hui	man participants, their data, or biological material	
olicy information about studies w nd sexual orientation and <u>race, et</u>	vith <u>human participants or human data</u> . See also policy information about <u>sex, gender (identity/presentation), thnicity and <u>racism</u>.</u>	
Reporting on sex and gender	N/A	
Reporting on race, ethnicity, or other socially relevant groupings	N/A	
Population characteristics	N/A	
Recruitment	N/A	
Ethics oversight	N/A	

Note that full information on the approval of the study protocol must also be provided in the manuscript.

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. X 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>

Life sciences study design

All studies must disclose on these points even when the disclosure is negative.		
Sample size	We considered already processed data from published articles. We considered all samples provided in such studies.	

Data exclusions No data exclusions were performed.

Sample size

Replication	We considered already processed data from published articles.
Randomization	We considered already processed data from published articles.
Blinding	We considered already processed data from published articles.

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		Methods	
n/a	Involved in the study	n/a Involved in the study	
\boxtimes	Antibodies	ChIP-seq	
\boxtimes	Eukaryotic cell lines	Flow cytometry	
\boxtimes	Palaeontology and archaeology	MRI-based neuroimaging	
\boxtimes	Animals and other organisms	•	
\boxtimes	Clinical data		
\boxtimes	Dual use research of concern		
\boxtimes	Plants		