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 <u>Editorial Policies</u> and the <u>Editorial Policy Checklist</u>.

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For	all statistical analyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.
n/a	Confirmed
	\square The exact sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement
	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.
	A description of all covariates tested
	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 <u>statistics for biologists</u> contains articles on many of the points above.

Software and code

Policy information about availability of computer code

Data collection

no software was used for data collection

Data analysis

Provide a description of all commercial, open source and custom code used to analyse the data in this study, specifying the version used OR state that no software was used.

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 <u>guidelines for submitting code & software</u> for further information.

Data

Policy information about availability of data

All manuscripts must include a <u>data availability statement</u>. 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 data from this paper are freely available

Human resea	arch parti	cipants	
Policy information a	about <u>studies ir</u>	nvolving human research participants and Sex and Gender in Research.	
Reporting on sex a	and gender	N/A)
Population characteristics		N/A)
Recruitment		N/A	
Ethics oversight		N/A	
Note that full informat	tion on the appro	oval of the study protocol must also be provided in the manuscript.	
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Field-spe	<u>cific re</u>	porting	_
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Life sciences	_	ehavioural & social sciences	
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Life scien	ices sti	udy design	
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/		ndicated for each experiment and was chosen to calculate statistical significant	
Data exclusions	no data exclusio	on	
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		was used in this paper and therefore no blinding was necessary)
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Reporting	g for sp	pecific materials, systems and methods	
		about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.	
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Materials & exp		$\frac{\text{Methods}}{\text{n/a} \text{Involved in the study}}$	
☐ ☐ Antibodies ☐ ChIP-seq			
□ □ Eukaryotic cell lines □ Flow cytometry			
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	d other organism	IS .	
Clinical data			
Dual use les	search of concer		
Antibodies			
Antibodies used	all anti	bodies used are described in Supplemental Table 1	
Validation Anti-Actin mouse monoclonal, Abcam ab 8245			
		/www.abcam.com/gapdh-antibody-6c5-loading-control-ab8245.html ed by the company and following publication : Wang, H., et al., Cell Death Diff 13 :435 (2022)	
		PD mouse monoclonal, Cosmo Bio CAC-NM-DND-001 //www.cosmobiousa.com/products/anti-cpds-mab-clone-tdm-2	

Validated by the company and following publication: Mori, T., et al., Photochem. Photobiol. 54, 225-232 (1991)

Anti-ATP5A mouse monoclonal, Proteintech 66037-1

https://www.ptglab.com/products/ATP5A1-Antibody-66037-1-Ig.htm

Validated by the company and following publication: Lu, YT., et al., Cell Death Dis, 9(6):672 (2018)

Anti-EXD2 rabbit polyclonal, SIGMA, HPA00548

https://www.sigmaaldrich.com/FR/fr/product/sigma/hpa005848

Validated by the company and following publication: Biehs, R., et al., Molecular Cell, 65(4):671-684 (2017)

Anti-Flag M2 mouse monoclonal, SIGMA, F1804

https://www.sigmaaldrich.com/FR/fr/product/sigma/f1804?gclsrc=aw.ds&gclid=EAlalQobChMlv_mRn8-

F_AIVkc3VCh0Y0wN1EAAYASAAEgJiyvD_BwE

Validated by the company and following publication: Srivastava, M., et al., Nature Communications, 6, 6253 (2015)

Anti-GAPDH mouse monoclonal, Abcam ab 8245

https://www.abcam.com/gapdh-antibody-6c5-loading-control-ab8245.html

Validated by the company and following publication: Wang, H., et al., Cell Death Diff 13:435 (2022)

Anti-GFP rabbit polyclonal, Amsbio TP401

https://www.amsbio.com/rabbit-anti-gfp-pab-tp401

Validated by the company and following publication: Kosar, M., et al., Nature Communications, 12, 3937 (2021)

Anti-GFP mouse monoclonal, Roche 11814460001

https://www.sigmaaldrich.com/FR/fr/product/roche/11814460001?

 $gclsrc=aw.ds\&gclid=EAIaIQobChMltNGEgNuF_AIVh6ztCh1uaQljEAAYASAAEgL-hvD_BwE$

Validated by the company and following publication: Chalfie, M., Science, 263, 802-805 (1994)

Anti-GST rabbit monoclonal, Abcam ab19256

https://www.abcam.com/gst-antibody-ab19256.htmlValidated by the company and following publication : Joset, A., J Cell Biol 188 :271-85 (2010).

Anti-gH2AX mouse monoclonal, Abcam ab22551

https://www.abcam.com/gamma-h2ax-phospho-s139-antibody-3f2-ab22551.html

Validated by the company and following publication: Ochs, F., Nature 574:571-574 (2019)

Anti-Histone H3 rabbit polyclonal, Abcam ab1791

https://www.abcam.com/histone-h3-antibody-nuclear-marker-and-chip-grade-ab1791.html Validated by the company and following publication: Yuan, L., NAR, 49:98-113 (2021)

Anti-RPB1 mouse monoclonal, 7C2 IGBMC Home made

Validated following publication: Boulon, S., Mol Cell, 39(6): 912-924

Anti-RPB1 rabbit polyclonal, Bethyl A300-653A

https://www.fortislife.com/products/primary-antibodies/rabbit-anti-rna-polymerase-ii-antibody/BETHYL-A300-653#dummy-cit Validated following publication : Sharifnia, W., Nat Med 292-300 (2019)

Anti-TFIIEa mouse monoclonal, 2A1 IGBMC Home made

Validated following publication: Compe, E., Nature Communications, 10:2084 (2019)

Anti-Vinculin mouse monoclonal, Sigma V9131

https://www.sigmaaldrich.com/FR/fr/sds/sigma/v9131

Validated following publication: Berico, P., Embo Rep, 22(9) 51683

Anti-XPA rabbit polyclonal, SCBT sc-853

https://www.scbt.com/fr/p/xpa-antibody-fl-273?requestFrom=search

 $Validated\ following\ publication: Fischer, J.M.,\ The\ FEBS\ journal,\ 281:3625-41$

Anti-(6-4PPs) mouse monoclonal, Clone 64M-2

https://www.cosmobiousa.com/products/anti-6-4pps-mab-clone-64m-2

Validated following publication: Yamamoto, A., et al., DNA Repair, 6, 649-657 (2007).

Eukaryotic cell lines

Policy information about <u>cell lines and Sex and Gender in Research</u>

Cell line source(s) U-2 OS, Hela, CS1ANSV, CS1ANSV+CSB

Authentication no authentification

no contamination certified by our cell culture departement

Mycoplasma contamination

Commonly misidentified lines

(See <u>ICLAC</u> register)

no commonly misidentified lines used in this work

Palaeontology and Archaeology

Specimen provenance	Provide provenance information for specimens and describe permits that were obtained for the work (including the name of the issuing authority, the date of issue, and any identifying information). Permits should encompass collection and, where applicable, export.
Specimen deposition	Indicate where the specimens have been deposited to permit free access by other researchers.
Dating methods	If new dates are provided, describe how they were obtained (e.g. collection, storage, sample pretreatment and measurement), where they were obtained (i.e. lab name), the calibration program and the protocol for quality assurance OR state that no new dates are provided.
Tick this box to confir	m that the raw and calibrated dates are available in the paper or in Supplementary Information.
Ethics oversight	Identify the organization(s) that approved or provided guidance on the study protocol, OR state that no ethical approval or guidance was required and explain why not.

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