

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](#).

Statistics

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

- ☒ ☐ 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. F , t , r) with confidence intervals, effect sizes, degrees of freedom and P value noted
Give P values as exact values whenever suitable.
- ☒ ☐ For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings
- ☒ ☐ For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes
- ☒ ☐ Estimates of effect sizes (e.g. Cohen's d , Pearson's r), indicating how they were calculated

Our web collection on [statistics for biologists](#) contains articles on many of the points above.

Software and code

Policy information about [availability of computer code](#)

Data collection Immunofluorescence images were acquired by ZEN (2.1, black); Electrophoretic mobility shift assay and nuclease assay images were acquired by Genesys (1.8.2); Immunoblotting images were acquired by Gelcap 5.6.

Data analysis Immunofluorescence images were analyzed by ZEN (3.1, blue); Images of immunoblotting, electrophoretic mobility shift assay, and nuclease assay were analyzed by ImageJ (1.8.0).

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

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](#)

Data availability

All data supporting the findings of this study are available from the corresponding author. Source data are provided with this paper.

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 [nature.com/documents/nr-reporting-summary-flat.pdf](https://www.nature.com/documents/nr-reporting-summary-flat.pdf)

Life sciences study design

All studies must disclose on these points even when the disclosure is negative.

Sample size	All sample sizes were based on experimental methods and our previous study (Bai et al. 2019. Mol Cell. PMID: 31353207).
Data exclusions	Data exclusions were not applied.
Replication	All experiments were independently replicated at least 3 times.
Randomization	All cells were randomly assigned to experimental groups.
Blinding	The investigators were blinded to experimental conditions before investigators finished data analysis.

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

n/a	Involved in the study
<input type="checkbox"/>	<input checked="" type="checkbox"/> Antibodies
<input type="checkbox"/>	<input checked="" type="checkbox"/> Eukaryotic cell lines
<input checked="" type="checkbox"/>	<input type="checkbox"/> Palaeontology and archaeology
<input checked="" type="checkbox"/>	<input type="checkbox"/> Animals and other organisms
<input checked="" type="checkbox"/>	<input type="checkbox"/> Human research participants
<input checked="" type="checkbox"/>	<input type="checkbox"/> Clinical data
<input checked="" type="checkbox"/>	<input type="checkbox"/> Dual use research of concern

Methods

n/a	Involved in the study
<input checked="" type="checkbox"/>	<input type="checkbox"/> ChIP-seq
<input checked="" type="checkbox"/>	<input type="checkbox"/> Flow cytometry
<input checked="" type="checkbox"/>	<input type="checkbox"/> MRI-based neuroimaging

Antibodies

Antibodies used

Immunoblotting (IB), Immunofluorescence (IF):
 anti-Flag-HRP (Sigma-Aldrich, A8592, 1:2000) for IB;
 anti-MBP-HRP (NEB, E8038, 1:3000) for IB;
 anti-MRE11 (Proteintech, 10744-1-AP, 1:1000) for IB;
 anti-MRE11 (Abcam, ab33125, 1:200) for IF;
 anti-NBS1 (Proteintech, 55025-1-AP, 1:1000) for IB;
 anti-RAD50 (ABclonal, A3078, 1:1000) for IB;
 anti-RNF4 (Proteintech, 17810-1-AP, 1:1000) for IB;
 anti-UBC9 (Proteintech, 51018-2-AP, 1:2000) for IB;
 anti-PIAS1 (Proteintech, 23395-1-AP, 1:1000) for IB;
 anti-phospho-RPA32 (S4/S8) (Abcam, ab87277) for IB (1:1000) and IF (1:500);
 anti-RPA32 (Proteintech, 10412-1-AP, 1:1000) for IB;
 anti-phospho-CHK1 (S345) (Cell Signaling Technology, 2348S, 1:1000) for IB;
 anti-CHK1 (Santa Cruz, sc-8408, 1:1000) for IB;
 anti-phospho-H2AX (S139) (Cell Signaling Technology, 9718, 1:1000) for IB;
 anti-phospho-ATM (S1981) (Cell Signaling Technology, 5883S, 1:1000) for IB;
 anti-phospho-NBS1 (S343) (Cell Signaling Technology, 3001S, 1:1000) for IB;
 anti-RAD51 (Abcam, ab133534, 1:250) for IF;
 anti-SUMO2/3 (Santa Cruz, sc32873, 1:500) for IB;
 anti-Histone H3 (Biodragon, B1055, 1:2000) for IB;
 anti-Cyclin A (Santa Cruz, sc-271682, 1:250) for IF;
 anti-c-Myc (Santa Cruz, sc-40) for IB (1:1000) and IF (1:250);
 anti-HA (BioLegend, 901503, 1:1000) for IB;
 anti-His (Santa Cruz, sc-803, 1:1000) for IB;
 anti-GAPDH (SUNGENE, KM9002, 1:5000) for IB;

anti- β -Actin (SUNGENE, KM9001, 1:5000) for IB.

Validation

anti-Flag-HRP for IB, <https://www.sigmaaldrich.cn/CN/zh/product/sigma/a8592>;
 anti-MBP-HRP for IB, <http://www.neb-china.com/pshow.asp?id=3613>;
 anti-MRE11 for IB, <https://www.ptglab.com/Products/MRE11A-Antibody-10744-1-AP.htm>;
 anti-MRE11 for IF, <https://www.abcam.cn/mre11-antibody-ab33125.html>;
 anti-NBS1 for IB, <https://www.ptglab.com/products/NBN-Antibody-55025-1-AP.htm>;
 anti-RAD50 for IB, <https://abclonal.com.cn/catalog/A3078>;
 anti-RNF4 for IB, <https://www.ptglab.com/products/RNF4-Antibody-17810-1-AP.htm>;
 anti-UBC9 for IB, www.ptglab.com/products/UBC9-Antibody-51018-2-AP.htm;
 anti-PIAS1 for IB, <https://www.ptglab.com/products/PIAS1-Antibody-23395-1-AP.htm>;
 anti-phospho-RPA32 (S4/S8) for IB and IF, <https://www.abcam.cn/rpa32rpa2-phospho-s4--s8-antibody-ab87277.html>;
 anti-RPA32 for IB, <https://www.ptglab.com/products/RPA2-Antibody-10412-1-Ap.htm>;
 anti-phospho-CHK1 (S345) for IB, https://www.cellsignal.cn/products/primary-antibodies/phospho-chk1-ser345-133d3-rabbit-mab/2348?site-search-type=Products&N=4294956287&Ntt=2348s&fromPage=plp&_requestid=1841901;
 anti-CHK1 for IB, <https://www.scbt.com/p/chk1-antibody-g-4>;
 anti-phospho-H2AX (S139) for IB, <https://www.cellsignal.cn/products/primary-antibodies/phospho-histone-h2a-x-ser139-20e3-rabbit-mab/9718?site-search-type=Products&N=4294956287&Ntt=h2ax&fromPage=plp>;
 anti-phospho-ATM (S1981) for IB, https://www.cellsignal.cn/products/primary-antibodies/phospho-atm-ser1981-d6h9-rabbit-mab/5883?site-search-type=Products&N=4294956287&Ntt=5883s&fromPage=plp&_requestid=1842322;
 anti-phospho-NBS1 (S343) for IB, https://www.cellsignal.cn/products/primary-antibodies/phospho-p95-nbs1-ser343-antibody/3001?_=1646978164176&Ntt=3001s&tahead=true;
 anti-RAD51 for IF, <https://www.abcam.cn/rad51-antibody-epr40303-ab133534.html>;
 anti-SUMO2/3 for IB, <https://www.scbt.com/p/sumo-2-3-antibody-fl-103?requestFrom=search>;
 anti-Histone H3 for IB, <https://www.biodragon.cn/nckt/73664.html>;
 anti-Cyclin A for IF, <https://www.scbt.com/zh/p/cyclin-a-antibody-b-8?requestFrom=search>;
 anti-c-Myc for IB and IF, <https://www.scbt.com/p/sumo-2-3-antibody-fl-103?requestFrom=search>;
 anti-HA for IB, <https://www.biolegend.com/en-us/products/purified-anti-ha-11-epitope-tag-antibody-11374>;
 anti-His for IB, <https://www.scbt.com/p/his-probe-antibody-h-15?requestFrom=search>;
 anti-GAPDH for IB, http://www.sungenebiotech.com/index.php?m=Product&a=product_xq&catid=2&proid=53&prid=290&pid=720&id=1557;
 anti- β -Actin for IB, http://www.sungenebiotech.com/index.php?m=Product&a=product_xq&catid=2&proid=53&prid=289&pid=717&id=1548.

Eukaryotic cell lines

Policy information about [cell lines](#)

Cell line source(s)

HEK293T and HeLa cell lines were purchased from ATCC.

Authentication

Cell lines authentication was identified by short tandem repeat profiling.

Mycoplasma contamination

All cell lines were confirmed without mycoplasma contamination.

Commonly misidentified lines
(See [ICLAC](#) register)

No commonly misidentified cell lines were used in this study.