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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 | For all statistical analyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section. | | | | | |
|-----|---|---|--|--|--|--|
| n/a | Cor | nfirmed | | | | |
| | x | 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. | | | | |
| X | | 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 Give <i>P</i> values as exact values whenever suitable. | | | | |
| X | | For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings | | | | |
| X | | 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 <u>statistics for biologists</u> contains articles on many of the points above. | | | | |

Software and code

| Policy information about <u>availability of computer code</u> | | | | | | |
|---|-----|--|--|--|--|--|
| Data collection | N/A | | | | | |
| Data analysis | N/A | | | | | |

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 <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 list of figures that have associated raw data

- A description of any restrictions on data availability

The data that support the findings of this study are available from the corresponding author upon reasonable request.

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>

Life sciences study design

| All studies must disclose on these points even when the disclosure is negative. | | | | | |
|---|--|--|--|--|--|
| Sample size | Sample sizes we based on previous published literature | | | | |
| Data exclusions | Outliers were removed based on a Grubbs' test, or if technical errors were apparent. | | | | |
| Replication | Major findings were replicated in 2 or more cohorts | | | | |
| Randomization | Mice were allocated to groups based on genotypes (littermates matched). | | | | |
| Blinding | The investigator were not blinded to the conditions. Genotype was known to allocate mouse groupings. | | | | |

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 Involved in the study Involved in the study n/a n/a × Antibodies X ChIP-seq × Eukaryotic cell lines × Flow cytometry X Palaeontology × MRI-based neuroimaging ✗ Animals and other organisms X Human research participants Clinical data × Antibodies All antibodies details, including supplier, dilution and catalog numbers are given in supplementary table 1 of manuscript Antibodies used Validation All antibodies were purchased from commercial suppliers. Catalog numbers are given in supplementary table 1 of manuscript and suppliers web-page contain validation experiments as well as references to previous publications using these antibodies test. All antibodies produced bands at or close to the predicted molecular weight and insulin receptor antibody was validated with

Animals and other organisms

Policy information about studies involving animals; ARRIVE guidelines recommended for reporting animal research

| Laboratory animals | Male PerIRKO mice were used. Strain, age and housing conditions are detailed in the methods section of the manuscript |
|-------------------------|---|
| Wild animals | N/A |
| Field-collected samples | N/A |
| Ethics oversight | N/A |

knockout mice tissue. pAkt antibody was validated with insulin induced increase in phosphorylation levels.

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