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.

For all statistical analyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.

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Data collection

Data analysis

n/a	Cor	nfirmed
	X	The exact sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement
	x	A statement on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly
	X	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)
	x	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>
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.
So ⁻	tw	vare and code
Poli	v in	formation about availability of computer code

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.

RNA Seq was done by the Illumina Hiseq 4000 system. Microscopic imaging was done by Leica DM750 microscope.

RNA-seq: STAR RNA-weq aligner sorfware (version STAR_2.5.0a) Gene Set Enrichment Analysis (GSEA, Broad Institute, version 4.1.0)

Plotting and Statistics: SPSS software (version 20.0) and Graphpad Prism (version 9.4)

Gene interaction network (Cytoscape, version 3.8.2) Western blot analysis: Las-4000 imager (version 1.2)

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

The source data for all box plots, bars, and line graphs as well as the original uncropped Western blots can be found in the online Source Data File. Raw and processed RNA-seq datasets were deposited into the NCBI GEO database under the accession number GSE214442.

Research involving human participants, their data, or biological material

Policy information about studies with <u>human participants or human data</u>. See also policy information about <u>sex, gender (identity/presentation)</u>, <u>and sexual orientation</u> and <u>race</u>, ethnicity and racism.

Reporting on sex and gender

Reporting on race, ethnicity, or other socially relevant groupings

Population characteristics

Blood and liver tissues are obtained from healthy volunteer, NAFLD patients or hepatocellular carcinoma patients.

Recruitment

The biospecimens and data used in this study were provided by the Biobank of Jeonbuk National University Hospital. participants were selected randomly and without self-selection.

Ethics oversight

Jeonbuk National University Hospital (IRB number CUH 2021-05-029, approved on 18 June 2021)

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 be	fore 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

Life sciences study design

Sample size

Blinding

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

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For in vitro and in vivo experiments, sample size selection was based on literature (PMID: 30530497, 21179166, 22183976, 25998209) and the lab's previous experience. We usually observed standard deviation of around 20% and therefore average of 6 mice per group is sufficient to detect 33% change with 80% power. For signaling assays, we typically performed at least three independent experiment repeats to allow statistical analysis and robust conclusions to be drawn. For RNA-seq analysis, the sample size was determined by minimum requirement for

statistical analysis (three samples).

Data exclusions No data exclusions in this manuscript.

Replication

For all experiments we used multiple biological replicates, as indicated in the figure legends. For quantitative measurements, three or more independent experiments were carried out and statistical analysis performed. For non-quantitative experiments, each experimental condition

was repeated at least one more time to confirm the observation.

Randomization Investigators were blinded to group allocation during data collection and data analyses.

Describe whether the investigators were blinded to group allocation during data collection and/or analysis. If blinding was not possible, describe why OR explain why blinding was not relevant to your study.

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	
	x Antibodies	x	ChIP-seq	
	x Eukaryotic cell lines	x	Flow cytometry	
×	Palaeontology and archaeology	x	MRI-based neuroimaging	
	🗶 Animals and other organisms		•	
×	Clinical data			
×	Dual use research of concern			
×	Plants			

Antibodies

Antibodies used

All antibodies used for Western blotting, immunofluorescence, and chromatin immunoprecipitation analyses are provided as follows:

Western blot

PAK4 (G222) (62690, 1:2500 dilution, Cell Signaling Technology)

Phospho-PAK4 (Ser474) (3241, 1:2500 dilution, Cell Signaling Technology)

HSP90 (C45G5) (4877,1:2500 dilution, Cell Signaling Technology)

Ubiquitin (E4I2J) (43124,1:2500 dilution, Cell Signaling Technology)

HMGCS2 (D3U1A) (20940, 1:2500 dilution, Cell Signaling Technology)

Phospho-PKA Substrate (100G7E) (9624, 1:2500 dilution, Cell Signaling Technology)

Phospho-(Ser/Thr) (9631, 1:2500 dilution, Cell Signaling Technology)

Sirt1 (2310, 1:2500 dilution, Cell Signaling Technology)

Sirt6 (D8D12) (12486, 1:2500 dilution, Cell Signaling Technology)

Phospho-Akt (Ser473) (D9E) (4060, 1:2500 dilution, Cell Signaling Technology)

Akt (9272, 1:2500 dilution, Cell Signaling Technology)

Acetylated-Lysine (9441, 1:2500 dilution, Cell Signaling Technology)

CHOP (L63F7) (2895, 1:2500 dilution, Cell Signaling Technology)

Phospho-S6 Ribosomal Protein (Ser240/244) (2215, 1:2500 dilution, Cell Signaling Technology)

S6 Ribosomal Protein (5G10) (2217, 1:2500 dilution, Cell Signaling Technology)

NCoR1 (5948, 1:2500 dilution, Cell Signaling Technology)

NEDD4 (2740, 1:2500 dilution, Cell Signaling Technology)

ATF-6 (D4Z8V) (65880, 1:2500 dilution, Cell Signaling Technology)

Phospho-PERK (Thr980) (16F8) (3179, 1:2500 dilution, Cell Signaling Technology)

PAK4 (B-3) (SC-390507, 1:2500 dilution, Santa Cruz Biotechnology)

FGF21 (H-105) (SC-292879, 1:2500 dilution, Santa Cruz Biotechnology)

PPARα (H-2) (SC-398394, 1:2500 dilution, Santa Cruz Biotechnology)

Sirt4 (3224, 1:2500 dilution, Bioworld Technology)

Lamin B1 (L75) (BS3547, 1:2500 dilution, Bioworld Technology)

GAPDH (A531) (AP0066, 1:2500 dilution, Bioworld Technology)

Sirt5 (aa30-46) (LS-B2060, 1:2500 dilution, LifeSpan Biosciences)

Sirt7 (aa317-366) (LS-B1566, 1:2500 dilution, LifeSpan Biosciences)

FoxO1 (Acetyl-Lys294) (LS-C800723, 1:2500 dilution, LifeSpan Biosciences)

Sirt2 (EPR20411-105) (ab211033, 1:2500 dilution, Abcam)

Sirt3 (ab189860, 1:2500 dilution, Abcam)

GRP78 BiP (ab21685, 1:2500 dilution, Abcam)

NCOR2/SMRT (ab24551, 1:2500 dilution, Abcam)

CPT1A (8F6AE9) (ab128568, 1:2500 dilution, Abcam)

T-OXPHOS (ab110413, 1:2500 dilution, Abcam)

THR β (ab53170, 1:2500 dilution, Abcam)

p-IRE1α (ab48187, 1:2500 dilution, Abcam)

MDM2 (2A10) (ab16895, 1:2500 dilution, Abcam)

LXRα (14351-1-AP, 1:2500 dilution, Proteintech)

Proximity Ligation Assay

NCoR1 (5948, 1:100 dilution, Cell Signaling Technology)

p300 (D8Z4E) (86377, 1:100 dilution, Cell Signaling Technology)

PPARα (H-2) (SC-398394, 1:100 dilution, Santa Cruz Biotechnology)

Immunofluorescence

NCoR1 (5948, 1:100 dilution, Cell Signaling Technology)

PPARα(H-2) (SC-398394, 1:100 dilution, Santa Cruz Biotechnology)

Alexa Fluor 488-conjugated goat anti-mouse IgG1 (11001, 1:100 dilution, Thermo Fisher Scientific)

Alexa Fluor 594-conjugated goat anti-rabbit IgM (11012, 1:100 dilution, Thermo Fisher Scientific)

Validation

All antibodies used in this study were commercially developed and validated by the companies.

PAK4 (G222), https://www.cellsignal.com/products/primary-antibodies/pak4-g222-antibody/62690

Phospho-PAK4 (Ser474), https://www.cellsignal.com/products/primary-antibodies/phospho-pak4-ser474-pak5-ser602-pak6-ser560-antibody/3241

HSP90 (C45G5), https://www.cellsignal.com/products/primary-antibodies/hsp90-c45g5-rabbit-mab/4877

Ubiquitin (E412J), https://www.cellsignal.com/products/primary-antibodies/ubiquitin-e4i2j-rabbit-mab/43124? site-search-type=Products & N=4294956287 & Ntt=ubiquitin & from Page=plp & Ntt=ubiquitin & Ntt=u

HMGCS2(D3U1A), https://www.cellsignal.com/products/primary-antibodies/hmgcs2-d3u1a-rabbit-mab/20940?site-search-

type=Products&N=4294956287&Ntt=hmgcs2%28d3u1a%29&fromPage=plp&_requestid=464612

Phospho-PKA Substrate(100G7E), https://www.cellsignal.com/products/primary-antibodies/phospho-pka-substrate-rrxs-t-100g7e-rabbit-mab/9624? =1664199480543&Ntt=p-PKA%20substrate&tahead=true

Phospho-(Ser/Thr) Phe, https://www.cellsignal.com/product/productDetail.jsp?productId=9631

Sirt1, https://www.cellsignal.com/product/productDetail.jsp?productId=2310

Sirt6(D8D12), https://www.cellsignal.com/product/productDetail.jsp?productId=12486

Phospho-Akt (Ser473) (D9E), https://www.cellsignal.com/product/productDetail.jsp?productId=4060

Akt, https://www.cellsignal.com/product/productDetail.jsp?productId=9272

Acety lated-Lysine, https://www.cellsignal.com/products/primary-antibodies/acety lated-lysine-antibody/9441? site-search-lysine-antibody lateral products and the search-lysine and the search-lysin

type=Products&N=4294956287&Ntt=acetyl+lysine&fromPage=plp

CHOP(L63F7), https://www.cellsignal.com/products/primary-antibodies/chop-l63f7-mouse-mab/2895?site-search-

type=Products&N=4294956287&Ntt=chop&fromPage=plp

Phospho-S6 Ribosomal Protein (Ser240/244), https://www.cellsignal.com/product/productDetail.jsp?productId=2215

S6 Ribosomal Protein (5G10), https://www.cellsignal.com/product/productDetail.jsp?productId=2217

NCoR1, https://www.cellsignal.com/products/primary-antibodies/ncor1-antibody/5948? site-search-action of the control of the

type=Products&N=4294956287&Ntt=ncor1&fromPage=plp

NEDD4,https://www.cellsignal.com/products/primary-antibodies/nedd4-antibody/2740

ATF-6 (D4Z8V), https://www.cellsignal.com/products/primary-antibodies/atf-6-d4z8v-rabbit-mab/65880?site-search-

type=Products&N=4294956287&Ntt=atf6&fromPage=plp

Phospho-PERK (Thr980) (16F8), https://www.cellsignal.com/products/primary-antibodies/phospho-perk-thr980-16f8-rabbit-

mab/3179? site-search-type=Products & N=4294956287 & Ntt=p-perk & from Page=plp-perk & from

PAK4(B-3).https://www.scbt.com/p/pak4-antibody-b-3

FGF21(H-105),chttps://www.citeab.com/antibodies/793317-sc-292879-fgf-21-antibody-h-10https://www.citeab.com/antibodies/793317-sc-292879-fgf-21-antibody-h-105

PPARα(H-2), https://www.scbt.com/ko/p/pparalpha-antibody-h-2

Sirt4, https://www.biovision.com/sirt4-antibody.html

Lamin B1 (L75), https://www.antibodypedia.com/gene/3937/LMNB1/antibody/1562122/BS3547

 ${\sf GAPDH (A531)}, https://www.citeab.com/antibodies/2207025-ap0066-gapdh-a531-polyclonal-antibody-n-apolyclonal-$

Sirt7, https://www.lsbio.com/antibodies/ihc-plus-sirt7-antibody-sirtuin-7-antibody-aa317-366-ihc-wb-western-ls-b1566/45501

FoxO1(Acetyl-Lys294), https://www.lsbio.com/antibodies/foxo1-antibody-fkhr-antibody-acetyl-lys294-ihc-wb-western-ls-c800723/827288

Sirt2 (EPR20411-105), https://www.abcam.com/sirt2-antibody-epr20411-105-ab211033.html

Sirt3, https://www.abcam.com/sirt3-antibody-ab189860.html

GRP78 BiP,https://www.abcam.com/products/primary-antibodies/grp78-bip-antibody-ab21685.html

NCOR2/SMRT, https://www.abcam.com/ncor2smrt-antibody-ab24551.html

CPT1A(8F6AE9), https://www.abcam.com/cpt1a-antibody-8f6ae9-ab128568.html

T-OXPHOS, https://www.abcam.com/products/panels/total-oxphos-rodent-wb-antibody-cocktail-ab110413.html

 $THR\beta, https://www.abcam.com/products/primary-antibodies/thyroid-hormone-receptor-beta-antibody-ab53170.html. Alternative and the state of the product of t$

 $p\text{-IRE1}\alpha, https://www.abcam.com/products/primary-antibodies/ire1-phospho-s724-antibody-ab48187.html}$

MDM2(2A10),https://www.abcam.com/mdm2-antibody-2a10-ab16895.html

 $LXR\alpha \ , \ https://www.ptglab.com/products/NR1H3-Antibody-14351-1-AP.htm$

P300, https://www.cellsignal.com/product/productDetail.jsp?productId=86377

Alexa Fluor 488-conjugated goat anti-mouse IgG1, https://www.thermofisher.com/antibody/product/Goat-anti-Mouse-IgG1-Cross-Adsorbed-Secondary-Antibody-Polyclonal/A-21121

Alexa Fluor 594-conjugated goat anti-mouse IgM, https://www.thermofisher.com/antibody/product/Goat-anti-Mouse-IgM-Heavy-chain-Cross-Adsorbed-Secondary-Antibody-Polyclonal/A-21044

Eukaryotic cell lines

Authentication

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

Cell line source(s) AML12 cells (CRL-2254), HEK293

AML12 cells (CRL-2254), HEK293T cells (CRL-3216) and Hepa 1-6 cells (CRL-1830) were purchased from ATCC.

Cell lines were authenticated by morphology. The isolation and authentication of primary hepatocytes have been reported in

our previous manuscript (10.1002/hep.32384).

Mycoplasma contamination All cells tested negative for mycoplasma.

Commonly misidentified lines (See ICLAC register)

No commonly misidentified cell lines were used in the study.

Animals and other research organisms

Policy information about <u>studies involving animals</u>; <u>ARRIVE guidelines</u> recommended for reporting animal research, and <u>Sex and Gender in Research</u>

Laboratory animals 8 week-old C57BL/6 male mice were purchased from Damul Science (Daejeon, Korea).

8 week-old ob/+, ob/ob, db/+, and db/db male mice were obtained from Jackson Laboratory (Bar Harbor, ME, USA).

Pak4 LKO mice were generated by mating PAK4Pak4flox/flox mice (B6.129S2-Pak4tm2.1Amin/J) and Albumin-cre mice (B6.Cg-Speer6-ps1Tg(Alb-cre)21Mgn/J). Male and female LKO mice were used at 8 week to 10 week of age.

Wild animals No wild animals were used in the study.

Reporting on sex | Indicate if findings apply to only one sex; describe whether sex was considered in study design, methods used for assigning sex. Provide data disaggregated for sex where this information has been collected in the source data as appropriate; provide overall numbers in

this Reporting Summary. Please state if this information has not been collected. Report sex-based analyses where performed, justify

reasons for lack of sex-based analysis.

Field-collected samples No filed-collected samples were used in the study.

Ethics oversight

All animal experiments were performed in accordance with the Guide for the Care and Use of Laboratory Animals published by the US National Institutes of Health (NIH Publication No. 85-23, revised 2011). The study protocol was approved by the Institutional

Animal Care and Use Committee of Jeonbuk National University (permit number: JBNU-2019-00122).

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