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Reporting Summary

X Life sciences

Behavioural & social sciences

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>.

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 coefficien 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.
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 <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 <u>availability of computer code</u>
Data collection No custom algorithms or new softwares were used.
Data analysis No custom algorithms or new softwares were 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/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 data availability statement. 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 source data underlying most figures are provided as a Source Data file. Other datasets generated and/or analyzed in the current 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.

Ecological, evolutionary & environmental sciences

Life sciences study design

All studies must dis	close on these	points even when the disclosure is negative.	
Sample size	No sample size	calculation in this study	
Data exclusions	No data exclusion	on in this study	
Replication	All experiments were repeated at least three times and similar results were observed.		
Randomization	No randomization in this study		
Blinding	No blinding in the	his study	
Reportin	g for sr	pecific materials, systems and methods	
Ne require information	on from authors a	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.	
Materials & exp		<u> </u>	
n/a Involved in the study n/a Involved in the study			
	Antibodies ChIP-seq Eukaryotic cell lines Flow cytometry		
Palaeontolo			
	੦੪) d other organism		
	earch participant		
Clinical dat			
Antibodies			
Antibodies used	(al PL PL (Co fro	atibodies against OGT (ab96718), OGA (MGEA5, ab124807), O-GlcNAc (RL2, ab2739), PLIN1 (ab3526), SNAP23 (ab3340), ATGL b99532), DGAT1 (11561-1-AP), Fsp27 (CIDE C, ab77115), and p-Ser (Phosphoserine, PSR-45) (Abcam); p492 phosphorylated IN1 (4855) and p517 phosphorylated PLIN1 (4856) (Vala Sciences); HA (H3663) (Sigma-Aldrich); CGI-58 (ABHD5, 12201-1-AP), IN2 (15294-1-AP), and PLIN3 (TIP47, 10694-1-AP) (Proteintech); p563-HSL (4139), phospho-Akt (Ser473, 9271), and Akt (9272) ell Signaling Technology); Myc (sc-40), DGAT2 (sc-66859), and β-actin (sc-8432) (Santa Cruz Biotechnology) were purchased om the indicated sources. Horseradish peroxidase-conjugated secondary antibodies were from Santa Cruz Biotechnology. Alexa year 594-conjugated secondary antibodies were obtained from Thermo Fisher Scientific.	
Validation	All	primary antibodies are commercially available and validation results can be found at manufacturer's websites.	
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Eukaryotic c			
Policy information a			
Cell line source(s))	C3H/10T1/2, HeLa, and 293T cells were from the American Type Culture Collection (ATCC). OGA-Tet off HeLa stable cell line was established in our laboratory.	
Authentication		Cell lines were not authenticated.	
Mycoplasma con	tamination	C3H/10T1/2, HeLa, and 293T cells were tested mycoplasma-free.	
Commonly miside		No commonly misidentified lines in this study	

Animals and other organisms

 $Policy\ information\ about\ \underline{studies\ involving\ animals;}\ \underline{ARRIVE\ guidelines}\ recommended\ for\ reporting\ animal\ research$

Laboratory animals

Ogt-flox, Adipoq-CreER and Rosa26-STOPflox-rOGT mice in the C57BL/6 background were used. Male mice were used unless otherwise stated. Experiments were performed in $4^{\sim}12$ weeks old mice used unless otherwise stated.

Wild animals

No wild animals were used.

Field-collected samples

No field-collected samples were used.

Ethics oversight

All relevant ethical regulations for animal testing and research have been complied with. All animal studies received ethical

approval from Yale University's Institutional Animal Care and Use Committee.

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