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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			
	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.			
X		A description of all covariates tested			
	x	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 statistics for biologists contains articles on many of the points above.			

### Software and code

Policy information at	pout availability of computer code
Data collection	Leica LAS X Version 2.0.0.14332; Micro-Manager1.4; 7500 Fast Software v2.0.6; Evos Software (EVOS FL Auto2, Life Technologies), EthoVision®XT Noldus, Camera OneView 4K 16 bit (Gatan), DigitalMicrograph (Gatan, JEOL JEM2100PLUS)
Data analysis	FJJI ImageJ 1.52p; KymoMaker (http://www.pharm.hokudai.ac.jp/shinkei/Kymomaker.html); Microsoft Excel 2016; WAVE version 2.6.0. (Seahorse assay); Maxquant version 1.5.3.8 (Mass spectrometry data), GraphPad Pris 8.0.1.

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

All data supporting the findings described in this study are available from the corresponding author upon reasonable request. Source data from all representative blots shown in this study as well as data underlying all quantitative analysis performed in this study are provided with the manuscript as a Source Data file. Data generated from mass spectrometry have been deposited in the ProteomeXchange Consortium via the PRIDE (PubMed ID: 26527722) partner repository with the dataset identifier PXD011279.

## Field-specific reporting

**X** Life sciences

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.

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

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

Sample size	Sample sizes were not chosen based on pre-specified effect size. Instead, multiple independent experiments were carried out using several sample replicates as detailed in the figure legends. For all experiments there was enough statistical power to detect the corresponding effect size.
Data exclusions	Data was only excluded when the quality of the sample was not optimal. Predefined quality criteria were: in living samples- vacuolization or other signs of cellular degeneration, normal cell morphology and protein transport (which is usually hampered in unhealthy cells) and image streams that are in focus; proper sample preparation and mounting in fixed samples.
Replication	Each experiment was replicated a minimum of three times and data was reliably reproduced with each replication attempt.
Randomization	Mice were genotyped for allocation into control (Cre negative) or experimental (Cre positive) groups. Within each group, all animals were generally used for experimental procedures, so no randomization was applied.
Blinding	Animals were genotyped prior to experiments, i.e. no blinding was used to allocate experimental groups. Immunofluorescence image data collection and analysis of axons was not blinded since a axonal swellings phenotype differentiated controls from KO groups. Data analysis of indistinguishable cellular structures, such as dendrites and soma of the neuron, were performed blinded. Western blot data were not blindly analysed since either KO protein or siRNA treatment were assessed for each experiment.

## 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         involved in the study       n/a         involved in the study       involved in the study         invol				
Image: Second state sta	n/a	Involved in the study	n/a	Involved in the study
Image: Second control logy   Image: Second		X Antibodies	×	ChIP-seq
Image: Sector of the sector		Eukaryotic cell lines	×	Flow cytometry
Image: Second	×	Palaeontology	×	MRI-based neuroimaging
		🗶 Animals and other organisms		•
Clinical data	×	Human research participants		
	×	Clinical data		

### Antibodies

Antibodies used Rabbit polyclonal anti-ATG16L1 MBL Cat# PM040, Lot. 017 Rabbit monoclonal (clone EPR1755(2)) anti-ATG5 abcam Cat# ab108327, Lot. GR312199-7 Rabbit polyclonal anti-ATG9 Novus Biologicals Cat# NB110-56893 Rabbit polyclonal anti-ATG13 Arigo (Biomol GmbH) Cat# ARG55112.050, Lot. 81210 Guinea pig polyclonal anti-Bassoon Synaptic Systems Cat# 141 004, Lot. 1-19 Rabbit polyclonal anti-CASPASE-3 cleaved (Asp175) Cell Signaling Cat# 9661, Lot. 45 Rabbit polyclonal anti-c-FOS Synaptic Systems Cat# 226 003, Lot. 22600313-41 Rabbit polyclonal anti-CLASP2 Millipore Cat# ABT263, Lot. 3040386 Goat polyclonal anti-DNCT1 abcam Cat# ab11806 Rabbit polyclonal anti-ELKS1 Novus Biologicals Cat# NBP1-88178 Rabbit monoclonal anti-FIP200 Cell Signaling Cat# 12436, Lot. 1 Mouse monoclonal anti-GAPDH (clone 71.1) Sigma-Aldrich Cat# G8795 Chicken polyclonal anti-GFP abcam Cat# ab13970, Lot. GR3190550-15 Mouse monoclonal (clone is not provided) anti-GFP Takara Bio Clontech Cat# 632375

Mouse monoclonal	ti-LC3B Novus Biologicals Cat# NB600-1384SS, Lot. CK-1
	(clone 4E12) anti-LC3A/B MBL Cat# M152-3, Lot. 051
Mouse monoclonal	anti-MAP2 (clone HM-2) Sigma-Aldrich Cat# M9942, Lot. 036M4804V
Mouse monoclonal	anti-MAPK activated (cloane MAPK-YT) Sigma-Aldrich Cat# M8159, Lot. 065M4806V
	ti-mCherry abcam Cat# ab167453, Lot. GR3265215-2
	anti-mCherry (clone 1C51) Novus Biologicals Cat# NBP1-96752
	ti-P-S6 Ribosomal Protein (Ser235/236) Cell Signaling Cat# 2211, Lot. 23
	al anti-p62 Progen Cat# GP62-C, Lot. 703241-02
	anti-P-AKT (clone D9E) Cell Signaling Cat# 4060, Lot.19
	anti-RAB7 (clone D95F2) Cell Signaling Cat#9367S
	ti-SYB2 Synaptic Systems Cat# 104 202
	anti-Synapsin 1 (clone 46.1) Synaptic Systems Cat# 106 001
	tti-P-TRKB (Y816) abcam Cat# ab75173
	anti-α-Tubulin (clone 3A2) Synaptic Systems Cat# 302 211, Lot. 1-6
	anti- beta-3-Tubulin (clone 2G10-TB3) eBioscience Cat# 14-4510-80, Lot. 4315809
	anti-α-Tubulin acetylated (Lys40, clone D20G3) Cell Signaling Cat# 5335, Lot. 5
	anti-α-Tubulin tyrosinated (clone TUB-1A2) Sigma-Aldrich Cat# T9028
	ti-α-Tubulin detyrosinated Millipore Cat# AB3201, Lot: 2886375
	ti-α-Tubulin Δ2 Synaptic Systems Cat# 302 213, Lot: 30221313
	ti-Ubiquitin Sigma-Aldrich Cat# U5379
	anti-WIPI2 (clone MCA5780GA) BioRad Cat# AB_10845951, Lot. 1801
0	G (H+L) peroxidase-conjugated Jackson ImmunoResearch Cat# 115-035-003
-	G, light chain specific, peroxidase conjugated Jackson ImmunoResearch Cat# 115-035-174
-	G (H+L) peroxidase-conjugated Jackson ImmunoResearch Cat# 111-035-003
	g IgG (H+L) peroxidase-conjugated Jackson ImmunoResearch Cat# 106-035-003, Lot. 124397 Cell Signaling Cat# 2729, Lot. 8
•	Villipore Cat# 12-371, Lot. 2757162
	t anti-Mouse IgG Thermo Fisher Sci Cat# A-11029; Lot. 1942237
	t anti-Rabbit IgG Thermo Fisher Sci Cat# A 11034, Lot. 1937195
	t anti-Chicken IgG Thermo Fisher Sci Cat# A-11039, Lot. 1691381
	t anti-Mouse IgG Thermo Fisher Sci Cat# A-11031, Lot. 1946341
	t anti-Rabbit IgG Thermo Fisher Sci Cat# A-11036, Lot. 1705911
	key anti-Goat IgG Thermo Fisher Sci Cat# A-11057, Lot. 2044862
	t anti-Mouse IgG Thermo Fisher Sci Cat# A-21236, Lot. 1705800 t anti-Rabbit IgG Thermo Fisher Sci Cat# A-21245, Lot. 2098544
	t anti-Guinea Pig IgG Thermo Fisher Sci Cat# A-21450, Lot. 2110845
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	t anti-Rat IgG Thermo Fisher Sci Cat# A-21247, Lot. 2005938 Goat anti-mouse IgG Abberior Cat#ST635P
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Abberior STAR 635P Goat anti-mouse IgG Abberior Cat#ST635P: manufacturer validation (https://www.abberior.com/jtl-shop/ Abberior-STAR-635P). Mazkereth, N., Rocca, F., Schubert, J.R., et al. Immunobiology 221 (12), 1395-1406 (2016).

### Eukaryotic cell lines

Policy information about <u>cell lines</u>	i
Cell line source(s)	HEK293T were obtained from Prof. M. Lammers (CECAD, Cologne) and were originally from ATCC, NSC34 cells (CLU140, Cedarlane) were obtained from Prof. B. Wirth (University Hospital, Cologne), MEF cells were obtained form Prof. T. Langer (MPI for Biology of Ageing, Cologne)
Authentication	Cell lines were authenticated by morphology
Mycoplasma contamination	Cell lines were not tested for mycoplasma contamination
Commonly misidentified lines (See <u>ICLAC</u> register)	No commonly misidentified cell lines were used in the study.

### Animals and other organisms

Policy information about <u>stud</u>	dies involving animals; ARRIVE guidelines recommended for reporting animal research				
Laboratory animals	Mus Musculus, C57BI/6, both sexes, postnatal days P0-P5, 12-13 weeks old				
Wild animals	This study did not involve wild animals				
Field-collected samples	This study did not involve samples collected from the field				
Ethics oversight	All animal experiments were reviewed and approved by the ethics committee of the "Landesamtes für Natur, Umwelt- und Verbraucherschutz des Landes Nordrhein-Westfalen", Cologne (2015.A578, 2016.A041, 2016.A451)				

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