

DEC 21, 2022

## OPEN ACCESS

### DOI:

[dx.doi.org/10.17504/protocols.io.5jyl89648v2w/v1](https://doi.org/10.17504/protocols.io.5jyl89648v2w/v1)

### External link:

<https://doi.org/10.1093/brain/awac325>

**Protocol Citation:** Marc P.M. Soutar, Daniela Melandri, Benjamin O'Callaghan, Emily Annuario, Amy E. Monaghan, Paul J. Whiting, Helene Plun-Favreau 2022. Regulation of mitophagy by the NSL complex underlies genetic risk for Parkinson's disease: Cell-Based *in vitro* Assays.

### protocols.io

<https://dx.doi.org/10.17504/protocols.io.5jyl89648v2w/v1>

### MANUSCRIPT CITATION:

Soutar MPM, Melandri D, O'Callaghan B, Annuario E, Monaghan AE, Welsh NJ, D'Sa K, Guelfi S, Zhang D, Pittman A, Trabzuni D, Verboven AHA, Pan KS, Kia DA, Bictash M, Gandhi S, Houlden H, Cookson MR, Kasri NN, Wood NW, Singleton AB, Hardy J, Whiting PJ, Blauwendraat C, Whitworth AJ, Manzoni C, Ryten M, Lewis PA, Plun-Favreau H, Regulation of mitophagy by the NSL complex underlies genetic risk for Parkinson's disease at 16q11.2 and MAPT H1 loci. *Brain* 145(12). doi: [10.1093/brain/awac325](https://doi.org/10.1093/brain/awac325)

**License:** This is an open access protocol distributed under the terms of the [Creative Commons Attribution License](https://creativecommons.org/licenses/by/4.0/), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited

# Regulation of mitophagy by the NSL complex underlies genetic risk for Parkinson's disease: Cell-Based *in vitro* Assays

Marc P.M. Soutar<sup>1</sup>, Daniela Melandri<sup>1</sup>, Benjamin O'Callaghan<sup>1</sup>, Emily Annuario<sup>2</sup>, Amy E. Monaghan<sup>3,4</sup>, Paul J. Whiting<sup>3,4</sup>, Helene Plun-Favreau<sup>5,6</sup>

<sup>1</sup>Department of Neurodegenerative Diseases, UCL Queen Square Institute of Neurology, London, UK;

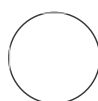
<sup>2</sup>Kings College, London, UK;

<sup>3</sup>UCL Alzheimer's Research UK, Drug Discovery Institute, London, UK;

<sup>4</sup>UCL Dementia Research Institute, London, UK;

<sup>5</sup>UCL Queen Square Institute of Neurology, London, UK;

<sup>6</sup>Aligning Science Across Parkinson's (ASAP) Collaborative Research Network, Chevy Chase, MD, 20815.



Benjamin O'Callaghan

## ABSTRACT

Impaired mitophagy is a key causative pathway in familial Parkinson's disease, but its relevance to idiopathic Parkinson's disease is unclear. We used a mitophagy screening assay to evaluate the functional significance of risk genes identified through genome-wide association studies and describe the cell-based *in vitro* assays used.

## MATERIALS TEXT

### Mitochondrial Toxins

Oligomycin (mitochondrial complex V inhibitor) was purchased from Cayman Chemicals (11341) and from Sigma-Aldrich (O4876)

Antimycin A (mitochondrial complex III inhibitor) were purchased from Sigma-Aldrich (A8674).

### All siRNAs were purchased as pre-designed siGENOME SMARTpools from Dharmacon:

non-targeting (D-001206-13)

PINK1 (M-004030-02), PLK1 (L-003290-00)

KIF-11 (L-003317-00), KAT8 (M-014800-00)

KANSL1 (M-031748-00)

KANSL2 (M-020816-01)

KANSL3 (M-016928-01), HCFC1 (M-019953-01)

MCRS1 (M-018557-00)

OGT (M-019111-00)

PHF20 (M-015234-02)

WDR5 (M-013383-01).

### The following antibodies were used for immunocytochemistry:

mouse anti TOM20 (Santa Cruz, sc-17764, 1:1000),

rabbit anti phospho-ubiquitin (Ser65) (Cell Signaling, 37642, 1:1000),

rabbit anti phospho-Parkin (Ser65) (Abcam/Michael J. Fox Foundation, MJF17, 1:250)

rabbit anti FLAG (Sigma-Aldrich, F7425, 1:500),

AlexaFluor 488 goat anti rabbit (Invitrogen, A11008, 1:2000),

AlexaFluor 568 goat anti mouse (Invitrogen, A11004, 1:2000).

### The following antibodies were used for immunoblotting:

**Protocol status:** Working

**Created:** May 17, 2022

**Last Modified:** Dec 21, 2022

**PROTOCOL integer ID:** 62752

**Keywords:** Parkinson's disease, Genetic Risk, Mitophagy, NSL Complex, GWAS, KANSL1, KAT8, mitophagy, MAPT locus, H1 Haplotype, HCS, ASAPCRN

mouse anti TIM23 (BD Biosciences, 611223, 1:1000),  
 rabbit anti TOM20 (Santa Cruz, sc-11415, 1:1000),  
 rabbit anti phospho-ubiquitin (Ser65) (Merck Millipore, ABS1513-I, 1:1000, and Cell Signaling, 37642, 1:1000),  
 mouse anti GAPDH (Abcam, ab110305, 1:10000),  
 rabbit anti KAT8 (Abcam, ab200600, 1:1000),  
 rabbit anti total-Tau (DAKO, 1:10000),  
 mouse anti V5 tag (Invitrogen, R960-25, 1:1000),  
 rabbit anti KANSL1 (Sigma-Aldrich, HPA006874, 1:500),  
 rabbit anti pParkin(Ser65) (Abcam/Michael J. Fox Foundation, MJF17, 1:1000),  
 mouse anti FLAG M2 (Sigma-Aldrich, F3165, 1:1000),  
 sheep anti pRab8A(Ser111) (MRC Protein Phosphorylation and Ubiquitylation Unit, University of Dundee, 1ug/ml preblocked with 10ug/ml non-phosphorylated peptide),  
 rabbit anti total Rab8 (Cell Signaling, 6975, 1:1000),  
 IRDye 680LT donkey anti mouse (LI-COR Biosciences, 925-68022, 1:20000),  
 IRDye 800CW donkey anti rabbit (LI-COR Biosciences, 925-32213, 1:20000),  
 IRDye 800CW Donkey anti-Goat (LI-COR Biosciences, 926-32214, 1:20000),  
 IRDye® 680RD Goat anti-Rabbit (LI-COR Biosciences, 926-68071, 1:20000).

The generation of rabbit monoclonal anti PINK1 antibody has been described in the following manuscript:


*Sci Rep.* 2015. DOI: <https://doi.org/10.1038/s41598-018-26949-6>

The generation and use of sheep anti pRab8A(Ser111) antibody has been described in the following manuscript:

*EMBO J.* 2015. DOI: <https://doi.org/10.15252/emboj.201591593>

## Cell Culture and siRNA Transfection

1

Culture cells in Dulbecco's Modified Eagle Medium (DMEM, Gibco, 11995-0 65) supplemented with 10% heat-inactivated foetal bovine serum (FBS, Gibco) in a humidified chamber at  37 °C with 5% CO<sub>2</sub>.

### Note

Parkin over-expressing (POE) SHSY5Y cells are a kind gift from H. Ardley and have been described in the following manuscript.

*Mol Biol Cell.* 2003. DOI: <https://doi.org/10.1091/mbc.e03-02-0078>

Mt-Keima POE SHSY5Y cells were a kind gift of C. Luft and have been described in the following manuscript.

*Autophagy.* 2019. DOI: <https://doi.org/10.1080/15548627.2019.1603549>

PINK1-HA overexpressing SHSY5Y cells were a kind gift from E. Deas and have been described in the following manuscript.


*Hum Mol Genet.* 2011. DOI: <https://doi.org/10.1093/hmg/ddq526>

- 2 For siRNA transfection, transfect cells using DharmaFECT1 transfection reagent (Dharmacon, T-2001-03) according to the manufacturer's instructions.


## High Content siRNA Screen

### 3 Cell Plating and siRNA Transfection



Dispense siRNA into Geltrex-coated 96-well CellCarrier Ultra plates (Perkin Elmer) at a final concentration of [M] 30 nanomolar (nM) using the Echo 555 acoustic liquid handler (Labcyte).

- 4 Add  25 µL of DMEM containing 0.48% of DharmaFECT1 transfection reagent to each well and incubate for

30m

 00:30:00

- 5 Seed POE SHSY5Y cells using the CyBio SELMA (Analytik Jena) at 15,000 cells per well, 100 µl per well in DMEM + [M] 10 % (v/v) FBS.



- 6 Incubate cells for  72:00:00 in a humidified chamber at  37 °C with 5% CO<sub>2</sub>.

3d

- 7 Treat cells with [M] 10 micromolar (µM) oligomycin/ [M] 10 micromolar (µM) antimycin for  03:00:00 to induce mitophagy.

3h

### 8 Immunofluorescence and Image Capture Analysis

Fix cells with [M] 4 % (w/v) PFA (Sigma-Aldrich,F8775),  00:15:00,  Room temperature

15m

- 9 Remove PFA solution and wash cells 3x with PBS.

- 10 Block and permeabilise cells with [M] 10 % (v/v) FBS, [M] 0.25 Mass / % volume Triton X-100 in PBS, 1 h, RT.

- 11 Immunostain cells with rabbit anti pUb(Ser65) (Cell Signaling, 37642, 1:1000) and mouse anti TOM20 (Santa Cruz, sc-17764, 1:1000) primary antibodies in [M] 10 % (v/v) FBS/PBS, ⌚ 02:00:00 ,  
🌡 Room temperature .
- 12 After 3x PBS washes, add AlexaFluor 568 anti-mouse and 488 anti-rabbit secondary antibodies and Hoechst 33342 (Thermo Scientific, 62249) in [M] 10 % (v/v) FBS/PBS, (1:2000 dilution for all) and incubate for ⌚ 01:00:00 , 🌡 Room temperature .
- 13 Remove secondary antibodies and wash cells 3x with PBS.
- 14 Image plates using the Opera Phenix (Perkin Elmer).  
Acquire 5x fields of view and 4x 1 µm Z-planes per well, using the 40X water objective, NA1.1.
- 15 Analyse images in an automated way using the Columbus 2.8 analysis system (Perkin Elmer) to measure the integrated intensity of pUb(Ser65) within the whole cell.
- 16 First of all, load the image as a maximum projection, before segmenting to identify the nuclei using the Hoechst 33342 channel (method B).
- 17 To identify the cytoplasm, use the “Find Cytoplasm” building block (method B) on the sum of the Hoechst and Alexa 568 channels.
- 18 Identify pUb(Ser65) as spots (method B) on the Alexa 488 channel, before measuring their integrated intensity.
- 19 **Screen Quality Control, Data Processing and Candidate Selection**  
Quality control screen plates based on the efficacy of the PINK1 siRNA control and O/A treatment window (minimum 3-fold).  
Check data for edge effects using Dotmatics Vortex visualization software.
- 20 Quality control raw data using robust Z prime > 0.5.

2h

1h



Process data using Python for Z score calculation before visualization in Dotmatics Vortex. Candidates should be considered a hit where Z score is  $\geq 2$  or  $\leq -2$ , and where replication of efficacy is seen both within and across plates.

## 21 siRNA Libraries

Resuspend siRNAs in RNase-free water for a final concentration of 20 micromolar ( $\mu\text{M}$ ).

### Note

siRNA libraries were purchased from Dharmacon as an ON-TARGETplus SMARTpool Cherry-pick siRNA library, 0.25 nmol in a 384-well plate.

22 Add SCR, PINK1 and PLK1 or KIF11 controls to the 384-well plate at a concentration of 20 micromolar ( $\mu\text{M}$ ) before dispensing into barcoded assay-ready plates.

## KANSL1 iNeuron Culture and Differentiation

23 Culture human induced pluripotent stem cells (hiPSCs) on Geltrex (Thermofisher) coated culture dishes in mTeSR1 (StemCell Technologies) and maintained in a humidified 37 °C incubator, 5% CO<sub>2</sub>.

### Note

Isogenic hiPSC lines with/without a heterozygous loss of function (LoF) frameshift mutation in Exon2 of the KANSL1 gene (c.531insT), which have also being stable transduced with transgenes permitting doxycycline-inducible overexpression of murine Ngn2 were a kind gift from the lab of N. Nadif Kasri and have been described in the following manuscript

*Autophagy*. 2022. DOI: <https://doi.org/10.1080/15548627.2021.1936777>

24 Differentiate isogenic KANSL1<sup>+/+</sup> and KANSL1<sup>+/-</sup> hiPSCs into excitatory cortical neurons by doxycycline induced overexpression of murine Ngn2.

### Note

iNeuron differentiation protocol adapted from that described in the following manuscripts.

*Autophagy*. 2022. DOI: <https://doi.org/10.1080/15548627.2021.1936777>

*J Vis Exp*. 2017. DOI: <https://doi.org/10.3791/54900>

- 25 On d0 dissociate hiPSCs into a single cell suspension using accutase (Sigma).
- 26 Plate cells onto geltrex coated dishes in induction medium consisting of DMEM/F12 supplemented with 1x Glutamax, 1x non-essential amino acids (NEAAs), 1x N2-supplement (all Thermofisher) and **100 nM 4 μg/ml** doxycycline (Sigma).  
Supplement induction media with **100 nM 10 micromolar (μM)** Y-27632 Rho Kinase inhibitor (ROCKi, Peprotech) during initial seeding.  
Seed  $7.5 \times 10^5$  cells per 6-well.
- 27 **24:00:00** later (d1) and **48:00:00** later (d2) perform a full medium change with freshly prepared induction media without Y-27632 ROCKi. 2d
- 28 On d3 perform a full medium change with freshly prepared N2-B27 media consisting of a 1:1 mixture of DMEM/F12:Neurobasal supplemented with 0.5x N2-supplement, 0.5x B27 supplement, 0.5x NEAAs, 0.5x Glutamax, **100 nM 45 micromolar (μM)** 2-Mercaptoethanol (all Thermofisher), **100 nM 2.7 μg/ml** insulin (Sigma).  
Supplement N2-B27 media with **100 nM 2 micromolar (μM)** Cytosine β-D-arabinofuranoside (ARA-C) (Sigma) on d3. 1d
- 29 Every 3-4 days thereafter, perform a half media change with N2-B27 lacking ARA-C. 1w 5d
- 30 Perform a half media change with N2-B27 on d16 and collect cells for experimental assays 24 h later on d17. 1d

## CRISPRi-3N iNeuron Culture and Differentiation

- 31 hiPSCs are cultured as outlined above for KANSL1 hiPSCs in Step 23 **⇒ go to step #23**

### Note

A hiPSC line stably transduced with transgenes permitting doxycycline-inducible overexpression of murine Ngn2 at the AAVS1 safe-harbor locus, and stably transduced with constitutively expressed enzymatically dead Cas9 (dCas9)-KRAB transcriptional repressor fusion protein at the CLYBL promoter safe-harbor locus (CRISPRi-3N hiPSCs) was a kind gift from the labs of M. E. Ward and M. Kampmann, and have been described in the following manuscripts.

*Curr Protoc cell Biol.* 2018. DOI: <https://doi.org/10.1002/cpcb.51>

*Neuron.* 2019. DOI: <https://doi.org/10.1016/j.neuron.2019.07.014>

- 32 First transduce CRISPRi-i3N hiPSCs with lentiviral particles encoding a mCherry-reporter and sgRNA sequences targeting the promoter regions of *KANSL1*, *KAT8* or *PINK1*, or a non-targeting sgRNA control. Then differentiated transduced cells into excitatory cortical neurons by doxycycline induced overexpression of murine Ngn2.

#### Note

CRISPRi-i3N-iNeuron differentiation protocol adapted from the following manuscripts.



*Curr Protoc cell Biol.* 2018. DOI: <https://doi.org/10.1002/cpcb.51>



*Neuron.* 2019. DOI: <https://doi.org/10.1016/j.neuron.2019.07.014>


#### Note

See Steps 39-41 for lentiviral particle generation and encoded sgRNA sequences.

- 33 On d-1 dissociate CRISPRi-i3N hiPSCs into a single cell suspension using accutase and reverse transduce with sgRNA lentiviral supernatant in mTeSR1 supplemented with 5ug/ml polybrene and **[M] 10 micromolar ( $\mu$ M)** Y-27632 ROCKi. Seed  $4.5 \times 10^5$  cells onto Geltrex-coated 6-well plates. 1d

- 34  24:00:00 later (d0) change media to induction media (composition outlined above in Step 26  go to step #26 for KANSL1 iNeurons). 1d

- 35  24:00:00 later (d1) and  48:00:00 later (d2) perform a full medium change with freshly prepared induction media. 3d

- 36 On d3, dissociate differentiating iNeurons into a single cell suspension using accutase (Sigma) and seed in N2-B27 media (composition outlined above in Step 28  go to step #28 for KANSL1 iNeurons) into Geltrex-coated 96-well CellCarrier Ultra plates for immunofluorescence (IF) ( $3 \times 10^4$  cells per well) and 12-well plates ( $5 \times 10^5$  cells per well) for biochemistry purposes. 1d

- 37 Perform a half media change with N2-B27 the following day (d4) and every 3-4 days thereafter. 1w 5d

38 Perform a half media change with N2-B27 on d16 and collect cells for experimental assays 24 h later on d17.

1d

## Lentiviral Particle Generation

39

1d

Transfect 70-90% confluent Lenti-X human embryonic kidney 293T cells (HEK293T) cultured in DMEM 10% FBS media with pMD2.G and pCMVR8.74 alongside appropriate delivery plasmids: pLV[Exp]-U6>sgRNA-hPGK>mApple (Vectorbuilder) plasmids (for sgRNA), empty pLVX-EF1α-IRES-Puro (Clontech, Takara Bio), V5-KANSL1 pLVX-EF1α-IRES-Puro, or V5-KAT8 pLV[Exp]-EF1α-IRES-Puro at a 1:1:2 molar mass ratio using Lipofectamine 3000 (Invitrogen).


### Note

pMD2.G (Addgene plasmid #12259) and pCMVR8.74 (Addgene plasmid # 12259) were gifts from Didier Trono. KANSL1 cDNA (ENST00000432791.7) with N-terminal V5 tag was cloned into the pLVX-EF1α-IRES-Puro plasmid using SpeI and NotI restriction sites. See Table below for sgRNA plasmids and V5-KAT8 pLV[Exp]-EF1α-IRES-Puro plasmids (Vectorbuilder)

A	B	C	D
Gene Target	sgRNA Sequence (5'-3')	Plasmid	Plasmid Map
<i>Non-Targeting</i>	gctgcatggggcgcgcaatca	pLV[Exp]-U6>NT-Seq1-hPGK>mApple	<a href="https://en.vectorbuilder.com/vector/VB210129-1165psy.html">https://en.vectorbuilder.com/vector/VB210129-1165psy.html</a>
<i>KANSL1</i>	gccgccgcggcgagacgagt	pLV[Exp]-U6>KANSL1_P1_Seq1-hPGK>mApple	<a href="https://en.vectorbuilder.com/vector/VB210126-1227jjz.html">https://en.vectorbuilder.com/vector/VB210126-1227jjz.html</a>
<i>KAT8</i>	ggttgccggcggggacttcag	pLV[Exp]-U6>KAT8_P1_Seq1-hPGK>mApple	<a href="https://en.vectorbuilder.com/vector/VB210129-1139enp.html">https://en.vectorbuilder.com/vector/VB210129-1139enp.html</a>
<i>PINK1</i>	ggcctgtcgaccgccatgg	pLV[Exp]-U6>PINK1_P1_Seq1-hPGK>mApple	<a href="https://en.vectorbuilder.com/vector/VB210129-1160qka.html">https://en.vectorbuilder.com/vector/VB210129-1160qka.html</a>
N/A	N/A	pLV[Exp]-EF1A>V5/hKAT8[NM_032188.3]:IRES-Puro	<a href="https://en.vectorbuilder.com/vector/VB220307-1149nfg.html">https://en.vectorbuilder.com/vector/VB220307-1149nfg.html</a>

sgRNA sequences and encoding plasmids, and V5-KAT8 encoding plasmid.





40 The next day, perform a full media change with mTeSR1 (for sgRNA lentivirus) or DMEM [M] 10 % (v/v) FBS media and culture cells for  24:00:00 .


1d

41 Collect the lentivirus containing mTeSR1 / DMEM [M] 10 % (v/v) FBS and dilute 1:2 with fresh mTeSR1 or [M] 10 % (v/v) FBS before filtering through 0.44 µm PES filters.

## Mitochondrial Enrichment and Western Blotting

42  Transfect POE SHSY5Y and H4 cells with [M] 100 nanomolar (nM) siRNA and incubate for  72:00:00 .

3d

43 Culture KANSL1 iNeurons as detailed above in Steps 23-30  go to step #23 .


44 Use whole cell lysates from POE SHSY5Y cells, H4 cells, and KANSL1 iNeurons. For some experiments, POE SHSY5Y lysates you can first fractionate into cytoplasmic and mitochondria-enriched preparations to facilitate detection of mitochondrial localised proteins of interest.

### Note

Mitochondrial enrichment and Western blotting protocols have been described previously in the following manuscript.

*Sci Rep.* 2018. DOI: <https://doi.org/10.1038/s41598-018-26949-6>

## KANSL1 and KAT8 siRNA KD Rescue

45  Transfect POE SHSY5Ys with 25 nM siRNA (d0) and incubate for 48 h.



46 Transduce siRNA KD cells with lentivirus in the presence of 10 µg/ml polybrene (d2).

- 47 Perform a full media change the following day (d3) and collect 4 days post siRNA transfection (and 2 days post lentivirus transduction) (d4).

## pRab8A(Ser111) Measurements

- 48
- Transfect SHSY5Y cells stably overexpressing PINK1-HA with 100 nM siRNA and incubate for 72 h.
- 49 Immunoprecipitate 200ug of protein (whole cell lysate) with Protein A Dynabeads™ (Invitrogen) prebound with 0.5ug of rabbit anti total-Rab8 antibody (Cell Signaling, 6975) at 4 °C overnight.
- 50 Elute samples from the beads by heating at 95 °C in 2x LDS supplemented with 50mM DTT for 5min.

## Immunofluorescence

- 51
-  Reverse transfect SHSY5Y cells with 50 nM siRNA in 96-well CellCarrier Ultra plates according to the manufacturer's instructions and incubate for 72 h.
- 52 Culture CRISPRi-i3N iNeurons as described above in Steps 31-38 [⇒ go to step #31](#) .
- 53 Treat, fix and stain cells as per the screening protocol detailed above in Steps 8-13 [⇒ go to step #8](#) .
- 54
-  For visualisation purposes, select brightness and contrast settings on the SCR (siRNA KD SHSY5Y) or no transduction (No TD) (CRISPRi-i3N iNeurons) controls and apply the same settings to all other conditions. Present representative images as maximum projections of the channels for one field of view.

55

Extract total RNA from cells using the Monarch Total RNA Miniprep Kit (New England Bioscience) with inclusion of the optional on-column DNase treatment.

56

Reverse transcribe RNA with SuperScript™ IV reverse transcriptase and random hexamers (Invitrogen).

57

Dilute the cDNA product then subject to quantitative real-time PCR (qPCR) using Fast SYBR™ Green Master Mix (Applied Biosystems) and gene specific primer pairs (See Table Below) on a QuantStudio™ 7 Flex Real-Time PCR System (Applied Biosystems).





List of primer pairs used for RT-qPCR of target genes

- 58 Calculate relative mRNA expression levels using the  $2^{-\Delta\Delta C_t}$  method and *RPL 18A* (SHSY5Y) or *UBC* (iNeurons) as the house-keeping gene.

## Mitophagy Measurement Using mt-Keima Reporter

59



Reverse transfect stable mt-Keima expressing POE SHSY5Y cells with 50 nanomolar (nM) siRNA in 96-well CellCarrier Ultra plates according to the manufacturer's instructions and incubate for 72:00:00.

60

For the assay, replace the cell medium with phenol-free DMEM + 10% FBS containing Hoechst 33342 (1:10000) and either DMSO or 1 micromolar ( $\mu$ M) oligomycin/ 1 micromolar ( $\mu$ M) antimycin to induce mitophagy.

61

Image cells immediately on the Opera Phenix (PerkinElmer) at 37 °C with 5% CO<sub>2</sub>, acquiring 15x single plane fields of view, using the 63X water objective, NA1.15.

### Note

The following excitation wavelengths and emission filters were used: cytoplasmic Keima: 488 nm, 650–760 nm; lysosomal Keima: 561 nm, 570–630 nm; Hoechst 33342: 375 nm, 435–480 nm.

- 62 Images were analysed in an automated way using the Columbus 2.8 analysis system (Perkin Elmer) to measure the mitophagy index.
- 63 Identify cells using the nuclear signal of the Hoechst 33342 channel, before segmenting and measuring the area of the cytoplasmic and lysosomal mt-Keima.
- 64 Calculate the mitophagy index as the ratio between the total area of lysosomal mitochondria and the total area of mt-Keima (sum of the cytoplasmic and lysosomal mtKeima areas) per well.

## Mitochondrial Membrane Potential Measurements using TMRM

- 65 Transfect POE SHSY5Ys with **100 50 nanomolar (nM)** siRNA in 96-well CellCarrier Ultra plates using Dharmafect1 and incubate for **72:00:00** . 3d
- 66 Perform Tetramethylrhodamine, Methyl Ester, Perchlorate (TMRM) measurements in redistribution mode where a decrease in TMRM signal intensity is associated with mitochondrial membrane potential ( $\psi_m$ ) depolarisation (confirmed post-imaging by treatment with **100 1 micromolar ( $\mu$ M)** Carbonyl cyanide m-chlorophenyl hydrazone, CCCP,  $\psi_m$  depolarising agent).
- 67 Incubate live cells in **100 25 nanomolar (nM)** TMRM (Sigma Aldrich) diluted in Hanks' Balanced Salt Solution (HBSS, Gibco) buffered with **100 10 millimolar (mM)** HEPES (Sigma Aldrich), **pH 7.4** for **00:40:00** at **37 °C** . 40m
- 68 Image cells on the Opera Phenix (PerkinElmer) at **37 °C** , acquiring confocal z-stacks for 27 fields of view across 3 individual wells per experimental condition using the 40X water objective, NA1.1.

### Note

Excite TMRM with the 561 nm laser (50% power) and collect signal with 570-630nm emission filter.

- 69 Analyse images in an automated way using the Columbus 2.8 analysis system (Perkin Elmer) to measure the TMRM signal intensity of maximum intensity projections.

## Statistical Analysis

- 70 Normalise intensity measurements from imaging experiments for each experiment. N numbers shown in figure legends should refer to the number of independent, replicate experiments. Within each experiment, calculate the mean values of every condition from a minimum of 3 technical replicates.
- 71 Normalise integrated density measurements from Western blot experiments to control conditions. Wherever possible, avoid normalisation to conditions for statistical comparisons in order to maintain experimental error associated.
- 72 Subject Data to either one-way or two-way ANOVA with Dunnett's post-hoc analysis for multiple comparisons, or other appropriate statistical analysis. Present data with error bars showing mean  $\pm$  standard deviation (SD) from replicate experiments.

### Note

We use GraphPad Prism 9 (La Jolla, California, USA) for statistical analyses and graph production.