

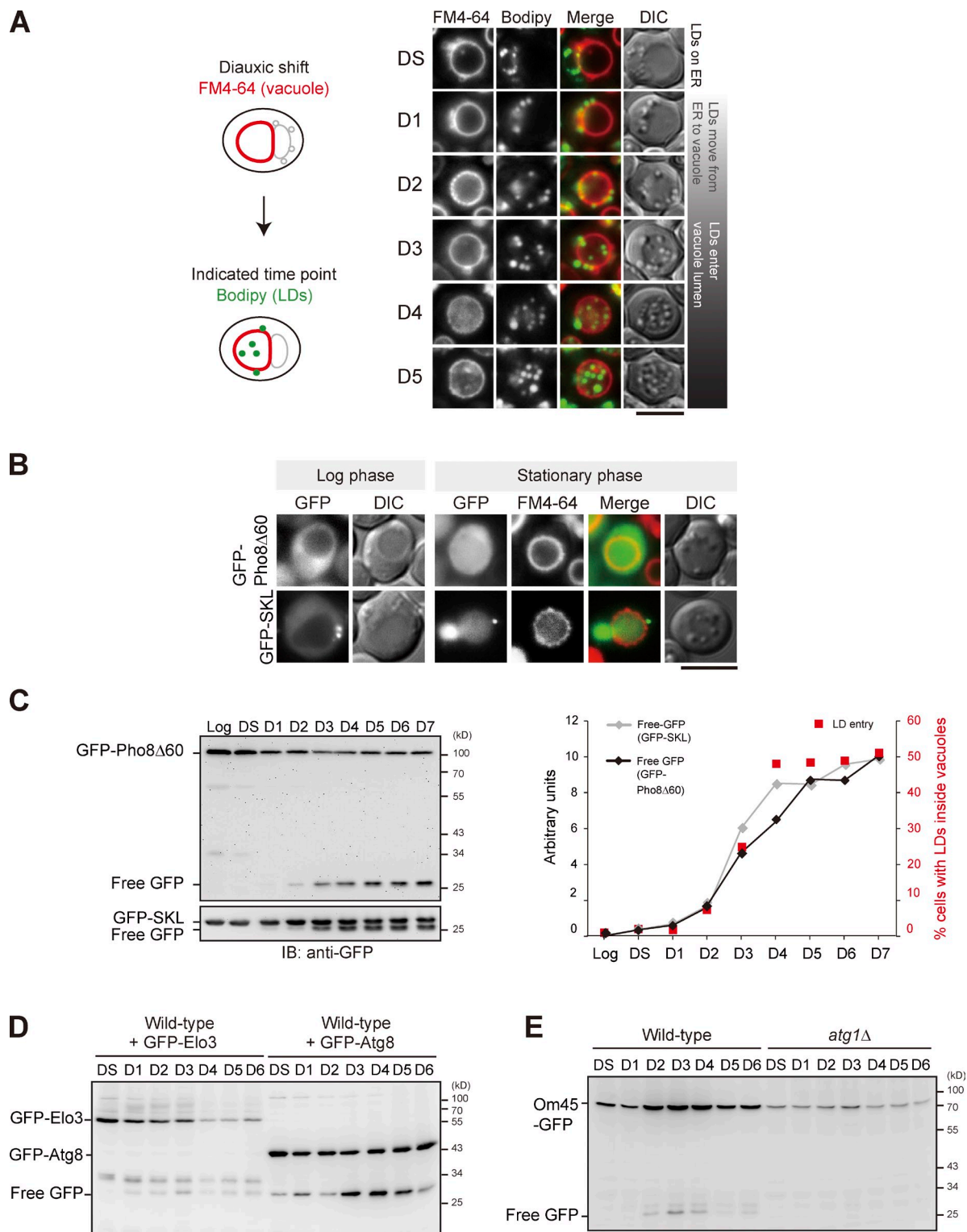
Wang et al., <http://www.jcb.org/cgi/content/full/jcb.201404115/DC1>

Figure S1. **Various forms of autophagy are activated during stat-phase.** (A, left) The scheme of vacuole and LD staining. (right) Representative images of LD association with vacuoles in wild-type cells grown from DS to D5. (B) Cells expressing GFP-Pho8 Δ 60 and GFP-SKL were imaged by fluorescence microscopy under growth conditions as indicated. (C, left) Cells expressing GFP-Pho8 Δ 60 and GFP-SKL under growth conditions as indicated were lysed, and the lysates were analyzed by immunoblotting (IB) with the anti-GFP antibody. (right) The free GFP signals were quantified and plotted. The percentage of cells containing LDs inside vacuole lumen in A on the same days was compared. The data shown are from a single representative experiment out of three repeats. (D) Wild-type cells expressing GFP-Elo3 or GFP-Atg8 under growth conditions as indicated were lysed, and the lysates were analyzed by immunoblotting with anti-GFP antibody. (E) Wild-type and *atg1* Δ cells harboring endogenous Om45-GFP under growth conditions as indicated were lysed, and the lysates were analyzed by immunoblotting with the anti-GFP antibody. DIC, differential interference contrast. Bars, 5 μ m.

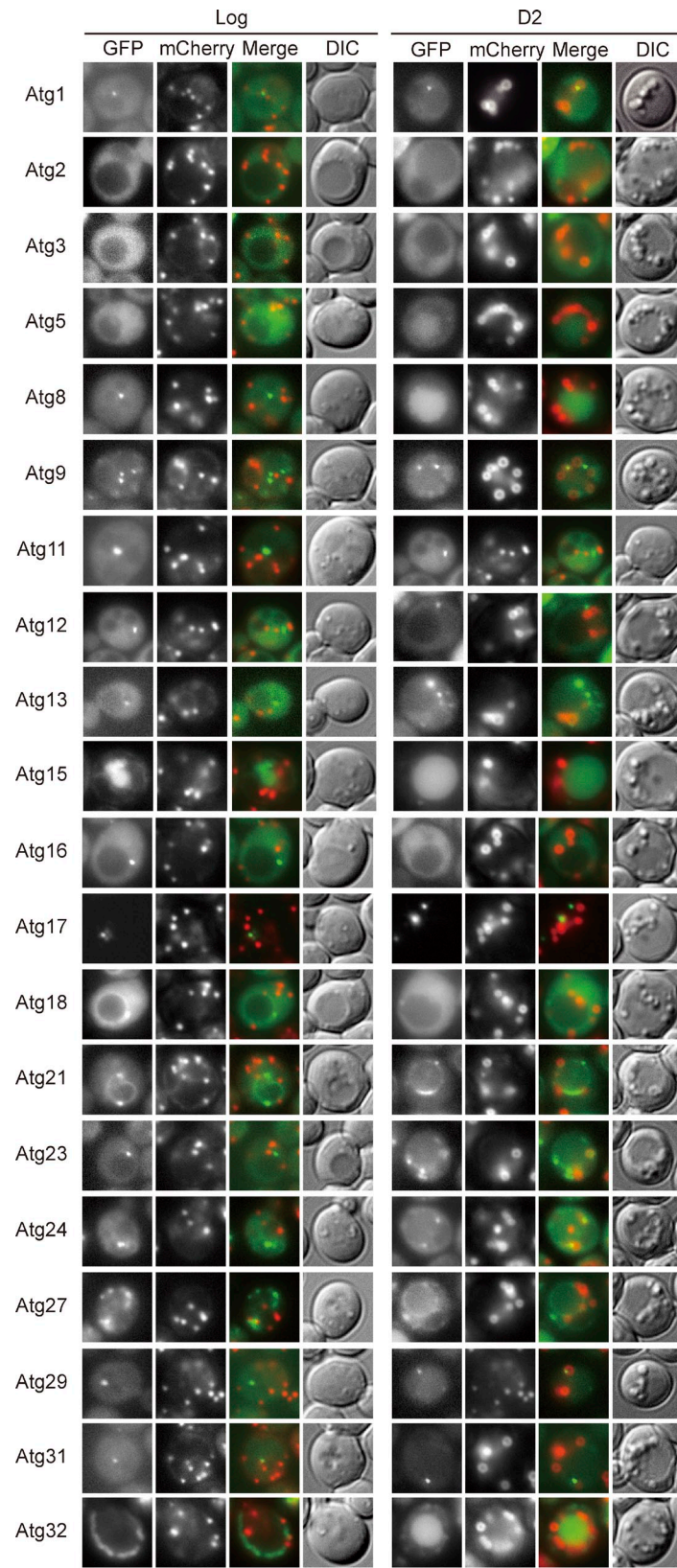


Figure S2. **Localization of various GFP-tagged Atg proteins during log and stat-phase.** Cells expressing Erg6-mCherry and various GFP-tagged Atg proteins as indicated grown in SC medium to log phase or D2 were imaged by fluorescence microscopy. DIC, differential interference contrast. Bar, 5 μ m.

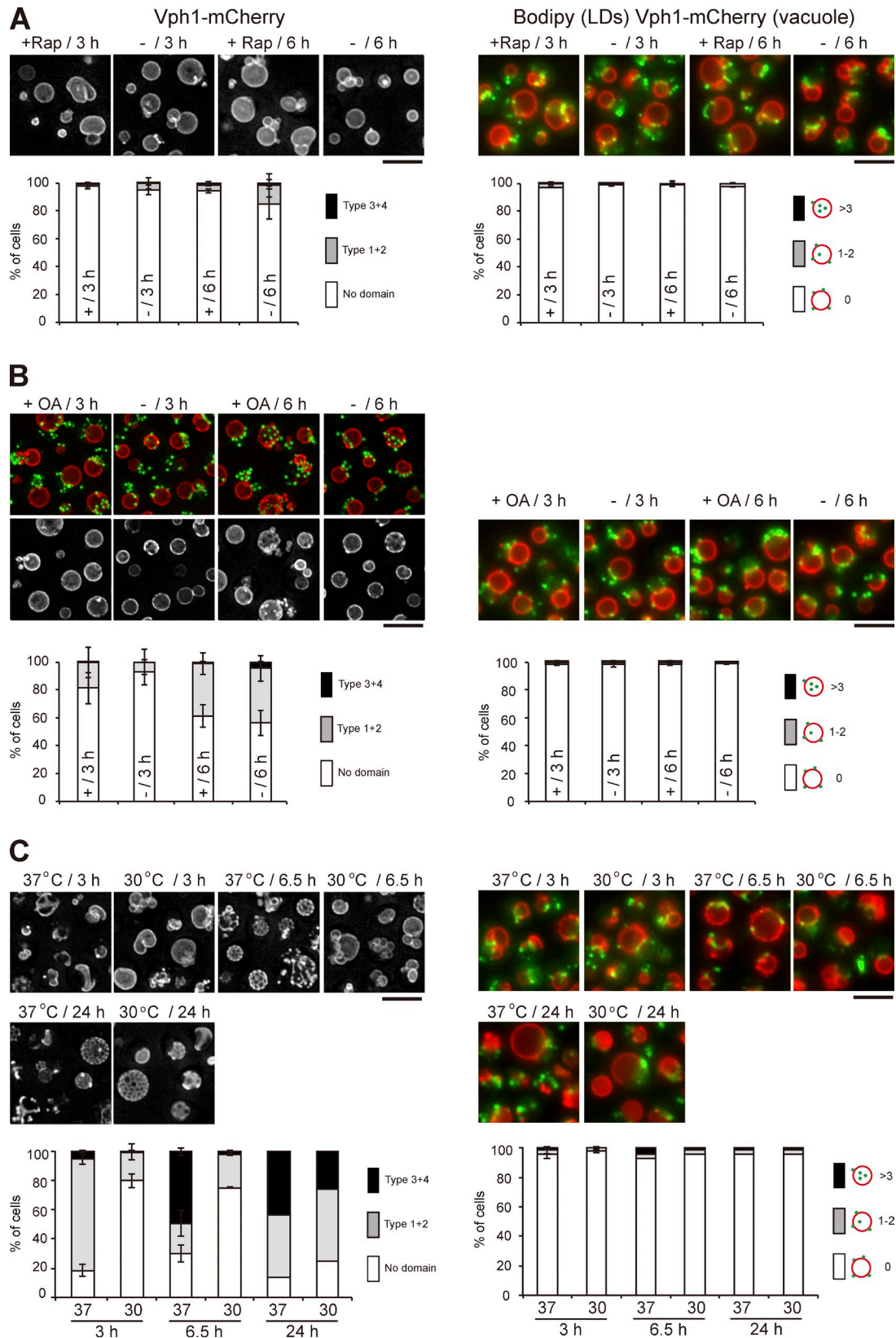


Figure S3. **Rapamycin, oleate, and higher temperature shift did not induce LD translocation into vacuoles.** (A) Cells expressing Vph1-mCherry grown in SC medium at 30°C to log phase were treated with (+) or without (-) 0.2 μ g/ml rapamycin (Rap) for 3 or 6 h. (left) The representative images of Vph1-mCherry processed by deconvolution and maximal projection. The data were quantified based on the three indicated patterns. (right) Cells were stained with BODIPY (LDs) and imaged by fluorescence microscopy. The data were quantified based on the three indicated patterns. (B) Same as A, except that cells were grown in SC medium with 1% Brij to log phase and treated with (+) or without (-) 1% oleic acids (OA). (left) The merged Vph1-GFP and LDs (BODIPY) images were also shown for the comparison of LD numbers. (C) Same as A, except that cells were grown in SC medium to DS followed by 1:1 diluted into SC medium prewarmed to 30 or 37°C and incubated for 3, 6.5, or 24 h. Data are means \pm SEM. Bars, 5 μ m.

Table S1. Yeast strains and plasmids used in this study

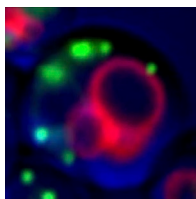
Strain	Description	Source
BY4742	<i>MATα leu2Δ his3Δ ura3Δ lys2Δ</i>	Laboratory collection
CWY6290	<i>ELO3-VENUS::LEU HIS::GPD-CFP-PHO8 ERG6-mCherry::KAN BY4742</i>	This study
CWY6302	<i>atg1Δ::HYG BY4742</i>	This study
CWY6304	<i>atg2Δ::HYG BY4742</i>	This study
Laboratory collection	<i>atg3Δ::KAN BY4742</i>	Invitrogen
CWY6469	<i>atg4Δ::KAN BY4742</i>	This study
Laboratory collection	<i>atg5Δ::KAN BY4742</i>	Invitrogen
CWY6306	<i>atg6Δ::HYG BY4742</i>	This study
CWY6308	<i>atg7Δ::HYG BY4742</i>	This study
CWY6471	<i>atg8Δ::KAN BY4742</i>	This study
Laboratory collection	<i>atg9Δ::KAN BY4742</i>	Invitrogen
CWY6473	<i>atg10Δ::KAN BY4742</i>	This study
Laboratory collection	<i>atg11Δ::KAN BY4742</i>	Invitrogen
CWY6475	<i>atg12Δ::KAN BY4742</i>	This study
Laboratory collection	<i>atg13Δ::KAN BY4742</i>	Invitrogen
CWY6497	<i>atg14Δ::LEU BY4742</i>	This study
Laboratory collection	<i>atg15Δ::LEU BY4742</i>	Invitrogen
Laboratory collection	<i>atg16Δ::KAN BY4742</i>	Invitrogen.
CWY6499	<i>atg17Δ::LEU BY4742</i>	This study
Laboratory collection	<i>atg18Δ::KAN BY4742</i>	Invitrogen
Laboratory collection	<i>atg19Δ::KAN BY4742</i>	Invitrogen
Laboratory collection	<i>atg20Δ::KAN BY4742</i>	Invitrogen
CWY6501	<i>atg21Δ::LEU BY4742</i>	This study
Laboratory collection	<i>atg22Δ::KAN BY4742</i>	Invitrogen
Laboratory collection	<i>atg23Δ::KAN BY4742</i>	Invitrogen
Laboratory collection	<i>atg24Δ::KAN BY4742</i>	Invitrogen
Laboratory collection	<i>atg26Δ::KAN BY4742</i>	Invitrogen
Laboratory collection	<i>atg27Δ::KAN BY4742</i>	Invitrogen
Laboratory collection	<i>atg29Δ::KAN BY4742</i>	Invitrogen
Laboratory collection	<i>atg31Δ::KAN BY4742</i>	Invitrogen
Laboratory collection	<i>atg32Δ::KAN BY4742</i>	Invitrogen
Laboratory collection	<i>atg33Δ::KAN BY4742</i>	Invitrogen
CWY6503	<i>atg34Δ::LEU BY4742</i>	This study
Laboratory collection	<i>atg36Δ::KAN BY4742</i>	Invitrogen
CWY7336	<i>FAA4-GFP::HIS BY4742</i>	This study
CWY7464	<i>atg1Δ::LEU FAA4-GFP::HIS BY4742</i>	This study
CWY7466	<i>atg6Δ::LEU FAA4-GFP::HIS BY4742</i>	This study
CWY6294	<i>atg1Δ::HYG ELO3-VENUS::LEU HIS::GPD-CFP-PHO8 ERG6-mCherry::KAN BY4742</i>	This study
CWY6312	<i>atg2Δ::HYG ELO3-VENUS::LEU HIS::GPD-CFP-PHO8 ERG6-mCherry::KAN BY4742</i>	This study
CWY6298	<i>atg7Δ::HYG ELO3-VENUS::LEU HIS::GPD-CFP-PHO8 ERG6-mCherry::KAN BY4742</i>	This study
CWY6758	<i>VPH1-GFP::LEU ERG6-mCherry::KAN BY4742</i>	This study
CWY7085	<i>GTR2-3xGFP::HIS ERG6-mCherry::LEU BY4742</i>	This study
CWY6760	<i>IVY1-GFP::LEU ERG6-mCherry::KAN BY4742</i>	This study
CWY7129	<i>VPH1-mCherry::LEU BY4742</i>	This study
CWY7183	<i>ATG6-3xGFP::HIS VPH1-mCherry::LEU BY4742</i>	This study
CWY7325	<i>ATG6-3xGFP::HIS GTR2-mCherry::LEU BY4742</i>	This study
CWY7226	<i>ATG14-3xGFP::HIS VPH1-mCherry::LEU BY4742</i>	This study
CWY7327	<i>ATG14-3xGFP::HIS IVY1-mCherry::LEU BY4742</i>	This study
CWY7169	<i>VPH1-mCherry::LEU fab1Δ::KAN BY4742</i>	This study
CWY7161	<i>VPH1-mCherry::LEU vps4Δ::KAN BY4742</i>	This study
CWY7159	<i>VPH1-mCherry::LEU nem1Δ::KAN BY4742</i>	This study
Laboratory collection	<i>fab1Δ::KAN BY4742</i>	Invitrogen
Laboratory collection	<i>vps4Δ::KAN BY4742</i>	Invitrogen
Laboratory collection	<i>nem1Δ::KAN BY4742</i>	Invitrogen
Laboratory collection	<i>pep4Δ::KAN BY4742</i>	Invitrogen
CWY7266	<i>VPH1-mCherry::LEU pep4Δ::KAN BY4742</i>	This study
CWY7204	<i>VPH1-mCherry::LEU atg1Δ::HYG BY4742</i>	This study
CWY7212	<i>VPH1-mCherry::LEU atg6Δ::HYG BY4742</i>	This study

Table S1. Yeast strains and plasmids used in this study (Continued)

Strain	Description	Source
CWY7216	VPH1-mCherry::LEU atg7Δ::HYG BY4742	This study
CWY7220	VPH1-mCherry::LEU atg8Δ::KAN BY4742	This study
CWY7234	VPH1-mCherry::LEU atg11Δ::KAN BY4742	This study
CWY7250	VPH1-mCherry::HIS atg14Δ::LEU BY4742	This study
CWY7238	VPH1-mCherry::LEU atg19Δ::KAN BY4742	This study
CWY7254	VPH1-mCherry::HIS atg21Δ::LEU BY4742	This study
CWY7242	VPH1-mCherry::LEU atg32Δ::KAN BY4742	This study
CWY7246	VPH1-mCherry::LEU atg33Δ::KAN BY4742	This study
CWY4836	are1Δ::KAN are2Δ::HIS BY4742	This study
CWY3626	dga1Δ::KAN Iro1Δ::HIS BY4742	Wang and Lee, 2012
CWY7167	VPH1-mCherry::LEU are1Δ::KAN are2Δ::HIS BY4742	This study
CWY7442	OM45-GFP::LEU BY4742	This study
CWY7444	atg1Δ::HYG OM45-GFP::LEU BY4742	This study
CWY6555	ATG1-GFP::LEU ERG6-mCherry::KAN BY4742	This study
CWY6558	ATG3-GFP::LEU ERG6-mCherry::KAN BY4742	This study
CWY6592	ATG9-GFP::LEU ERG6-mCherry::KAN BY4742	This study
CWY6572	ATG13-GFP::LEU ERG6-mCherry::KAN BY4742	This study
CWY6580	ATG21-GFP::LEU ERG6-mCherry::KAN BY4742	This study
CWY6582	ATG23-GFP::LEU ERG6-mCherry::KAN BY4742	This study
CWY6584	ATG24-GFP::LEU ERG6-mCherry::KAN BY4742	This study
CWY6586	ATG27-GFP::LEU ERG6-mCherry::KAN BY4742	This study
CWY6679	ATG29-GFP::HIS ERG6-mCherry::KAN BY4742	This study

Table S2. Plasmids used in this study

Plasmid	Description/reference
pRS416-P _{GPD} -GFP-Pho8Δ60-T _{CYC1}	GPD promoter–GFP-PHO8Δ60-CYC1 terminator cloned into pRS416
pRS416-P _{GPD} -GFP-SKL-T _{CYC1}	GPD promoter–GFP-SKL-CYC1 terminator cloned into pRS416
pBS-GTR2-3xGFP-His3	GTR2 ORF residues 778–1,025 and the residues 50–411 after stop codon were cloned into HindIII–BamHI and SacI sites, respectively, of pBS-3xGFP-His3 (Toshima et al., 2006)
pBS-ATG6-3xGFP-His3	ATG6 ORF residues 1,248–1,672 and the residues 57–572 after stop codon were cloned into HindIII–BamHI and SacI sites, respectively, of pBS-3xGFP-His3 (Toshima et al., 2006)
pBS-ATG14-3xGFP-His3	ATG14 ORF residues 763–1,033 and the residues 44–455 after stop codon were cloned into HindIII–BamHI and SacI sites, respectively, of pBS-3xGFP-His3 (Toshima et al., 2006)
pFA6α-His3-GPD-CFP	GPD promoter–N-terminal CFP integration plasmid with a His3 marker
pFA6α-VENUS-LEU2	C-terminal VENUS integration plasmid with a LEU2 marker
pFA6α-mCherry-KanMX6	C-terminal mCherry integration plasmid with a KanMX6 marker
pFA6α-mCherry-LEU2	C-terminal mCherry integration plasmid with a LEU2 marker
pFA6α-mCherry-His3	C-terminal mCherry integration plasmid with a His3 marker
pFA6α-GFP-LEU2	C-terminal GFP integration plasmid with a LEU2 marker
pFA6α-GFP-His3MX6	Longtine et al., 1998
pFA6α-His3MX6	Longtine et al., 1998
pFA6α-HYG	Deletion plasmid with a Hygromycin resistance gene marker
pFA6α-LEU2	Deletion plasmid with a LEU2 marker
pFA6α-KanMX6	Longtine et al., 1998
pRS416-P _{GPD} -GFP-Elo3-T _{CYC1}	GPD promoter–GFP-ELO3-CYC1 terminator cloned into pRS416
pRS416-P _{Cu} -GFP-Atg8 (Aut7)	Kim et al., 2002
pRS416-P _{ADH1} -ATG2-GFP-T _{CYC1}	ADH1 promoter–ATG2-GFP-CYC1 terminator cloned into pRS416
pRS416-P _{ADH1} -ATG5-GFP-T _{CYC1}	ADH1 promoter–ATG5-GFP-CYC1 terminator cloned into pRS416
pRS416-P _{Cu} -GFP-Atg11 (Cvt9)	Kim et al., 2001
pRS416-P _{ADH1} -GFP-ATG12-T _{CYC1}	ADH1 promoter–GFP-ATG12-CYC1 terminator cloned into pRS416
pRS416-P _{ADH1} -GFP-ATG15-T _{CYC1}	ADH1 promoter–GFP-ATG15-CYC1 terminator cloned into pRS416
pRS416-P _{ADH1} -ATG16-GFP-T _{CYC1}	ADH1 promoter–ATG16-GFP-CYC1 terminator cloned into pRS416
pRS426-P _{Cu} -GFP-ATG17	Cheong et al., 2005
pRS426-ATG18-GFP	ATG18-GFP cloned into pRS426
pRS416-P _{CYC1} -GFP-Atg32-T _{CYC1}	CYC1 promoter–GFP-ATG32-CYC1 terminator cloned into pRS416



Video 1. **Localization of LDs in wild-type cells during stat-phase.** Wild-type cells stained with FM4-64 (vacuole) at DS and with BODIPY (LDs) at the indicated time points. Images were analyzed by time-lapse microscopy using the Delta Vision system (Applied Precision). Frames were taken every 30 s for 30 min. Cells in D4 were also imaged for z sections. Frames from top to bottom were taken every 0.2 μm for 5 μm . The dotted circles outline the boundaries of yeast cells. N, nucleus; V, vacuole.

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