

Supplementary Information

Trs130 participates in autophagy through GTPases Ypt31/32 in

Saccharomyces cerevisiae

Zou *et al.*,

Table S1. Yeast strains and plasmids used in this study

A. Strains

Strain	Alias	Genotype	Source
NSY991	VSY459, TRS130	<i>MATa leu2-3,112 his3-Δ200 trp1-Δ901 lys2-801 suc2-Δ9 ura3-52 TRS130-HA::HIS3MX6</i>	(1)
NSY992	<i>trs130ts</i>	<i>MATa leu2-3,112 his3-Δ200 trp1-Δ901 lys2-801 suc2-Δ9 ura3-52 trs130-(33 aa truncation)-HA^{ts}::HIS3MX6</i>	(1)
YLY2360	GFP-Atg8 TRS130	NSY991, <i>GFP-ATG8::URA3</i>	This study
YLY2361	GFP-Atg8 <i>trs130ts</i>	NSY992, <i>GFP-ATG8::URA3</i>	This study
YLY2741	GFP-Atg8 RFP-Ape1 TRS130	NSY991, <i>GFP-Atg8::URA3,RFP-APE1::LEU2</i>	This study
YLY2742	GFP-Atg8 RFP-Ape1 <i>trs130ts</i>	NSY992, <i>GFP-ATG8::URA3,RFP-APE1::LEU2</i>	This study
YLY4081	TRS130 <i>trs85Δ</i>	YLY2741, <i>trs85Δ::Kan</i>	This study
YLY4083	<i>trs130ts</i> <i>trs85Δ</i>	YLY2742, <i>trs85Δ::Kan</i>	This study
YLY3661	TRS130 <i>atg1Δ</i>	YLY2741, <i>atg1Δ::Kan</i>	This study
YLY3662	<i>trs130ts</i> <i>atg1Δ</i>	YLY2742, <i>atg1Δ::Kan</i>	This study
NSY1176	Trs130-HA Trs120-myc	<i>MATa leu2-3,112 his3-Δ200 trp1-Δ901 lys2-801 suc2-Δ9 ura3-52 TRS130-HA::HIS3MX6 TRS120-myc::TRP1</i>	(2)
NSY1177	<i>trs65ts</i>	NSY1176, <i>trs65Δ::Kan</i>	(2)
YLY2483	GFP-Atg8 TRS65	NSY1176, <i>GFP-ATG8::URA3</i>	This study

YLY2484	GFP-Atg8 <i>trs65ts</i>	NSY1177, <i>GFP-ATG8::URA3</i>	This study
YLY2851	GFP-Atg8 RFP-Ape1 TRS65	NSY1176, <i>GFP-ATG8::URA3,RFP-APE1::LEU2</i>	This study
YLY2852	GFP-Atg8 RFP-Ape1 <i>trs65ts</i>	NSY1177, <i>GFP-ATG8::URA3,RFP-APE1::LEU2</i>	This study
YLY4504	GFP-Atg8 RFP-Ape1 <i>tca17ts</i>	YLY2851, <i>tca17Δ::Kan</i>	This study
YLY3362	TRS130 <i>atg11Δ</i>	YLY2741, <i>atg11Δ::Hyg</i>	This study
YLY3365	<i>trs130ts</i> <i>atg11Δ</i>	YLY2742, <i>atg11Δ::Hyg</i>	This study
YLY3331	TRS130 <i>atg17Δ</i>	YLY2741, <i>atg17Δ::Kan</i>	This study
YLY3334	<i>trs130ts</i> <i>atg17Δ</i>	YLY2742, <i>atg17Δ::Kan</i>	This study
YLY3126	TRS130 <i>atg5Δ</i>	YLY2741, <i>atg5Δ::Hyg</i>	This study
YLY3127	<i>trs130ts</i> <i>atg5Δ</i>	YLY2742, <i>atg5Δ::Hyg</i>	This study
YLY4205	TRS130 <i>Atg13Δ</i>	YLY2741, <i>atg13Δ::Hyg</i>	This study
YLY4208	<i>trs130ts</i> <i>atg13Δ</i>	YLY2742, <i>atg13Δ::Hyg</i>	This study
YLY4068	TRS130 <i>Atg9Δ</i>	YLY2741, <i>atg9Δ::Hyg</i>	This study
YLY4071	<i>trs130ts</i> <i>atg9Δ</i>	YLY2742, <i>atg9Δ::Hyg</i>	This study
YLY3212	TRS130 <i>atg14Δ</i>	YLY2741, <i>atg14Δ::Hyg</i>	This study
YLY3213	<i>trs130ts</i> <i>atg14Δ</i>	YLY2742, <i>atg14Δ::Hyg</i>	This study
YLY4462	TRS33	<i>MAT a ura3-52 leu2-3,112 trp1-Δ901 lys2-801 his3-Δ200, TRS130-HA::HIS3MX6, GFP-Atg8::URA3,RFP-APE1::LEU2</i>	This study
YLY4464	<i>trs33Δ</i>	YLY4462, <i>trs33Δ::Hyg</i>	This study
YLY4431	TRS120	SEY6210, <i>TRS130-myc::TRP1 + pRS425-YPT31, GFP-Atg8::URA3,RFP-APE1::LEU2</i>	This study

YLY4021	<i>trs120Δ</i>	SEY6210, <i>trs120::HIS3MX6</i> <i>TRS130-myc::TRP1</i> + pRS425-YPT31, <i>GFP-Atg8::URA3,RFP-APE1::LEU2</i>	This study
YLY2705	Atg9-GFP RFP-Ape1 TRS130	NSY991, <i>ATG9-3XGFP::URA3,RFP-APE1::LEU2</i>	This study
YLY2704	Atg9-GFP RFP-Ape1 <i>trs130ts</i>	NSY992, <i>ATG9-3XGFP::URA3,RFP-APE1::LEU2</i>	This study
YLY2978	Atg9-GFP RFP-Ape1 TRS130 <i>atg1Δ</i>	YLY2705, <i>atg1Δ::Kan</i>	This study
YLY2979	Atg9-GFP RFP-Ape1 <i>trs130ts</i> <i>atg1Δ</i>	YLY2704, <i>atg1Δ::Kan</i>	This study
YLY2710	Atg9-GFP RFP-Ape1 TRS65	NSY1176, <i>ATG9-3XGFP::URA3,RFP-APE1::LEU2</i>	This study
YLY2870	Atg9-GFP RFP-Ape1 <i>trs65ts</i>	YLY2710, <i>trs65Δ::Hyg</i>	This study
YLY2746	Atg9-GFP RFP-Ape1 TRS65 <i>atg1Δ</i>	YLY2710, <i>atg1Δ::Kan</i>	This study
YLY2872	Atg9-GFP RFP-Ape1 <i>trs65ts</i> <i>atg1Δ</i>	YLY2746, <i>trs65Δ::Hyg</i>	This study
YLY2540	TRS130 Sec7-DsRed	NSY991, <i>SEC7-DsRed</i>	This study
YLY2541	<i>trs130ts</i> Sec7-DsRed	NSY992, <i>SEC7-DsRed</i>	This study
YLY2817	Sec7-DsRed Atg9-3XGFP T RS130	YLY2540, <i>ATG9-3XGFP::URA3</i>	This study
YLY2818	Sec7-DsRed Atg9-3XGFP <i>trs130ts</i>	YLY2541, <i>ATG9-3XGFP::URA3</i>	This study

YLY3091	Sec7-DsRed Atg9-3XGFP TRS130 <i>atg1Δ</i>	YLY2817, <i>atg1Δ::Kan</i>	This study
YLY3092	Sec7-DsRed Atg9-3XGFP <i>trs130ts</i> <i>atg1Δ</i>	YLY2818, <i>atg1Δ::Kan</i>	This study
YLY2914	Sec7-DsRed GFP-Atg8 TRS130	YLY2540, <i>GFP-ATG8::URA3</i>	This study
YLY2917	Sec7-DsRed GFP-Atg8 <i>trs130ts</i>	YLY2541, <i>GFP-ATG8::URA3</i>	This study
YLY3337	Sec7-DsRed GFP-Atg8 TRS130 <i>atg1Δ</i>	YLY2914, <i>atg1Δ::Kan</i>	This study
YLY3339	Sec7-DsRed GFP-Atg8 <i>trs130ts</i> <i>atg1Δ</i>	YLY2917, <i>atg1Δ::Kan</i>	This study
YLY3359	TN124	<i>MATα leu2-3,112 trp1 ura3-52 pho8::pho8Δ60 pho13::LEU2</i>	(3)
YLY3799	TRS130 (TN124)	TN124, <i>HIS3Δ::Hyg,</i> <i>TRS130-HA::HIS3MX6</i>	This study
YLY3800	<i>trs130ts</i> (TN124)	TN124, <i>HIS3Δ::Hyg, trs130-(33 aa truncation)-HA^{ts}::HIS3MX6</i>	This study
YLY3956	<i>atg1Δ</i> (TN124)	YLY3799, <i>atg1Δ::Kan</i>	This study
YLY4060	TRS65 (TN124)	TN124, <i>HIS3Δ::Hyg,</i> <i>TRS130-HA::HIS3MX6 TRS120-myc::TRP1</i>	This study
YLY4140	<i>trs65ts</i> (TN124)	YLY4060, <i>trs65Δ::Kan</i>	This study
NSY1081	YPT1	<i>MAT a ura3-52 Leu2 his3</i>	(2)
NSY1082	<i>ypt1ts</i>	<i>MAT a ura3-52 Leu2 his3 ypt1A136D</i>	(2)
YLY2366	GFP-Atg8 YPT1	NSY1081, <i>GFP-ATG8::URA3</i>	This study
YLY2367	GFP-Atg8 <i>ypt1ts</i>	NSY1082, <i>GFP-ATG8::URA3</i>	This study

YLY2417	GFP-Snc1 Sec7-DsRed	<i>MAT a leu2-3,112 lys2-801</i> <i>TRS130-HA::HIS3MX6, GFP-Snc1::URA3,</i> <i>SEC7-DsRed</i>	This study
YLY2418	GFP-Snc1 Sec7-DsRed	<i>MAT a leu2-3,112 lys2-801 trs130-(33 aa</i> <i>truncation)-HA^{ts}::HIS3MX6,</i> <i>GFP-Snc1::URA3, SEC7-DsRed</i>	This study
RSY367	SEC7	<i>MATa ade2-1 ura3-1 his3-11,15</i> <i>leu2-3,112 trp1-Δ1 can1-100</i>	R. Scheckman lab
RSY301	<i>sec7-4</i>	<i>MATa ura3-1 his3-11,15 Leu2-3,112</i> <i>trp1-Δ1</i>	R. Scheckman lab
YLY4478	SEC7 GFP-Atg8	RSY367, <i>TRS130-HA::HIS3MX6,</i> <i>GFP-ATG8::URA3</i>	This study
YLY4479	<i>trs130ts</i> GFP-Atg8	RSY367, <i>trs130-(33 aa truncation)-</i> <i>HA^{ts}::HIS3MX6, GFP-ATG8::URA3</i>	This study
YLY4480	<i>sec7-4</i> GFP-Atg8	RSY301, <i>TRS130-HA::HIS3MX6,</i> <i>GFP-ATG8::URA3</i>	This study
YLY4481	<i>sec7-4</i> <i>trs130ts</i> GFP-Atg8	RSY301, <i>trs130-(33 aa truncation)-</i> <i>HA^{ts}::HIS3MX6, GFP-ATG8::URA3</i>	This study
YLY40	NSY128	<i>MATa ade2 his3-200 leu2-3,112 lys2-801</i> <i>ura3-52</i>	(4)
YLY44	NSY340 <i>ypt31Δ</i> <i>ypt32ts</i>	<i>MATa leu2-3,112 lys2-801 ura3-52</i> <i>ypt31Δ::HIS3 ypt32A141D</i>	(4)

B. Plasmids

Plasmid	Alias	Genotype	Source
p1K-GFP- <i>ATG8-406</i>	pYL233		(5)
<i>RFP-APE1-305</i>	pYL238		(6)
<i>ATG9-3XG FP-PG5</i>	pYL242		(7)
Cu-Cherry- <i>ATG8-415</i>	pYL254		(8)
pNS180	pRS425	2μ, <i>LEU3</i> , Amp ^r	(9)
pNS781	Ypt31	pRS425- <i>YPT31</i>	(1)
pYL316	Ypt31S	pRS425- <i>YPT31</i> S27N	This study
pYL313	Ypt31Q	pRS425- <i>YPT31</i> Q72L	This study
pYL317	Ypt1	pRS425- <i>YPT1</i>	This study
pYL120	pRS426	2μ, <i>URA3</i> , Amp ^r	(9)

pYL335		pRS426-Ypt31	This study
pYL338		pRS426- <i>YPT31</i> S27N	This study
pYL336		pRS426- <i>YPT31</i> Q72L	This study
pYL339		pRS426- <i>YPT1</i>	This study
pYL151	YEp351	2 μ , <i>LEU2</i>	(10)
pYL152		YEp351- <i>TRS130</i>	This study
pYL283		YEp351- <i>YPT32</i>	This study
pYL225		YEp351- <i>TRS65</i>	This study
pYL342		YEp351- <i>TCA17</i>	This study

Supplementary Materials and Methods

Growth assay of genetic interaction

For genetic interaction assays, cultures were grown overnight at 26°C in minimal medium (SD-Leu), normalized to the same density and spotted onto SD-Leu plates in serial dilutions of 1:10. Plates were incubated at various temperatures for different numbers of days and photographed.

References

1. Sciorra VA, Audhya A, Parsons AB, Segev N, Boone C, Emr SD. Synthetic genetic array analysis of the PtdIns 4-kinase Pik1p identifies components in a Golgi-specific Ypt31/rab-GTPase signaling pathway. *Mol Biol Cell* 2005;16(2):776-793.
2. Liang Y, Morozova N, Tokarev AA, Mulholland JW, Segev N. The role of Trs65 in the Ypt/Rab guanine nucleotide exchange factor function of the TRAPP II complex. *Mol Biol Cell* 2007;18(7):2533-2541.
3. Noda T, Matsuura A, Wada Y, Ohsumi Y. Novel system for monitoring autophagy in the yeast *Saccharomyces cerevisiae*. *Biochem Biophys Res Commun* 1995;210(1):126-132.
4. Jones S, Jedd G, Kahn RA, Franzusoff A, Bartolini F, Segev N. Genetic interactions in yeast between Ypt GTPases and Arf guanine nucleotide exchangers. *Genetics* 1999;152(4):1543-1556.
5. Xie Z, Nair U, Klionsky DJ. Atg8 controls phagophore expansion during autophagosome formation. *Mol Biol Cell* 2008;19(8):3290-3298.
6. Shintani T, Reggiori F. Fluorescence microscopy-based assays for monitoring yeast Atg protein trafficking. *Methods Enzymol* 2008;451:43-56.
7. Monastyrska I, He C, Geng J, Hoppe AD, Li Z, Klionsky DJ. Arp2 links autophagic machinery with the actin cytoskeleton. *Mol Biol Cell* 2008;19(5):1962-1975.
8. Mari M, Griffith J, Rieter E, Krishnappa L, Klionsky DJ, Reggiori F. An Atg9-containing compartment that functions in the early steps of autophagosome biogenesis. *J Cell Biol* 2010;190(6):1005-1022.
9. Christianson TW, Sikorski RS, Dante M, Shero JH, Hieter P. Multifunctional yeast high-copy-number shuttle vectors. *Gene* 1992;110(1):119-122.
10. Yamamoto K, Jigami Y. Mutation of TRS130, which encodes a component of the TRAPP II complex, activates transcription of OCH1 in *Saccharomyces cerevisiae*. *Curr Genet* 2002;42(2):85-93.

Figure S1

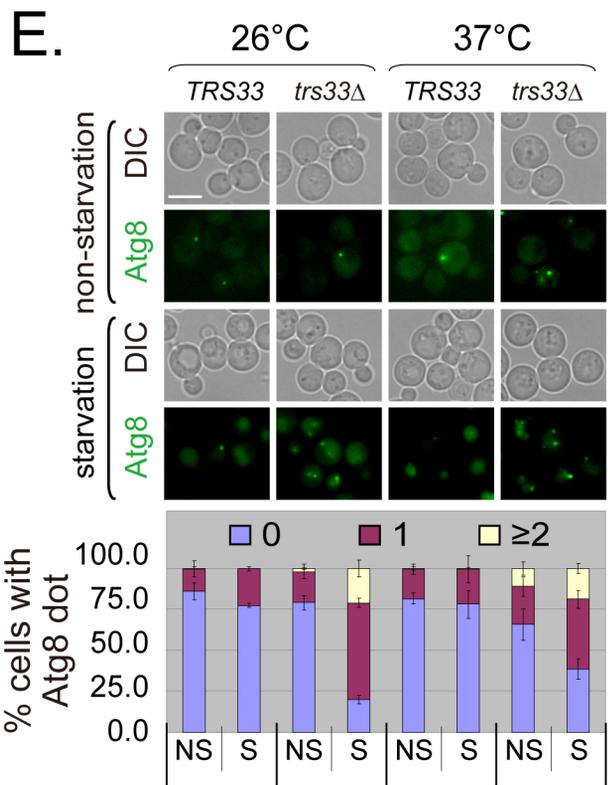
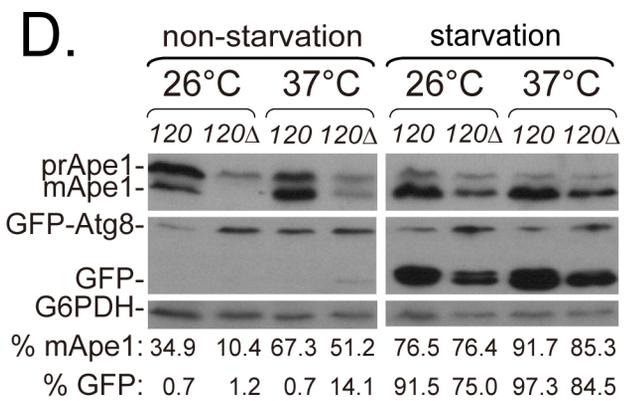
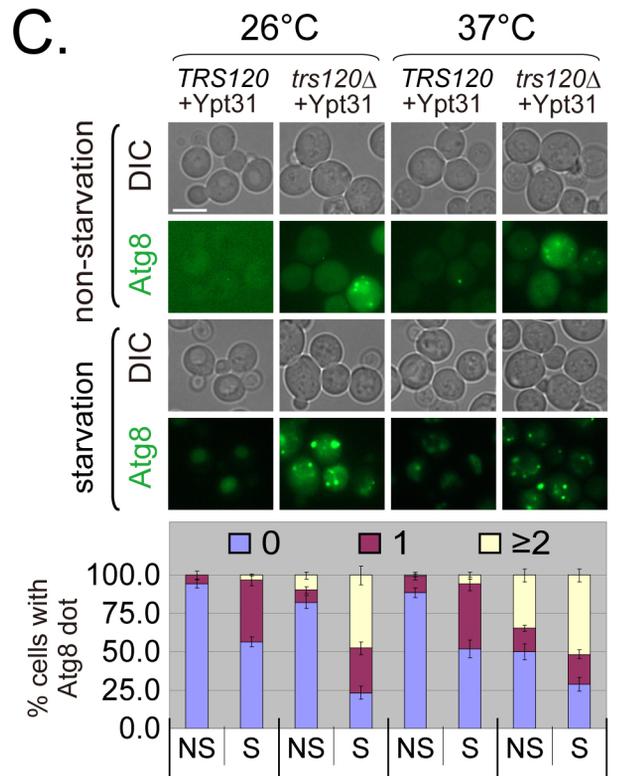
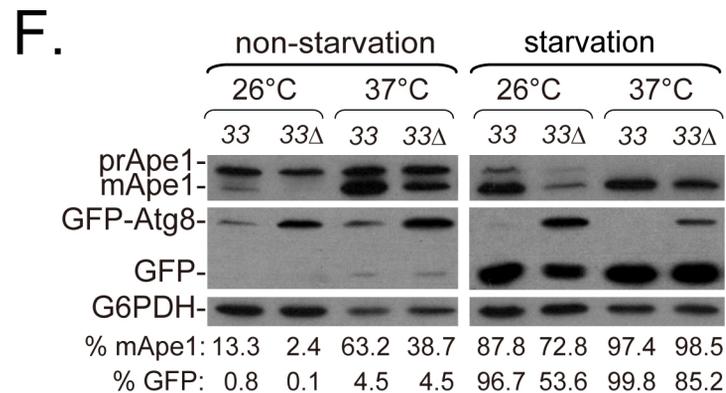
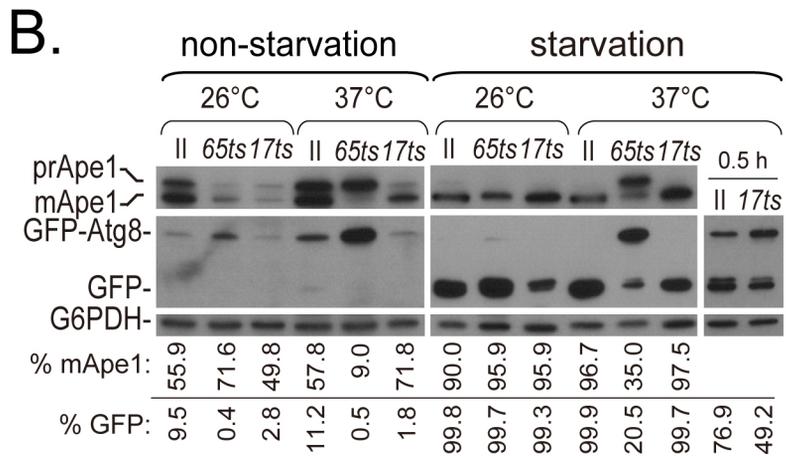
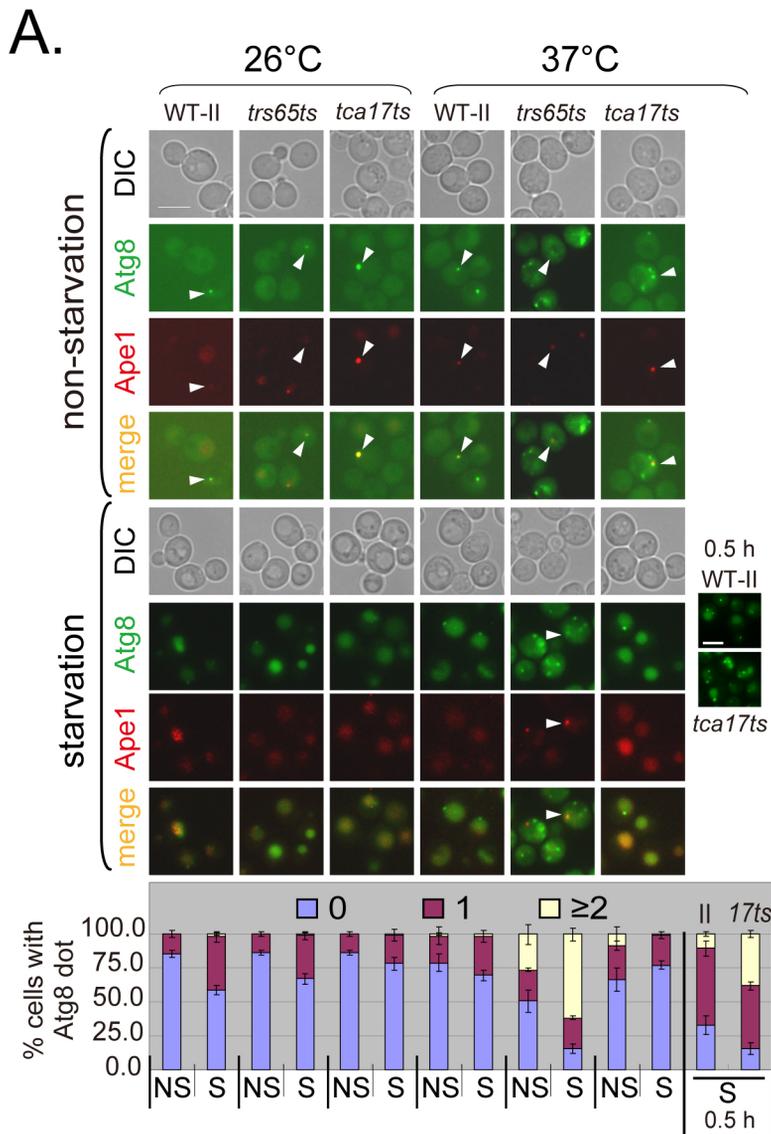
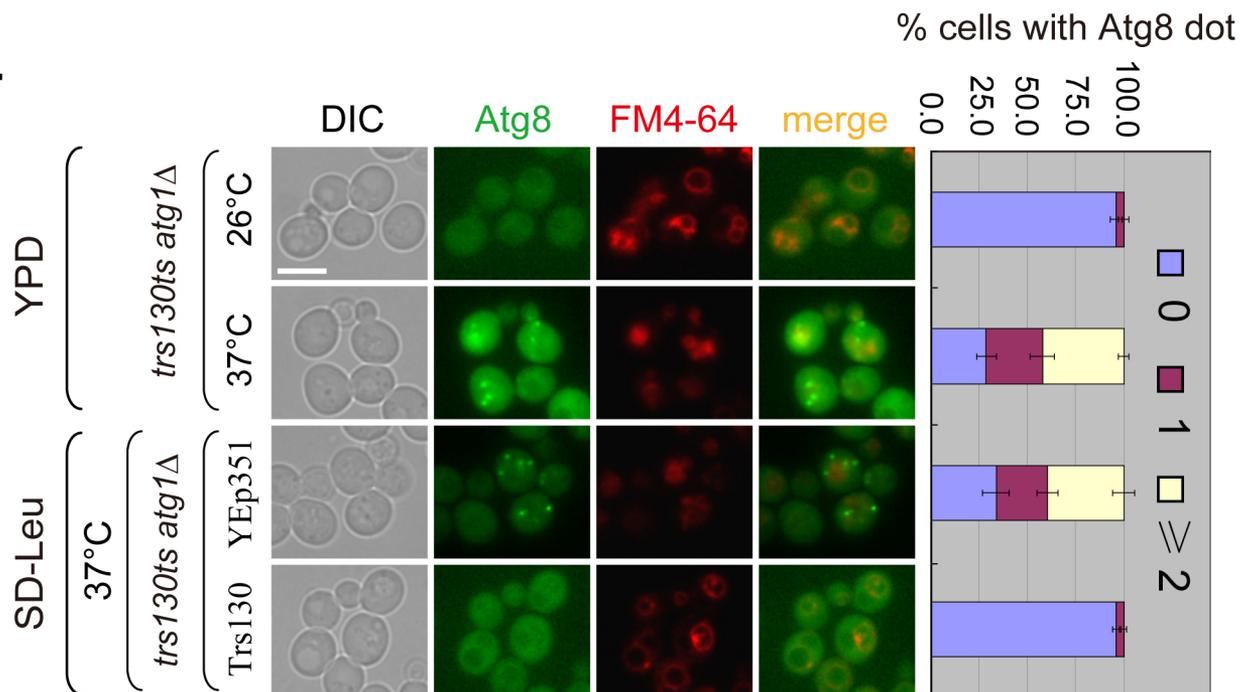


Figure S1. Cvt pathway and starvation-induced autophagy are impaired in other TRAPP II-specific subunit mutants at the restrictive temperature. (A) Atg8 and Ape1 localization in *trs65ts* and *tca17ts* mutant cells. WT-II (for indicating TRAPP II-specific subunits Trs130 and Trs120 are tagged as Trs130-HA Trs120-myc) and mutant cells were tagged using GFP-Atg8 and RFP-Ape1 integration plasmids. Cells were grown and treated as in [Figure 1A](#). GFP-Atg8 morphology of WT-II and *tca17ts* cells starved for 0.5 hour at 37°C is at right. Arrowheads indicate colocalization of Atg8 and Ape1. Bar, 5 μm. The percentage of cells with Atg8 dots was quantitated and presented as in [Figure 1A](#). **(B)** Ape1 maturation was blocked under non-starvation conditions and GFP-Atg8 degradation was reduced under starvation conditions in *trs65ts* mutant cells at 37°C. Cells grown as in (A) but in YPD were subjected to immunoblot assay, quantitated and presented as in [Figure 1B](#). II for WT-II as in (A). **(C)-(F)**: GFP-Atg8 localization in the *trs120Δ* mutant (*trs120Δ*+Ypt31) **(C)** and the *trs33Δ* mutant (*trs33Δ* in Trs130-HA background) (Trs33 was a candidate for TRAPP II-specific subunit in this study) **(E)**; immunoblot assay for autophagy defects in *trs120Δ* mutant **(D)** and *trs33Δ* mutant **(F)**. Experiments were done as in (A) and (B).

Figure S2

A.



B. SD-Ura, 37°C

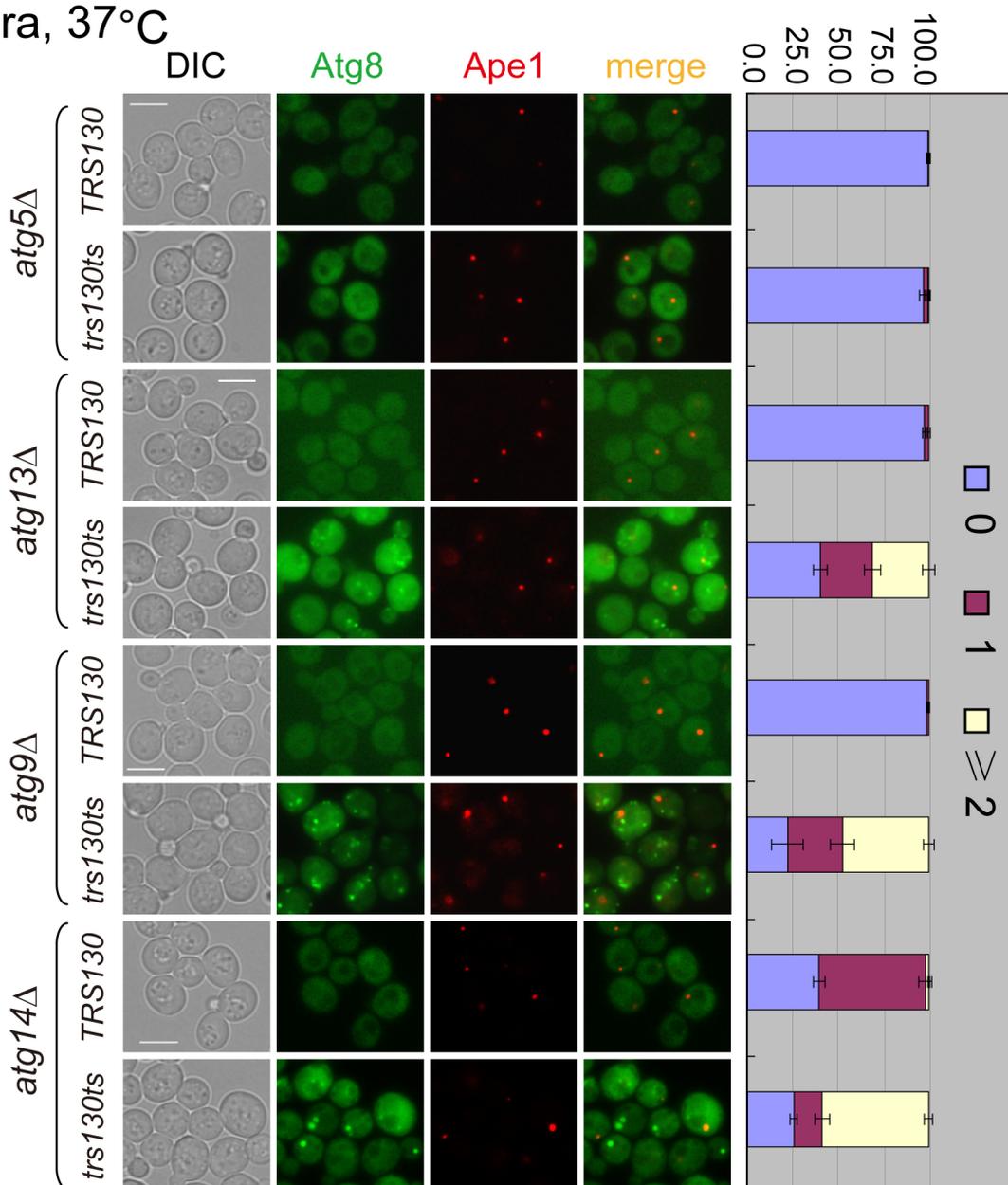
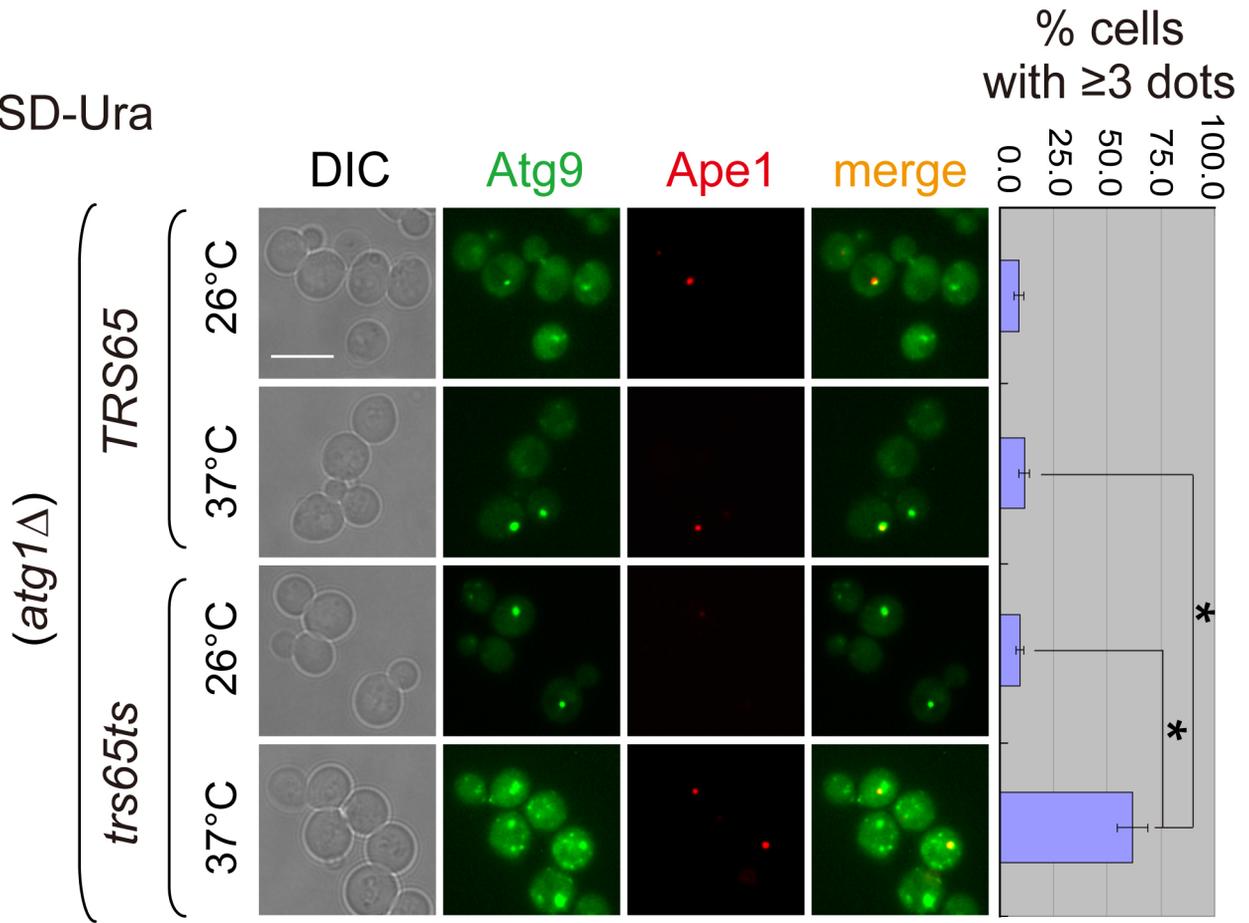


Figure S2. Localization of Atg8 in *trs130ts* mutants with deletion of *ATG* genes without starvation. (A) Multiple GFP-Atg8 dots were formed in *trs130ts atg1Δ* without starvation at the restrictive temperature; this was rescued by Trs130. Cells grown in rich medium (YPD or SD-Leu) to mid-log phase at 26°C were treated as in [Figure 1A](#) for non-starvation and stained with FM4-64 during the last hour to detect the vacuole. Bar, 5 μm. (B) Yeast strains were grown and treated as described in [Figure 3B](#) except that incubation was in SD-Ura for 1.5 hours at 37°C without starvation. Cells were examined by live microscopy imaging for changes in GFP-Atg8 and RFP-Ape1. Bar, 5 μm. The percentage of cells with Atg8 dots was quantitated and presented as in [Figure 1A](#).

Figure S3

A. SD-Ura



B. SD-N

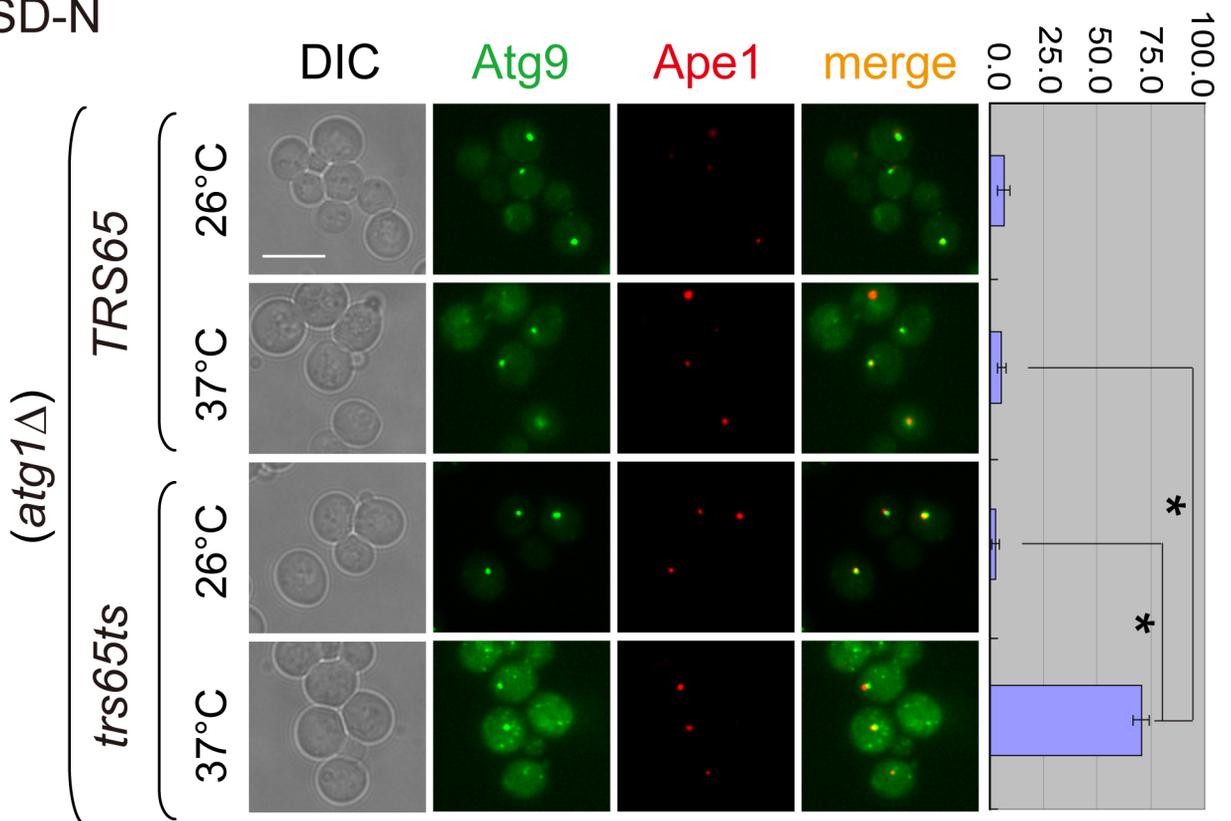


Figure S3. Anterograde transport of Atg9 to the PAS is altered in *trs65ts* mutant cells at the restrictive temperature. WT and *trs65ts* mutant cells tagged with Atg9-3XGFP and RFP-Ape1 in an *atg1Δ* background were incubated as described in [Figure 4](#) and examined for fluorescence. **(A)** Non-starvation. **(B)** Starvation. Bar, 7 μ m. Data are presented as in [Figure 4](#). Asterisks indicate $P < 0.001$ as highly significant.

Figure S4

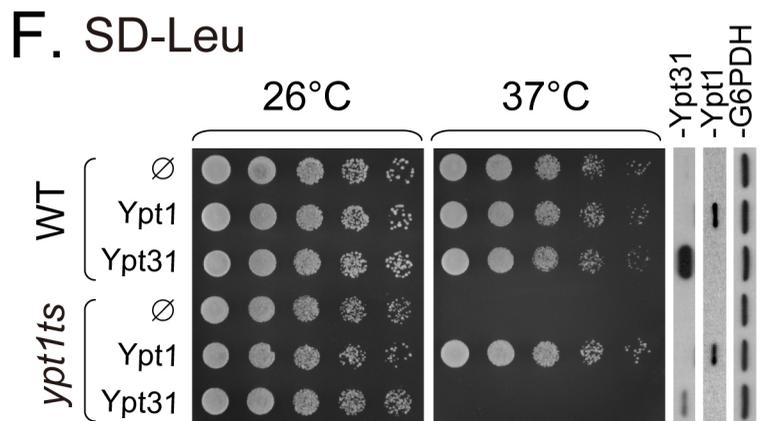
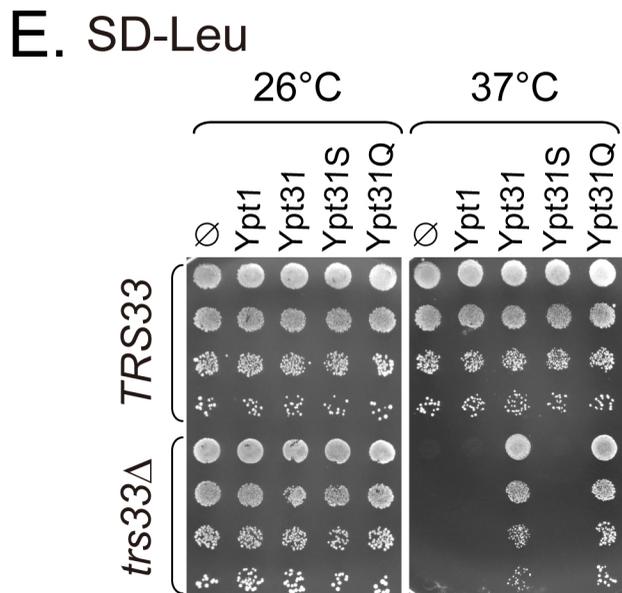
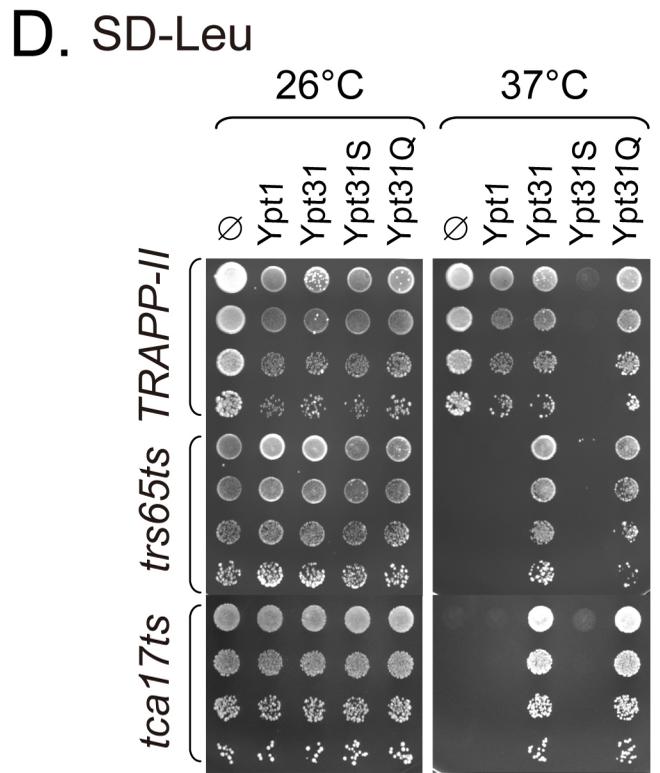
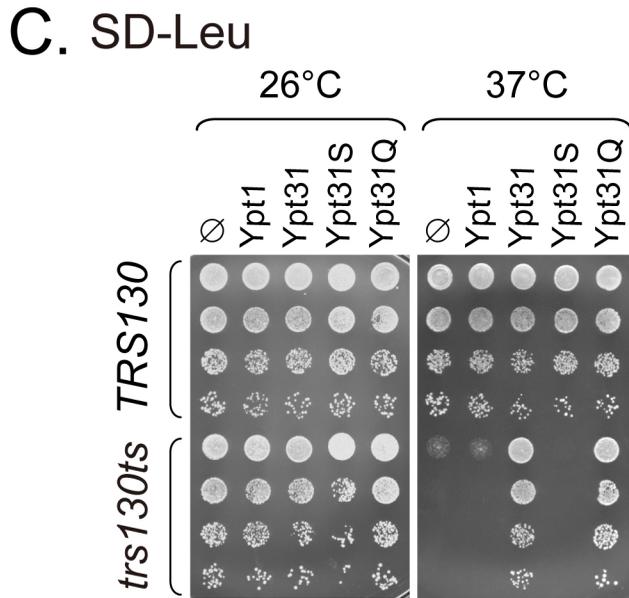
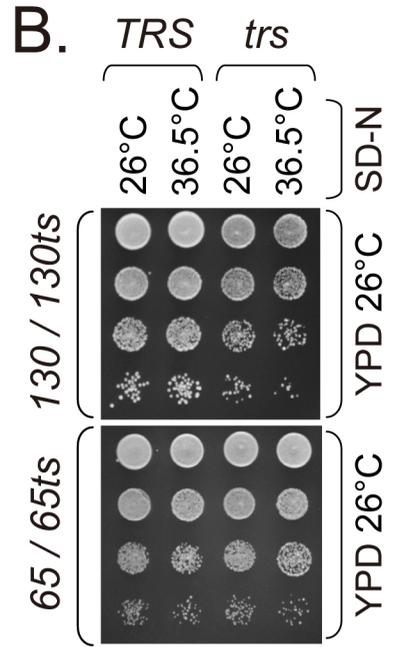
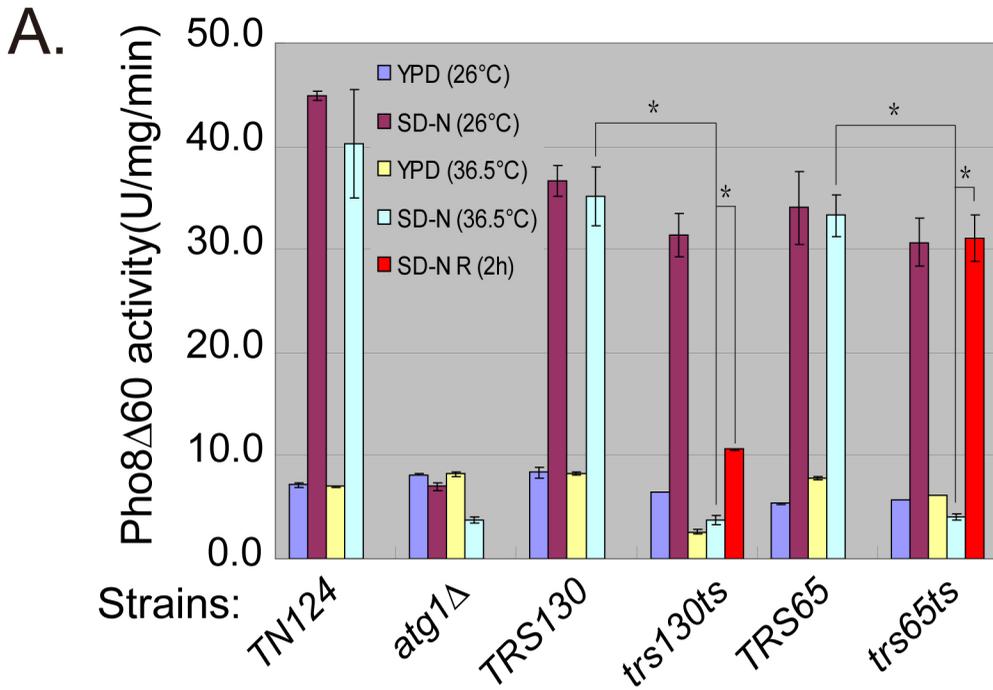
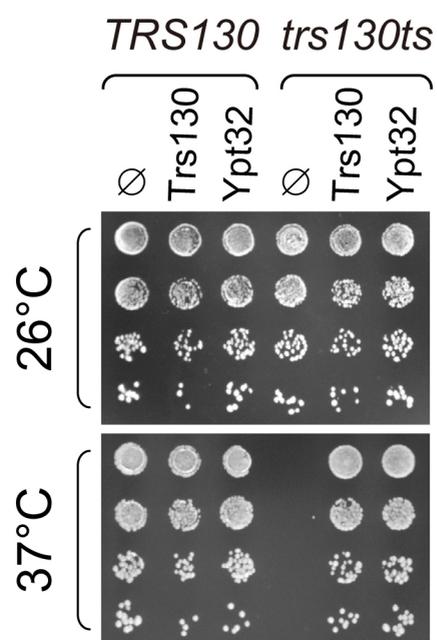


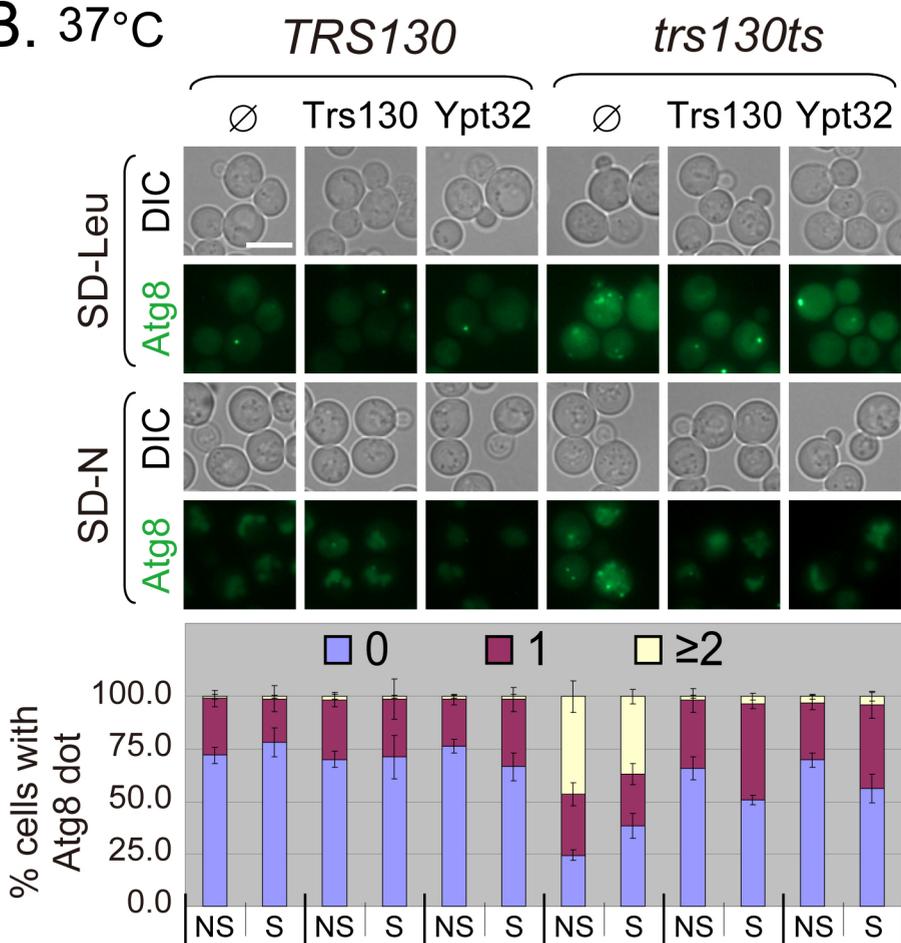
Figure S4. Changes in autophagic flux in some TRAPP II-specific subunit mutants and suppression of growth defects in mutants by Ypt31. (A) Pho8 Δ 60 activity in *trs130ts* and *trs65ts* mutants. WT (TN124) and the *atg1 Δ* (in *TRS130* background) strains were used as positive and negative controls. All cells were treated as in [Figures 1B](#) except that 36.5°C was used to guarantee that the *trs130ts* cells would be viable after treatment since the mutants in the TN124 background were more temperature sensitive than the strains in [Figures 1A and S1A](#). The *trs130ts* and *trs65ts* mutants were transferred from 36.5°C in SD-N to 26°C in SD-N for 2 hours recovery and designated SD-N R (2h). Pho8 Δ 60 assay data are presented as in [Figure 7B](#) except with absolute values because of slightly lower Pho8 Δ 60 activity in *TRS130* than in TN124. Asterisks indicate $P < 0.001$ as highly significant. **(B)** The *trs130ts* and *trs65ts* mutants were viable after treatment at 36.5°C in SD-N for 2 hours. WT and the mutants incubated in SD-N at 26°C and 36.5°C for 2 hours were plated on YPD plates with tenfold serial dilution from top to bottom and grown at 26°C for 2 days. **(C)-(E)** Cells as in [Figures 7 and S1](#) but without RFP-Ape1 tagging were transformed with the indicated plasmids to examine growth. Cells were spotted onto SD-Leu medium with tenfold serial dilution from top to bottom and grown at various temperatures. Ypt31 (WT and GTP-bound form) but not Ypt1 suppressed the growth defect of *trs130ts* mutant cells **(C)**, *trs65ts* and *tca17ts* mutant cells **(D)**, and *trs33 Δ* mutant cells **(E)** at 37°C. **(F)** The Ypt1 plasmid in (C)-(E) was used to overexpress Ypt1 in *ypt1ts* to confirm expression and function. Ypt31 served as a negative control. Growth was checked as in (C) and protein expression was examined as in [Figure 7C](#).

Figure S5

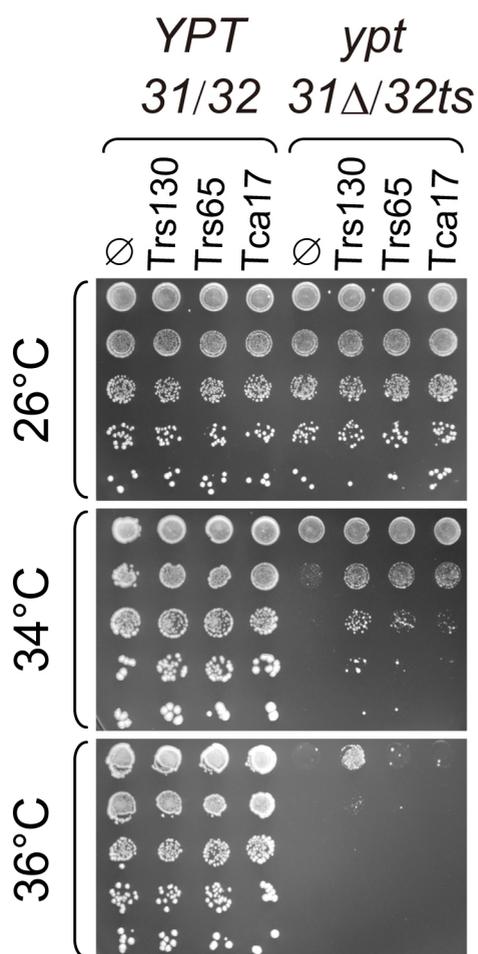
A. SD-Leu



B. 37°C



C. SD-Leu



D.

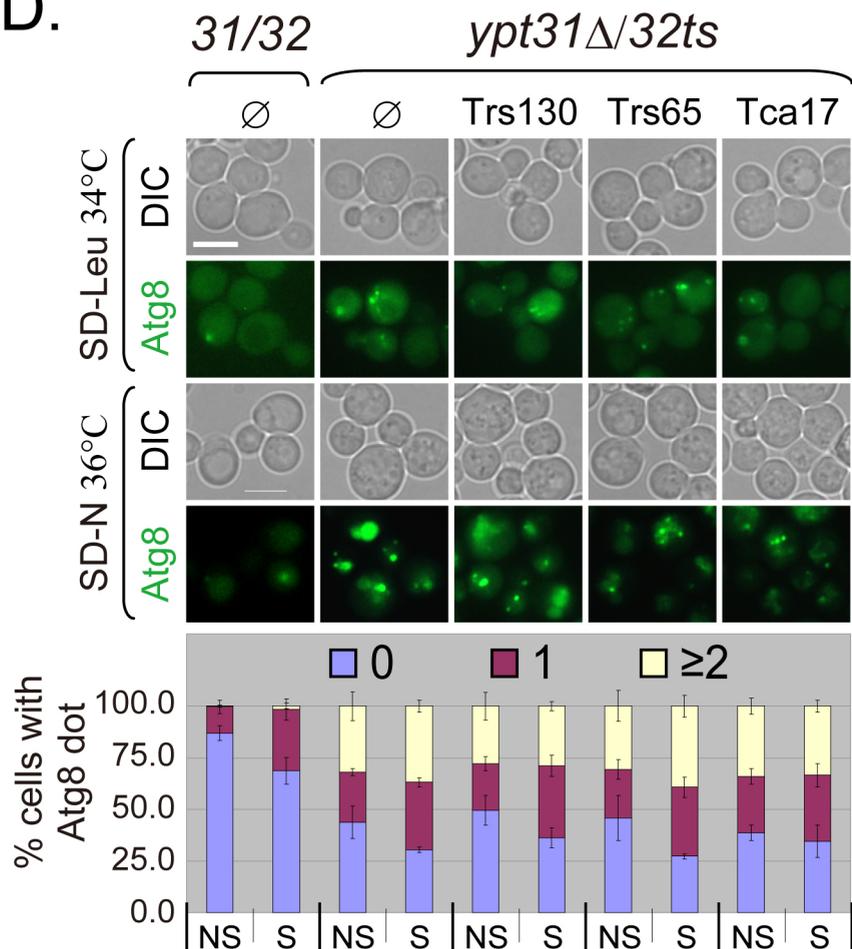


Figure S5. Ypt32 fully suppresses growth and autophagy defects in *trs130ts* mutant cells, but TRAPP II subunits suppress *ypt31Δ/32ts* growth defects weakly. (A) Trs130 and Ypt32 suppress the growth defect in *trs130ts* mutant cells at the restrictive temperature. Trs130 and Ypt32 in YEp351 were transformed into WT and *trs130ts* mutant cells tagged with GFP-Atg8 and grown as in [Figure S4C](#). **(B)** Trs130 and Ypt32 suppressed the GFP-Atg8 transport defect in *trs130ts* cells. Yeast transformants as in (A) were investigated to determine whether Trs130 and Ypt32 suppressed the GFP-Atg8 transport defect in *trs130ts* cells in synthetic minimal medium (SD-Leu) or starvation medium (SD-N) at the restrictive temperature. Bar, 5 μm. The percentage of cells with Atg8 dots was quantitated and presented as in [Figure 1A](#). **(C)** TRAPP II-specific subunits (Trs130, Trs65 and Tca17) weakly suppressed the growth defect in *ypt31Δ/32ts* mutant cells at the non-permissive temperature (34°C). Trs130, Trs65 and Tca17 in YEp351 (all are functional) were transformed and grown as in [Figure S4C](#). **(D)** TRAPP II-specific subunits (Trs130, Trs65 and Tca17) did not obviously suppress the GFP-Atg8 transport defect in *ypt31Δ/32ts* cells. Yeast transformants as in (C) were investigated to determine whether TRAPP II-specific subunits suppressed the GFP-Atg8 transport defect in *ypt31Δ/32ts* cells in synthetic minimal medium (SD-Leu) at a growth suppression temperature (34°C) or in starvation medium (SD-N) at a temperature at which GFP-Atg8 transport starts to show defects (36°C). Bar, 5 μm. The percentage of cells with Atg8 dots was quantitated and presented as in [Figure 1A](#).