

SUPPLEMENTAL MATERIALS

Requirement for Golgi-localized PI(4)P in fusion of COPII vesicles with Golgi compartments

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SUPPLEMENTAL MATERIALS AND METHODS

Determination of Sac1^{WT} phosphatase activity on semi-intact cell membranes under *in vitro* pre-treatment conditions

Washed semi-intact cell membranes were treated with 50 μ M Sac1^{WT} or 50 μ M Sac1^{C392S} as in Figure 4. Membrane lipids were extracted as described (Gray *et al.*, 2003) with minor modifications: 1.5 A₂₈₀ units (~1.03 OD₆₀₀ units) of treated membranes were independently precipitated in quadruplicate with 5% TCA, 1 mM EDTA in 1.5 ml on ice for 15 min followed by centrifugation. The pellet was washed twice with 1.5 ml ice-cold 5% TCA, 1 mM EDTA, and acidic lipids were extracted and dried. Lipids were reconstituted with 10 μ l 2:1:0.8 CH₃OH:CHCl₃:H₂O by vortexing and sonication for 5 min at 30°C on a B2200R-1 Branson ultrasonic bath (Branson Ultrasonic Corporation, Danbury, CT). Lipids (4 μ l) were spotted on a PI(4)P Mass Strip KitTM membrane as indicated by the manufacturer (Echelon Biosciences, Salt Lake City, UT), and developed by chemiluminescence and densitometric analysis as described for immunoblots. A Mann-Whitney-Wilcoxon test to compare the medians was performed on the observed results.

NEM inactivation of Sac1^{WT} phosphatase activity

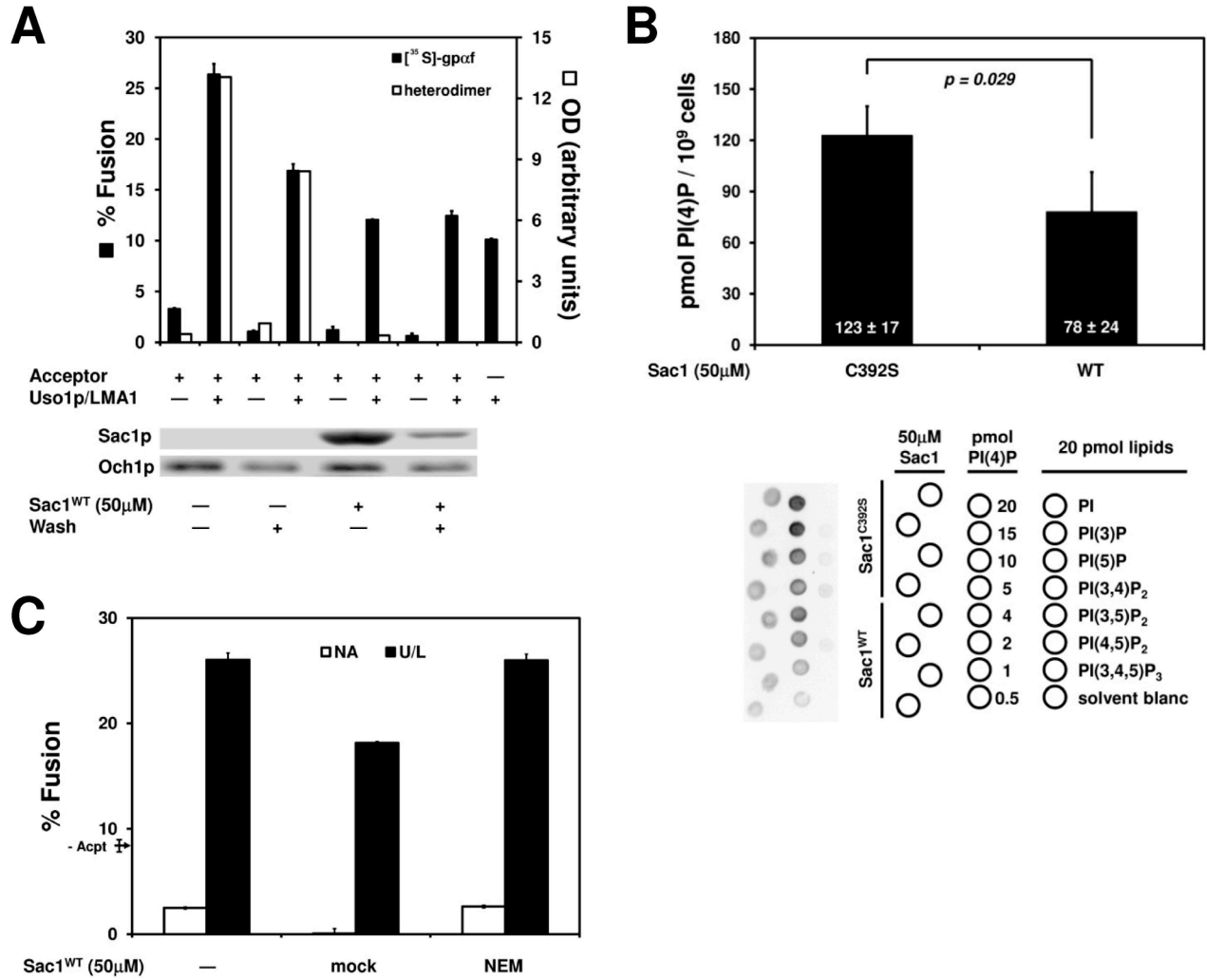
To inactivate Sac1^{WT}, protein (600 μ g) was incubated at 8°C for 10 min with 50 mM NEM (Maehama *et al.*, 2000). Excess NEM was quenched with 100 mM DTT at 8°C for 10 min, and then dialyzed overnight against buffer 88 without sorbitol containing 0.5 mM DTT. A mock inactivation was performed in parallel using ethanol carrier in place of NEM.

SUPPLEMENTAL TABLE S1. CANDIDATE STRAINS TESTED FOR FUSION DEFECTS

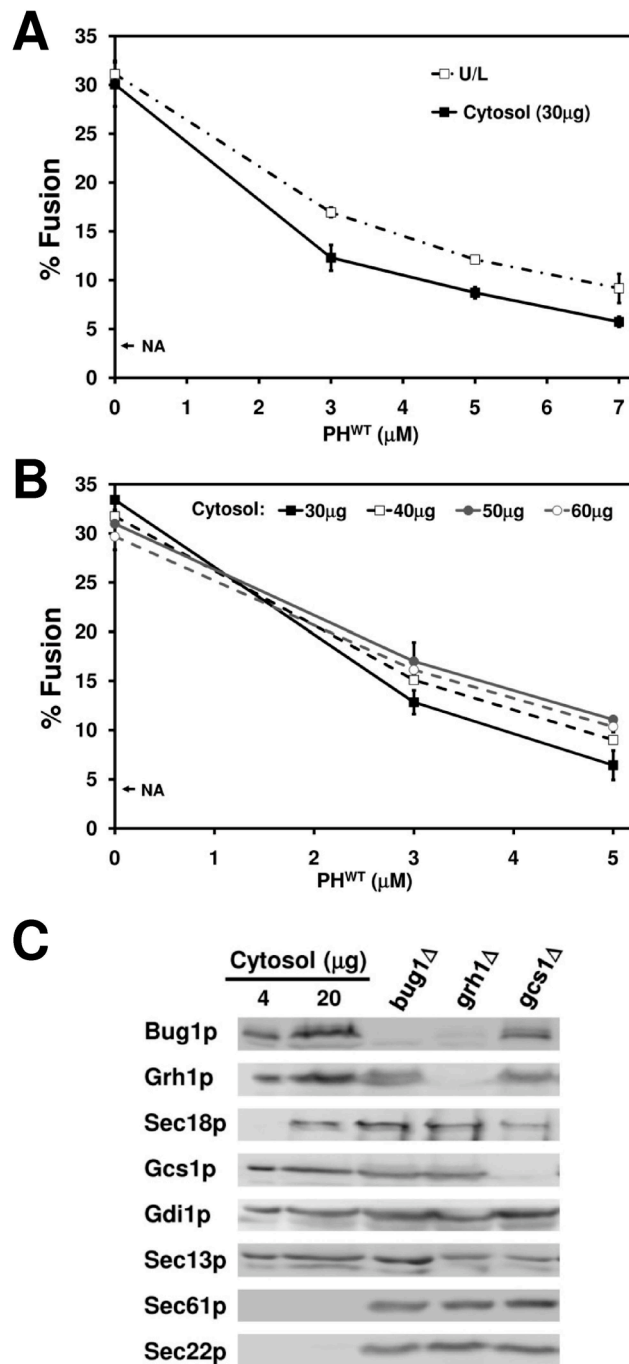
Strain	Genotype	Study reporting PI(4)P binding ¹
<i>atg26Δ</i>	BY4742; <i>atg26Δ::KAN^R</i>	(Yu <i>et al.</i> , 2004)
<i>bem3Δ</i>	BY4742; <i>bem3Δ::KAN^R</i>	(Zhu <i>et al.</i> , 2001)
<i>caf120Δ</i>	BY4742; <i>caf120Δ::KAN^R</i>	(Yu <i>et al.</i> , 2004)
<i>cla4Δ</i>	BY4742; <i>cla4Δ::KAN^R</i>	(Yu <i>et al.</i> , 2004)
<i>cog5Δ</i>	BY4742; <i>cog5Δ::KAN^R</i>	(Zhu <i>et al.</i> , 2001)
<i>gcs1Δ</i>	BY4742; <i>gcs1Δ::KAN^R</i>	(Zhu <i>et al.</i> , 2001)
<i>osh1Δ</i>	BY4742; <i>osh1Δ::KAN^R</i>	(Yu <i>et al.</i> , 2004)
<i>osh2Δ</i>	BY4742; <i>osh2Δ::KAN^R</i>	(Yu <i>et al.</i> , 2004)
<i>osh3Δ</i>	BY4742; <i>osh3Δ::KAN^R</i>	(Yu <i>et al.</i> , 2004)
<i>osh1Δ2Δ3Δ</i>	SEY6210; <i>osh1Δ::URA3 osh2Δ::URA3 osh3Δ::LYS2</i>	
<i>osh4Δ</i>	BY4742; <i>osh4Δ::KAN^R</i>	(Zhu <i>et al.</i> , 2001)
<i>osh6Δ</i>	BY4742; <i>osh6Δ::KAN^R</i>	(Zhu <i>et al.</i> , 2001)
<i>pdr16Δ</i>	BY4742; <i>pdr16Δ::KAN^R</i>	(Zhu <i>et al.</i> , 2001)
<i>pdr17Δ</i>	BY4742; <i>pdr17Δ::KAN^R</i>	(Zhu <i>et al.</i> , 2001)
<i>rgc1Δ</i>	BY4742; <i>rgc1Δ::KAN^R</i>	(Yu <i>et al.</i> , 2004)
<i>scs2Δ</i>	BY4742; <i>scs2Δ::KAN^R</i>	(Zhu <i>et al.</i> , 2001)
<i>skg3Δ</i>	BY4742; <i>skg3Δ::KAN^R</i>	(Yu <i>et al.</i> , 2004)
<i>spo14Δ</i>	BY4742; <i>spo14Δ::KAN^R</i>	(Yu <i>et al.</i> , 2004)
<i>syt1Δ</i>	BY4742; <i>syt1Δ::KAN^R</i>	(Yu <i>et al.</i> , 2004)
<i>tus1Δ</i>	BY4742; <i>tus1Δ::KAN^R</i>	(Yu <i>et al.</i> , 2004)
<i>vtc4Δ</i>	BY4742; <i>vtc4Δ::KAN^R</i>	(Zhu <i>et al.</i> , 2001)
<i>yel1Δ</i>	BY4742; <i>yel1Δ::KAN^R</i>	(Yu <i>et al.</i> , 2004)
<i>yhr131cΔ</i>	BY4742; <i>yhr131cΔ::KAN^R</i>	(Yu <i>et al.</i> , 2004)
<i>ypr091cΔ</i>	BY4742; <i>ypr091cΔ::KAN^R</i>	(Yu <i>et al.</i> , 2004)

¹Refers to the original study that reported binding of the indicated protein to PI(4)P.

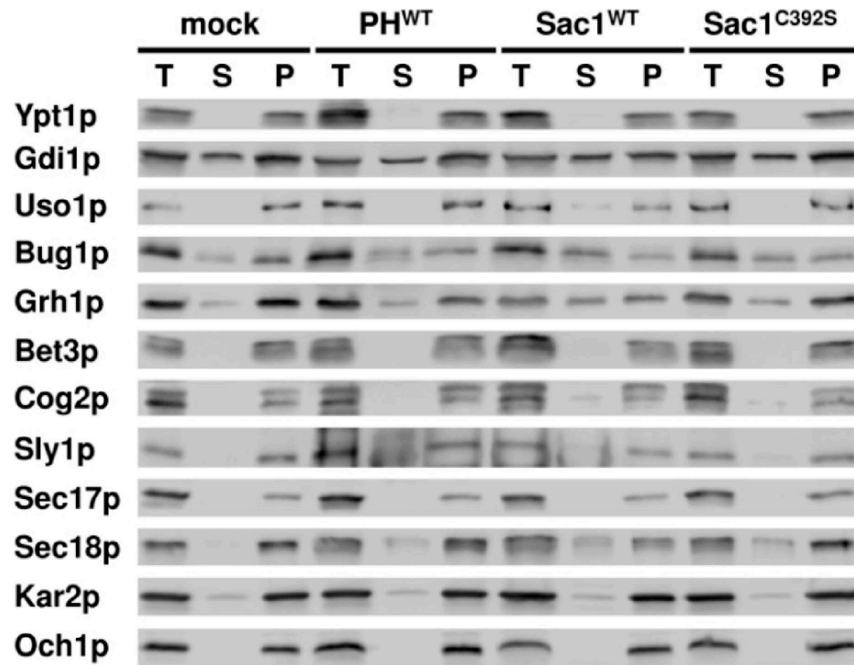
All of the above strains are from the Research Genetics Collection (Brachmann *et al.*, 1998), except for the *osh1Δ2Δ3Δ* strain (Beh *et al.*, 2001). Because our results showed that PI(4)P was required on Golgi membranes for fusion with COPII vesicles, we performed two-stage fusion assays chasing wild type vesicles into acceptor semi-intact cell membranes prepared from the indicated deletion mutants compared to their isogenic wild type. All deletion strains tested above displayed normal fusion levels except for the *gcs1Δ* mutant. Semi-intact cell membranes from the *gcs1Δ* strain exhibited a 47% decrease in fusion efficiency but displayed normal levels of COPII vesicle budding and tethering *in vitro*.



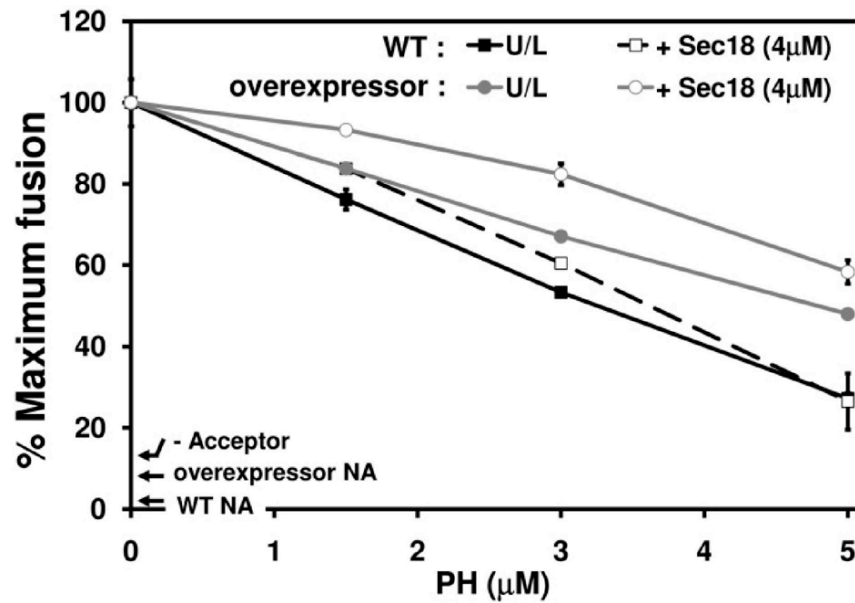
Supplemental Figure S1. The Sac1^{WT} phosphatase activity is required for inhibition of fusion. (A) Washing excess Sac1^{WT} does not rescue fusion inhibition or formation of cross-linked trans-SNARE complexes. Experiment as in Figure 4D monitored formation of cross-linked Bet1p^{I83C}-Sec22p^{D153C} heterodimers (open bars) and Golgi-modified [³⁵S]gpαf fusion (black bars). Ice controls were subtracted as in Figure 4D. Aliquots of treated membranes were immunoblotted for Sac1p using an anti-penta-His antibody and for the Golgi membrane protein Och1p (loading control). Note that the wash removed ~78% Sac1^{WT}. (B) Sac1^{WT} is catalytically active on semi-intact cell membranes *in vitro* under pre-treatment conditions. Washed semi-intact cell membranes were treated with 50 μM Sac1^{WT} or 50 μM Sac1^{C392S} as in Figure 4, lipids extracted and levels of PI(4)P/cell quantified using a lipid overlay assay (shown below). Note that PI(4)P levels are reduced by ~37% ($p=0.029$). (C) NEM inactivation of the catalytic phosphatase activity in Sac1^{WT} reverses inhibition. Semi-intact cell acceptor membranes pretreated with buffer, 50 μM mock-inactivated Sac1^{WT} or 50 μM NEM-inactivated Sac1^{WT} as in Figure 4 were used in two-stage fusion reactions as described in Figure 4A. Note that the mock treated Sac1^{WT} inhibited fusion while the NEM-inactivated Sac1^{WT} did not. The arrow indicates background fusion in the absence of added acceptor membranes.



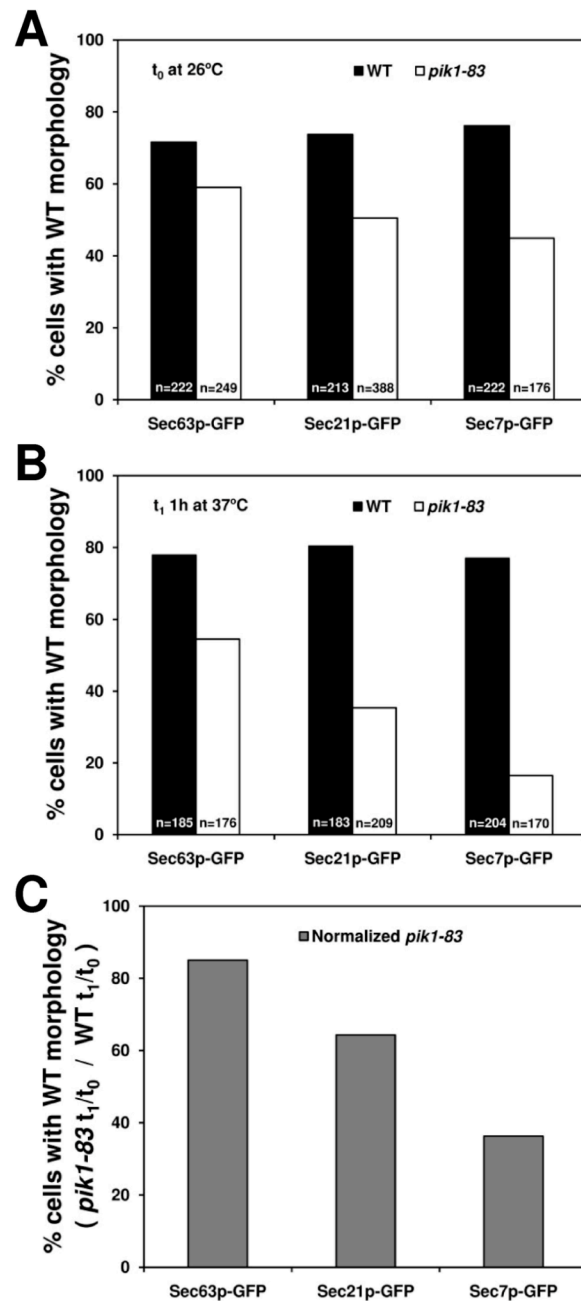
Supplemental Figure S2. Cytosol does not reverse PH^{WT} inhibition of transport. (A) Two-stage fusion reactions reconstituted with cytosol are sensitive to inhibition with PH^{WT} domain. Semi-intact cell acceptor membranes pre-treated with indicated amounts of PH^{WT} for 20 min at 4°C were used in two-stage fusion reactions as in Figure 1B. (B) Increasing concentrations of cytosol do not reverse inhibition of fusion by PH^{WT} domain. Two-stage fusion reactions using increasing concentrations of cytosol were performed as in panel A. (C) Immunoblot analysis of cytosol. Two different amounts of cytosol were compared to aliquots of semi-intact cell membranes prepared from the indicated deletion strains (Brachmann *et al.*, 1998). Note that Grh1p, Bug1p, Sec18p and Gcs1p are enriched in this cytosol. Antibodies against Gcs1p have been described (Lewis *et al.*, 2004). Cytosol was prepared as described (Barlowe, 1997).



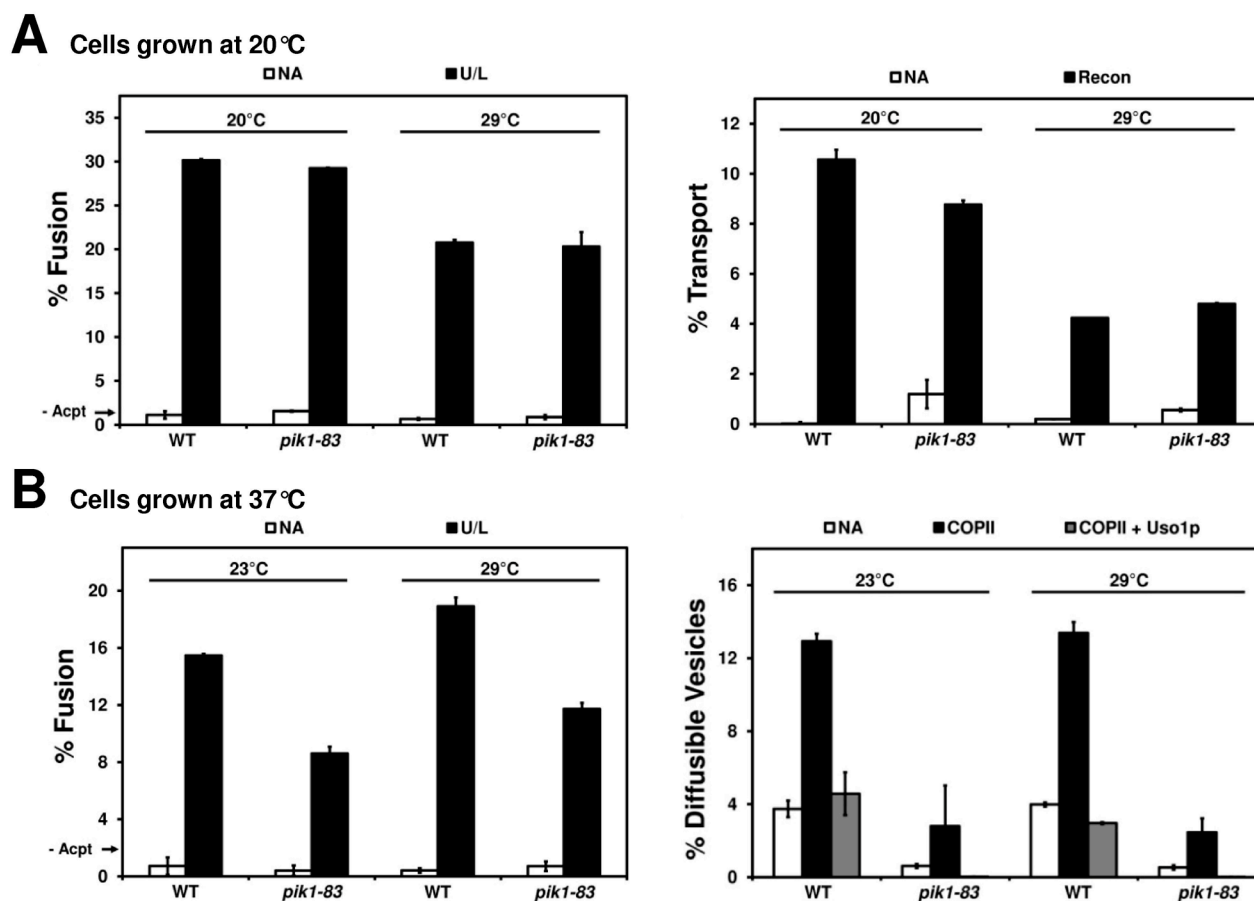
Supplemental Figure S3. Membrane association of transport factors after treatment of semi-intact cells with PH^{WT}, Sac1^{WT} or Sac1^{C392S}. Semi-intact cells (65 A₂₈₀ units/ml) were pretreated with 50 μM Sac1^{WT} for 30 min at 12°C and then diluted in buffer 88 to a final concentration of 12.5 A₂₈₀ units/ml. In parallel, semi-intact cells (12.5 A₂₈₀ units/ml) were incubated on ice for 20 min with buffer, 7 μM PH^{WT} or 5 μM Sac1^{C392S}. All reactions were then incubated for 30 min at 23°C and separated into soluble (S100) and pellet (P100) fractions for immunoblot analysis of the Rab GTPase Ypt1p and Rab GDP dissociation inhibitor Gdi1p; the tethering factors Uso1p, Bug1p, Grh1p, Bet3p (TRAPP complex), and Cog2p (COG complex); the fusion factors Sly1p (SM protein), Sec17 (α-SNAP) and Sec18 (NSF); and the control proteins Kar2p (soluble luminal ER chaperone) and Och1p (Golgi membrane marker). Note that these treatments shifted a low level of Sec18p, Grh1p and Bug1p into the soluble fraction whereas other factors were not affected. In the Sly1p immunoblot PH^{WT} was detected as a cross-reactive species.



Supplemental Figure S4. Overexpression of anterograde ER-Golgi fusion factors partially rescues PH^{WT} inhibition. Data in Figure 7C plotted relative to the control transport reaction in the absence of inhibitor. NA is background fusion in the absence of fusion factors. The - Acceptor condition indicates the background fusion in the absence of added semi-intact cell acceptor membranes.



Supplemental Figure S5. Quantification of ER and Golgi morphologies in wild type and *pik1-83* cells from Figure 8. (A) ER and Golgi morphologies in GFP-tagged strains after growth at 26°C and (B) after shift to 37°C for 1 h. To quantify ER morphology, cells expressing Sec63p-GFP were scored for the appearance of elaborated and gapped ER structures compared to wild type cells, which exhibited a typical perinuclear and peripheral ER morphology (Prinz *et al.*, 2000). Cis-Golgi structures were assessed in cells expressing Sec21p-GFP where wild type morphology corresponded to 3-8 medium-sized puncta per cell; and trans-Golgi structures in Sec7p-GFP expressing wild type cells corresponded to 2-6 larger puncta per cell (Rossanese *et al.*, 2001). Abnormal Golgi morphologies included smaller and more numerous puncta, as well as ring-like structures that may correspond to Berkeley bodies. (C) Data from panel B was normalized to panel A (t_1/t_0), and the *pik1-83* t_1/t_0 data was then normalized to WT t_1/t_0 to account for the defective morphologies at 26°C. The morphological defects in *pik1-83* cells shifted to the restrictive temperature of 37°C are severe at trans-Golgi compartments, moderate at cis-Golgi compartments and minor at ER compartments.



Supplemental Figure S6. Analysis of *pik1-83* membranes in cell free transport assays. (A) Semi-intact cells were prepared from wild-type (WT) or *pik1-83* cells grown at 20°C. LEFT: Two-stage fusion reactions performed as in Figure 1B using wild-type COPII vesicles and the indicated semi-intact cell acceptor membranes. The reactions were incubated at 20°C or 29°C (highest temperature at which the *in vitro* assay remains active). RIGHT: One-stage transport reactions as in Figure 1A were incubated at 20°C or 29°C. Note that there are no significant fusion or transport defects in *pik1-83* membranes compared to WT. NA is the background level of transport in the absence of transport factors. The “-Acpt” condition represents the background level of fusion in the absence of added acceptor membranes. (B) Semi-intact cells were prepared from WT or *pik1-83* cells grown at 26°C and then shifted to 37°C for 1 h. Two-stage fusion reactions (LEFT) and diffusible vesicle experiments (RIGHT) were conducted as in panel A and Figure 1C, respectively, and incubated at 23°C or 29°C. The *pik1-83* semi-intact cell membranes displayed defects in vesicle fusion (~44% inhibition) and budding (~80% inhibition) after cells were grown at 37°C for 1 h. (A-B) Note that the *pik1-83* membranes did not display thermosensitive defects *in vitro* and suggests that the *pik1-83* encoded enzyme remains active at 29°C *in vitro* or that there is a sufficient pool of PI(4)P on acceptor membranes such that continual synthesis is not needed under conditions of the cell free transport assay. However, after shifting cells to 37°C for 1 hr, the semi-intact cell membranes displayed defects in vesicle budding and fusion suggesting that depletion of PI(4)P *in vivo* compromises ER and Golgi membrane function in these cell free assays.

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