Inhibition of Endosome Function in CHO Cells Bearing a Temperature-sensitive Defect in the Coatomer (COPI) Component \( \epsilon \)-COP

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Abstract. Recent evidence has suggested that subunits of the coatomer protein (COPI) complexes are functionally associated with endosomes in mammalian cells. We now provide genetic evidence that COPI plays a role in endocytosis in intact cells. The ldlF mutant CHO cell line bears a temperature-sensitive defect in the COPI subunit \( \epsilon \)-COP. In addition to exhibiting conditional defects in the secretory pathway, we find that the cells are also defective at mediating endosome-associated functions. As found for cells microinjected with anti-COPI antibodies, ldlF cells at the restrictive temperature could not be infected by vesicular stomatitis (VSV) or Semliki Forest virus (SFV) that require delivery to acidic endosomes to penetrate into the cytosol. Although there was no temperature-sensitive defect in the internalization of receptor-bound transferrin (Tfn), Tfn recycling and accumulation of HRP were markedly inhibited at the restrictive temperature. Sorting of receptor-bound markers such as EGF to lysosomes was also reduced, although delivery of fluid-phase markers was only partially inhibited. In addition, lysosomes redistributed from their typical perinuclear location to the tips of the ldlF cells. Mutant phenotypes began to emerge within 2 h of temperature shift, the time required for the loss of detectable \( \epsilon \)-COP, suggesting that the endocytic defects were not secondary to a block in the secretory pathway. Importantly, the mutant phenotypes were also corrected by transfection of wild-type \( \epsilon \)-COP cDNA demonstrating that they directly or indirectly reflected the \( \epsilon \)-COP defect. Taken together, the results suggest that \( \epsilon \)-COP acts early in the endocytic pathway, most likely inhibiting the normal sorting and recycling functions of early endosomes.

Endocytosis is responsible for the internalization of extracellular fluid and macromolecules as well as plasma membrane proteins, receptors, and their bound ligands. Within early endosomes, this complex cargo must be sorted to ensure proper delivery to the appropriate intracellular destinations. Many receptors such as low density lipoprotein (LDL) receptor and some ligands such as transferrin (Tfn) are recycled back to the plasma membrane, whereas bulk fluid and ligands that dissociate from their receptors because of acidic endosomal pH are targeted to lysosomes (Gruenberg and Maxfield, 1995; Mellman, 1996). Other sorting events lead from endosomes back to the Golgi complex, specialized storage compartments, or in polarized cells, a plasma membrane domain other than the domain of entry (Mellman, 1996). The mechanisms responsible for sorting in early endosomes remain largely unknown, but seem likely to involve the formation of transport vesicles or tubules that selectively include or exclude individual components. As found for sorting or transport events in other organelles, endosomal sorting may directly or indirectly involve the assembly of cytosolic coatomer proteins (COP).

Several distinct coat complexes have been identified to be functionally important in molecular sorting and vesicle budding. These include clathrin and the clathrin adaptor proteins responsible for much of the transport of receptors from the plasma membrane and the trans-Golgi network to endosomes (Pearse and Robinson, 1990) and the COPI and COPII complexes, which are well known to play important roles in mediating transport through the secretory pathway. COPI was first identified as coating Golgi-derived transport vesicles (Orci et al., 1986, 1993) and shown to function in the transport between the ER and Golgi (Pepperkock et al., 1993). It is a multimeric complex termed
coatomer consisting of α (160), β (110), β′ (102), γ (98), δ (61), ε (31), and ζ (20 kD) subunits (Waters et al., 1991).

The COPI coatomer is thought to associate with the cytoplasmic face of membranes under the regulatory control of small GTPases known as ADP ribosylation factors (ARFs; Takizawa and Malhotra, 1993). Inhibition of ARF nucleotide exchange by the fungal metabolite brefeldin A (BFA) prevents the binding of coatomer to membranes, and thus the normal sorting and transport functions of the ER and the Golgi complex (Donaldson et al., 1992; Helms and Rothman, 1992). In the secretory pathway, COPI may function in vesicle bud formation, selection of cargo for transport, or both (Scheder and Orci, 1996). A role in cargo selection was suggested by findings that COPI subunits can interact with KKXX or aromatic amino acid-containing motifs on the cytoplasmic domains of many membrane proteins (Letourneur et al., 1994; Fiedler et al., 1996; Lowe and Kreis, 1996; Tisdale et al., 1997).

Both clathrin and coatomer have been identified as candidates for endocytic sorting and vesicle budding, largely because of biochemical and immunocytochemical evidence that they can associate with endosomes in vitro (Whitney et al., 1995; Aniento et al., 1996; Stoovogel et al., 1996; Takei et al., 1996). In the case of COPI, endosome-binding is specific and ARF dependent. Only a subcomplex of coatomer appears to bind, with γ- and δ-COP associating at lower than the normal 1:1 stoichiometry relative to the other subunits (Whitney et al., 1995; Aniento et al., 1996). Although the functional role for clathrin on endosomes has not yet been explored, several considerations have suggested that COPI may be associated with one or more endosomal activities. Initial hints came from observations that BFA caused a dramatic increase in long tubular elements of the endosomes reminiscent of those observed emanating from the Golgi complex in BFA-treated cells (Hunziker et al., 1991; Lippincott-Schwartz et al., 1991; Wood et al., 1991). Similarly, overexpression of inactive ARF mutants have been shown to alter endosome morphology or function (Zhang et al., 1994; D’Souza-Schorey et al., 1995; Peters et al., 1995). Although BFA did not greatly affect the uptake of receptor-bound or fluid-phase tracers, it did markedly alter endosomal sorting functions, altering recycling efficiency as well as the fidelity of basolateral to apical transcytosis in epithelial cells (Hunziker et al., 1991; Apodaca et al., 1993; Matter et al., 1993).

More direct evidence of a role for COPI in endosome function has come from experiments performed both in intact cells and in cell-free assays. Microinjection of antibodies to β-COP, previously shown to block transport between the ER and the Golgi, were found to block infection of vesicular stomatitis virus (VSV), an enveloped virus that must reach endosomes of a suitably acidic pH (<pH 6) to fuse with the endosome membrane and penetrate into the cytosol for replication (Whitney et al., 1995). A similar result was obtained for the low pH-dependent penetration of diaphragm toxin from endosomes (Lemichez et al., 1997). In vitro, β-COP antibodies were found to inhibit the formation of endosome-derived structures thought to represent vesicles that mediate transport from early to late endosomes (Aniento et al., 1996).

In addition, a mutant CHO cell line (ldlf) found to have a temperature-sensitive defect in ε-COP was found not only to have a defect in the transport of newly synthesized LDL receptors, but also to perturb the behavior of LDL receptors that had already reached the plasma membrane (Hobbe et al., 1994; Guo et al., 1994, 1996). At the restrictive temperature, surface LDL receptors were not recycled properly but prematurely and ectopically degraded.

If COPI subunits were to play a role in endosome function, a minimal expectation would be that cells defective in ε-COP would exhibit one or more defects in the endocytic pathway. To this end, we have now examined endocytosis in ldlf cells in some detail. Our results clearly show that these cells exhibit significant alterations in the handling of endocytic tracers internalized as bulk fluid or receptor-bound ligands. Although the experiments do not pinpoint the precise step(s) at which COPI might act in the endocytic pathway, they do suggest that alterations in COPI activity directly affect the normal sorting functions of early endosomes in CHO cells.

Materials and Methods

Cell Culture

All cells were maintained in α-MEM containing 10% FCS. Wild-type (wt) and mutant CHO cells were maintained at 34°C and shifted to 40°C for experiments, whereas complementation ldlF[LDLF] CHO cells were maintained under selection at the nonpermissive temperature, 40°C, and then shifted to 34°C for experiments to parallel wt and ldlf cells at this temperature. Cells were plated 2 d before all experiments into multiwell plates for [125]Tfn and HRP experiments or onto glass coverslips for immunofluorescence. Cells were used at 50–75% confluence.

Detection of ε-COP by Western Blot

After incubation at either 34°C or 40°C for 12 h, cells were washed extensively in PBS/900 μM Ca2+ per 500 μM pMg2+ and lysed in PBS containing 1% Triton X-100. Samples were then normalized for total cell protein using a modified biuret assay (BCA assay; Pierce Chemical Co., Rockford, IL), which was insensitive to the concentrations of Triton X-100 used. Samples were then subjected SDS-PAGE and then transferred to 0.2-μm nitrocellulose for Western blotting. A rabbit polyclonal anti-ε-COP (gift from F. Wieland, University of Heidelberg, Heidelberg, Germany) was used at 1:1,000 and detected using HRP-coupled goat anti–rabbit secondary antibody diluted at 1:2,000, followed by enhanced chemiluminescence (Amer sham Corp., Arlington Heights, IL).

Virus Infection of Cells

Endocytosis and infection of Semliki Forest virus (SFV) or tsO45 VSV was performed essentially as described (Kreis, 1986). Briefly, before infection, cells were preincubated at 34°C or 40°C for <10 h. After 1 h of preabsorption, cells were incubated with virus for 2.5 h. Cells were then fixed and extracted with methanol at −20°C for 4 min. Viral infection was detected by immunofluorescence using monoclonal antibody, PSD4, directed against VSV G protein (Kreis, 1986), or E2-1 (Kielian et al., 1990) directed against SFV E1/E2 spike glycoprotein. Delivery of SFV to an acidic compartment was monitored by immunofluorescence using the monoclonal antibody, E1-al (Kielian et al., 1990), directed against the acid-specific conformation of the E1 subunit of the SFV viral spike glycoprotein. For these experiments, endocytosis of SFV was carried out for 30 min in cells that were first preincubated at 34°C or 40°C for 10 h.

Steady-state Distribution of Tfn Receptors

Cells were first incubated at 34°C and 40°C for <12 h. To rid the cells and medium of unlabeled Tfn, a brief incubation in serum-free medium (30 min, α-MEM + 2 mg/ml BSA) was performed. The cells were then labeled for 60 min at 34°C or 40°C with 2.5 μg/ml [125]Tfn and were then transferred to ice to prevent any further endocytic traffic. Iodination of Tfn was performed as previously described (Podbielczew and Mellman, 1990). External [125]Tfn was removed by extensive washing with PBS at
4°C. Surface-bound [125I]Tfn was determined by cycling several times between acid (α-MEM + 2 mg/ml BSA, pH 2.6) and neutral (PBS/900 μM CaCl₂ per 500 μM MgCl₂) to disassociate the ligand from its receptor. The combined washes were assayed to determine total amount of ligand from the cell surface. After removal of cell surface [125I]Tfn, the internal fraction was assayed by lysing cells in 1% Triton X-100 and γ counting. Values are presented as percent total of bound [125I]Tfn. Total [125I]Tfn was calculated as the sum of cell surface and internal [125I]Tfn.

Internalization of [125I]Tfn
After incubation at 34°C or 40°C for 0–8 min. The cells were then washed extensively and surface-bound Tfn was removed as described above. The cells were then lysed in 1% Triton X-100 total and cell protein was determined by a modified biuret assay (BCA assay; Pierce Chemical Co.) that was insensitive to the concentrations of Triton X-100 used.

[125I]Tfn Recycling
After incubation at 34°C or 40°C for <12 h, the cells were labeled to steady state as described above. Cells were then washed extensively with ice-cold PBS/900 μM CaCl₂ per 500 μM MgCl₂. Recycling was then initiated by warming the cells simultaneously to 34°C with 2× MEM + 2 mg/ml BSA in the presence of 100-fold excess unlabeled Tfn. At each time point the medium was removed for γ counting and replaced with fresh medium. At the end of the recycling assay, cell-associated [125I]Tfn was obtained by lysing in 1% Triton X-100 and scraping the wells with a rubber policeman. Total [125I]Tfn was calculated as the sum of all recycled and cell-associated [125I]Tfn. Essentially the same kinetics were obtained when cells were labeled to steady state and surface-bound Tfn was acid stripped as described above before initiation of recycling (data not shown).

Specificity of [125I]Tfn
After incubation at 34°C and 40°C, all cell lines were labeled to steady state as described above in the absence or presence of 100-fold excess unlabeled Tfn. Under all conditions, noncompetitive uptake of [125I]Tfn represented ~2% of the total specific signal.

Fluid-phase Endocytosis
Cells were labeled at either 34°C or 40°C with 2.5 mg/ml HRP (Type VI; Sigma Chemical Co., St. Louis, MO) in α-MEM + 2 mg/ml BSA for various times <120 min after incubation at the permissive and nonpermissive temperatures for <12 h. The cells were then transferred to ice and washed three times with α-MEM + 5% BSA, three times with α-MEM + 2 mg/ml BSA, and three times with PBS/900 μM CaCl₂ per 500 μM MgCl₂. The cells were then lysed in 1% Triton X-100 and spun at 1,000 rpm for 10 min in an Eppendorf microcentrifuge (model 1514C; Brinkman Instruments, Westbury, NY). The supernatants were assayed for HRP activity as previously described (van der Sluijs et al., 1991) and normalized to total cell protein (BCA assay; Pierce Chemical Co.).

Density Gradient Fractionation
Cells were labeled either with [125I]Tfn or by uptake of HRP as described above. For each gradient, three confluent 10-cm culture dishes were scraped into 0.6 ml of cracking buffer (78 mM KCl, 4 mM MgCl₂, 8.4 mM CaCl₂, 10 mM EGTA, 50 mM Hepes/KOH, pH 7.0) plus 250 mM sucrose and passed four times through a ball-bearing cell homogenizer using an 0.2498" ball bearing. The homogenate was centrifuged at 1,000 g for 5 min to produce a postnuclear supernatant (PNS) that was loaded on top of a 12-ml preformed linear 5 to 20% gradient (optiprep: Nycomed, Oslo, Norway) made with cracking buffer. The gradients were developed at 100,000 g for 20 h and then decepted mechanically. Fractions were tested for β-hexosaminidase activity using 4-methylumbelliferyl-N-acetyl-β-D-glucosaminide (Sigma Chemical Co.) as a fluorescent substrate. Activity was assayed with an excitation wavelength of 365 nm and an emission wavelength of 450 nm.

Immunofluorescence
Cells grown on coverslips were first incubated at the permissive or non-permissive temperatures for up to 12 h. The cells were then fixed in 2.5% paraformaldehyde and subsequently processed for immunofluorescence. All steps were performed using PBS/2% BSA per 0.03% saponin. Fixed cells were permeabilized for 30 min, and then incubated in primary antibodies for 1 h (for lysosomes: a mouse monoclonal antibody against lysosomal glycoprotein (Igp)-B, diluted 1:10 [Harter and Mellman, 1992]). After incubation in the primary antibodies, the cells were washed for 30 min and incubated in secondary antibodies (Texas red or FITC goat anti-mouse or goat anti-rabbit; Jackson ImmunoResearch Laboratories, Inc., West Grove, PA). The cells were then washed for 30 min, rinsed briefly in PBS and then water, and mounted in Moviol (Calbiochem-Novabiochem Corp., La Jolla, CA) containing 2.5% 1,4-diazabicyclo(2.2.2)octane (Sigma Chemical Co.).

Labeling of Lysosomes by Lucifer Yellow, EGF, and lgp
For Lucifer yellow (LY), cells were first incubated for 6 h at either 34°C or 40°C and then labeled with 2 mg/ml LY (Molecular Probes Inc., Eugene, OR) for 90 min and washed for 90 min at either 34°C or 40°C. For EGF, cells were preincubated for ≤12 h at either 34°C or 40°C, and fed 100 μg/ml of biotinylated EGF complexed to Texas red-conjugated avidin (Molecular Probes Inc.) for 30 min at either 34°C or 40°C. The EGF or LY labeled cells were then rinsed briefly in PBS/900 μM CaCl₂ per 500 μM MgCl₂, fixed in 2.5% paraformaldehyde, and counterstained for late endosomes and lysosomes as described above. For internalization of the anti–lgp-B antibody, cells were first preincubated for 6 h at 34°C or 40°C and then incubated for 60 min at either temperature in antibody-containing culture supernatant before fixation in 2.5% paraformaldehyde. The cells were then permeabilized and the internalized primary antibody was detected using FITC goat anti-mouse secondary antibodies as described above. The lgp-β antibody concentration was sufficiently low to prevent detectable uptake of antibody by fluid-phase endocytosis.

Confocal Fluorescence Microscopy
Fluorescently labeled cells were examined using a modified confocal microscope (model MRC600; Bio-Rad Laboratories, Hercules, CA) attached to a microscope (model 451888 Axiotool; Carl Zeiss, Inc., Thornwood, NY) using separate filters for each fluorochrome viewed. FITC: exc = 490 nm; em = 515 nm; Texas red: exc = 565 nm; em = 585 nm. Control slides (single-labeled with each fluorophore) were examined to ensure that no cross-talk or bleed-through was occurring for the given confocal conditions. Images were imported into Adobe Photoshop and processed.

FITC–Tfn Labeling of Endosomes
Cells grown on coverslips were first incubated at the permissive or non-permissive temperatures and then serum-starved for 30 min at either 34°C or 40°C in α-MEM + 2 mg/ml BSA to deplete the cells and medium of unlabeled Tfn. The cells were then incubated for 90 min in 50 μg/ml FITC–Tfn (human holo form; Molecular Probes Inc.) in α-MEM + 2 mg/ml BSA per 20 mM Hepes, pH 7.4, at either 34°C or 40°C. The cells were then rinsed briefly with PBS/900 μM CaCl₂ per 500 μM MgCl₂, fixed in 2.5% paraformaldehyde, rinsed briefly in PBS and then water, and mounted in Moviol (Calbiochem-Novabiochem Corp.) containing 2.5% 1,4-diazabicyclo(2.2.2)octane (Sigma Chemical Co.).

Intracellular pH Measurement
Intracellular pH was determined essentially as described (Kaplan and Bon ron, 1994). Briefly, cells were grown on rectangular coverslips designed for use in a 2-ml cuvette. After an 8-h incubation at either 34°C or 40°C, cells were loaded for 15 min with 1 μM BCECF-AM (Molecular Probes Inc.) diluted in loading buffer (125 mM NaCl, 5 mM KCl, 1 mM CaCl₂, 1.2 mM MgSO₄, 7H₂O, 2 mM NaH₂PO₄, H₂O, 32 mM Hepes, 10.5 mM glucose, 40 mM HCl, 5.8 mM N-methyl-D-glucamine (Sigma Chemical Co.) at pH 7.0, 7.5, 7.0, 6.5, 6.0, and 5.5.

Endosomal pH Measurement
Endosomal pH was measured using a modification of the method of Yamashiro and Maxfield (Yamashiro and Maxfield, 1987). Cells were labeled...
with FITC–Tfn as described above. Coverslips were then transferred to a chamber to permit confocal imaging of live cells. The coverslips were then rinsed in loading buffer (see Intracellular pH Measurement) and images were acquired using the BHS filter set (ex = 488 nm, em = 515 LP) and the same confocal settings for all cell lines. The pH of the endosomes was then artificially raised to pH 7.4 using 1 μM nigericin (Sigma Chemical Co.) in calibration buffer described in intracellular pH measurement, and a second set of images were acquired using the same confocal settings. The percent increase in fluorescence intensity was calculated using NIH image. A greater percent increase in fluorescence intensity represents a lower endosomal pH.

Results

Idlf Cells Lack e-COP

The Idlf CHO cell mutant was identified in a screen intended to isolate cell lines with temperature-sensitive defects in LDL receptor activity. The cells were isolated because of their inability to transport newly synthesized LDL receptors to the plasma membrane and to maintain previously synthesized LDL receptors at the surface (Hobbie et al., 1994). cDNA transfection experiments and cloning of the mutant Idlf allele identified the site of the mutation as being a point mutation in the coding region of the gene encoding e-COP (Guo et al., 1994, 1996). To confirm that the cells used for these experiments were deficient in e-COP, Western blot analysis was performed on wt, mutant (Idlf cells), and mutant cells transfected with a cDNA expressing wt e-COP (Idlf[LDLF] cells) after incubation at either the permissive (34°C) or nonpermissive temperature (40°C) for 12 h. As shown in Fig. 1, Idlf cells have more than five-fold less e-COP than wt or Idlf[LDLF] cells even at 34°C and after incubation at 40°C for 12 h, e-COP was undetectable in the mutants, but barely affected in cells expressing wt e-COP. The time course of the loss of e-COP from Idlf cells indicated that at 40°C, the t1/2 of e-COP degradation was ~1–2 h (Guo et al., 1996).

Virus Infection Is Inhibited in Idlf Cells

In previous work, we found that microinjection of antibodies to β-COP, another COPII subunit that binds to endosomes, blocked infection by VSV (Whiteley et al., 1995). Microinjection of VSV nucleocapsids, however, circumvented this block. Since the normal infectious pathway of enveloped viruses such as VSV involves delivery to endosomes of sufficiently acidic pH to trigger fusion of the viral membrane with the endosomal membrane, we reasoned that the anti-β-COP antibody affected the delivery of the virus to endosomes capable of hosting viral penetration or inhibited some feature (perhaps acidification) of endosomes themselves. A minimal expectation, therefore, was that Idlf cells would be similarly unable to host virus infection. For these experiments, we used both VSV and a second enveloped virus, SFV. Whereas VSV has a fairly broad pH threshold for fusion (pH 5.3–6.3; Matlin et al., 1982), the SFV E1/E2 spike glycoproteins exhibit a pH-induced conformational change leading to fusion at pH < 6.2 (White et al., 1980). Thus, SFV may penetrate from an earlier or less acidic site on the endocytic pathway than VSV (Schmid et al., 1989).

Cells were incubated at 34° or 40°C for 10 h and then exposed to wt SFV or the tsO45 mutant of VSV and fixed, and then infection was monitored by immunofluorescence using antibodies to SFV E1/E2 or VSV G protein. As shown in Fig. 2, SFV was incapable of infecting the mutant cells at the restrictive temperature. However, clear synthesis of E1/E2 protein was observed in Idlf cells incubated at 34°C or in wt or complemented Idlf[LDLF] cells at either temperature. Similar results were obtained using tsO45 VSV (data not shown). These experiments were quantified by counting several hundred cells stained by immunofluorescence for the SFV E1/E2 or VSV G spike glycoproteins (Whiteley et al., 1995). Although at the permissive temperature, >90% of the Idlf cells were infected by VSV or SFV; after 12 h at 40°C the infection frequency was reduced to 5 and 10%, respectively.

The failure of VSV and SFV to infect CHO cells in the absence of functional e-COP suggested that virus particles were not delivered to endosomal compartments of sufficiently acidic pH to trigger fusion, or that the endosomes were rendered defective for some other secondary reason. To distinguish between these possibilities, we monitored the delivery of SFV to acidic compartments using a monoclonal antibody (E1a-1) specific for the acid conformation of the E1 spike glycoprotein (Kielian et al., 1990). After 30 min of uptake, E1a-1 reactive virus (i.e., virus particles that had reached pH < 6.1) was detected in wt cells and Idlf[LDLF] cells preincubated at either 34° and 40°C and in Idlf cells preincubated at 34°C (Fig. 3). The acid conformation was not detected after SFV internalization by Idlf cells incubated at the restrictive temperature. Thus, in the absence of e-COP, the virus was unable to reach a compartment of sufficiently low pH to trigger fusion. This seemed unlikely to reflect a global acidification defect in Idlf cells, since they accumulated the pH-sensitive probe acridine orange in a fashion similar to wt cells (see Endosomal pH Assessment). Additionally, Idlf cells were capable of viral infection when virus was microinjected directly into the cytosol. Thus, the inhibition of viral infection may reflect a block in intracellular transport.

Idlf Cells Have a Partial Defect in Tfn Endocytosis

To study transport along the endocytic pathway in Idlf

Figure 1. Idlf cells lack e-COP. After incubation of wt and mutant cells at either 34° or 40°C for 12 h, cell lysates were normalized for total cell protein and then subjected to SDS-PAGE and immunoblotting. At 34°C, Idlf cells have much less e-COP than wt cells, and after incubation at 40°C for 12 h, e-COP is undetectable in the mutant cells. The defect was complemented by transfection with wt e-COP in Idlf[LDLF] cells.

Figure 2. Idlf cells lack e-COP. After incubation of wt and mutant cells at either 34° or 40°C for 12 h, cell lysates were normalized for total cell protein and then subjected to SDS-PAGE and immunoblotting. At 34°C, Idlf cells have much less e-COP than wt cells, and after incubation at 40°C for 12 h, e-COP is undetectable in the mutant cells. The defect was complemented by transfection with wt e-COP in Idlf[LDLF] cells.
ldlF Cells Exhibit a Temperature-sensitive Defect in Tfn Recycling

The observation that ldlF cells did not exhibit an alteration in equilibrium distribution of TfRs suggested the existence of an inhibition of Tfn recycling to balance the slowed rate of endocytosis. To measure recycling rates, cells were first incubated at 34°C or 40°C for ≤12 h and then labeled to steady state with [125I]Tfn at either temperature. After washing at 4°C, recycling of internalized Tfn was initiated by warming the cells back to 34°C or 40°C in the presence of excess, unlabeled Tfn to ensure quantitative dissociation of recycled [125I]Tfn. The amount of [125I]Tfn recycled back to the medium was determined at various
wt and mutant cells at 40°C. Tfn internalization is inhibited in ldlF cells. After incubation of wt and mutant cells with [125I]Tfn, intracellular and cell surface receptors were saturated with [125I]Tfn by labeling for 60 min at 34°C and 40°C. Surface and internal Tfn were determined as described in Materials and Methods. Internalization of Tfn was inhibited in ldlF cells lacking e-COP. 

Importantly, expression of wt e-COP in ldlF cells almost completely complemented the temperature-sensitive defect in Tfn recycling (Fig. 5 A; ldlF[LDLF] cells) with an overall t1/2 of recycling of ~3.7 min at 34°C and ~3.4 min at 40°C. These results demonstrate that the temperature-sensitive recycling defect was a direct or indirect consequence of the loss of e-COP, but not because of a second site mutation in ldlF cells.

The fact that e-COP, as well as most other coatomer subunits, bind to endosomes suggests that the loss of e-COP function may directly affect the recycling activity of one or more populations of early endosomes. If e-COP was required for endosome function, the kinetics of the onset of the recycling defect should mirror the kinetics of loss of e-COP upon shifting to the restrictive temperature. To determine the time of onset of the recycling defect, ldlF cells were incubated at 40°C for 0–9 h before loading with [125I]Tfn. Recycling was assayed by measuring the amount of [125I]Tfn released into the medium during a 15-min chase (the time at which the difference between wt and mutant cells was maximal; see Fig. 5 A). As shown in Fig. 5 A, recycling activity began to decrease 1 h after shifting to the restrictive temperature and the defect was qualitatively half-maximal by 2 h. These data agree well with the kinetics of e-COP loss upon shifting ldlF cells to 40°C and with the onset of the transport defect in the secretory pathway, both of which appear with t1/2 of ~1–2 h (Hobbie et al., 1994). The simultaneous loss of recycling activity and e-COP is consistent with the possibility that e-COP plays a direct role in the endocytic pathway.

**Endosomal pH Assessment**

Since defects in endocytosis and recycling are sometimes observed after treatment with drugs that alter endosomal pH, we next asked if the early endosomes accessed by Tfn in ldlF cells exhibited an altered internal pH. These determinations were made by quantitative fluorescence microscopy of single cells using a modification of previously published methods (Yamishiro, 1987). After incubation at 34°C or 40°C for 8 h, wild-type, ldlF, or ldlF[LDLF] cells were

![Figure 4.]()
loaded with FITC–Tfn for 60 min, washed, and then viewed by confocal microscopy to establish baseline values. The percent increase in FITC fluorescence intensity was then monitored after nigericin addition and subsequent equilibration of endosomal pH to the pH of the medium (pH 7.4). The greater the percentage increase, the lower the starting endosomal pH. As shown in Fig. 6, within the limits of this technique, we were unable to detect any significant differences in the relative pH values of Tfn-containing endosomes in wt or mutant cells at the permissive or nonpermissive temperatures. Given the nature of these measurements, we did not attempt to assign specific pH values. Since wt and ldlF cells were able to deliver FITC–Tfn to early and/or recycling endosomes of comparable internal pH values, these results strongly suggest that the inability of SFV and VSV to infect ldlF cells at 40°C does not reflect a failure of endosomal acidification, but rather a failure of delivery of virus particles to acidic endosomes.

**Loss of ε-COP Inhibits Accumulation of Fluid-phase Endocytic Tracers**

We next measured the capacity of ldlF cells to accumulate the extracellular tracer, HRP by fluid-phase endocytosis. HRP is known to be internalized and delivered to lysosomes where it concentrates, because of its relative resistance to degradation. After incubation at either 34°C or 40°C for 8.5 h, cells were incubated with HRP for 90 min. As shown in Fig. 7, wt cells accumulated similar amounts of HRP at either temperature. However, at the restrictive temperature, mutant cells exhibited a marked defect in HRP accumulation. Uptake of HRP at early time points (0–10 min) was also inhibited in ldlF cells (data not shown). The defect was restored to wt levels by complementation with wt ε-COP in ldlF[LDLF] cells, indicating that the phenotype was because of the defective ε-COP allele in ldlF cells (Fig. 7).

**Endosomes and Lysosomes Are Redistributed in ldlF Cells**

Our analysis of endocytosis in ldlF cells revealed that loss...
of e-COP in IdlF cells resulted in inhibition of the initial rate of internalization of solutes and ligands, as well as a decrease in the ability of the cells to recycle ligands back to the cell surface or accumulate solutes in lysosomes. Together, these phenotypes suggested that the mutation disrupted transport or sorting in early endosomes. Since the Golgi complex exhibits an altered morphology in the mutant cells (Guo et al., 1994), we next asked if there was any alteration in the distribution or organization of endocytic organelles.

We examined the distribution of late endosomes and lysosomes by staining wild-type and mutant cells with an antibody directed against a major lysosomal membrane protein lgp-B (lgp 110, lamp-2; Harter and Mellman, 1992). As shown in Fig. 8A, wt cells at 34°C and 40°C exhibited the characteristic distribution of late endosomes and lysosomes throughout the cytoplasm with a more concentrated accumulation near the nucleus (large arrows). IdlF cells, on the other hand, appeared to exhibit somewhat less perinuclear clustering at 34°C. Strikingly, after incubation at 40°C for 5–12 h, the lysosomes in the mutant cells redistributed to the cell periphery, being concentrated at the periphery. This redistribution was not observed at the permissive temperature (34°C) (Fig. 8B, large arrowheads).

Figure 7. Loss of e-COP inhibits accumulation of fluid-phase markers. Cells were incubated for 8.5 h at either 34° or 40°C before labeling with the fluid-phase marker, HRP, for 90 min at either temperature. After extensive washing, cells were lysed and assayed for HRP activity and total cell protein. Loss of e-COP in IdlF cells at 40°C resulted in an inhibition of HRP accumulation. The defect was restored in IdlF[LDLF] cells which were complemented with wt e-COP. n = 12 for wt and IdlF cells and n = 6 for IdlF[LDLF] cells.

Figure 8. Lysosome, but not endosome, distribution is altered in IdlF cells. Cells were incubated at either 34°C or 40°C to compare the distribution of internalized FITC–Tfn and the lysosomal glycoprotein lgp-B. (A) Distribution of lysosomes. IdlF and wt cells were fixed and immunostained for lgp-B. Wt cells exhibited the characteristic clustering of lysosomes in the perinuclear region (large arrowheads), whereas the lysosomes in the mutant cells seem to be more randomly dispersed at the permissive temperature. Upon incubation at the nonpermissive temperature for ≤12 h, the lysosomes of the mutant cells seem to redistribute to the cell periphery (small arrowheads), extending to the very tips of the cells. Occasionally, mutant cells were observed to display both centrally (large arrowheads, bottom right) and peripherally (small arrowheads, bottom right) clustered lysosomes. Addition of wt e-COP in the IdlF[LDLF] cells restored this phenotype (large arrowheads). Similar results were obtained when cells were incubated at 40°C for only 5 h. wt and IdlF[LDLF] panels are representative of results obtained at either temperature. (B) Distribution of FITC–Tfn-containing early endosomes. Wt, IdlF, and IdlF[LDLF] cells were incubated at the permissive or nonpermissive temperature for 12 h and then allowed to internalize FITC–Tfn for 1 h. Most (>80%) wild-type and 30–60% of IdlF CHO cells exhibited the characteristic clustering of recycling endosomes at the pericentriolar region (arrows) at both temperatures. IdlF[LDLF] cells were similar to wt cells. Arrows indicate the position of the PNRC. There was no redistribution of Tfn-containing endosomes to the cell periphery.
very tips of the cells (small arrows). This is in contrast to the pattern observed with FITC–Tfn (Fig. 8 B) that labels the recycling pathway. Thus, the lgp-B–containing structures in the cell periphery represent redistributed lysosomes rather than mistargeting of lgp-B to peripheral early endosomes. Complementation with wt e-COP in ldlF[LDLF] cells restored the lysosomal distribution to the juxtanuclear region (Fig. 8 A, large arrows), indicating that the phenotype was indeed because of the loss of e-COP. Occasionally, mutant cells were observed to display both centrally (large arrow, bottom right) and peripherally clustered lysosomes (small arrows, bottom right).

It remained possible that the lysosomal marker enzymes have shifted from the lysosomes to an endosomal compartment. To address this possibility, we separated endosomes from lysosomes on an optiprep density gradient. Subcellular fractionation of ldlF and wt cells at both the permissive and nonpermissive temperatures revealed no difference in the distribution of the lysosomal marker β-hexosaminidase, suggesting that redistribution of this lysosomal marker was not occurring (data not shown).

The redistribution of lgp-B–positive structures to the periphery of ldlF cells at 40°C was not because of a reorganization of the microtubule network in these cells, since by immunofluorescence, there were no apparent differences in methanol-fixed microtubules in wt or ldlF cells grown at the permissive or restrictive temperatures (data not shown). Microtubule disruption has been shown to disperse endosomes (McGrath et al., 1993; Daro et al., 1996), late endosomes, and lysosomes (Gruenberg et al., 1989) from the perinuclear region of CHO and other cells to a more random distribution.

CHO cells exhibit a characteristic perinuclear accumulation of early endosomes containing recycling markers (Dunn and Maxfield, 1992; Hopkins and Trowbridge, 1983). These structures are thought to represent a distinct endosome population based on their unique rab protein composition (Daro et al., 1996; Ullrich et al., 1996), ability to accumulate recycling markers such as FITC–Tfn, and ability to be physically separated from early endosomes by centrifugation (Sheff, D., E. Daro, and I. Mellman, manuscript in preparation). On average, fewer (~60%) ldlF cells demonstrated the characteristic distribution of Tfn-containing endosomes as compared to wt cells (~80%; data not shown). However, this difference was observed at both 34°C and 40°C, making it unlikely that the altered distribution was responsible for or reflected any of the temperaturesensitive endocytic defects. Although the distribution of Tfn was slightly altered in ldlF cells, Tfn-containing structures did not redistribute to the periphery or tips of the cells (Fig. 8 B).

Acidification of the cytosol to pH < 6.9 has also been shown to redistribute endosomes and lysosomes to the cell periphery in a fashion remarkably similar to that exhibited by ldlF cells at 40°C (Heuser, 1989; Parton et al., 1991). We therefore measured the intracellular pH of wt, Idl, and ldlF[LDLF] cells after incubation at 34°C and 40°C for 8 h, and found that the pH of the mutant cells was ~pH 7.2 whereas that of wt and ldlF[LDLF] was ~pH 7.4 at either temperature (Table 1). Thus, it was unlikely that cytosol acidification was responsible for the organelle redistribution observed at 40°C. In any event, the fact that the redistribution was not observed in ldlF[LDLF] cells strongly suggested that the effect was at least indirectly the consequence of an inactivation of e-COP activity.

**Delivery of Markers to Lysosomes Is Inhibited in ldlF Cells**

Given the apparent defects in endosomal sorting combined with the dramatic redistribution of endocytic organelles, we next asked if internalized fluid-phase and receptor-bound tracers could reach their appropriate destinations using fluorescence microscopy.

To determine if any of the fluid-phase marker could reach the peripheral late endosomes or lysosomes, wt, Idl, or ldlF[LDLF], cells were incubated for 6 h at 34°C or 40°C and then labeled with LY for 90 min, washed, and then chased for 90 min at 34°C or 40°C in the absence of LY. The cells were then fixed and stained for lysosomes using the lgp-B antibody. wt and ldlF[LDLF] cells exhibited the characteristic perinuclear clustering of lysosomes together with a high degree of colocalization with internalized LY, indicating that this internalized marker reached late endosomes/lysosomes at either temperature in these cells. At 34°C, there was also a high degree of colocalization between the internalized LY and late endosomes/lysosomes in the mutant ldlF cells, although the perinuclear clustering was less pronounced. Upon incubation of ldlF cells at 40°C (Fig. 9), the lgp-B–positive late endosomes/lysosomes (red) redistributed to the tips of the cells (arrowheads), but these structures were still accessible to internalized LY (green; visualized using a higher excitation intensity than wt or ldlF[LDLF] cells because of the decreased amount of LY accumulated under these conditions). Interestingly, lgp-B–positive structures that remained more towards the center of the cell were relatively devoid of LY. Thus, at the restrictive temperature, ldlF cells remained capable of delivering LY to at least a subpopulation of lgp-positive structures. These peripheral lgp/LY-containing structures may correspond to late endosomes, or may simply represent a very low number of both late endosomes and lysosomes that receive a minimal amount of LY but become visually apparent because of their concentration in a small area of the cell. In any event, it was clear that at least a portion of the internalized LY could reach late endosomes/lysosomes.

We next examined the ability of the receptor-bound ligand EGF to be delivered to lysosomes. Cells were permitted to internalize Texas red–labeled biotinylated EGF for 30 min at 34°C or 40°C, fixed, and then stained using the

<table>
<thead>
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<th>pHpKα</th>
<th>34°C</th>
<th>40°C</th>
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<tr>
<td>wt</td>
<td>7.44±7.21</td>
<td>7.43±7.15</td>
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<tr>
<td>ldlF</td>
<td>7.18±7.30</td>
<td>7.19±7.35</td>
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<tr>
<td>ldlF[LDLF]</td>
<td>7.30±7.32</td>
<td>7.41±7.24</td>
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Acidification of cytosolic pH can cause the redistribution of endocytic structures. We therefore measured cytosolic pH using the fluorescent probe BCECF in each of the cell types after incubation for 8 h at either the permissive (34°C) or nonpermissive (40°C) temperatures. For each condition, a standard curve was generated to calculate both the pKα of the dye and the intracellular pH. The mutant cells were only slightly acidified. Values are from one experiment wherein a standard curve was prepared for each condition. Similar results were obtained for wt cells and ldlF cells at 40°C in several other experiments.
anti–lgp-B antibody. In wt cells at either temperature, EGF (red) was found to colocalize extensively with lgp-positive late endosomes and lysosomes (green) in the perinuclear region yielding yellow double-positive structures (Fig. 10, top two rows). In ldlF cells, however, colocalization was already limited at 34°C and even further reduced at 40°C (Fig. 10, lower two rows). EGF was strikingly absent from the lgp-positive structures at the cell tips which stained green (arrowheads). Thus, the delivery of EGF to late endosomes and lysosomes was inhibited far more strongly than the delivery of LY. This difference may in part reflect the fact that unlike fluid-phase tracers, EGF must dissociate from its receptor after reaching acidic endosomes to reach lysosomes efficiently.

An additional assay further demonstrated that the lysosomal pathway was, in general, very inefficient in ldlF cells at 40°C. A small fraction of lgp-B made in CHO cells reaches late endosomes and lysosomes after insertion into the plasma membrane (Harter and Mellman, 1992). By binding anti–lgp-B IgG to cell surface lgps, and incubating to permit uptake of the IgG–lgp-B complex, we were able to monitor delivery to late endosomes and lysosomes in cells that were preincubated at 34° or 40°C (data not shown). Delivery of internalized anti-lgp antibody to characteristic perinuclear lysosomal structures was found in wt and ldlF-[LDLF] cells at either temperature. However, in ldlF cells that were preincubated at 40°C, the antibody failed to reach the peripheral lysosomal structures found in the tips of cells. Thus, mutant cells even failed to deliver a resident lysosomal membrane protein to its intrinsic destination at 40°C.

**Discussion**

Our results demonstrate that multiple aspects of the endocytic pathway are disrupted in CHO cells expressing a temperature-sensitive point mutation in the COPII complex subunit e-COP. Although it is well established that COPII coatomer functions in early stages of the secretory pathway, our results provide genetic evidence of a role for at least one COPI subunit in endocytic transport. These data do not define the precise transport step or steps under the control of COPI in endosomes. Nor do they prove that COPI is directly required for endosome function, since it remains formally possible that COPI is required for a transport step that delivers a rapid turn over or recycling component from the Golgi complex to endosomes. However, given recent biochemical data showing that COPI components physically associate with endosomes and appear to inhibit certain endosome-related activities, the availability of the ldlF mutants provided a critical initial test for the concept that COPI indeed is required for endosome function: the failure of ldlF cells to exhibit any defect in the endocytic pathway would have dismissed a role for COPI in endosomes. Importantly, the fact that these defects could be complemented by transfection of a wt copy

**Figure 9.** Peripheral lysosomes are accessible to fluid-phase markers in ldlF cells. To determine if the peripheral lysosomes in Fig. 7 were accessible to a fluid-phase tracer, cells were first incubated for 6 h at 40°C and then labeled with LY for 90 min and washed for 90 min at 40°C. The cells were then fixed and stained for lysosomes using the lgp-B antibody shown in Fig. 7. In ldlF cells, the lysosomes (red) are found at the tips of the cells (arrowheads), but these structures are still accessible to internalized LY (green) as indicated by the yellow color in the merge panel.

**Figure 10.** EGF does not reach lysosomes in ldlF cells. To follow the intracellular destination of a lysosomally targeted ligand, Texas red–conjugated EGF was internalized for 30 min after the cells were first preincubated for <12 h at 34°C or 40°C. The cells were then fixed and counter-stained for lysosomes using the same lgp-B antibody shown in Figs. 7 and 8. In wt cells at either temperature, the internalized EGF (red) reached lgp-positive structures (green). Overlap is yellow. In ldlF cells, there is very little overlap between the red and green channels at either temperature, indicating that very little of the internalized EGF reached lgp-positive structures. In ldlF cells at 40°C, the lysosomes (green) are found at the tips of the cells (arrowheads), but EGF (red) is not found in these structures.
of the ε>COP gene proved that the mutant phenotype was directly or indirectly because of the loss of ε>COP.

Our analysis of endocytosis in ldIF cells revealed that temperature-dependent loss of ε>COP resulted in a complex but interconnected series of temperature-sensitive and ε-insensitive phenotypes, each appearing to inhibit some aspect of early endosome function. The first of these was a dramatic inhibition of virus infection at 40°C. In the case of SFV, the use of a conformation-specific antibody to the E1/E2 spike glycoprotein revealed that at the restrictive temperature, the virus never reached an endosome compartment of suitably low pH to acquire the acid conformation required for fusion of the viral envelope with the endosome membrane. Yet, there appeared to be no global acidification defect, since acridine orange was accumulated normally by ldIF cells at all temperatures. Moreover, quantitative fluorescence analysis of the pH within FITC–Tfn-containing endosomes indicated again the absence of obvious differences between ldIF and wt cells at either 34° or 40°C. Although early endosomes have generally been assumed to be the site of penetration of viruses such as SFV (which fuses at pH < 6.1), our previous results noted the presence of an early endocytic compartment that does not have sufficiently low pH to support even SFV entry (Schmid et al., 1989). Conceivably, in ldIF cells, incoming virus cannot be efficiently transferred from this compartment (which may reflect early endosomes or primary endocytic vesicles such as coated vesicles) to a compartment of suitably low pH to host fusion. In any event, resolving the actual site of virus entry will require some further exploration.

Also consistent with a defect in early endosome function was the observation that ldIF cells exhibit a marked temperature-sensitive inhibition of Tfn recycling. It is becoming clear that the pathway of Tfn recycling may be somewhat more complex than originally conceived, involving the existence of two parallel pathways: the faster, predominant one corresponding to a short circuit pathway that returns receptors back to the cell surface directly from early endosomes, and a second, slower pathway, which involves passage through (and transient residence within) the perinuclear population of recycling endosomes (Mellman, I., and D. Sheff, manuscript in preparation). The data are consistent with an inhibition of the short circuit pathway in ldIF cells lacking ε>COP. Conceivably, a greater fraction of Tfn in ldIF cells at the restrictive temperature passes through the perinuclear recycling endosome population (Fig. 11).

Finally, we observed a defect in transport from early to late endocytic compartments such as late endosomes and lysosomes. It is well established that the lysosomal accumulation of extracellular macromolecules internalized as fluid-phase tracers or receptor-bound ligands depends on the ability of early endosomes to sort recycling receptors from the fluid content of the endosome lumen (Mellman, 1996). To some extent, this is a nonspecific, geometric phenomenon, but signal-mediated targeting events must also play a role particularly in polarized cells. Thus, the inability of HRP and EGF to sufficiently reach lysosomes is consistent with an inhibition of early endosomes in ldIF cells to properly sort membrane from content. The fact that anti-lgp-B antibodies failed to reach the peripheral lysosomal structures in ldIF cells at 40°C suggests that the cells may also be defective at targeting even resident membrane components to late endocytic compartments. This would be consistent with previous suggestions that late endosome-like structures cannot be generated from early endosomes in vitro in the absence of β>COP (Aniento et al., 1996).

How might the loss of ε>COP affect recycling, sorting, and transport to lysosomes? It seems unlikely that COPI coats are responsible for each of these events individually. However, given that early endosomal sorting activities must be completed within 1–2 min after internalization (Mellman, 1996), any loss of function affecting any one early endosome function would be expected to have rapid consequences for the activities of compartments both upstream and downstream. A similar situation may prevail in the early secretory pathway, where inhibition of COPI function might block ER to Golgi transport as a rapid consequence of an arrest of retrograde transport back from the cis-Golgi. It should be pointed out, however, that the precise site(s) of action of COPI coats in the secretory pathway remains controversial, with evidence suggesting its involvement not just in retrograde Golgi to ER transport, but also anterograde transport through the Golgi stack itself (Schekman and Mellman, 1997). In this context, it is useful to conceptualize the endocytic pathway as being somewhat similar to the secretory pathway in the organization of transport steps (Fig. 11). The rapid bidirectional traffic of membrane between the ER and the Golgi complex is thought to involve a rapid turn over of an intermediate compartment that may occupy a position analogous to that occupied by early endosomes between the plasma membrane and late endosomes and/or the perinuclear recycling endosomes. As such, any perturbation of early en-
dosome function due to the action of COPI can be seen as causing both proximal and distal effects.

An important unknown remains whether COPI acts directly or indirectly to affect endosome function. Upon shifting to 40°C, the onset kinetics of the endocytosis and recycling defects are very similar to the onset of the secretory defect as well as to the loss of immunodetectable e-COP. This is consistent with the idea, suggested by the physical association of COPI components with endosomes, that COPI plays a direct role in endosome function. However, it remains possible the effect is indirect, perhaps blocking the delivery of a Golgi-derived component. Such an indirect effect is unlikely to involve the delivery of a newly synthesized component since inhibition of protein synthesis is without any inhibition of endocytosis in most mammalian cell types (Steinman et al., 1983).

Thus far, the participation of COPI in the formation of small diameter coated vesicles, analogous to those believed to function during intercisternal Golgi transport, has not been unambiguously detected in endosomes, although small Tfn-containing vesicles associated with COPI have been detected by immunocytochemistry (Whitney et al., 1995). Entire COPI-coated regions on larger endosomal structures have also been observed. Conceivably, COPI coats act to selectively include cargo as is thought to occur in the secretory pathway, where KXXX motif proteins are thought to be sequestered in COPI-coated vesicles for retrograde transport from the cis-Golgi back to the ER. The KXXX motif appears to interact with the α, β, and ε-COP subunits (Cosson and Letourneur, 1994; Fiedler et al., 1996; Lowe and Kreis, 1996), and indeed there is evidence that KXXX-containing proteins such as the KDEL receptor can act to retrieve ligands internalized from the plasma membrane (Miesenbock and Rothman, 1995). It is also of interest that COPI complexes can interact with a second, more degenerate sequence motif containing an aromatic amino acid (Fiedler et al., 1996). Conceivably, this second motif, which is not restricted to ER proteins, is the motif decoded by endosomal COPI.

It has been suggested that the regulated assembly of COPI coats may act to control vesicular transport in ways that do not always involve the formation of discrete COPI-coated vesicles (Klausner et al., 1992). According to this view, long membrane tubules reminiscent of the tubular extensions characteristic of early endosomes may form when coats are not allowed to bind. In the Golgi complex and in endosomes this is most obvious upon the inhibition of COPI binding by the drug brefeldin A (Hunziker et al., 1991; Lippincott-Schwartz et al., 1991; Wood et al., 1991). At 40°C, IdlF cells do not exhibit a brefeldin-like phenotype, but this is probably because of the fact that the ε-COP-free COPI complex can still bind to membranes, perhaps inhibiting tubule formation in a dominant negative fashion (Gomez, M., and T. Kreis, unpublished data). Were COPI complexes to regulate the formation of endosomal tubules, an IdlF-like phenotype might be easily explained.

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