The association of epsin with ubiquitinated cargo along the endocytic pathway is negatively regulated by its interaction with clathrin

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Monoubiguitination of plasma membrane proteins is a mechanism to control their endocytic trafficking by promoting their interaction with cytosolic adaptor proteins that contain ubiquitin (Ub)-binding domains. Epsin, which contains Ub interaction motifs (UIMs), as well as binding sites for the clathrin coat and clathrin accessory factors, is thought to function as one of such adaptors. The importance of clathrin in the internalization of ubiguitinated cargo, however, has been questioned. Here, we show that a GFP-Ub chimera directly targeted to the plasma membrane via a lipidbased interaction is efficiently taken up by endocytosis and delivered to the same endosomes that accumulate internalized EGF. Internalization of the chimera requires integrity of the UIM binding interface of Ub, but does not require clathrin. Surprisingly, WT epsin showed little colocalization with this chimera, whereas UIM-containing epsin constructs that lack the clathrin and AP2 binding region, strikingly colocalized with this chimera on endocytic vacuoles. In addition, extensive colocalization of WT epsin with the chimera on endocytic structures could be observed in cells where clathrin levels were drastically reduced by RNA interference. Our results reveal an important regulatory mechanism in epsin function. The mutually exclusive colocalization of epsin with membrane-bound Ub or clathrin may play a role in controlling the endocytic route taken by ubiquitinated cargo.

ubiquitin | endocytosis | Rab5 | endosome | FYVE domain

rotein ubiquitination plays pleotropic roles in cell physiology. Polyubiquitination of cytosolic proteins directs them for destruction by the proteasome. Reversible monoubiquitination regulates protein location, function, and interactions (1-3). In the case of membrane proteins, monoubiquitination triggers their interaction with adaptor proteins that control their intracellular traffic. More specifically, monoubiquitination of plasma membrane (PM) proteins tags them for internalization and subsequent sorting to late endosomes/lysosomes for degradation (2, 4). PM-associated ubiquitin (Ub) is recognized by endocytic adaptors, among which epsin has been extensively characterized (5-11). Epsin has a modular structure comprising an epsin N-terminal homology (ENTH) domain, which binds phosphoinositides, Ub interaction motifs (UIMs), and a long flexible region that includes binding sites for clathrin heavy chain, the clathrin adaptor AP-2, and EH domain-containing proteins such as Eps15, another UIM containing endocytic adaptor (8, 12–14). The UIM region of epsin, in addition to mediating binding to Ub, is required for the ubiquitination of epsin itself (6). This reaction, which occurs constitutively and is further stimulated by growth factors (6), inhibits the property of epsin to interact with the clathrin coat and the membrane (15). The importance of Ub metabolism in epsin function is further supported by genetic studies in Drosophila (16-19). The presence in epsin of binding sites for both Ub and core components of the clathrin coat, together with the localization of epsin at clathrin-coated pits, suggested that a function of this protein is to recruit ubiquitinated cargo to clathrin-coated pits, thus triggering its internalization (13). Surprisingly, however, the endocytosis of a chimeric protein composed of the extracellular and transmembrane domains of the EGF receptor fused to a cytoplasmically exposed Ub does not require clathrin, questioning the role of clathrin-mediated budding in the internalization of ubiquitinated cargo (20).

Here, we used a GFP-fusion protein of Ub that is directly targeted to the PM via lipid-based interactions, to further elucidate mechanisms in epsin-mediated sorting of PMassociated Ub.

Methods

Antibodies, Constructs, and Reagents. Antibody directed against epsin 1 was generated in our laboratory as described (5). Mouse mAbs against clathrin heavy chain, GFP, the hemagglutinin (HA) epitope, the Xpress epitope, and the Flag epitope were from Affinity BioReagents (Golden, CO), Clontech, Roche Applied Science, Invitrogen, and Sigma, respectively. Human antibody directed against EEA1 was a kind gift of Harald Stenmark (Norwegian Radium Hospital, Oslo). Rabbit antibody directed against Rabankyrin-5 was a kind gift of Marino Zerial (Max Planck Institute of Molecular Cell Biology and Genetics, Dresden, Germany). Xpress-tagged full-length epsin 1 and HAtagged amphiphysin A1 fragment have been described (5, 21). PM-GFP-Ub was made by PCR using PM-GFP in a pcDNA3 vector (a kind gift of Tobias Meyer, Stanford University, Stanford, CA) as a template, followed by subcloning in a pcDNA3-HA-Ub vector as described (15) except that the HA tag was removed in the final construct. PM-GFP-Ub^{∆GG} was generated by replacement of Ub in PM-GFP-Ub with a mutant Ub fragment obtained by PCR using a reverse primer that lacks the last two glycine residues in Ub cDNA sequence. GFP-Ub $^{\Delta GG}$ was generated by subcloning the mutant Ub lacking the last two glycine residues in pEGFP-C2 vector (Clontech). PM-GFP-Ub $^{\Delta GG-K48R}$ and PM-GFP-Ub $^{\Delta GG-144A}$ double mutants were obtained from PM-GFP-Ub $^{\Delta GG}$ by using the QuikChange Site-Directed Mutagenesis Kit (Stratagene) according to the manufacturer's instruction with primers harboring the corresponding mutation sequences. GFP-FYVE(Hrs)-Ub^{∆GG} was constructed by subcloning the mutant Ub lacking the last two glycine residues in a pEGFP-FYVE vector (a kind gift from Harald Stenmark). GFP-PH(FAPP1)-Ub was generated by subcloning Ub in a pEGFP-PH vector (a gift from Dario Alessi, University of Dundee, Dundee, Scotland). A mutant epsin 1 lacking the clathrin and AP-2 binding region was generated by ligating together two pieces of the rat epsin 1 cDNA sequence (5)

Abbreviations: Ub, ubiquitin; UIM, Ub interaction motif; HA, hemagglutinin; PM, plasma membrane; RNAi, RNA interference; siRNA, small interfering RNA; ENTH, epsin N-terminal homology.

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corresponding to amino acids 1–254 and 401–576. Blunt-end ligation was followed by subcloning in a pcDNA3-Flag vector. The ENTH-UIM construct, corresponding to amino acids 1–254 of epsin 1, was generated by PCR followed by subcloning in pcDNA3-Flag vector. The UIM construct, encoding amino acids 181–254 of epsin 1, was obtained by PCR followed by subcloning in pcDNA3-HA vector. All constructs were confirmed by DNA sequencing. Rhodamine-conjugated EGF and Alexa Fluor 594-conjugated dextran were purchased from Molecular Probes.

Cell Culture and cDNA Transfection. HeLa or CHO cells were grown at 37°C and 5% CO₂ in DMEM with 10% (vol/vol) FBS, 10 mM Hepes, 1 mM sodium pyruvate, 100 units/ml penicillin, and 100 μ g/ml streptomycin. cDNA transfection was performed with Lipofectamine (Invitrogen). Twenty-four hours later, cells were fixed in 4% formaldehyde and processed for immunofluorescence. In some experiments, cells were incubated with rhodamine-conjugated EGF or Alexa Fluor 594-conjugated dextrin for 20 min followed by rapid wash in PBS three times before fixation.

Clathrin Heavy Chain RNA Interference (RNAi). Small interfering RNA (siRNA) pairs corresponding to the sequences GAAA-GAATCTGTAGAGAAA and GCAATGAGCTGTTT-GAAGA of clathrin heavy chain (22, 23) were synthesized and purchased from Integrated DNA Technologies (Coralville, IA). HeLa cells were transfected with 50 nM each of the two siRNA pairs or with 50 nM of a control siRNA pair corresponding to nucleotides 695–715 of the firefly luciferase (U31240) (AA-GAATATTGTTGCACGATTT), using Oligofectamine (In-vitrogen). Three days later, cells were transfected with PM-GFP-Ub^{AGG} or Xpress-tagged epsin 1 cDNA in a pcDNA3 vector with Lipofectamine and processed for immunofluorescence studies.

Miscellaneous Procedures. SDS/PAGE, Western blotting, and immunoprecipitation were performed according to standard procedures. Immunofluorescence microscopy was performed with an Axioplan 2FS microscope (Zeiss) using $\times 63$ oilimmersion objectives. Images were acquired with a chargecoupled device camera (ORAC II, Hamamatsu, Middlesex, NJ) and analyzed with METAMORPH software (Universal Imaging, Downingtown, PA). Confocal images were obtained with a Zeiss 510 laser scanning confocal microscope.

Results

Monoubiquitin Targeted to the PM by a Lipid Anchor Is Efficiently Internalized. To determine mechanisms in the internalization of PM-bound Ub we generated a reporter construct (PM-GFP-Ub) represented by Ub fused in-frame to the COOH terminus of PM-targeted GFP (PM-GFP). PM-GFP is a modified form of GFP resulting from the addition to its NH2 terminus of the short amino acid sequence (10 aa) that directs the myristoylation and palmitoylation, and therefore PM targeting of Lyn kinase (24). We used a lipid-bound Ub chimera rather than Ub fused to a transmembrane protein to avoid potential effects caused by traffic of the chimera through biosynthetic compartments. As expected, when expressed in HeLa or CHO cells, PM-GFP accumulated selectively at the PM, consistent with a direct targeting to this membrane independent of the secretory pathway (Fig. 1A and data not shown). In contrast, PM-GFP-Ub also accumulated on internal vacuolar structures (Fig. 1A).

Anti-GFP Western blots of anti-GFP immunoprecipitates from cells transfected with PM-GFP-Ub revealed a smear characteristic of polyubiquitin-containing conjugates, in addition to the PM-GFP-Ub band, suggesting that this construct participates in the formation of polyubiquitin chains (Fig. 2). This smear was absent when WT Ub in PM-GFP-Ub was replaced by a mutant Ub that lacks the last two glycine residues (Ub^{Δ GG}). Ub^{Δ GG} cannot be conjugated to lysine residues of



Fig. 1. PM-GFP is selectively localized at the PM, whereas PM-GFP-Ub and PM-GFP-Ub^{ΔGG} are internalized and also accumulate on intracellular endocytic compartments. (A) Confocal microscopy images of HeLa cells. (B) HeLa cells transfected with PM-GFP-Ub^{ΔGG} were incubated with or without Alexa Fluor 594-conjugated dextran or rhodamine-conjugated EGF, then fixed and either processed by immunofluorescence for the endosomal proteins EEA1 and rabankyrin or examined directly for the internalized dye-conjugated endocytic probes. (C) Epifluorescence microscopy of HeLa cells transfected with the constructs indicated. Note the diffuse cytosolic localization of GFP-Ub^{ΔGG} that lacks the short amino acid sequence directing acylation and PM localization. (Bars: 12 μ m.)

ubiquitinated substrates because this conjugation requires a COOH-terminal glycine (25). Similar results were obtained with an Ub mutant that contained an additional lysine-to-alanine



Fig. 2. PM-GFP-Ub, but not PM-GFP-Ub^{Δ GG}, is incorporated into polyubiquitin chains. Anti-GFP immunoprecipitates from detergent extracts of untransfected cells and cells transfected with GFP constructs were processed by Western blotting for GFP immunoreactivity. IP, immunoprecipitation; WB, Western blot.

substitution at position 48, a major acceptor site for the COOHterminal glycine of another Ub (25). Like PM-GFP-Ub, both PM-GFP-Ub^{Δ GG} and PM-GFP-Ub^{Δ GG-K48R} were targeted to internal vacuoles (Fig. 1*A* and data not shown). PM-GFP-Ub^{Δ GG} was used for most subsequent experiments.

Extracellular applications of Alexa Fluor 594-conjugated dextran (100 μ g/ml for 20 min), a hydrophilic polysaccharide that does not cross the bilayer, and rhodamine-conjugated EGF (40 ng/ml for 20 min) resulted in the accumulation of these probes in the PM-GFP-Ub ΔGG -positive vacuoles, proving their endocytic nature. Furthermore, partial overlap (with some variability from cells to cells) was observed between these vacuoles and the endosomal proteins EEA1 and Rabankyrin-5, two Rab5 effectors (26, 27) (Fig. 1B). The large size of these endocytic structures may be related to the accumulation of the exogenous protein. To confirm that PM-GFP-Ub ΔGG found on endocytic vacuolar structures is not recruited directly to these membranes via endosomal-associated Ub-binding proteins, a GFP-Ub $^{\Delta GG}$ chimera was used as a control. This chimera, which lacks the sequence directing its acylation, and therefore PM targeting, had a diffuse localization in the cytosol (Fig. 1C).

Interactions with Endocytic Adaptors, but Not Clathrin, Are Required for the Endocytosis of Membrane-Targeted Ub. Studies in yeast and cells of higher eukaryotes have demonstrated that isoleucine 44 of Ub plays a critical role in endocytosis (28, 29), and biochemical and structural studies have confirmed the important role of this residue in the binding of Ub to the UIM motifs of endocytic adaptors (28–30). Accordingly, a PM-GFP-Ub chimera harboring an isoleucine-to-alanine substitution at position 44 (PM-GFP-Ub Δ GG-I44A) accumulated selectively at the PM and failed to be internalized (Fig. 3), thus implicating adaptors comprising UIMs or other Ub-binding domains in its endocytosis.

An EGF receptor-Ub chimera can be internalized in a clathrin-independent way (20). In agreement with this finding, disruption of clathrin function by siRNA or dominant negative interference did not block the internalization of PM-GFP-Ub^{Δ GG}. Thus, addition to HeLa cells of two different clathrin



Fig. 3. The internalization of PM-GFP-Ub^{Δ GG} is abolished by the substitution of isoleucine 44 with alanine as examined by epifluorescence microscopy of HeLa cells. (Magnification: \times 500.)

heavy chain-specific siRNA pairs, which alone were sufficient to nearly deplete clathrin, as shown by Western blotting (Fig. 4A) and immunofluorescence (Fig. 4B), blocked the internalization of transferrin as expected (data not shown), but did not prevent the accumulation of PM-GFP-Ub ΔGG on endosomal vacuoles (Fig. 4B). In contrast, PM-GFP-Ub^{Δ GG} still accumulated on endosomes in cells treated with a control siRNA pair (Fig. 4B). Furthermore, transfection of the fragment of amphiphysin 1 that comprises the binding sites for clathrin and the endocytic clathrin adaptor AP-2 (fragment A1) did not prevent accumulation of PM-GFP- $Ub^{\Delta GG}$ on endosomes (data not shown). This construct has a powerful inhibitory effect on clathrin-mediated endocytosis because it titrates out available clathrin and AP-2 (21). At variance with results obtained with the EGF receptor-Ub chimera (20), however, dynamin, a GTPase that mediates the fission steps in a variety of endocytic reactions (31), was also not required for the endocytosis of PM-GFP-Ub $^{\Delta GG}$. This reporter protein was found on endosomes in the presence of dynamin^{K44A} (32), a mutant dynamin that has a dominant negative effect on the function of endogenous dynamin, or in HeLa cells in which dynamin 2 had been knocked down by RNAi (data not shown).

Clathrin Affects the Colocalization of Epsin with Membrane-Targeted Ub. Given the putative role of epsin as an endocytic adaptor for ubiquitinated cargo, we investigated whether epsin colocalizes, at least partially, with PM-GFP-Ub or PM-GFP-Ub^{ΔGG}. To this aim, we cotransfected cells with Xpress-tagged epsin 1 (5) and



Fig. 4. Endocytosis of PM-GFP-Ub does not depend on clathrin. (*A*) Anticlathrin Western blot of extracts from cells treated with control siRNA and two distinct sets of clathrin siRNA duplexes. (*B*) HeLa cells treated with control siRNA or a mixture of the two siRNA duplexes used for *A* were transfected with PM-GFP-Ub^{Δ GG}, then fixed and analyzed for GFP fluorescence and clathrin immunoreactivity. (Bar: 12 μ m.)



Fig. 5. Full-length epsin 1 shows little colocalization with GFP-Ub constructs. CHO cells (*Top*) or HeLa cells (*Middle* and *Bottom*) cotransfected with epitopetagged full-length epsin 1 and GFP-Ub constructs were fixed and examined for GFP fluorescence and epitope tag immunoreactivity by epifluorescence microscopy. Note the typical punctate pattern of epsin, reflecting clathrincoated pits localization (5), regardless of the subcellular localization of GFP-Ub constructs. (Magnification: ×750.)

either PM-GFP-Ub or PM-GFP-Ub $^{\Delta GG}$. Epsin immunoreactivity showed little colocalization with either chimera, but displayed primarily the typical, previously reported punctate pattern characteristic of clathrin-coated pits (5, 11) (Fig. 5). This result may reflect the occurrence of regulatory mechanisms in the binding of epsin's UIMs to Ub or the inefficient interaction of epsin with Ub in the context of the cell cytosol.

To address these possibilities, fragments of epsin containing the UIMs, including a construct represented by the UIMs alone (Fig. 6), were coexpressed in HeLa cells together with PM-GFP-Ub^{ΔGG} or other Ub-containing GFP fusion proteins targeted to specific intracellular compartments by lipid-binding domains. Two tandem-arranged FYVE domains of the late endosomal protein Hrs (33), which binds PI(3)P, were used to target Ub to endosomes [GFP-FYVE(Hrs)-Ub^{ΔGG}], and the PH domain of FAPP1, which binds PI(4)P, was used to target Ub to the Golgi complex [GFP-PH(FAPP1)-Ub] (34, 35). Both of these fusion proteins behaved as expected, although they partially disrupted the morphology of the two organelles, possibly as a result of interactions, or functional interference, with the many intracellular Ub-binding proteins (36–38). GFP-FYVE(Hrs)-Ub^{ΔGG} accumulated on vacuolar endosome-like structures (often en-



Fig. 6. Domain cartoon of epsin and epsin constructs used in this study. Interacting regions for binding partners are indicated by arrows.



Fig. 7. Fragments of epsin containing the UIM (*A*) or the ENTH-UIM (*B*) region colocalize extensively with internalized PM-GFP-Ub^{Δ GG}. HeLa cells cotransfected with GFP-Ub constructs and epitope-tagged UIMs or ENTH-UIMs were fixed and examined for GFP fluorescence and epitope tag immunore-activity. Note colocalization of UIMs and ENTH-UIMs with GFP-Ub constructs irrespective of their subcellular localization. (Magnification: ×800.)

larged and/or clustered), whereas GFP-PH(FAPP1)-Ub accumulated on perinuclear particles and, at high levels of expression, disrupted the Golgi complex (Fig. 7*A*).

Coexpressed full-length epsin 1 did not colocalize with either FYVE(Hrs)- or PH(FAPP1)-containing constructs, and in all cases had the typical distribution of endocytic clathrin-coated pits (Fig. 5 and data not shown). In contrast, an epitope-tagged epsin 1 fragment composed only of its UIM domains (Fig. 6) strongly colocalized with membrane-targeted Ub fusion proteins irrespective of their localization (Fig. 7A). It colocalized with PM-GFP-Ub^{Δ GG} and GFP-FYVE(Hrs)-Ub^{Δ GG} on intracellular vacuoles and with GFP-PH(FAPP1)-Ub on the Golgi remnants observed in cells transfected with this domain (Fig. 7A). Thus, epsin's UIMs do interact efficiently with Ub and monoubiquitin in the cell cytosol. An epsin 1 construct including both the ENTH domain and the UIMs (Fig. 6) also colocalized with PM-GFP-Ub^{Δ GG} and GFP-FYVE(Hrs)-Ub^{Δ GG}, but not with GFP-PH(FAPP1)-Ub (Fig. 7B and data not shown), indicating that the ENTH domain contributes specificity to UIM binding.



Fig. 8. An epitope-tagged epsin mutant (Epsin^{Δ clathrin}) lacking the clathrin and AP-2 binding region colocalized with PM-GFP-Ub^{Δ GG} and GFP-FYVE(Hrs)-Ub^{Δ GG} on endocytic compartments. Cotransfected CHO cells (*Top*) or HeLa cells (*Middle* and *Bottom*) were fixed and examined for GFP fluorescence and epitope tag immunoreactivity. (Magnification: ×750.)

More important, even full-length epsin 1 with an internal deletion, i.e., deletion of the clathrin and AP-2 binding region (epsin^{$\Delta clathrin$}), a predicted unfolded region (Fig. 6), colocalized with PM-GFP-Ub^{ΔGG} and GFP-FYVE(Hrs)-Ub^{ΔGG} (Fig. 8). It would therefore appear that the presence of binding sites for core components of the clathrin coat is what prevents the colocalization of epsin with Ub-containing reporter fusions localized on endocytic membranes.

To determine whether this finding resulted from a masking effect of the clathrin-AP-2 binding region of epsin on its UIM domain region or an effect of the binding of clathrin to epsin, the localization of Xpress-tagged epsin was investigated in HeLa cells expressing PM-GFP-Ub^{Δ GG} after RNAi-mediated clathrin knockdown (see Fig. 4). Strikingly, in these cells, a significant fraction of WT epsin colocalized with PM-GFP-Ub^{Δ GG} on internal vacuoles (Fig. 9). Thus, binding of epsin to the clathrin coat either prevents its binding to Ub or makes this interaction very transient.

Discussion

Our results demonstrate the importance of Ub as an internalization signal at the PM (2) and of epsin in the recognition of this signal (2, 8, 13). In addition, a key conclusion of our study is that the interaction of epsin with Ub is critically regulated by the interaction of epsin with clathrin.

Epsin contains a multiplicity of binding sites, including binding sites for phosphoinositides (39), a transcription factor (40), Ub (6, 14), clathrin (41, 42), the clathrin adaptor AP-2 (5), and other signaling (43) and endocytic proteins (5, 44, 45), which implies the occurrence of regulatory mechanisms to control these interactions. As we have shown previously, some of the interactions of epsin are regulated by its phosphorylation and reversible ubiquitination (15, 46). Here, we report evidence for an additional control mechanism based on its binding to clathrin. Full-length epsin has the typical, previously described (5, 11) clathrin-coated pit localization even in cells expressing a PMtargeted, Ub-containing reporter protein. In contrast, an epsin construct lacking the clathrin and AP-2 binding region (epsin^{Δ clathrin}) shows a striking colocalization with this reporter protein throughout the endocytic pathway and also colocalizes



Fig. 9. A pool of full-length epsin colocalizes with PM-GFP-Ub^{Δ GG} on intracellular endocytic vacuoles after RNAi-mediated clathrin knockdown. HeLa cells pretreated with control siRNA or clathrin heavy chain-specific siRNA duplexes were transfected with epitope-tagged full-length epsin 1 construct and PM-GFP-Ub^{Δ GG}. Cells were then fixed and analyzed for GFP fluorescence and epitope tag immunoreactivity. (Magnification: ×1,000.)

with a Ub-reporter protein targeted directly to endosomes. This colocalization is observed for full-length epsin 1 as well, but only in cells where clathrin levels were drastically decreased by RNAi-mediated knockdown. It therefore would appear that the recruitment of epsin to clathrin coats promotes its dissociation from, or prevents its binding to, ubiquitinated cargo. The ubiquitination of epsin itself prevents its interaction with clathrin and AP-2, possibly as a result of an intramolecular interaction (15). It remains to be seen whether this "cis" interaction with an intramolecular Ub occludes clathrin binding in a way mechanistically similar to "trans" binding to another ubiquitinated protein. We note that our present results may not necessarily imply a direct, mutually exclusive interaction of epsin with either ubiquitinated cargo or clathrin, a possibility that, because of technical limitations, we could not test satisfactorily with in vitro biochemical experiments.

Clearly, as shown by Sigismund *et al.* (20) and our present results, ubiquitinated PM proteins can be internalized in a clathrin-independent way. The simple explanation is that epsin may function as an endocytic adaptor in two completely distinct internalization pathways, one clathrin-dependent but Ubindependent, and the other clathrin-independent but Ubdependent. However, this explanation seems unlikely, given the striking effect on epsin's localization of its ability to interact with clathrin in the context of the cell cytoplasm. Furthermore, binding sites for both Ub and clathrin are conserved in epsin from yeast to mammals. Thus, an interplay between Ub binding

and clathrin binding appears more plausible. A pool of epsin bound to a ubiquitinated cargo may be recruited to clathrincoated pits, for example, by EH domain endocytic proteins, but then dissociate from Ub, thus allowing other interactions of such cargo with coat components. Interestingly, a recent study demonstrated that epsin is critically needed for the function of the Notch receptor Delta (18, 19), and that this function involves not only the ubiquitination of Delta and its endocytosis, but also the targeting of internalized Delta to a specific endosomal subcompartment (19). Perhaps, epsin functions to recruit to clathrincoated pits and a recycling endosomal compartment, a pool of ubiquitinated Delta that is otherwise internalized and targeted to lysosomes through a clathrin-independent pathway. Alternatively, epsin may participate in coated-pit assembly (11), and the subsequent ubiquitination of cargo accumulated at coated pits may induce epsin separation form the coat. This possibility is

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consistent with the lack of enrichment of epsin in purified clathrin-coated vesicle fractions (5). It is also possible that the interplay between clathrin and Ub may occur on the surface of endosomes, at the specialized clathrin coat that has been described on these structure. This coat is thought to play a role in the progression of membrane cargo into luminal vesicles of multivesicular bodies (47, 48). Normally epsin is not detectable at these sites, but a small undetectable pool of epsin on endosomes may be greatly amplified if epsin cannot interact with this clathrin coat.

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