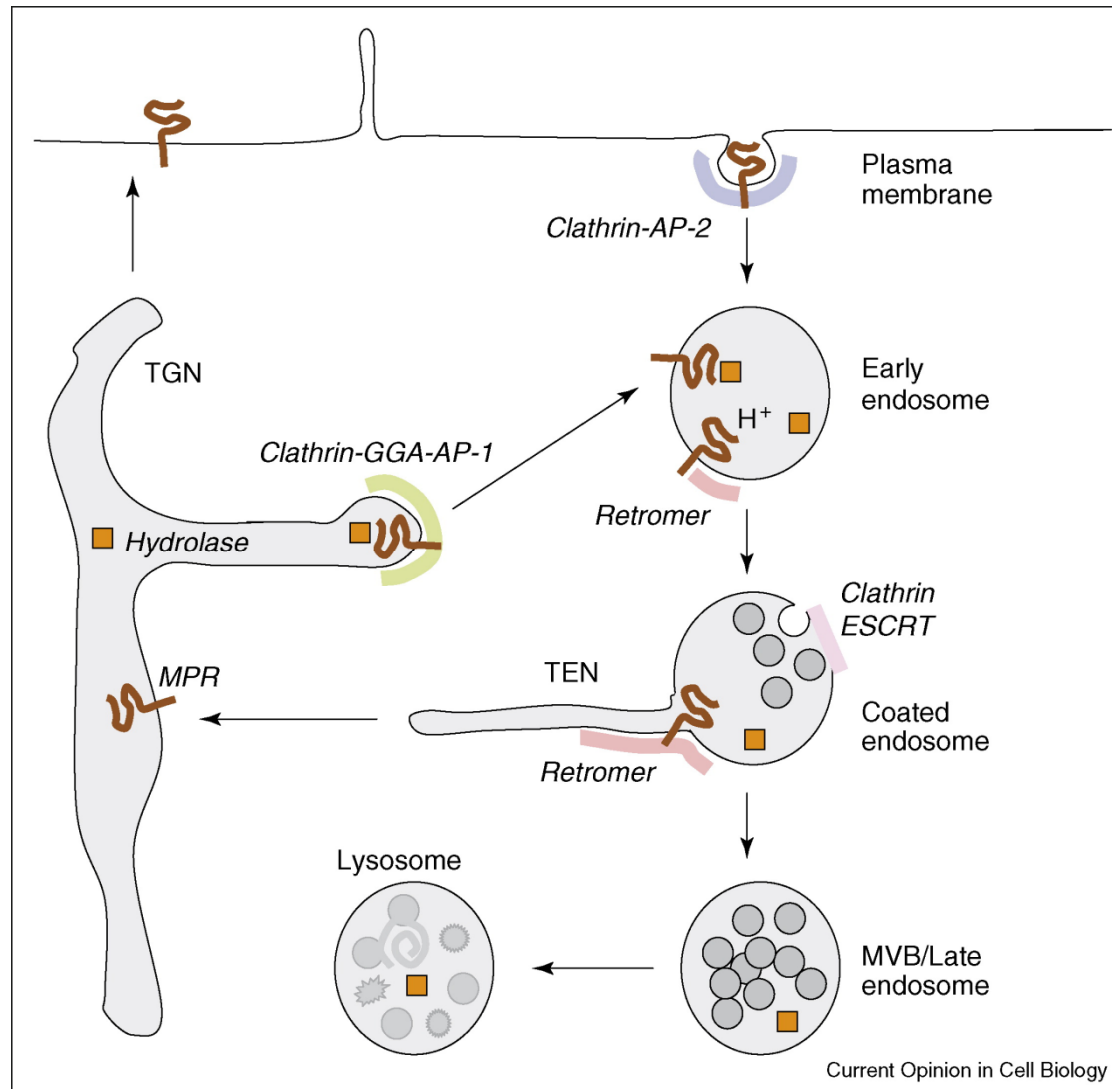
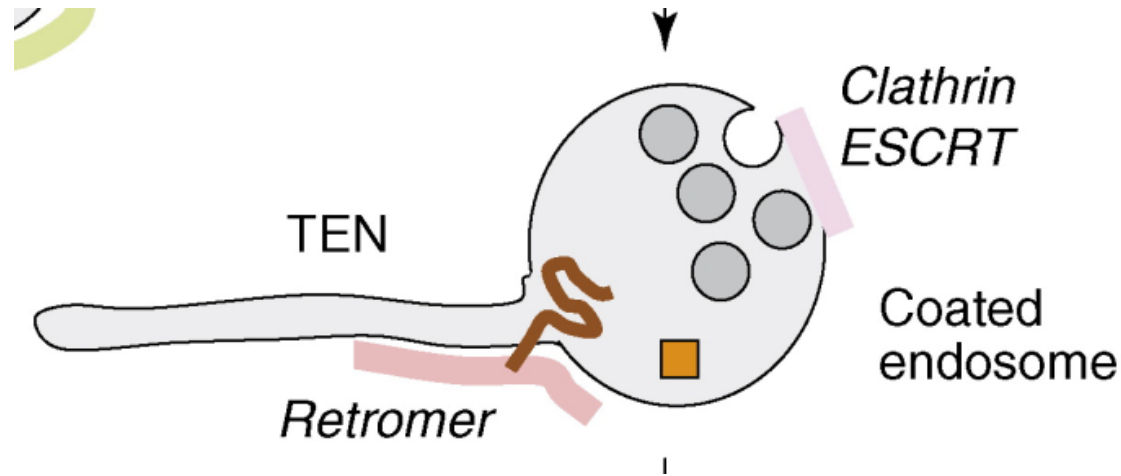


Retromer and ESCRTs

Coats of the late endosomal system: Retromer and ESCRT



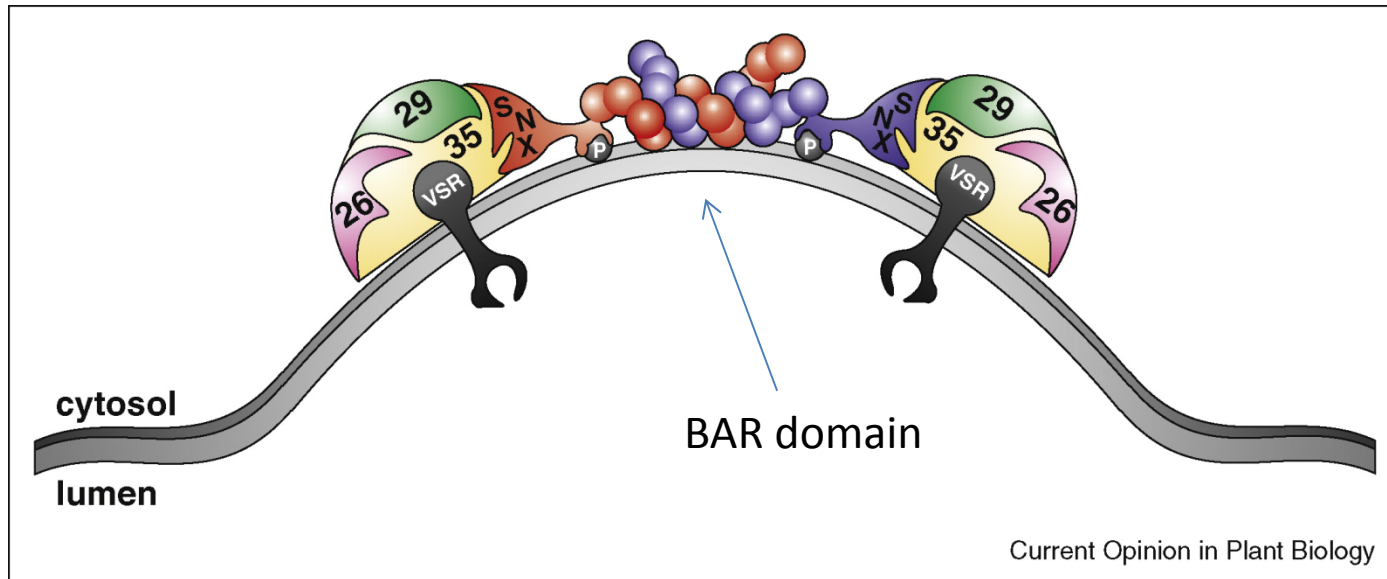
Pause to consider



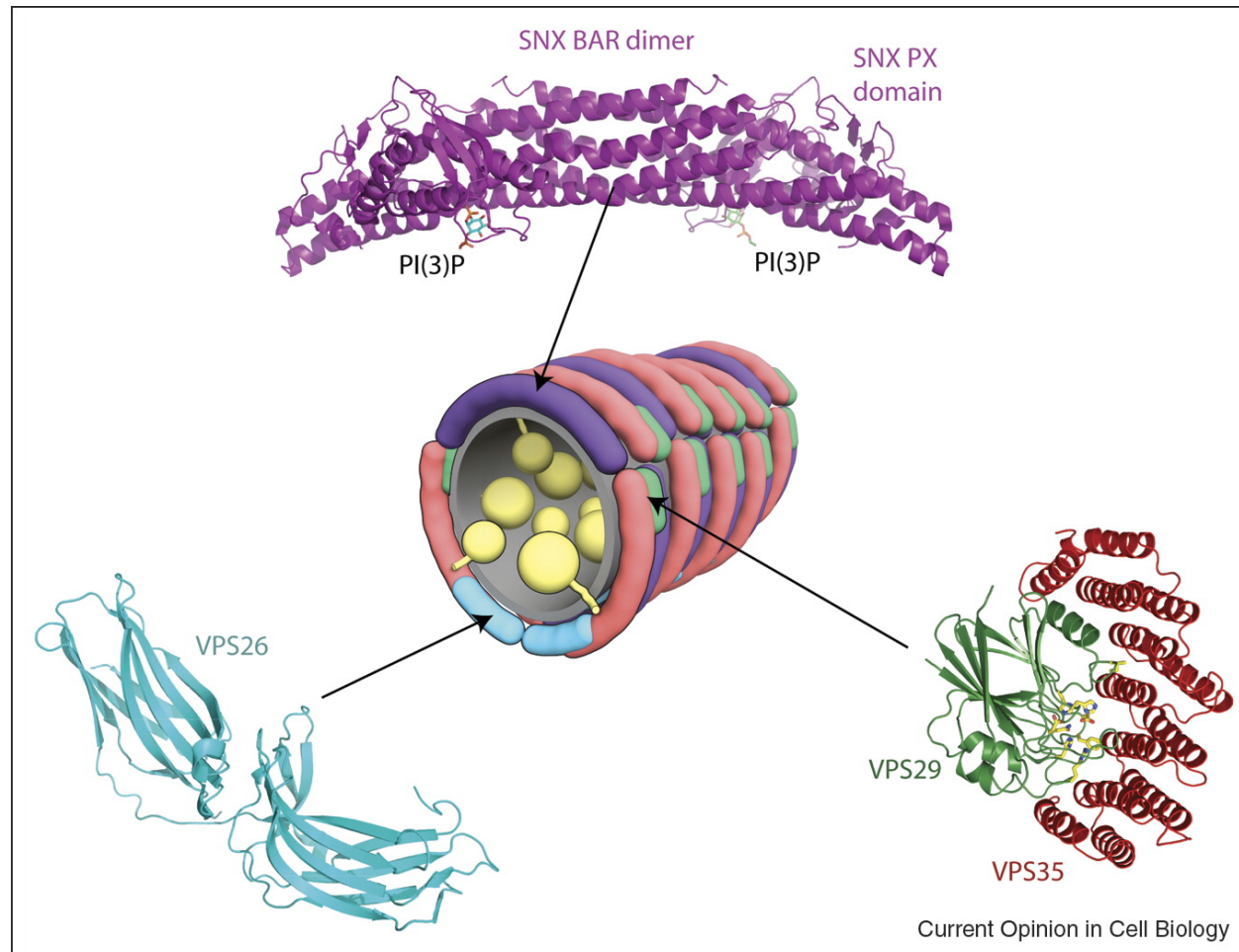
How do these two traffic pathways differ from ones we've seen before?

What are the architectural challenges for MVB formation?
for retromer tubules?

Retromer schematic

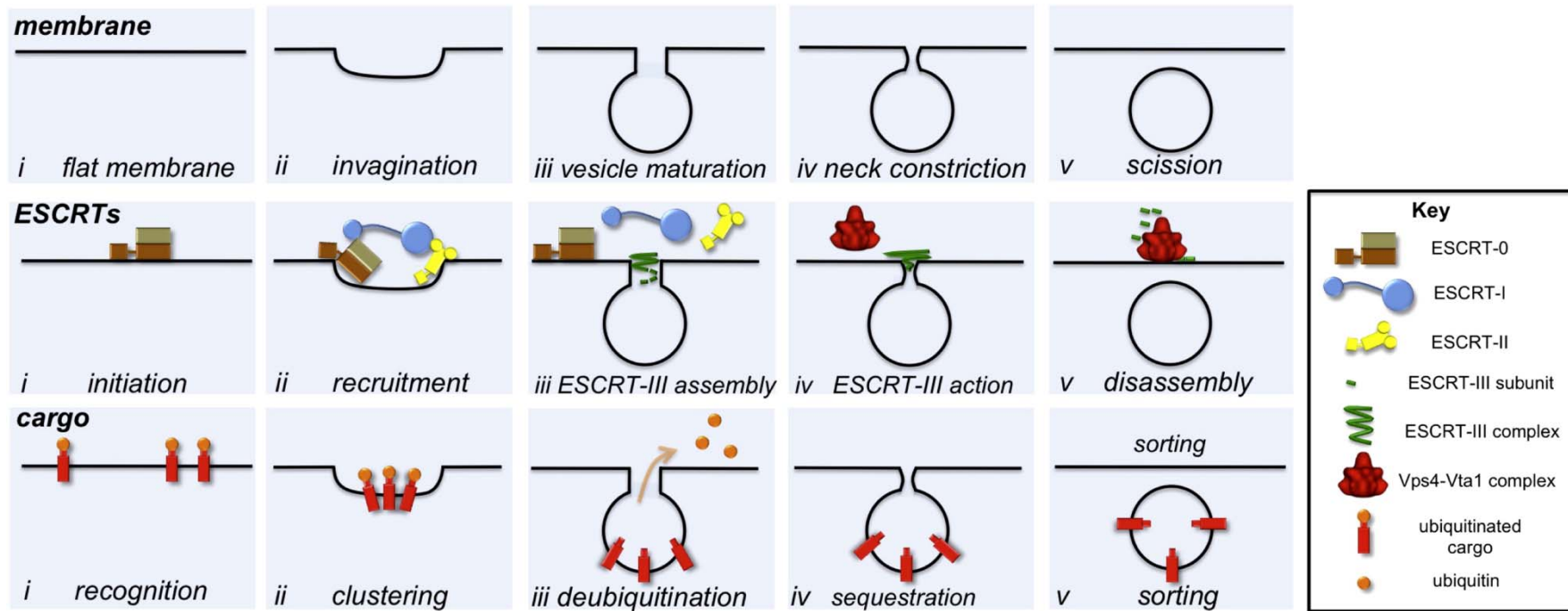


Retromer Schematic

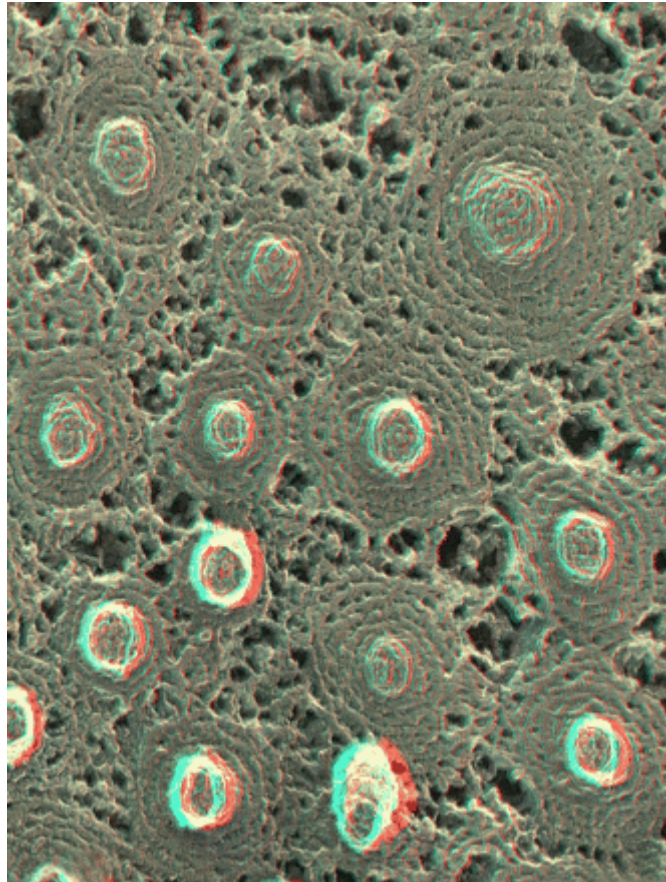


Bonifacino & Hurley, 2008

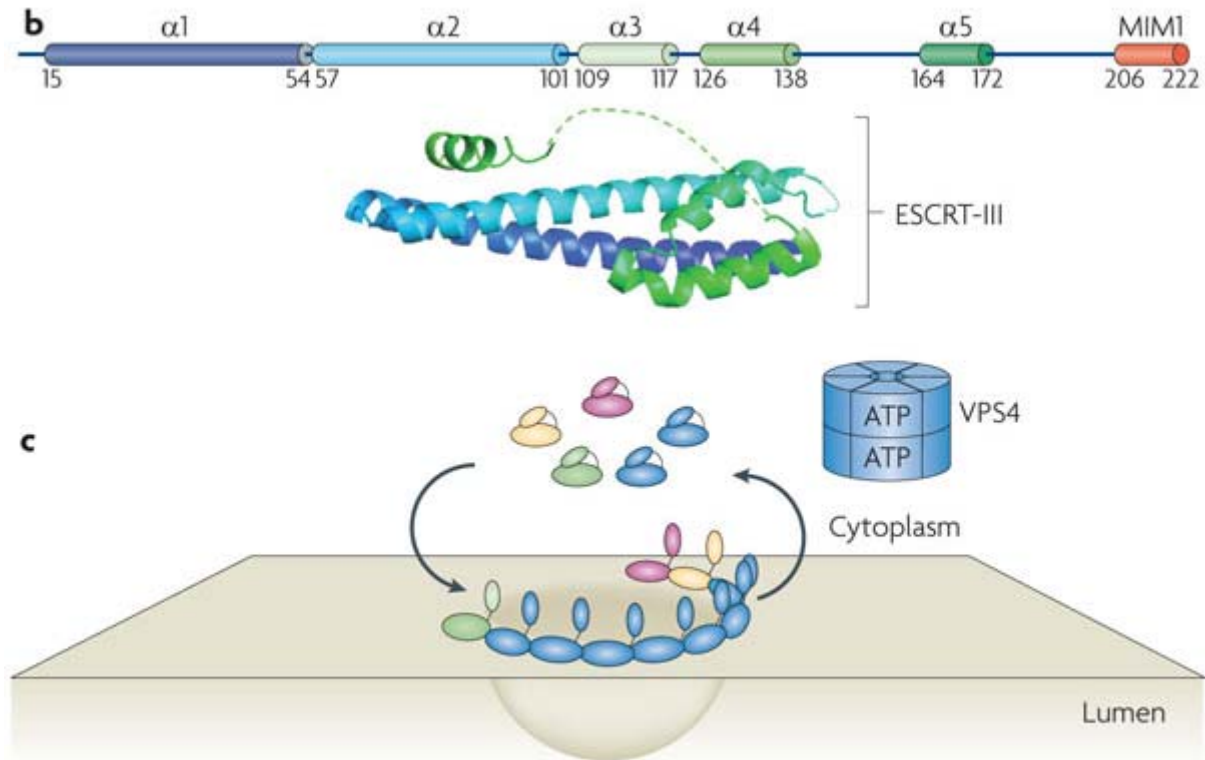
ESCRT



Top view of assembled ESCRTIII



ESCRT schematic



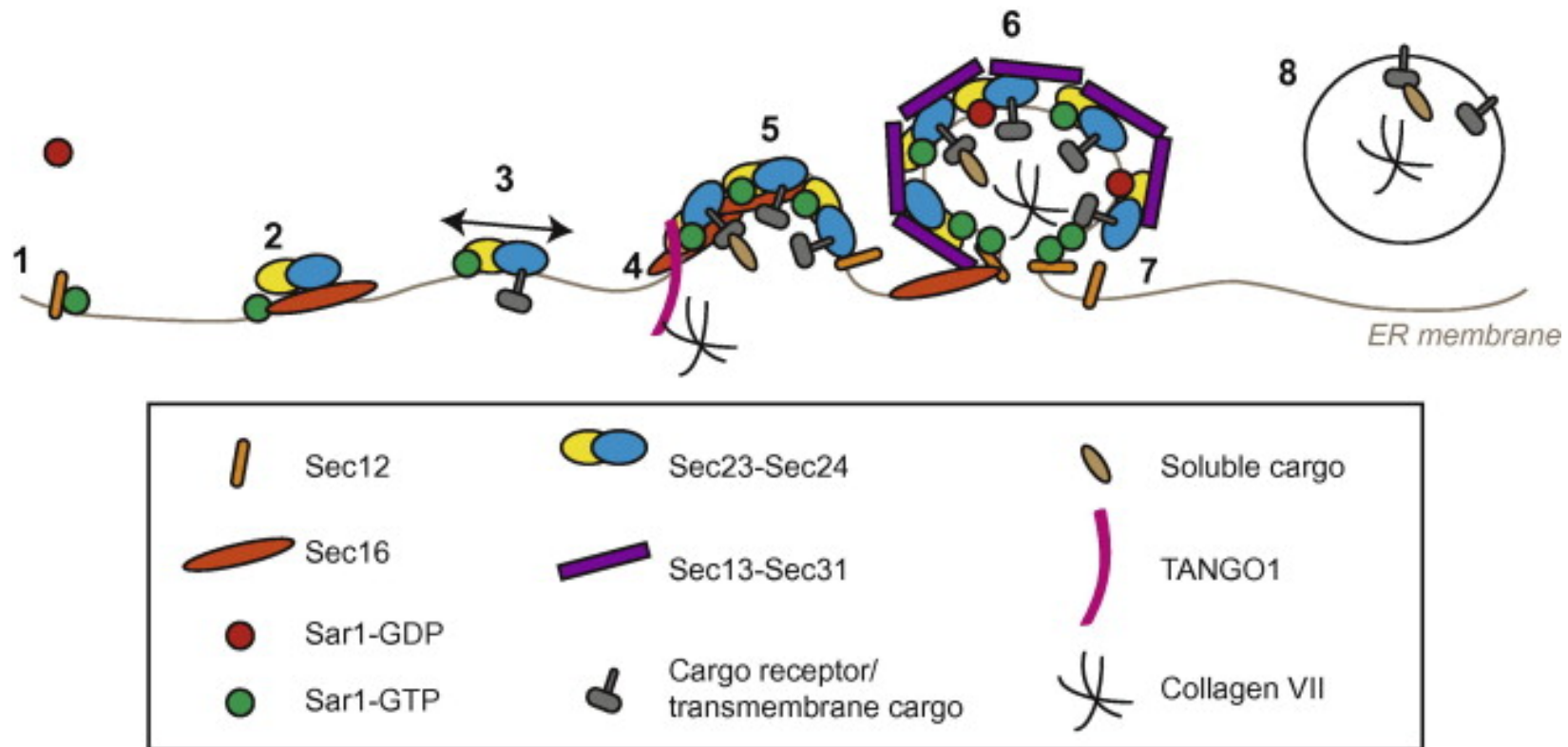
Nature Reviews | Molecular Cell Biology

Coat recruitment

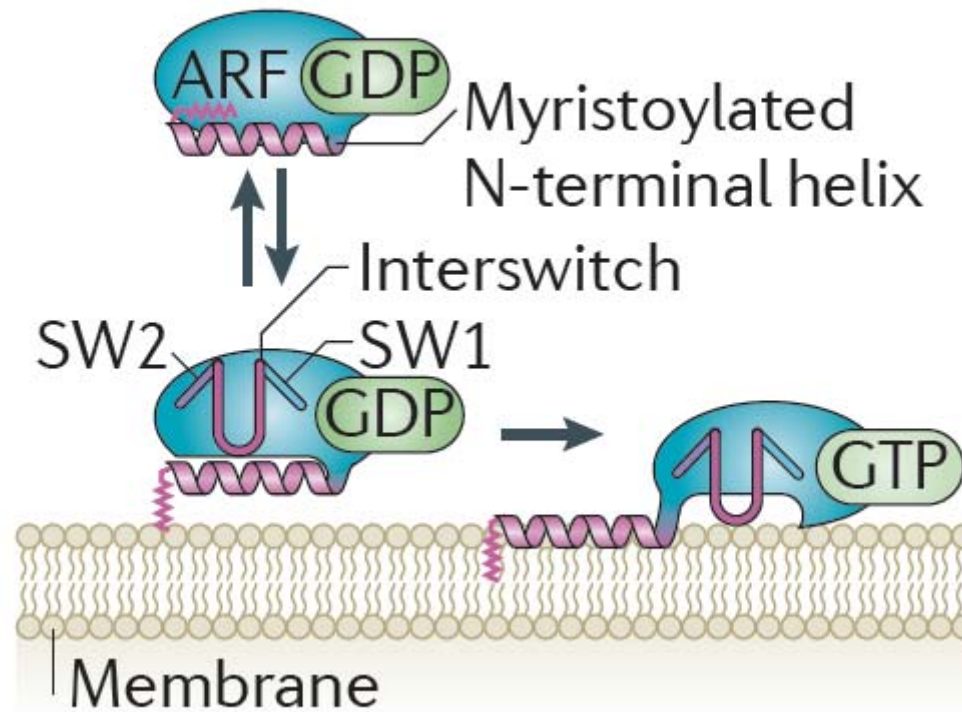
How are coats recruited?



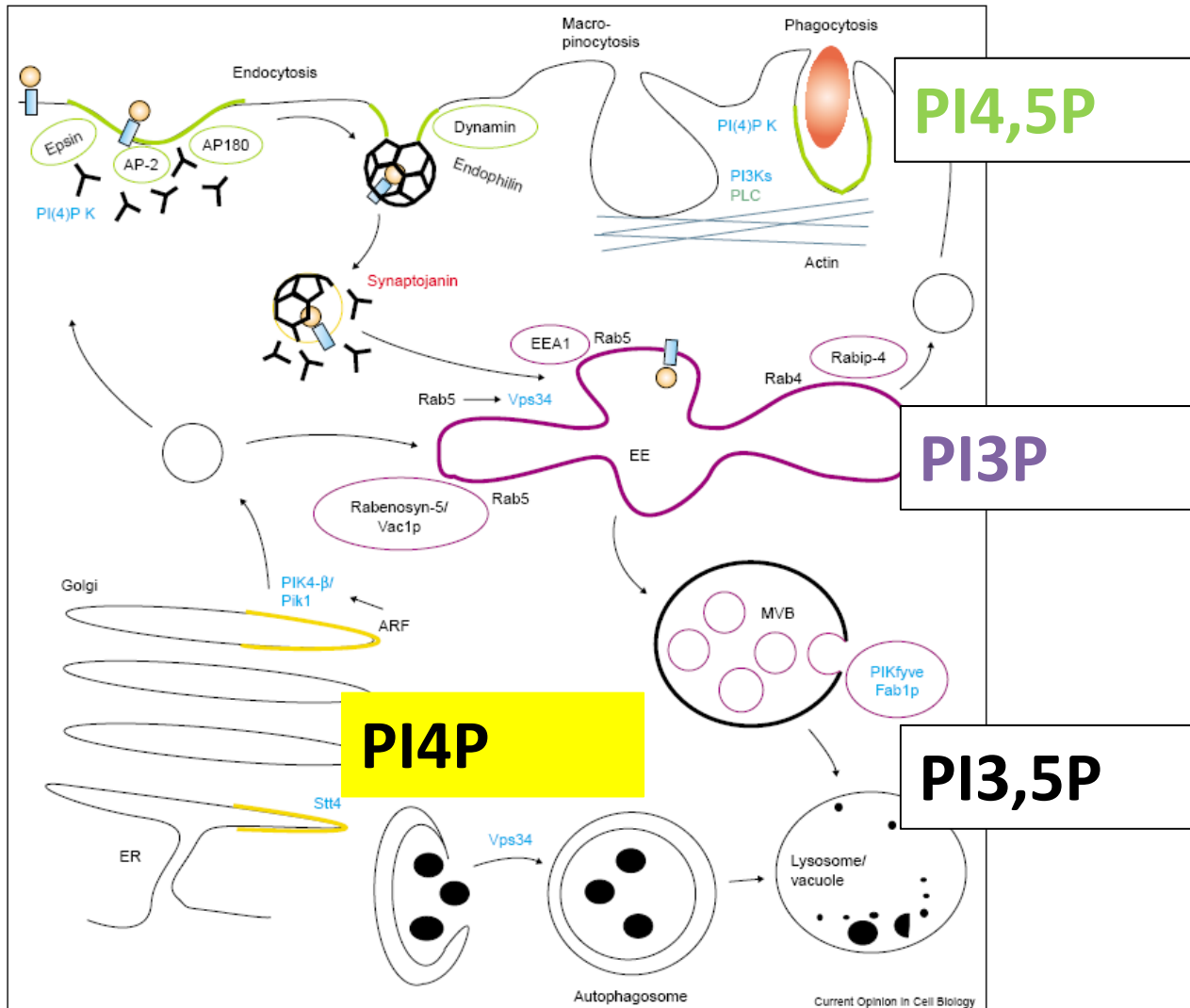
Recruitment of COPII by Sar1



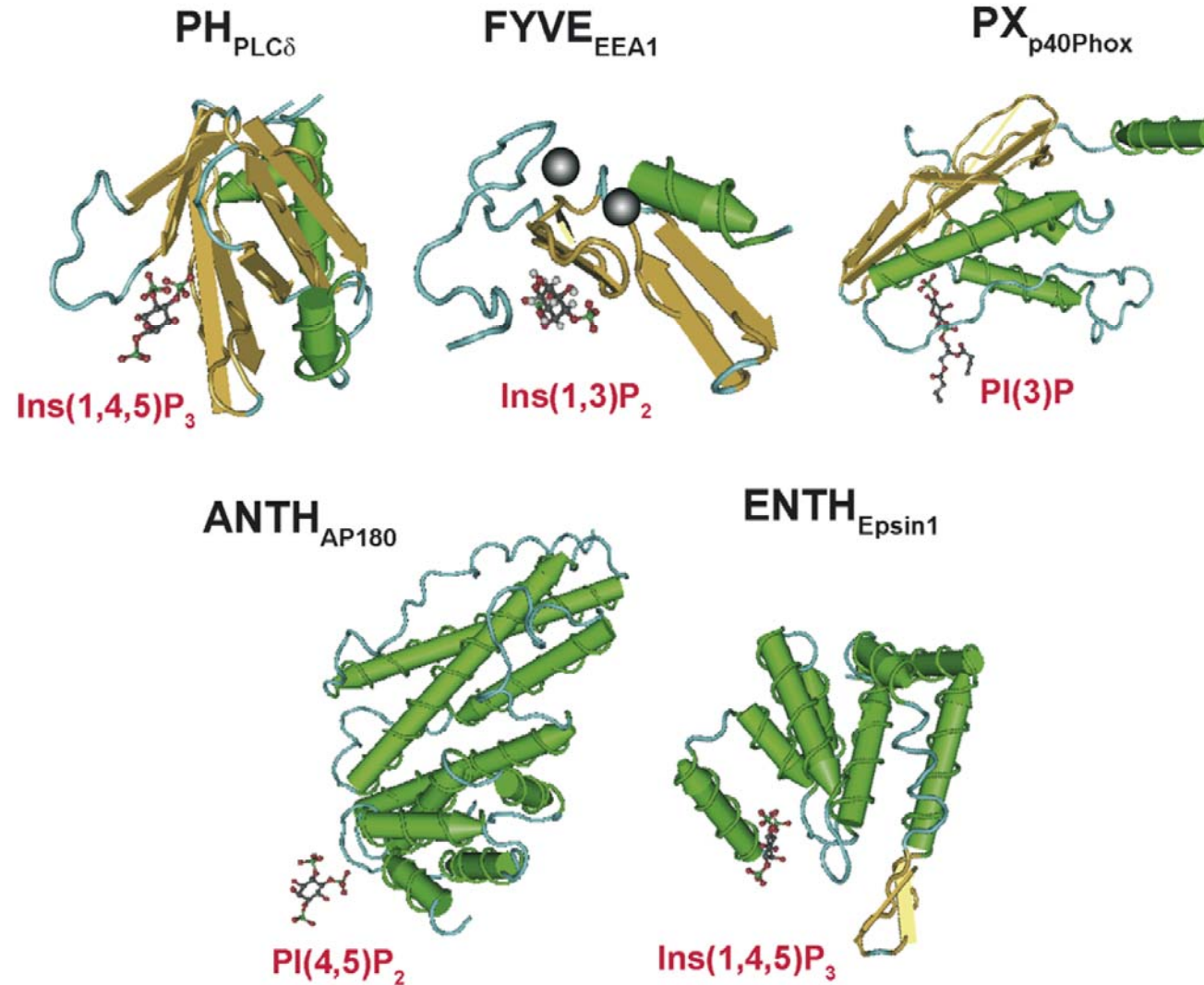
GTP-exchange factors allow membrane association of ARF family GTPases



Recruitment of coats by lipids

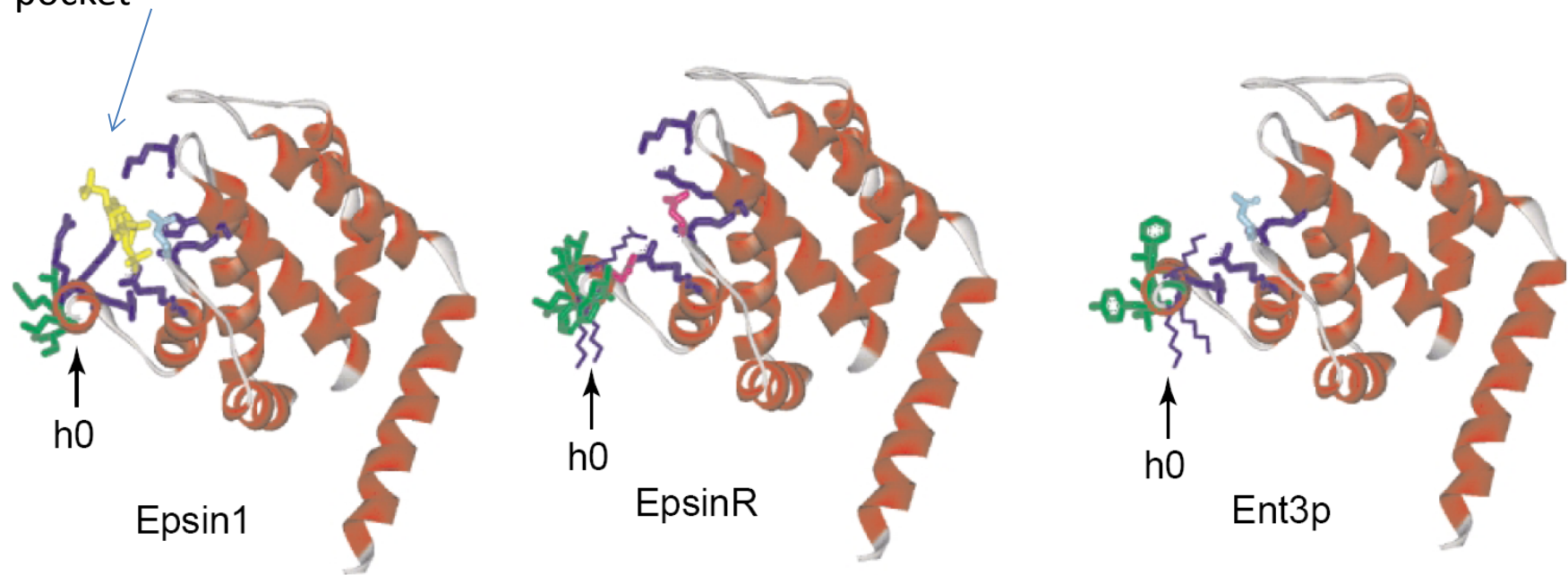


Common PIP binding domains

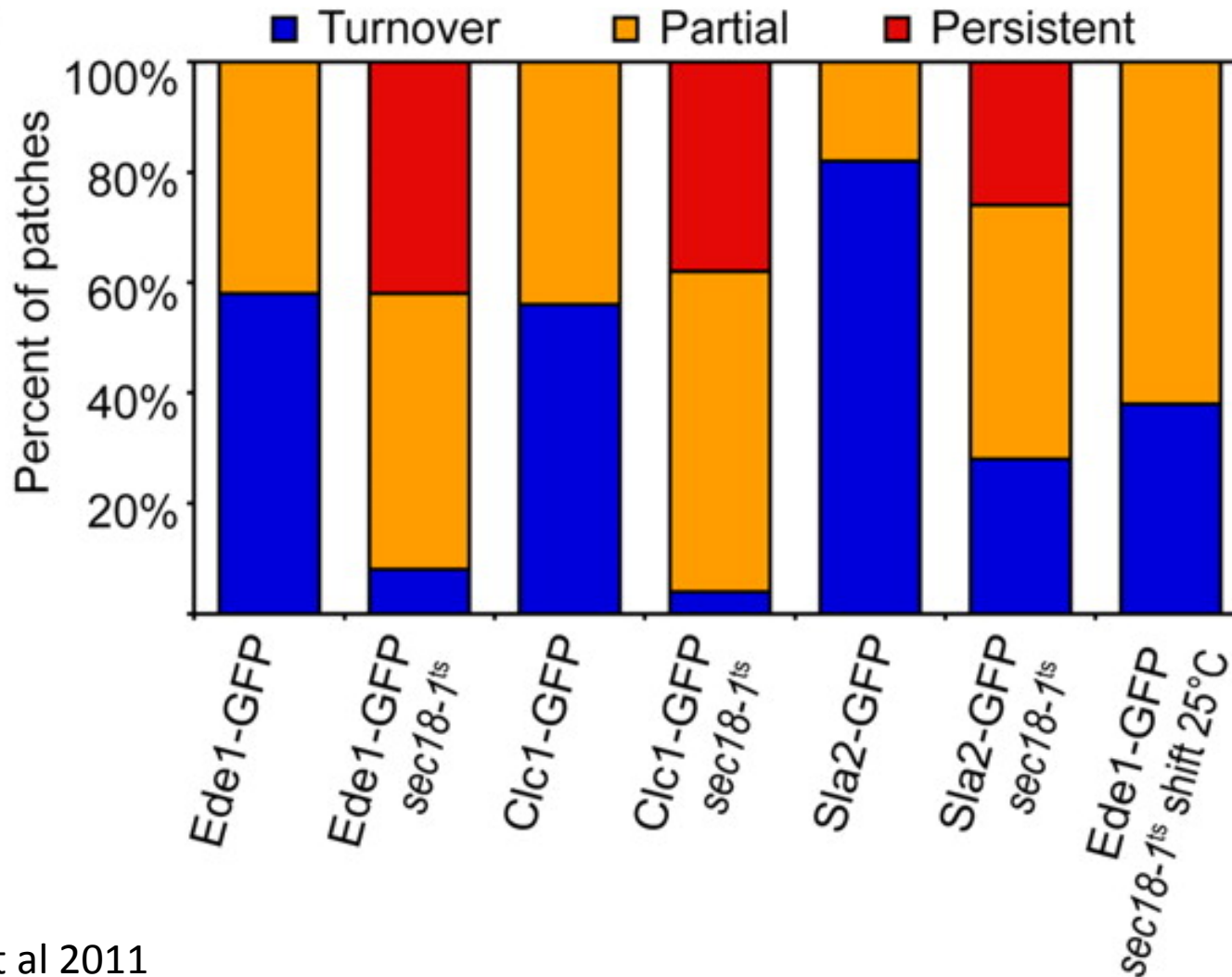


Individual domain families have diversified

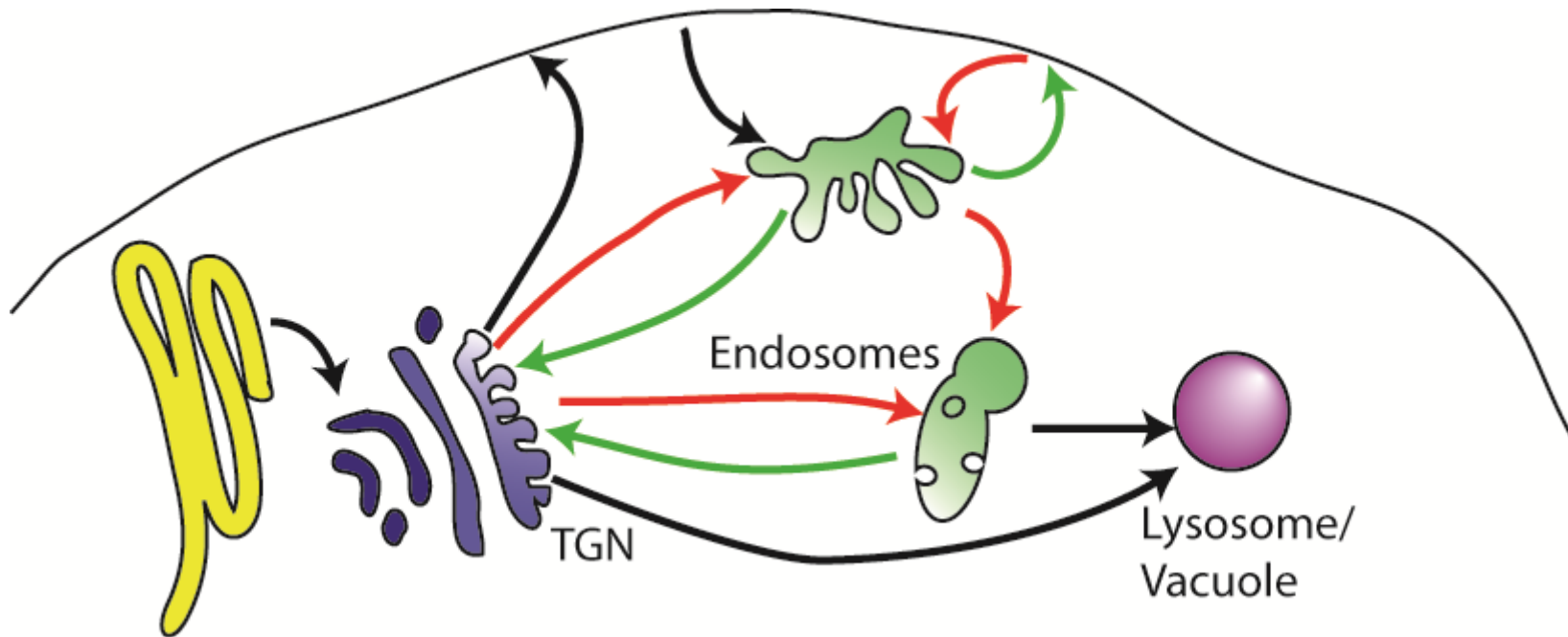
Lipid binding pocket



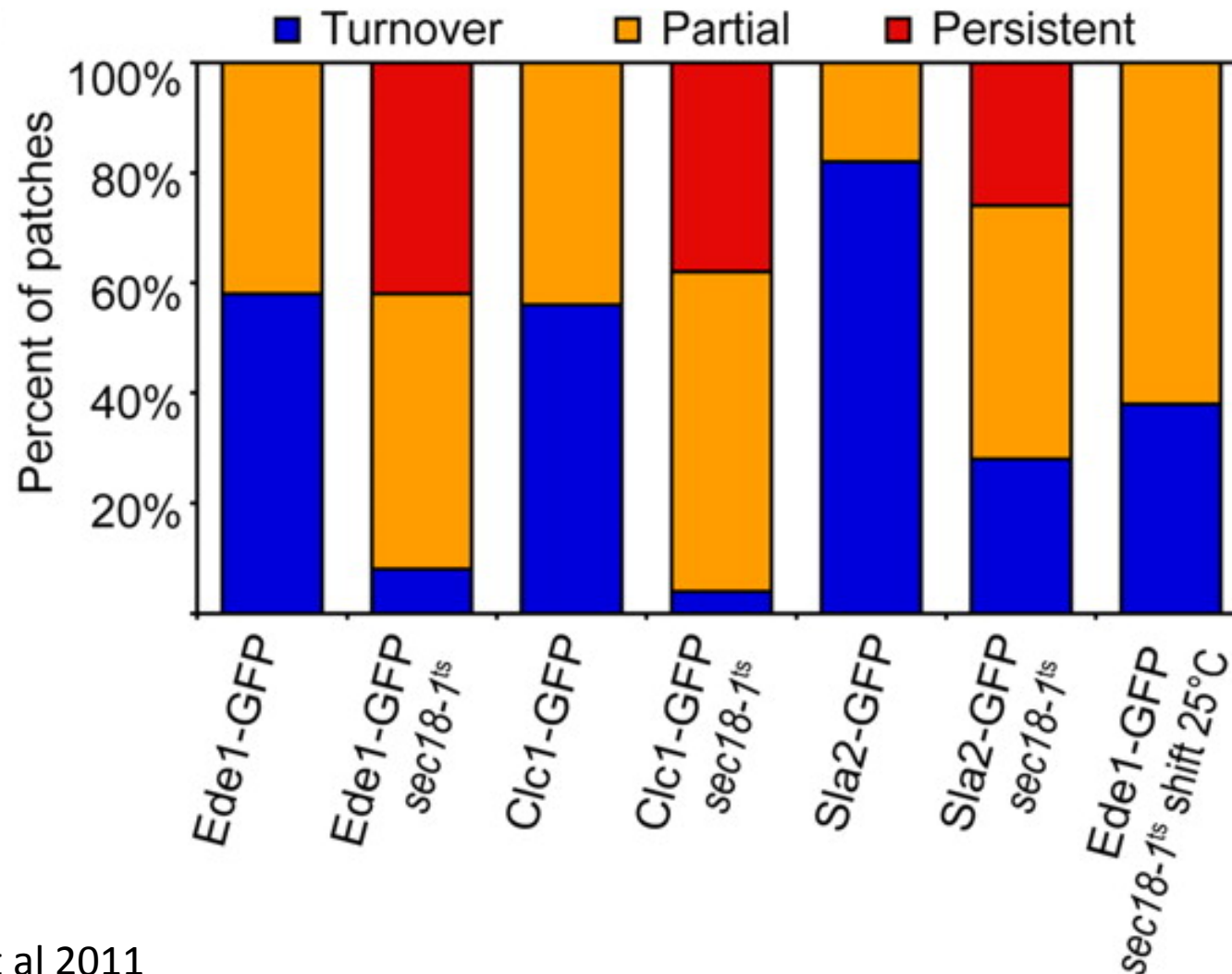
Recruitment of coats by cargo



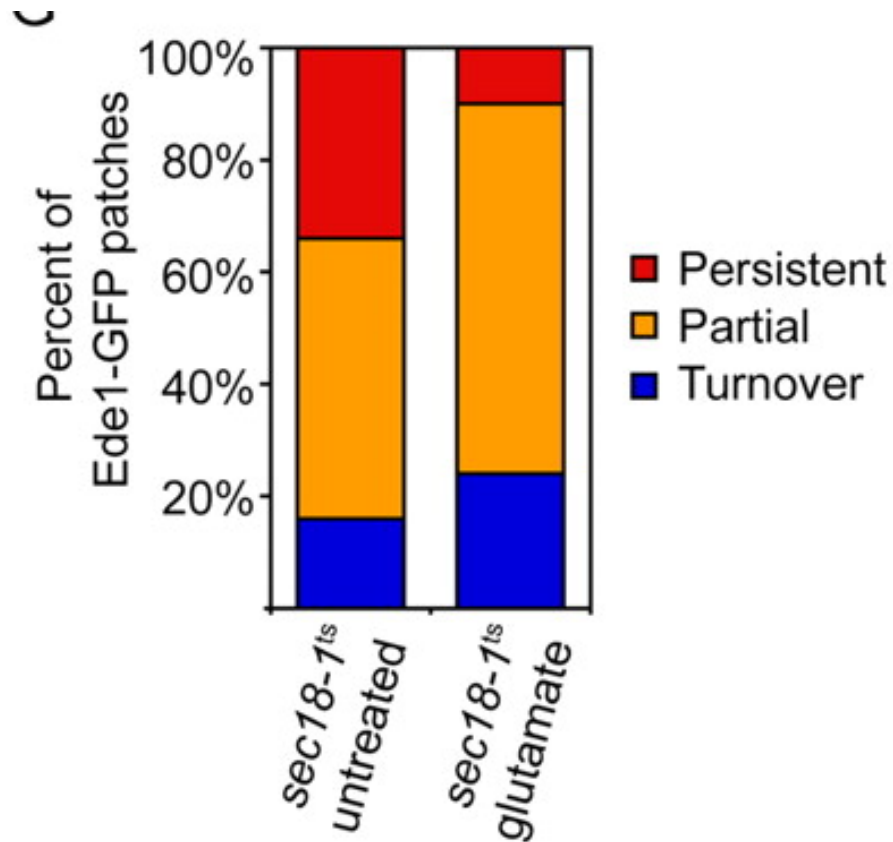
Sec18-ts depletes cargo



Cargo depletion reduces endocytic structures



Induction of endocytosis in depleted cells stimulates coat assembly



Common cargo motifs

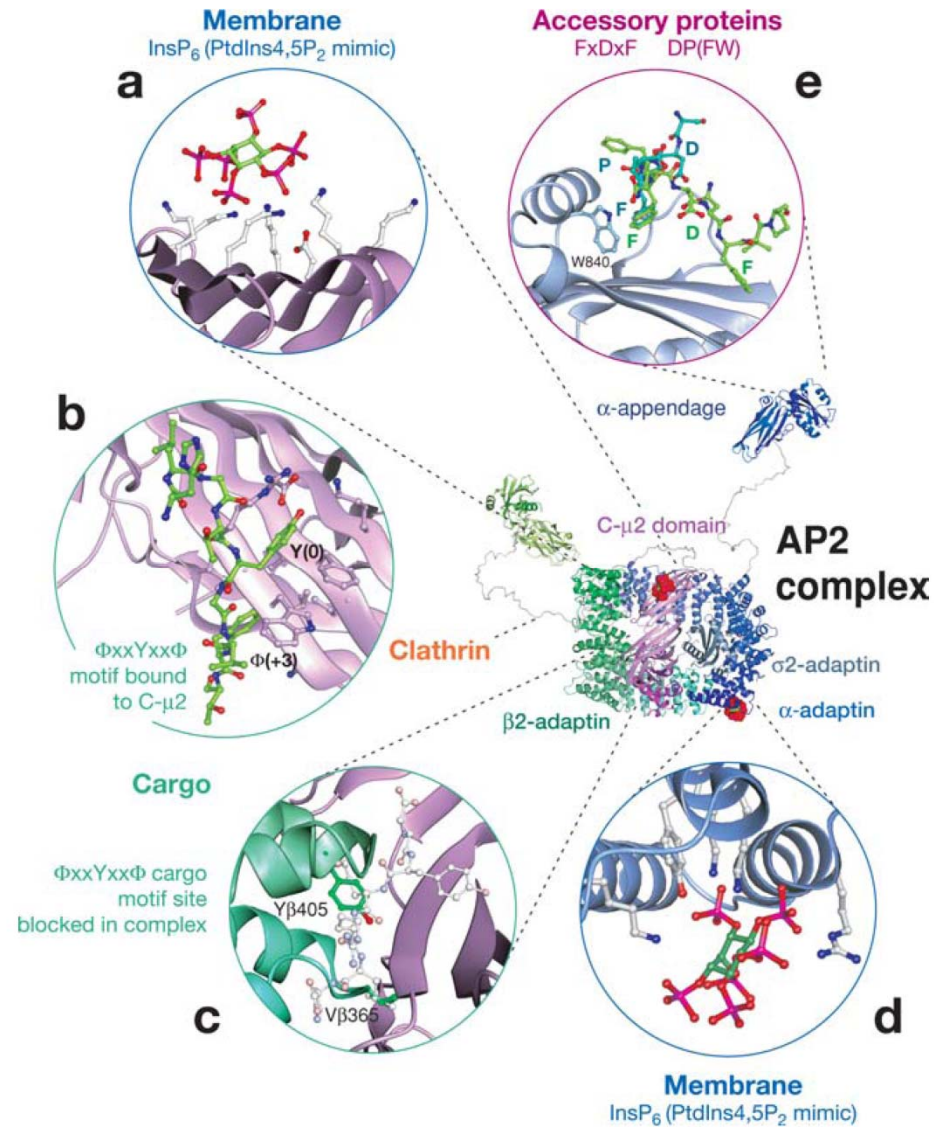
YXX ϕ -clathrin adaptors

Acidic dileucine [DE]xxxLL-clathrin adaptors

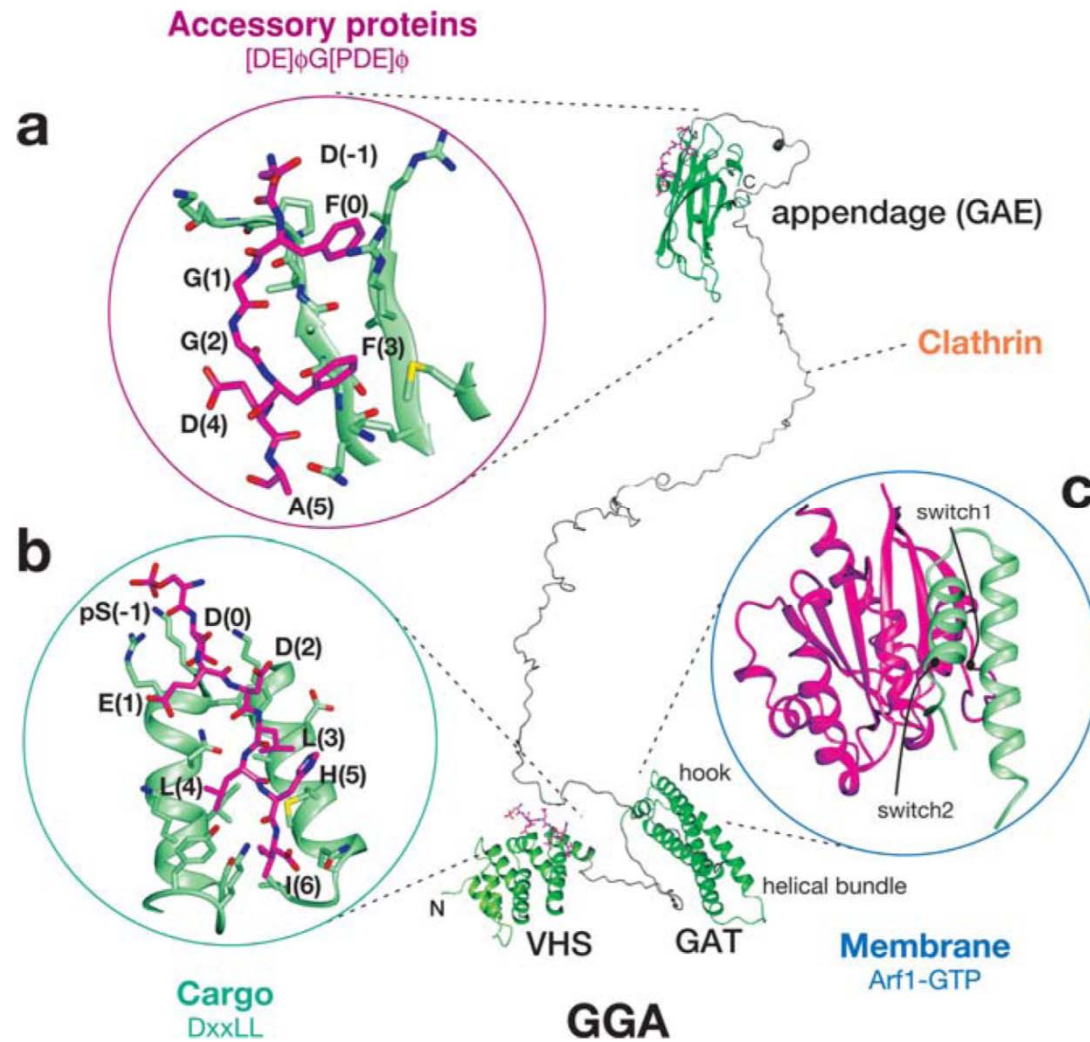
NPXY-endocytic clathrin adaptors

Ubiquitin-clathrin adaptors, ESCRTs

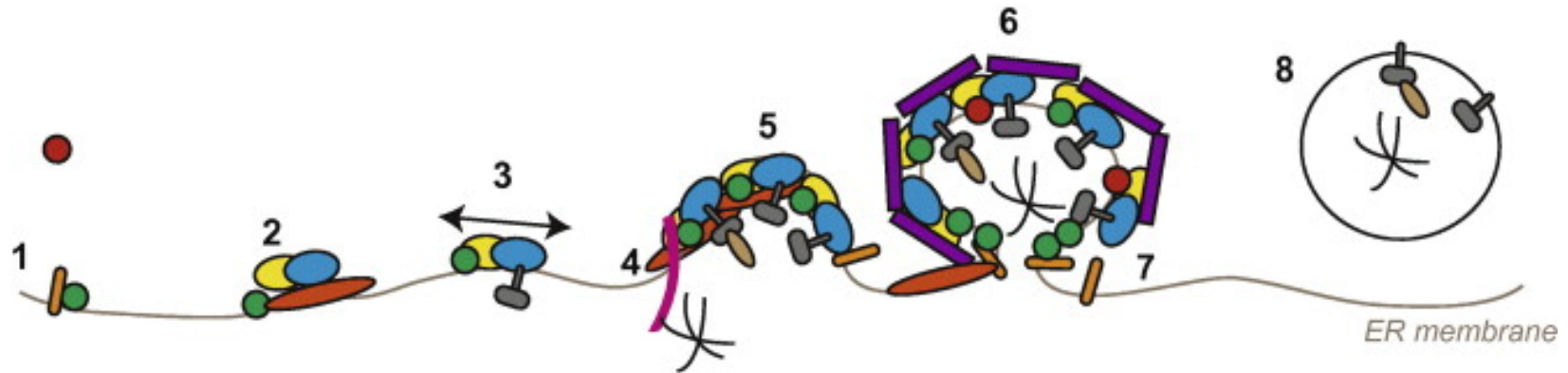
Cargo binding sites



Cargo binding sites

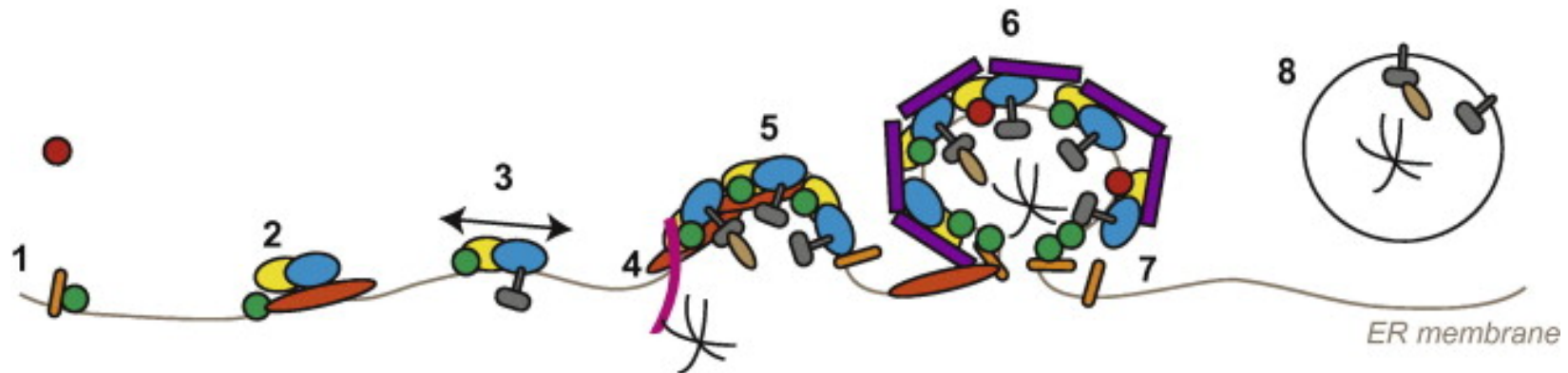


Pause to consider



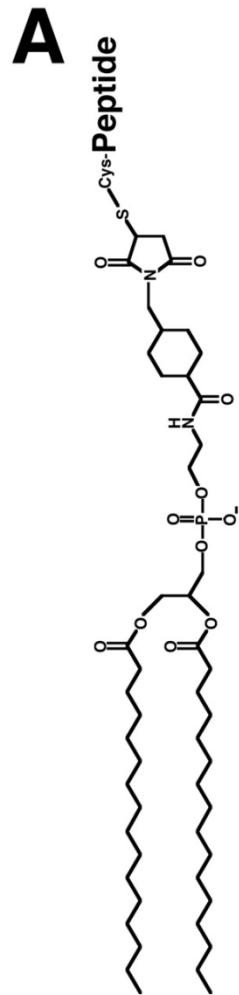
How can we experimentally determine recruiting factors?

Pause to consider



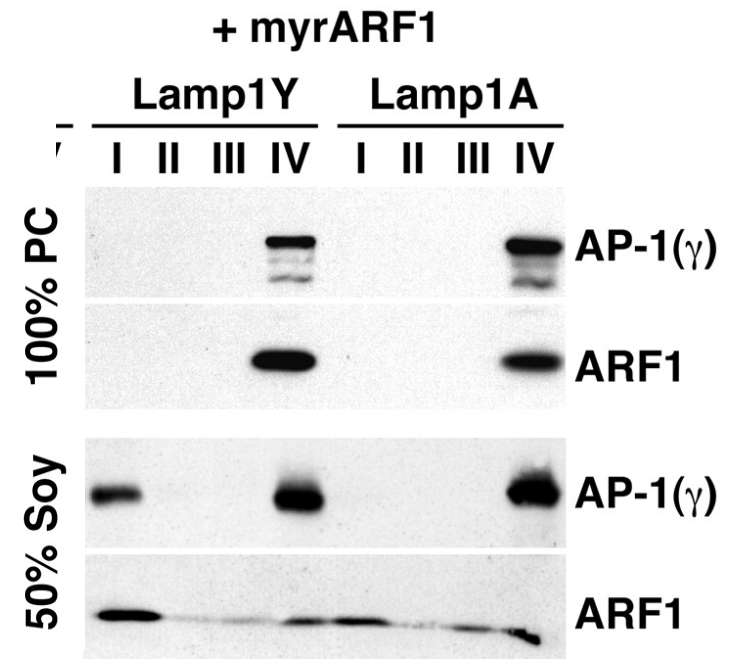
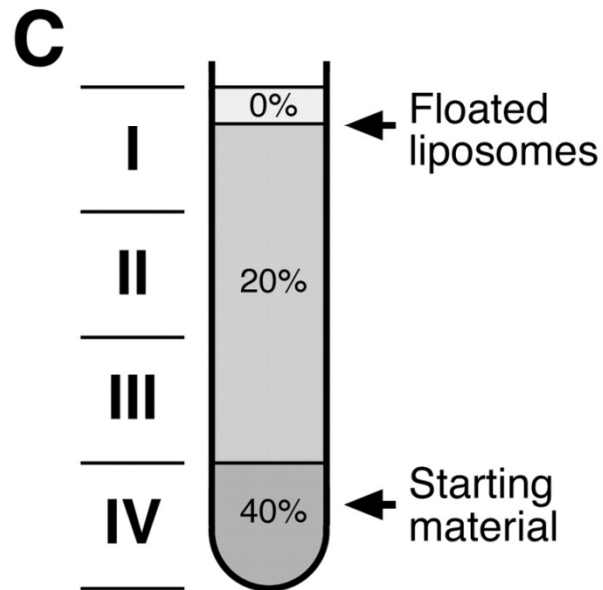
Trans-membrane proteins (sec12 & cargo) that act as initiators of coat assembly, end up in the target organelle. Why don't they recruit coats at the target organelle?

Cooperative recruitment

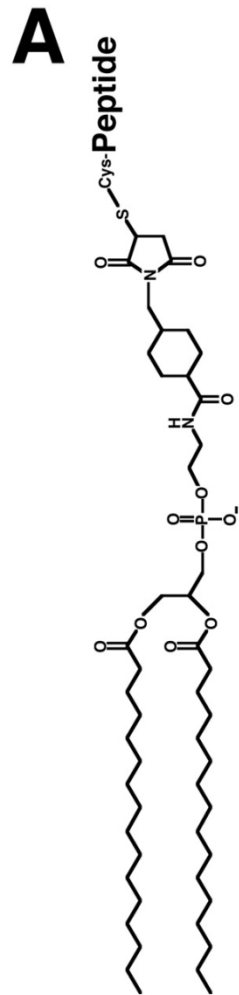


B

Lamp1Y	C-RKRSHAGYQTI
Lamp1A	C-RKRSHAGAQTI
TGN38Y	C-KVTRRPKASDYQRL
TGN38A	C-KVTRRPKASDAQRL

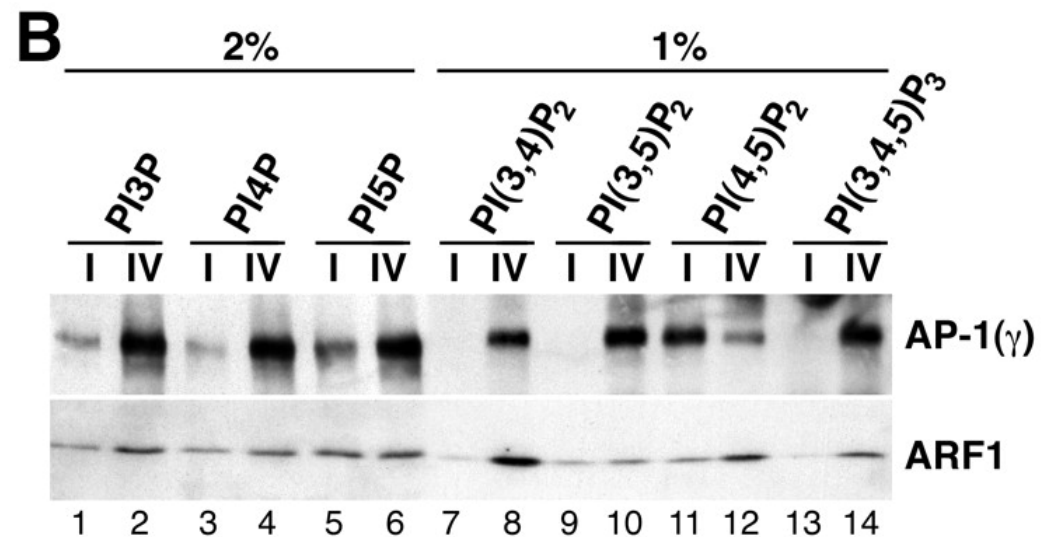


Cooperative recruitment

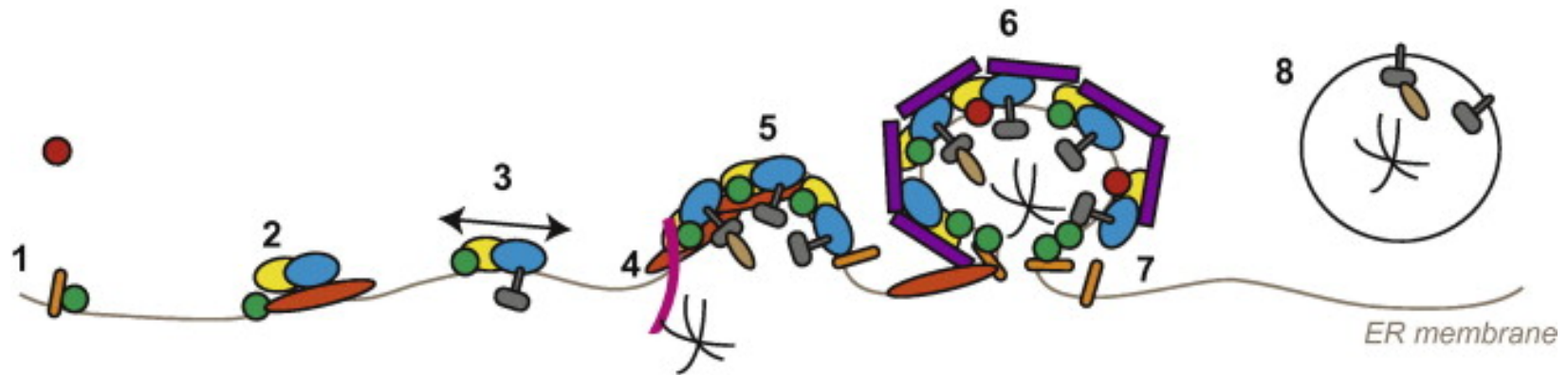


B

Lamp1Y	C-RKRSHAGYQTI
Lamp1A	C-RKRSHAGAQTI
TGN38Y	C-KVTRRPKASDYQRL
TGN38A	C-KVTRRPKASDAQRL



Pause to consider



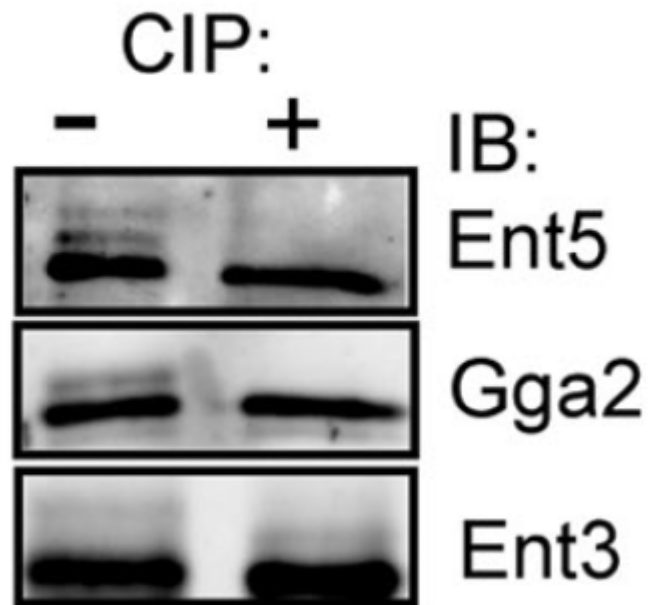
Does knowing how coats are recruited suggest mechanisms for uncoating?

Common mechanisms

- GTP hydrolysis
- Uncoating enzymes
- Lipid modifying enzymes
- Kinases/phosphatases

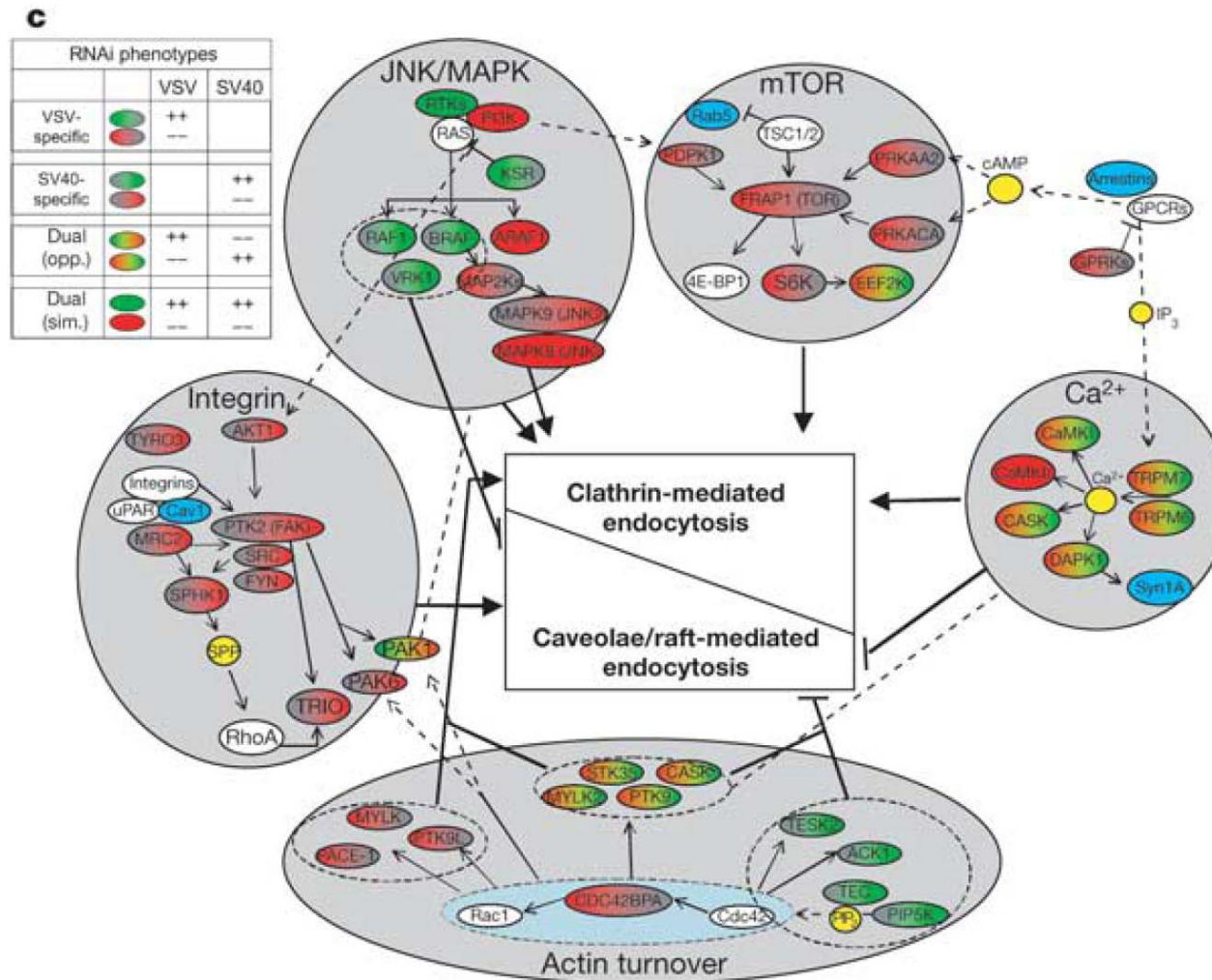
Phosphoregulation of coats

Pause to consider



Numerous proteins involved in membrane traffic are phosphorylated; how might this modulate function?

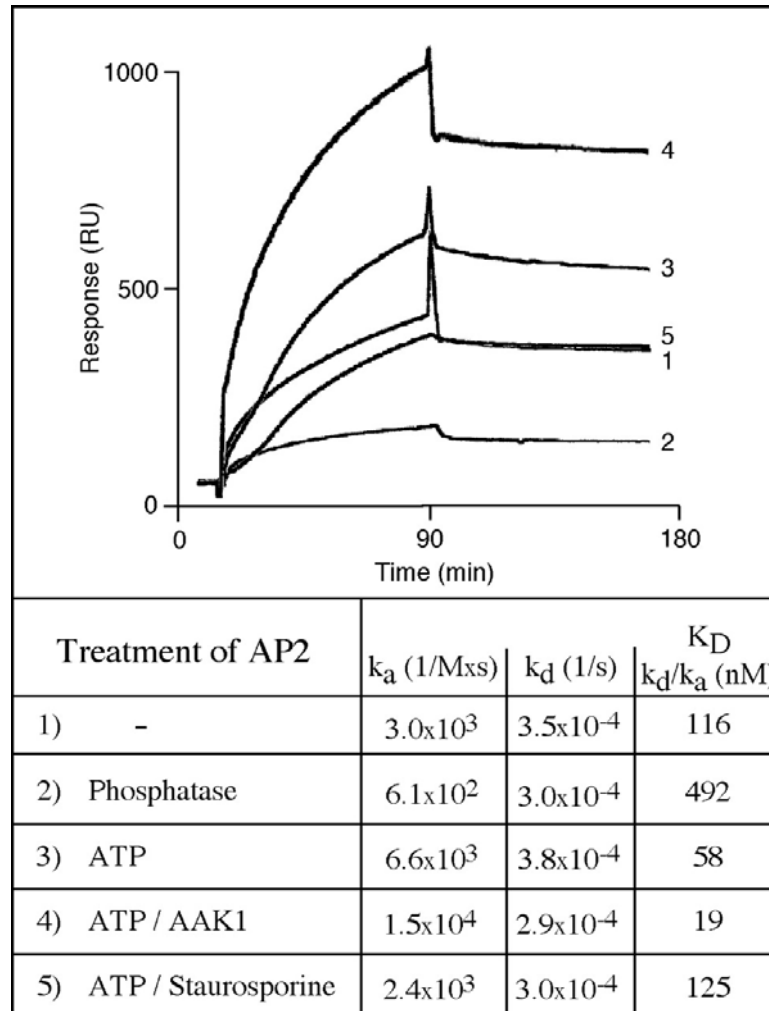
Multiple kinases regulate single traffic events



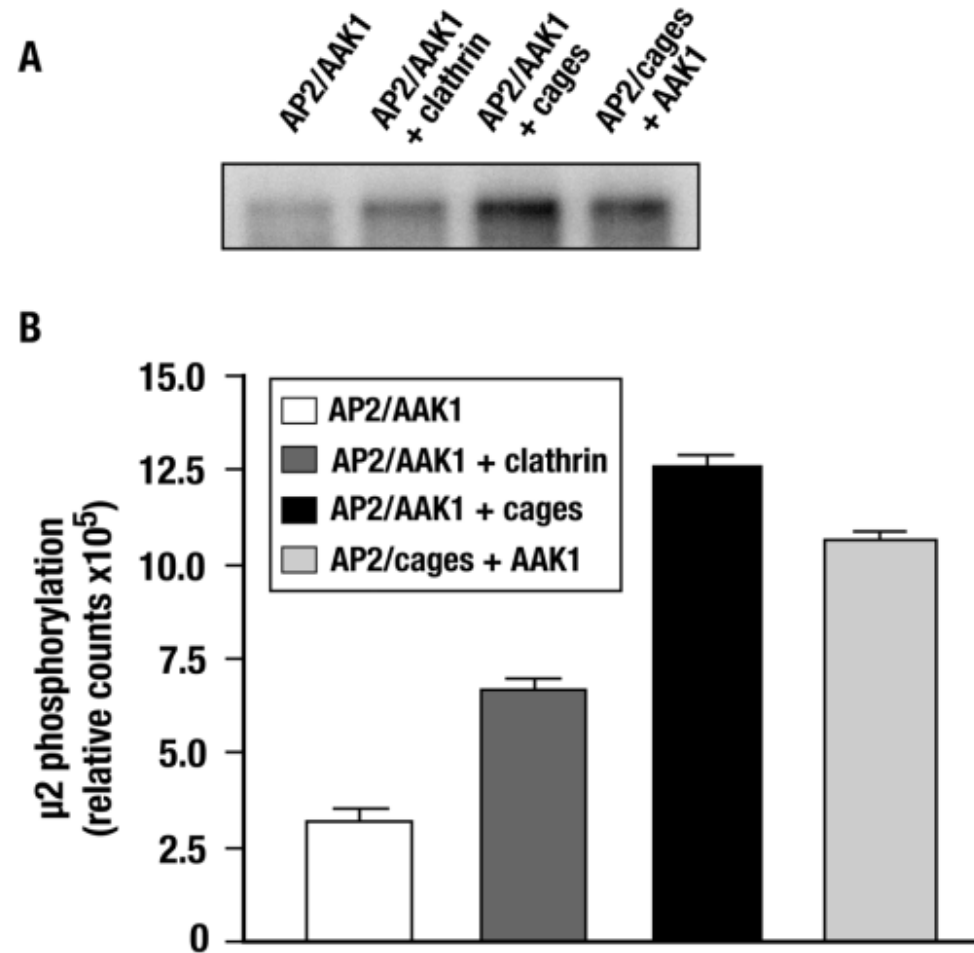
Common mechanisms

- Phosphorylation of tail promotes/inhibits coat binding
- Promote/inhibit autoregulation
- Promote/inhibit protein-protein interactions
- Activate/inhibit enzymatic activity

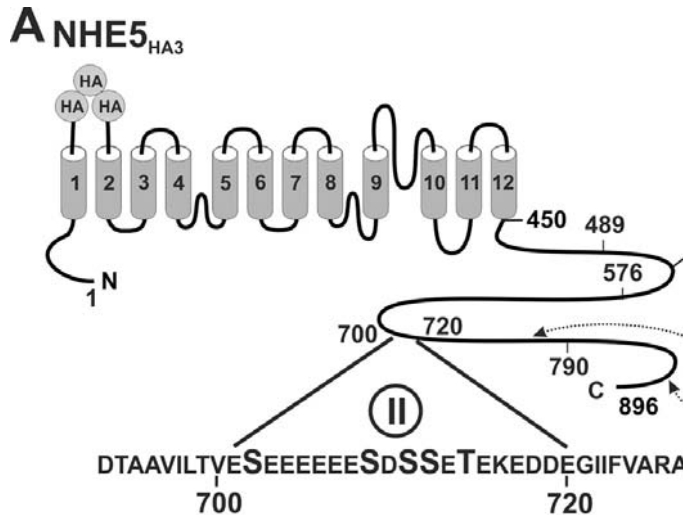
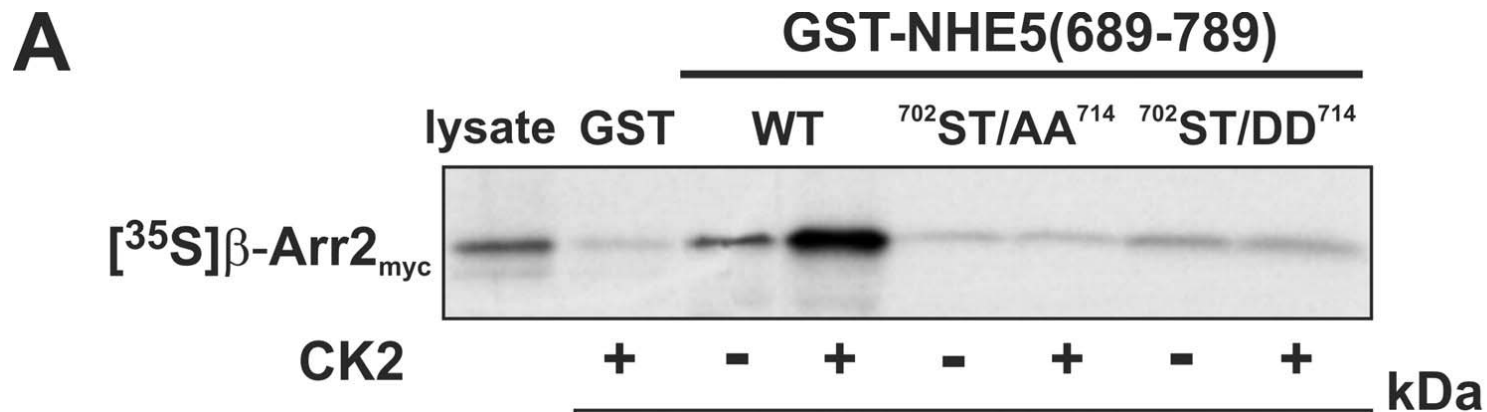
Phosphorylation of AP-2 enhances cargo binding



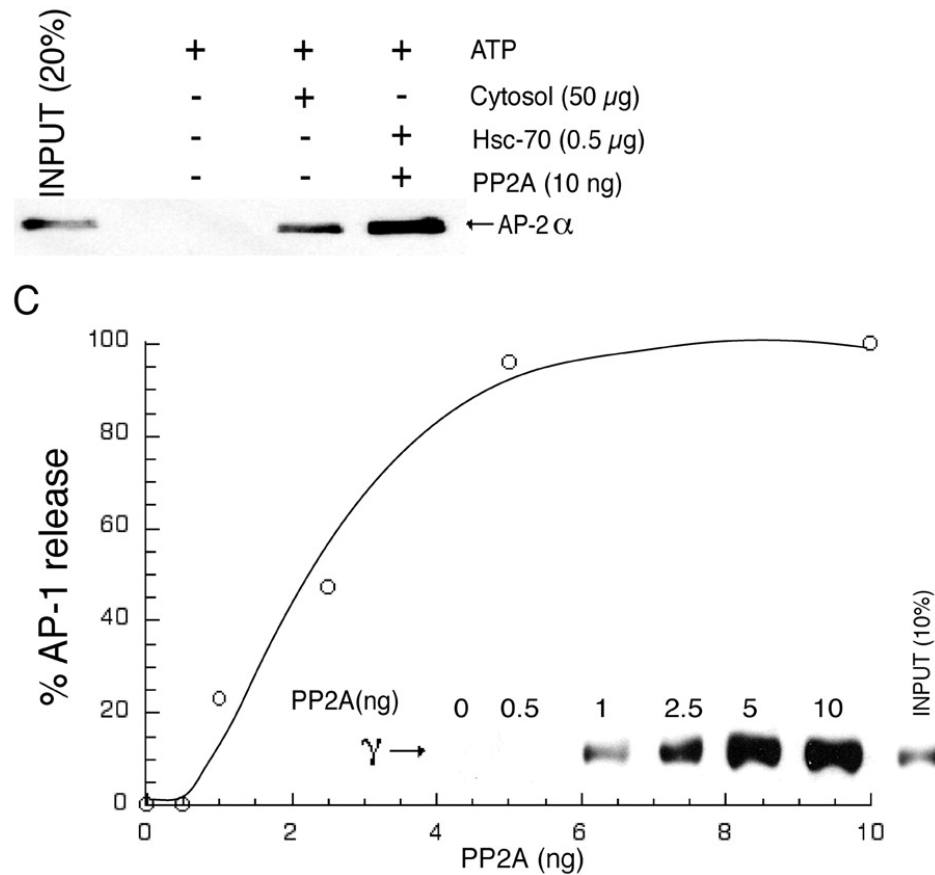
Clathrin stimulates AAK1 activity



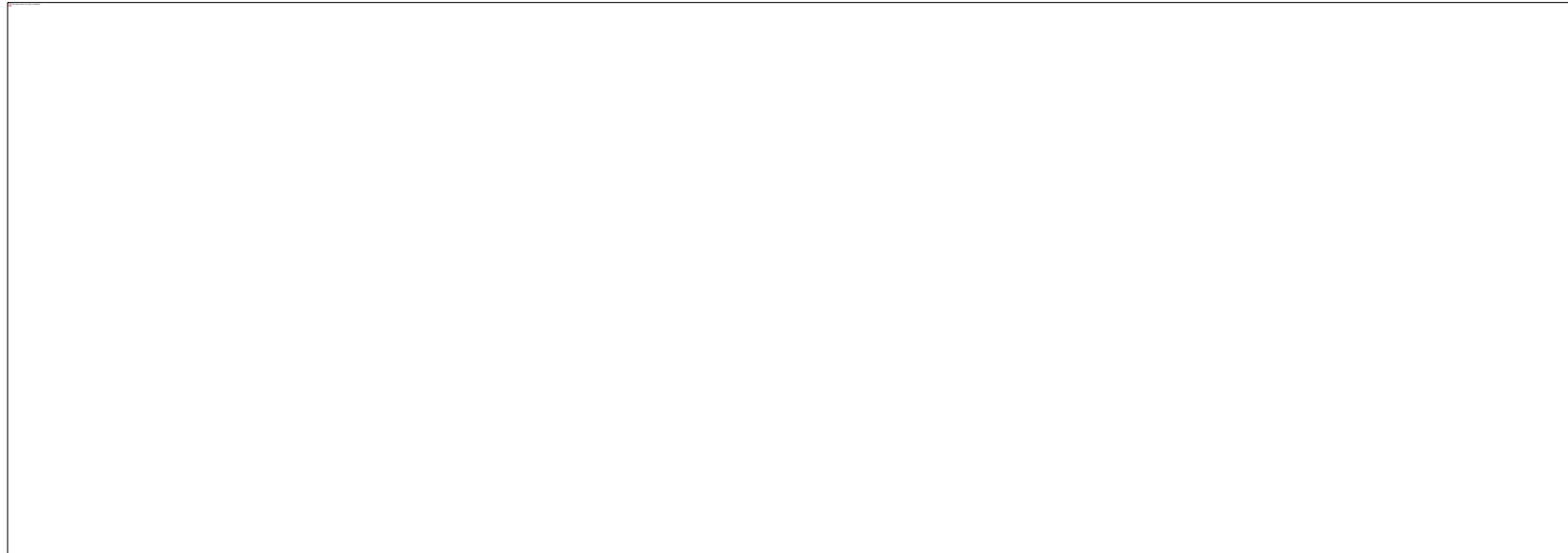
Phosphorylation of cargo enhances adaptor binding



Dephosphorylation of an adaptor reduces membrane association



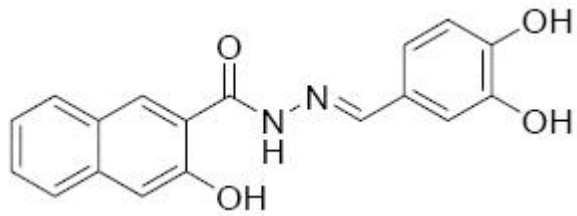
Phosphorylation of coat components promotes disassembly



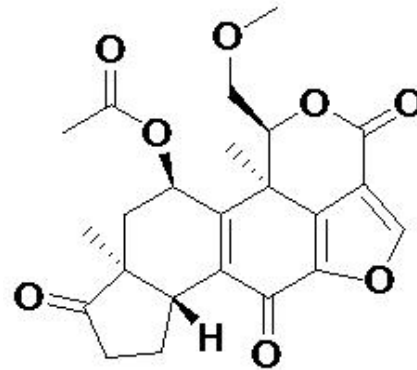
Chemical inhibitors of traffic

Chemicals that block coats

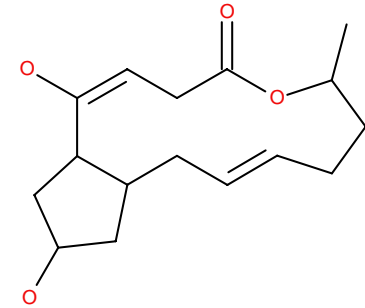
Dynasore



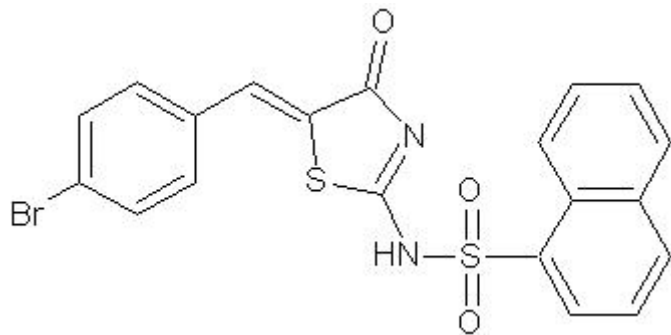
Wortmannin



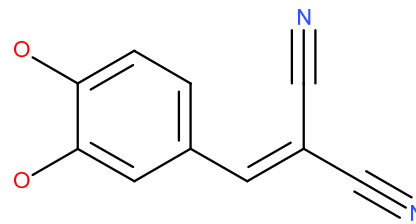
Brefeldin A



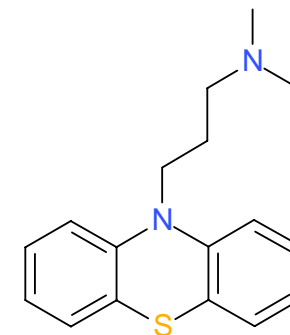
Pitstop



Tyrphostin A23

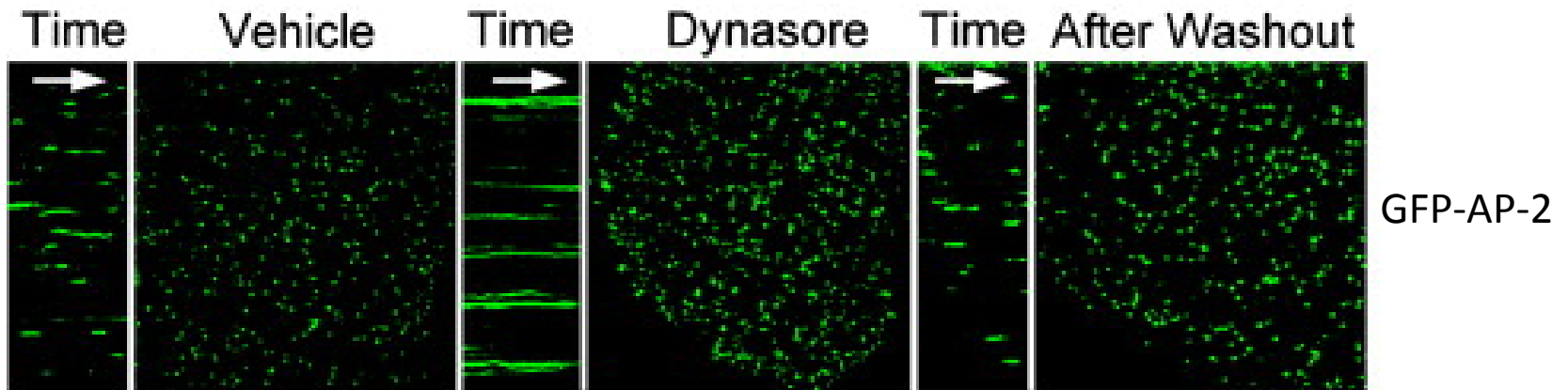
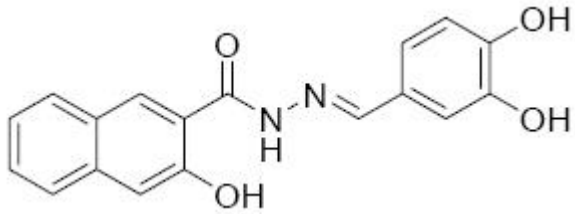


Chlorpromazine



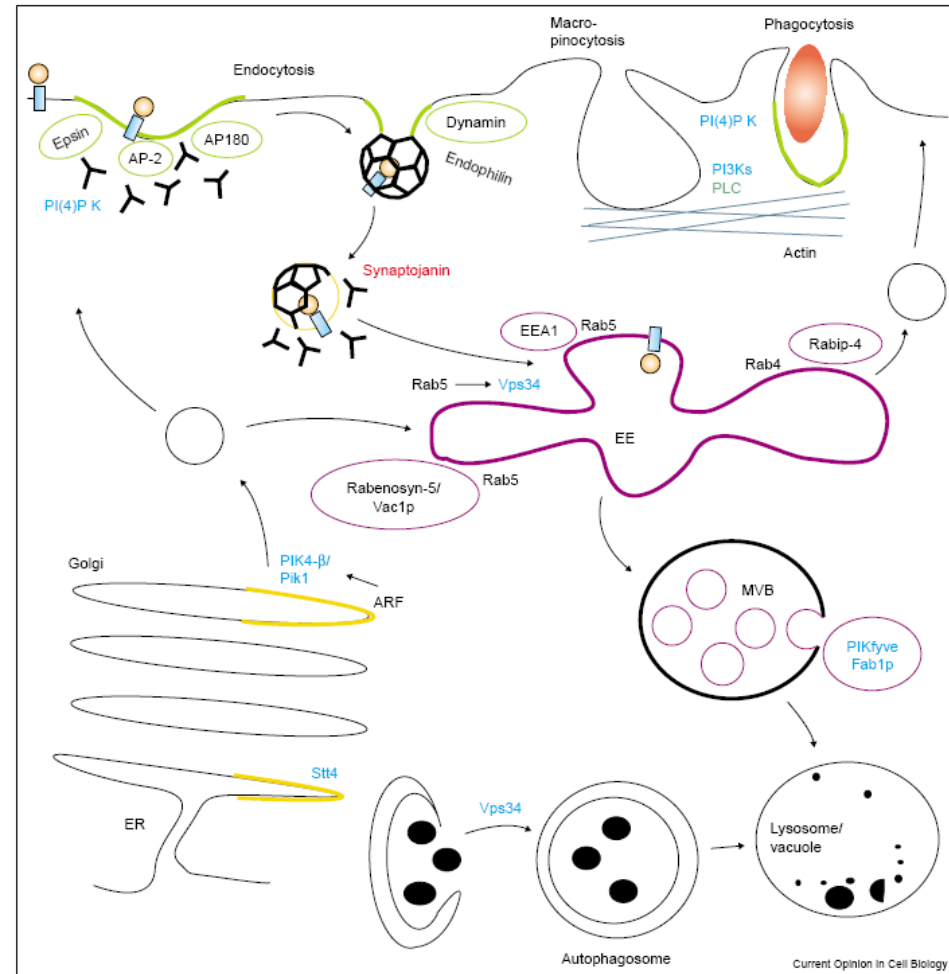
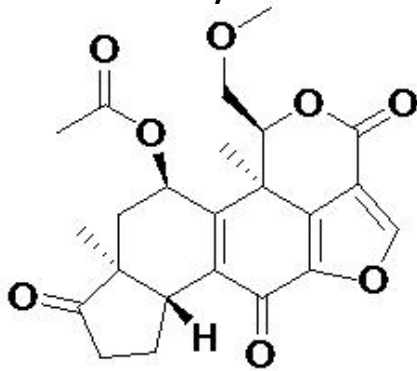
Chemicals that block coats

Dynasore: Inhibitor of Dynamin GTPase activity



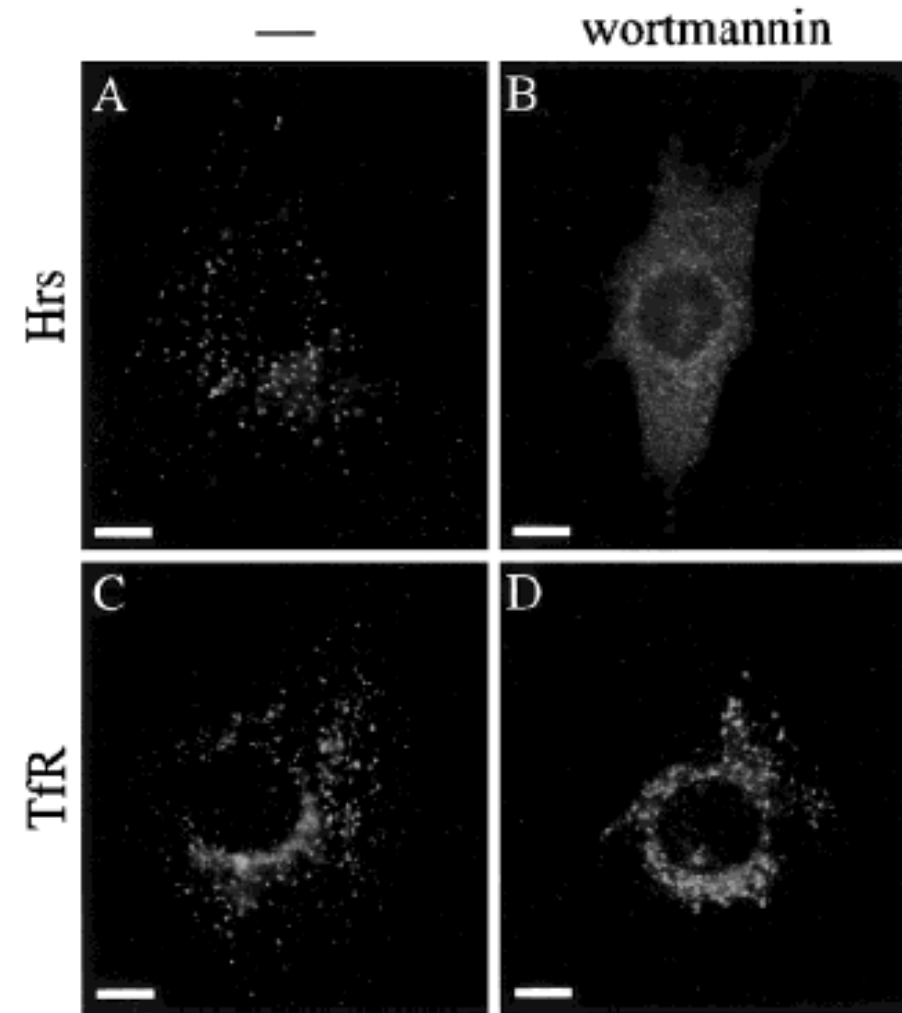
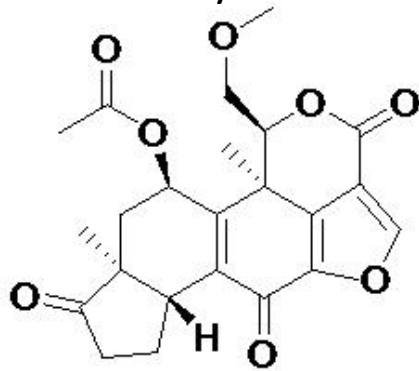
Chemicals that block coats

Wortmannin: Inhibitor of PI3 kinase activity



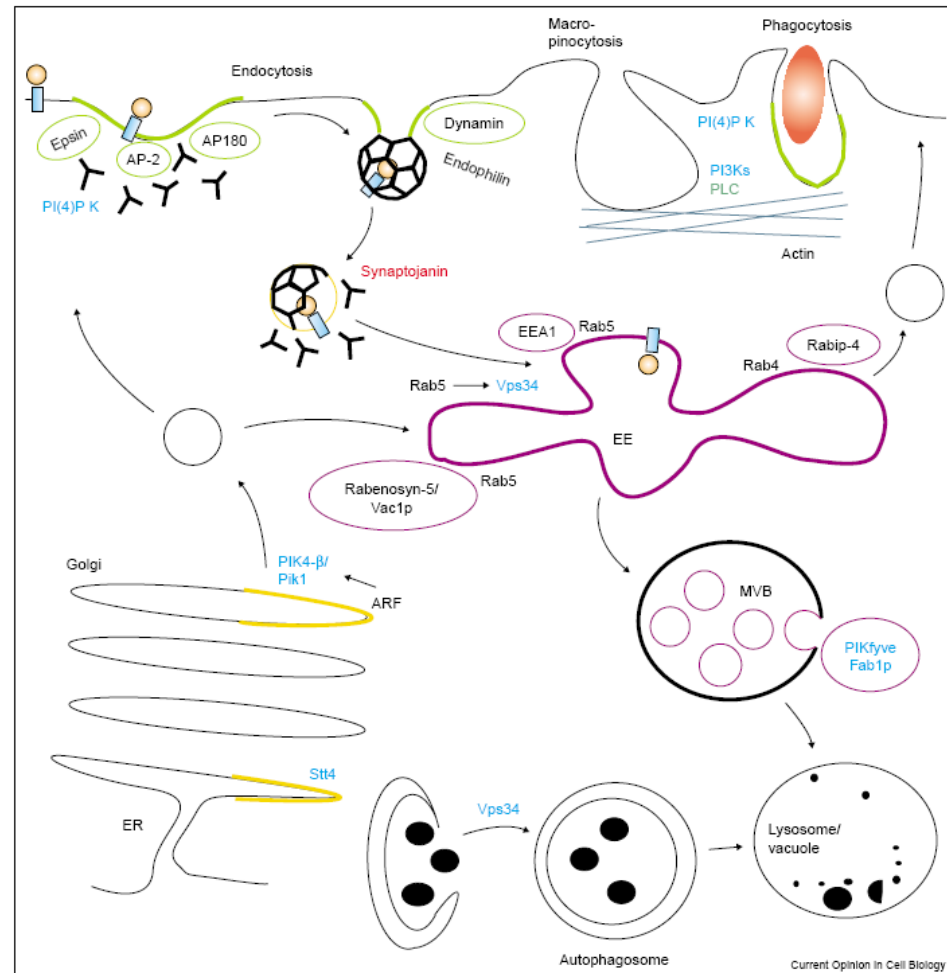
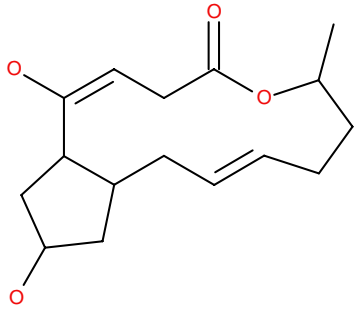
Chemicals that block coats

Wortmannin: Inhibitor of PI3 kinase activity



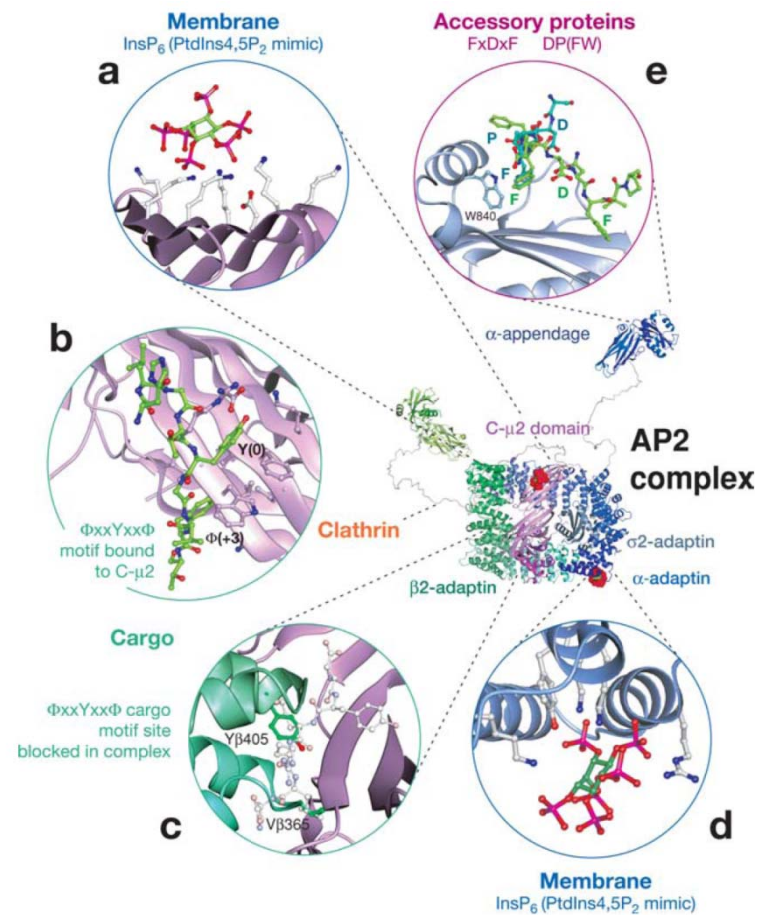
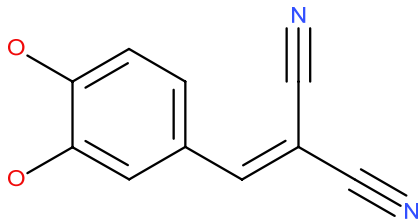
Chemicals that block coats

Brefeldin A: Inhibitor of many ARF GEFs



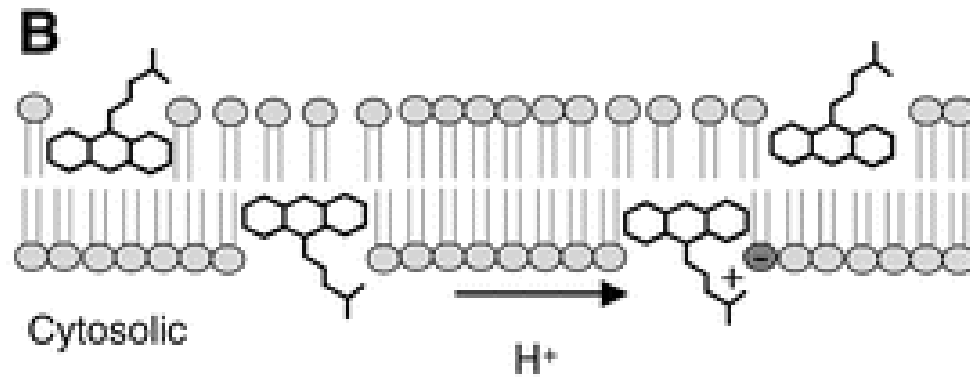
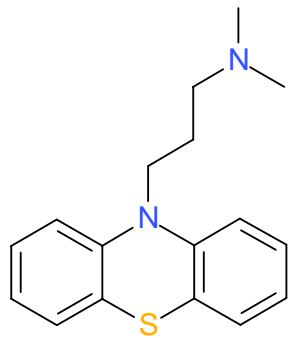
Chemicals that block coats

Tyrphostin A23-Mimics tyrosine sorting signal



Chemicals that block coats

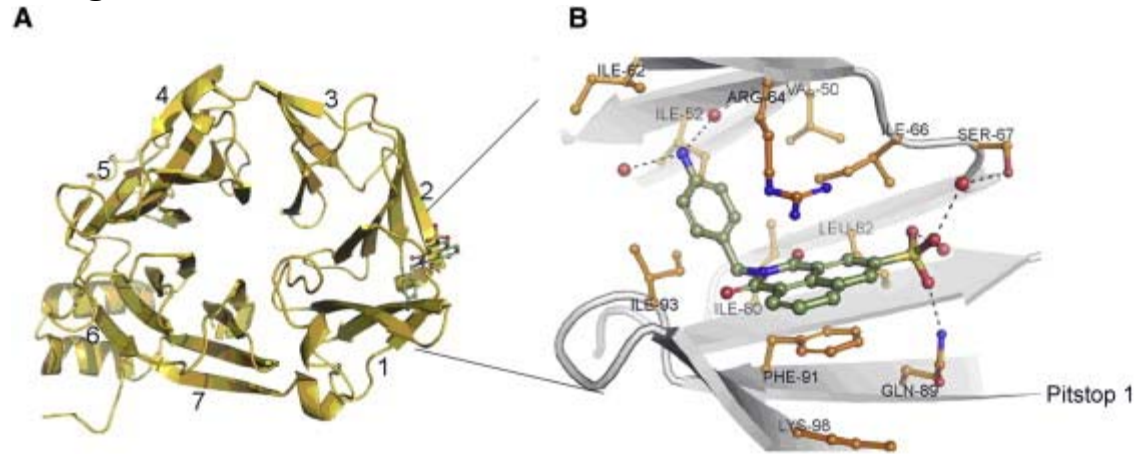
Chlorpromazine: Traps clathrin on membranes



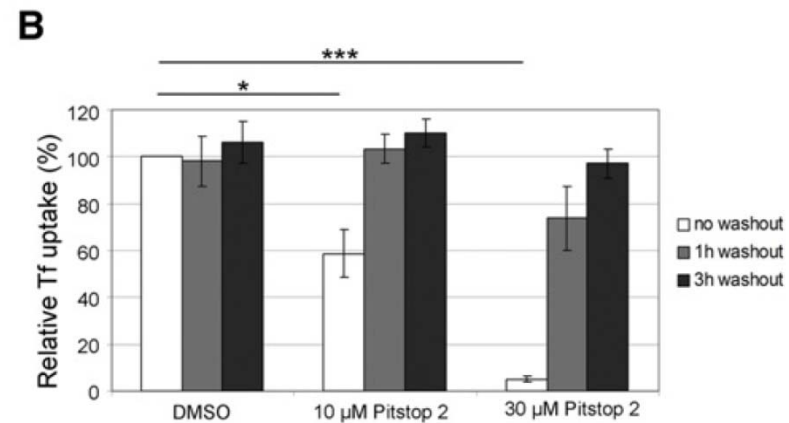
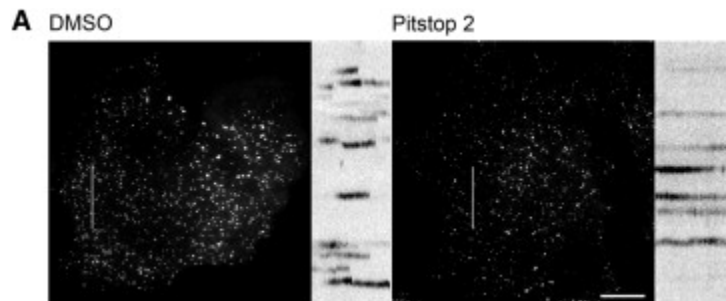
De Filippi L, et al . 2007

Chemicals that block coats

Pitstop-Blocks one adaptor binding site on the clathrin terminal domain

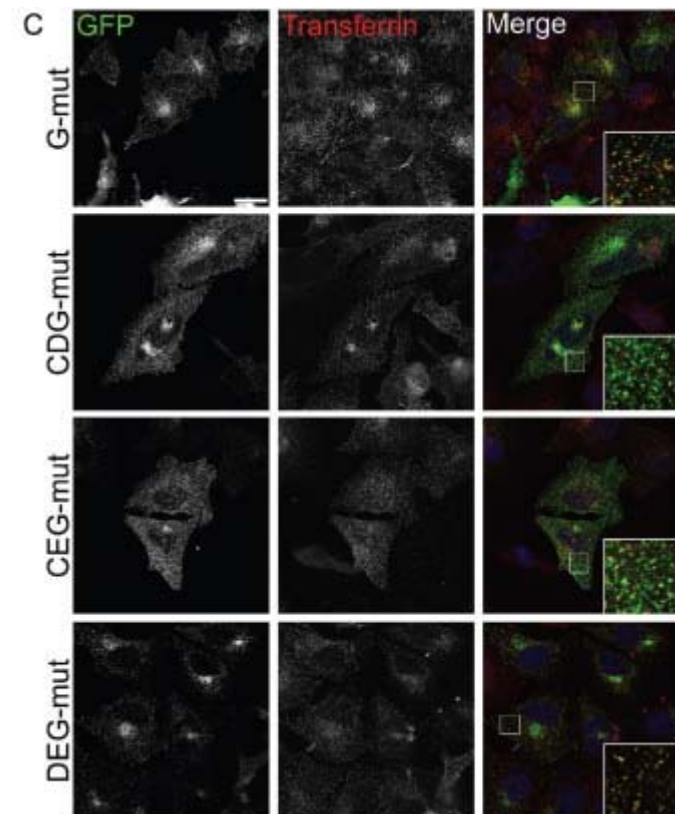
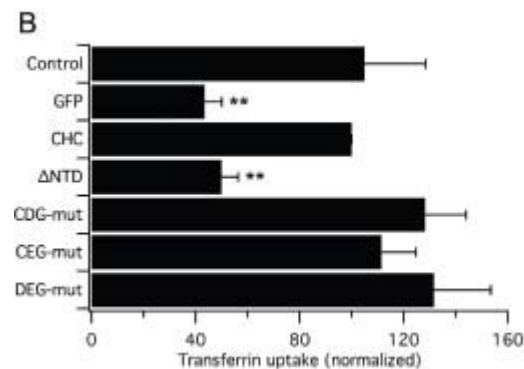
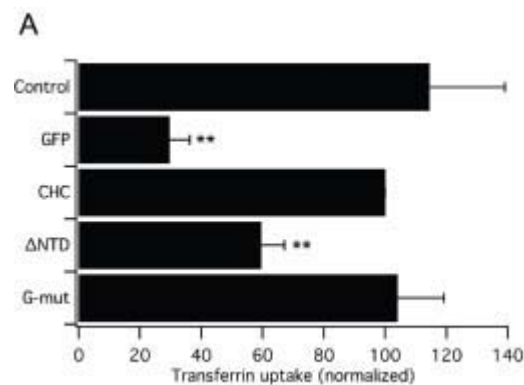


clc1EGFP



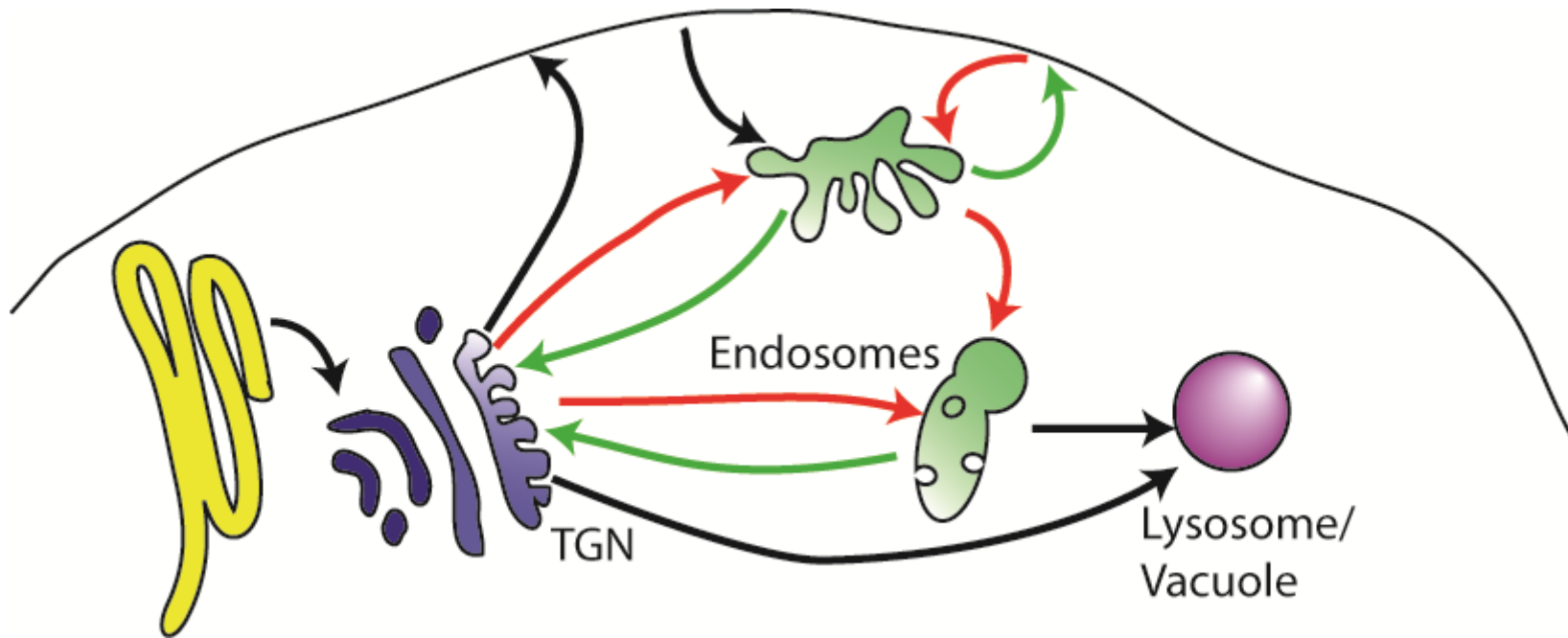
Controversies:

Pitstop arrests traffic but point mutations of its binding site do not? What are the possible explanations?



Controversies

Why is it so hard to determine directionality?



Selected reading and references used

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