

# II. Approaches/ Methods

1. Biochem: In vitro (reconstitution) assays to identify the players. by removing one at a time. (e.g. mutants, inhibitors, or physical removal), or adding one.

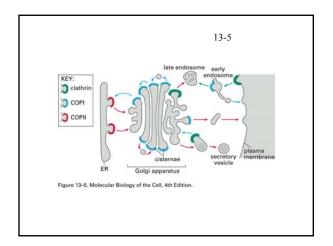
2. Genetic: Yeast *sec* mutants deficient in secretion. >25 genes Identify proteins that are essential for each step.

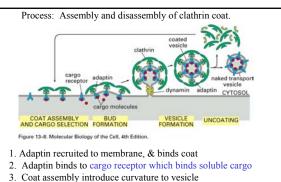
### 3. Cell biology.-

a. endocytosis of labeled ligand by a living cell, and visualize internalized ligand by Immuno-Gold or flourescence. [To visualize, cell is fixed] Follow path of endocytosis.

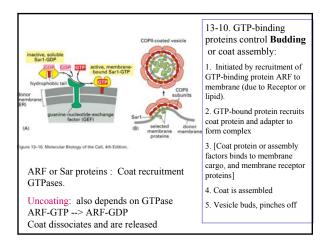
b. Transfection with GFP-tagged proteins! [living cell] Follow sorting in real time. Measure time it takes to move from ER-> G. Movie later

#### Budding with specificity. 17-51. Three types of coated vesicles Small GTP-Binding Vesicle Coat and Adapter Protein Transport Step Clathrin heavy and light chains; AP2 ARE Plasma membrane $\rightarrow$ endosome (endocytosis Clathrin Clathrin heavy and light chains; AP1 ARE Golgi → endosome $Golgi \rightarrow$ lysosome, vacuole, melanosome, or platelet vesicles Clathrin heavy and light chains; AP3 ARF COPI COP α, β, β', γ, δ, ε, ς ARE $Golgi \rightarrow ER$ Retrograde transport between Golgi cisternae Sec23/Sec24 complex; Sec13/Sec31 complex; Sec16 COPII Sar1 $ER \rightarrow Golgi$ Subclasses within each type: each specialized for a different transport step





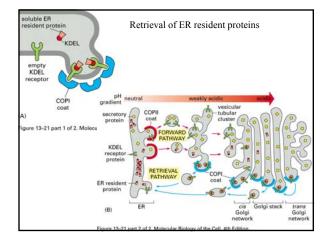
- 4. Coat is nearly assembled
- 5. Vesicle pinches off with help of dynamin
- 6. After budding, coat disassembles

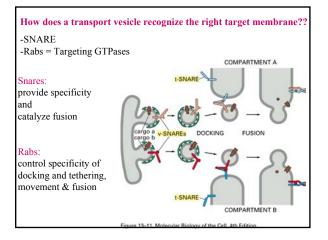


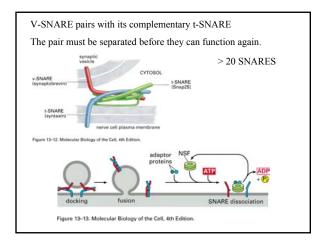
### How does sorting occur during budding??

Tab. 17-6. Sorting signals and receptors

| Signal Sequence*              | Type of Protein <sup>†</sup> | Transport Step   | Vesicle<br>Type | Signal Receptor  |
|-------------------------------|------------------------------|--|-----------------|--|
| Lys-Asp-Glu-Leu (KDEL)        | Secreted                     | Golgi to ER  | COP I           | KDEL receptor (ERD2<br>protein) in Golgi membrane                                |
| Lys-Lys-X-X (KKXX)            | Membrane                     | Golgi to ER  | COP I           | COP $\alpha$ and $\beta$ subunits  |
| Di-acidic (e.g., Asp-X-Glu)   | Membrane                     | ER to Golgi  | COP II          | Not known  |
| Mannose 6-phosphate (M6P)     | Secreted                     | Trans-Golgi and<br>plasma membrane to<br>late endosome | Clathrin        | M6P receptor in Golgi<br>and plasma membrane;<br>AP1 and AP2 adapter<br>proteins |
| Tyr-X-X- $\phi$ (YXX $\phi$ ) | Membrane                     | Plasma membrane to<br>endosome                         | Clathrin        | AP2 adapter proteins   |
| Leu-Leu (LL)                  | Membrane                     | Plasma membrane to<br>endosome                         | Clathrin        | AP2 adapter proteins   |

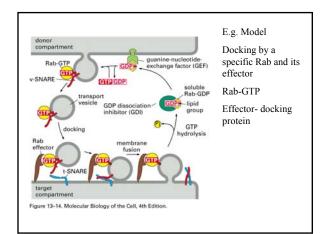


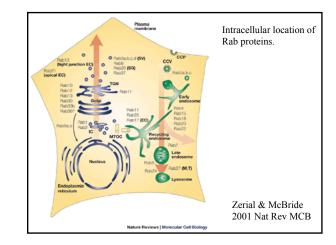


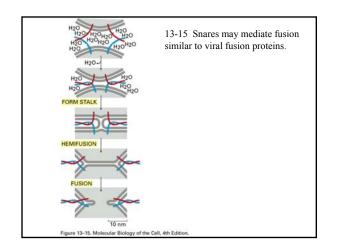


### Rab GTPases and their effectors determine compartment specificity Additional layer of regulation to ensure that t-SNARE fuse membranes only at the right time and in the correct place. Effectors: Protein(s) that bind a Rab-GTPase directly and is required for downstream function determined by that Rab. [e.g. tethering, motor proteins] Rabs Organelle Rab Effectors Function Rab1 ER-G p115 tethering Rab3a synaptic ves rabphilin 3 potentiates fusion Rab4 early endosome Rabaptin/NEF sorting, recycling Rab6 G -> ER, intra G Rabkinesin6 vesicle motility

RabGDP = soluble proteins that become attached to membranes when it is converted to Rab-GTP







### Summary

### How is vesicular transport specificity determined?

- 1. Budding
- a. Recognition of membrane proteins exposed to cytosolic side by
- ARF or Sar proteins
- b. Type of coat protein recruited and assembled

# 2. Transport

- motor -cargo recognition by motor receptor on the membrane
- 3. Docking and fusion
- a. Rab GTPases (distinct) and their specific effectors
- b. SNARE pairs

Arf-, Sar-, Rab-GTPases serve as membrane organizers