

# ***Coatopathies: Genetic Disorders of Protein Coats***

**Juan S. Bonifacino, PhD**

***Section on Intracellular Protein Trafficking***

***Cell Biology and Neurobiology Branch***



***Eunice Kennedy Shriver National Institute  
of Child Health and Human Development***



# ***Section on Intracellular Protein Trafficking***

---

# ***Section on Intracellular Protein Trafficking***

---

- **Molecular mechanisms of protein and organelle distribution within the cell**

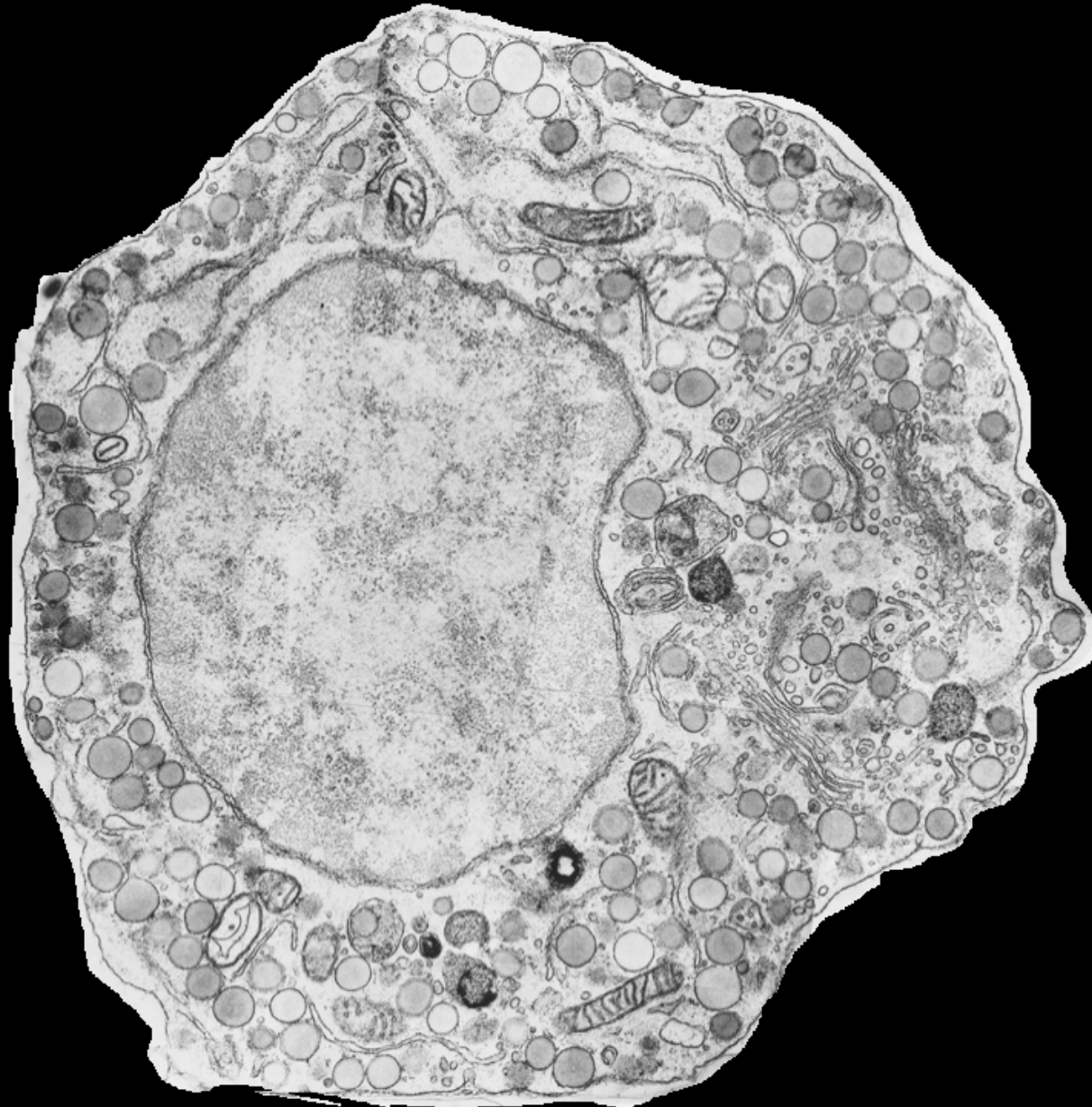
# ***Section on Intracellular Protein Trafficking***

---

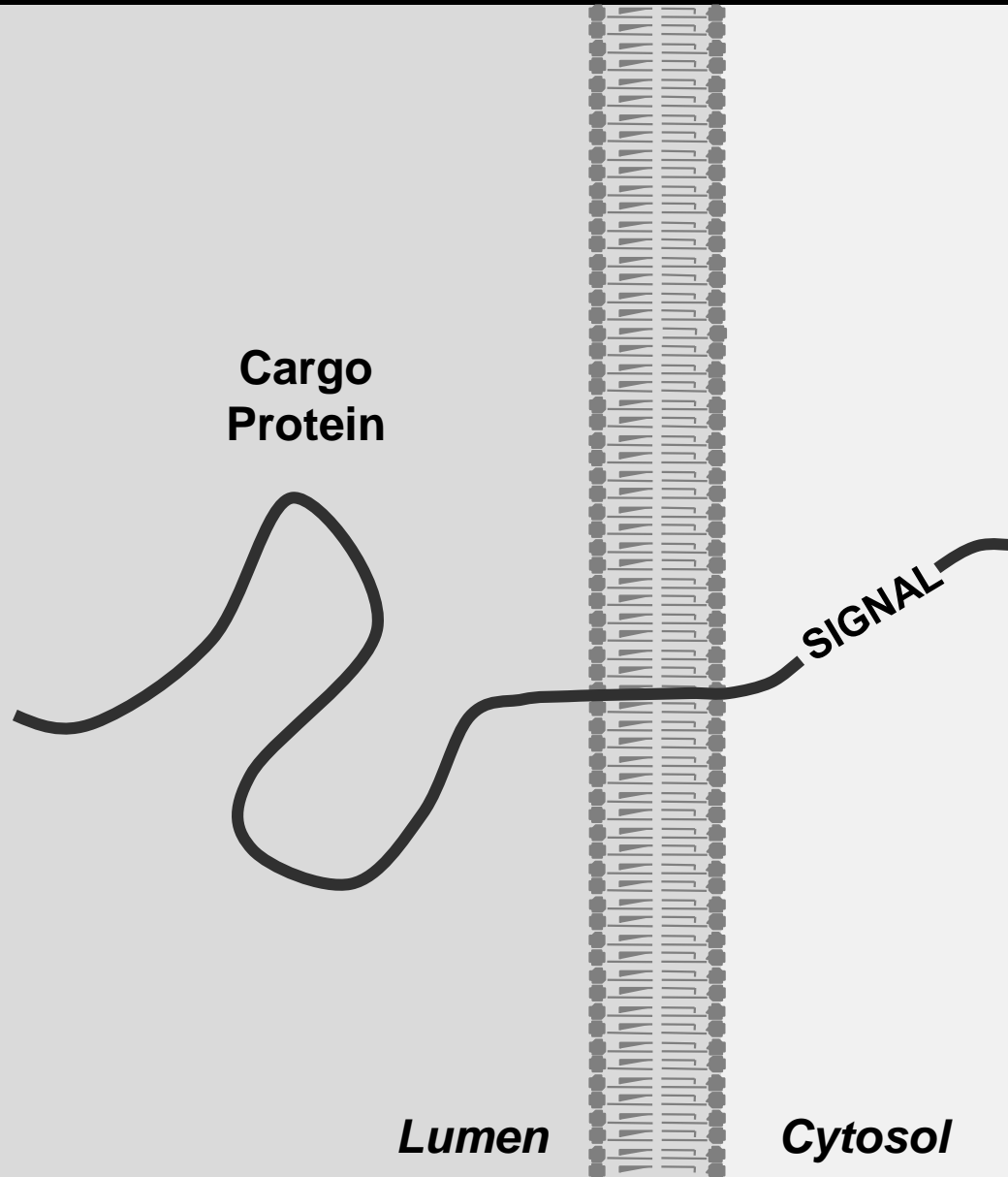
- **Molecular mechanisms of protein and organelle distribution within the cell**
- **Dysfunction in human disease**
  - Hermansky-Pudlak syndrome (HPS)**
  - MEDNIK syndrome**
  - Hereditary spastic paraplegias (HSP)**
  - Progressive cerebral cerebellar atrophy (PCCA)**

# *Electron Microscopy of a Pituitary Cell*

---



# Sorting Signals

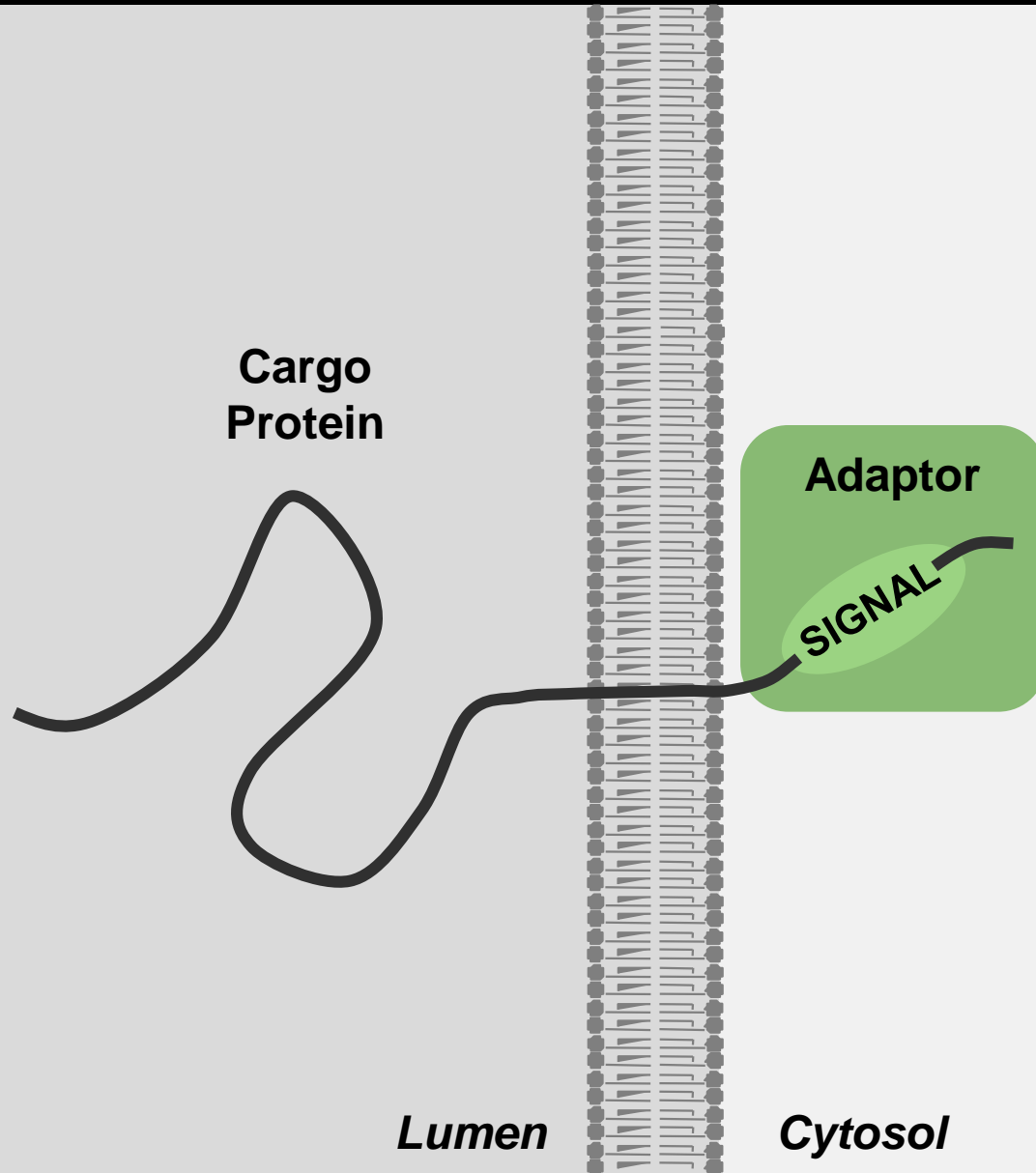




# Sorting Signals



# Adaptors

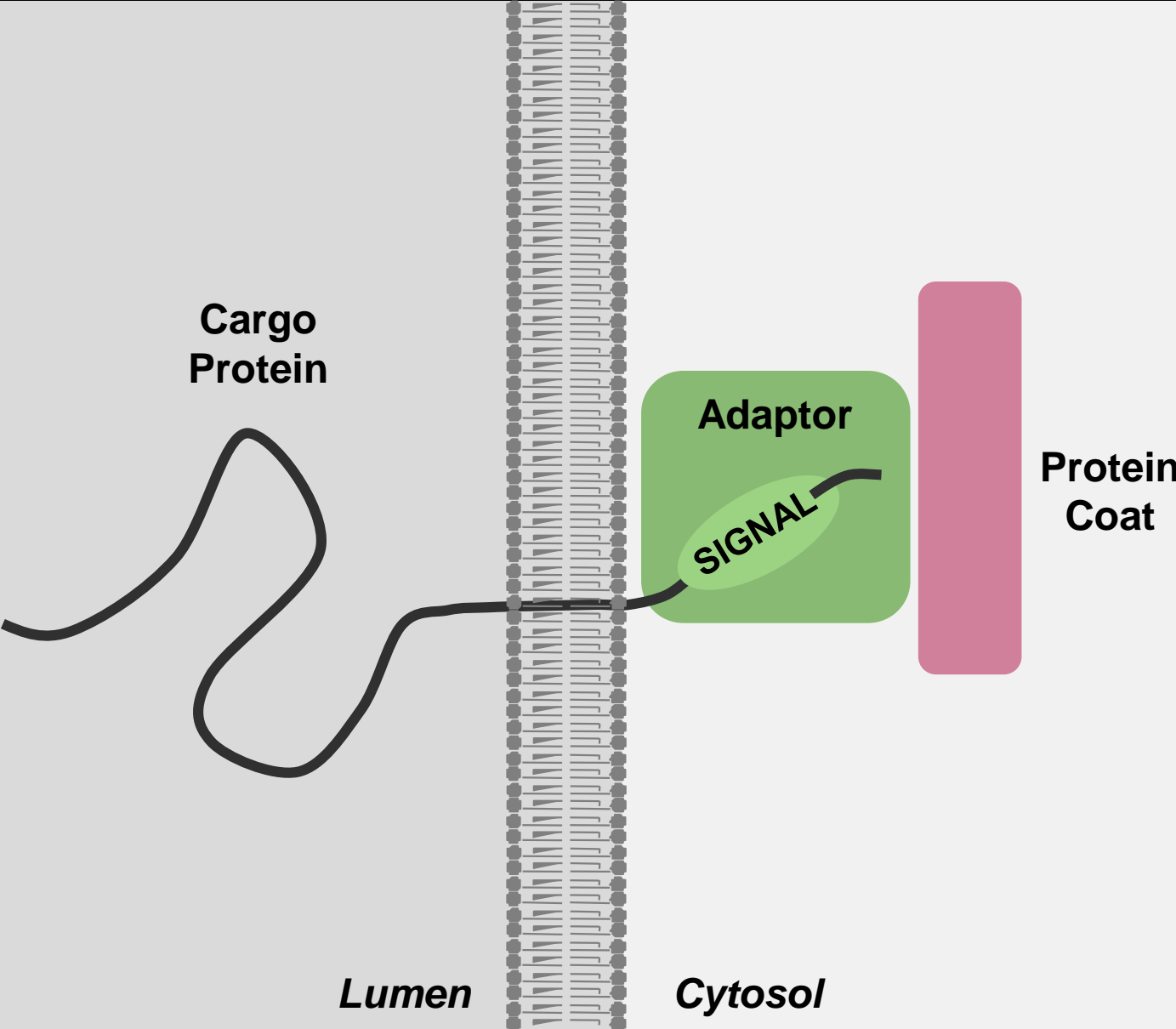




# Adaptors

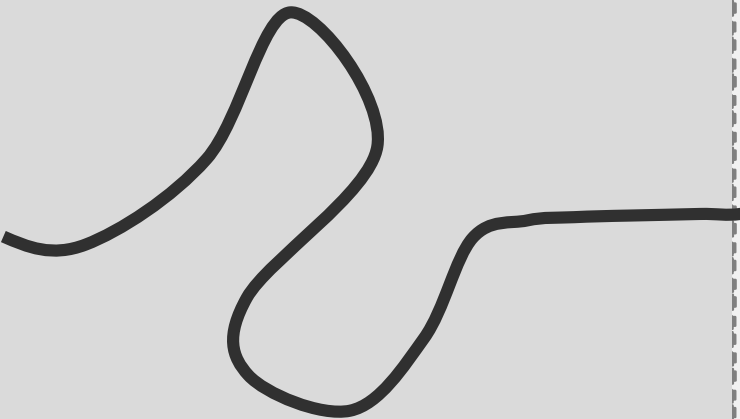


# Protein Coats

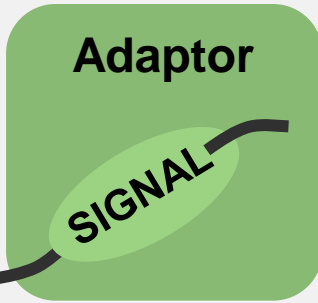


# Protein Coats

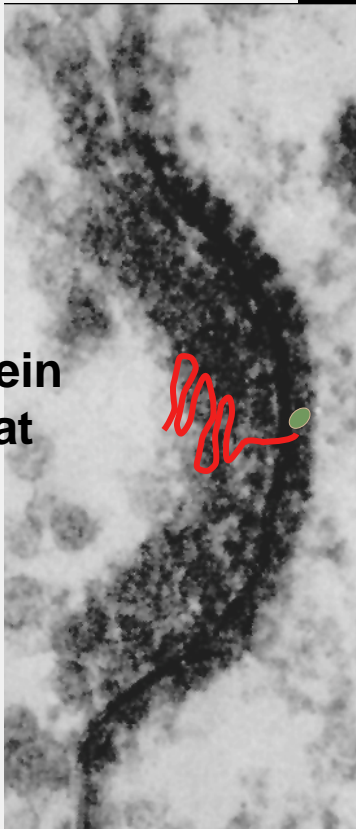
Cargo Protein



Adaptor



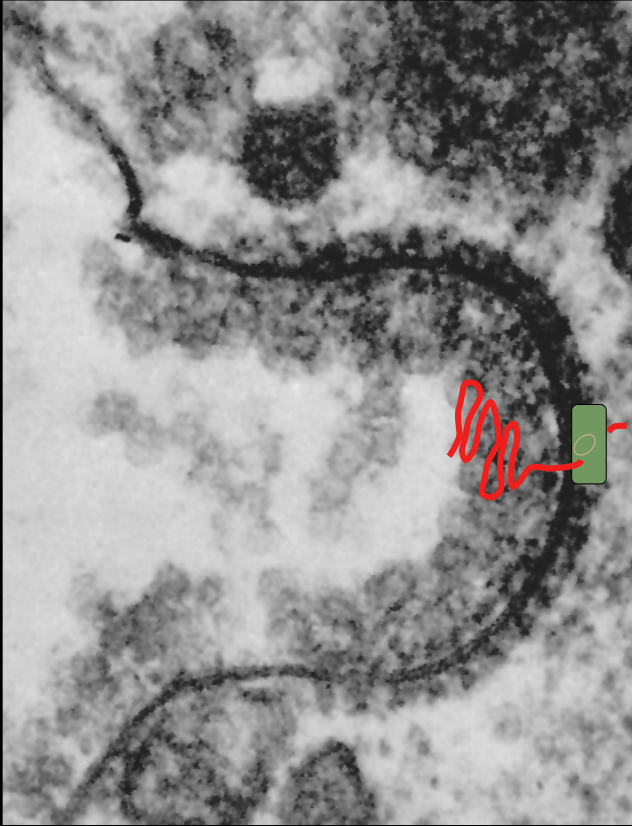
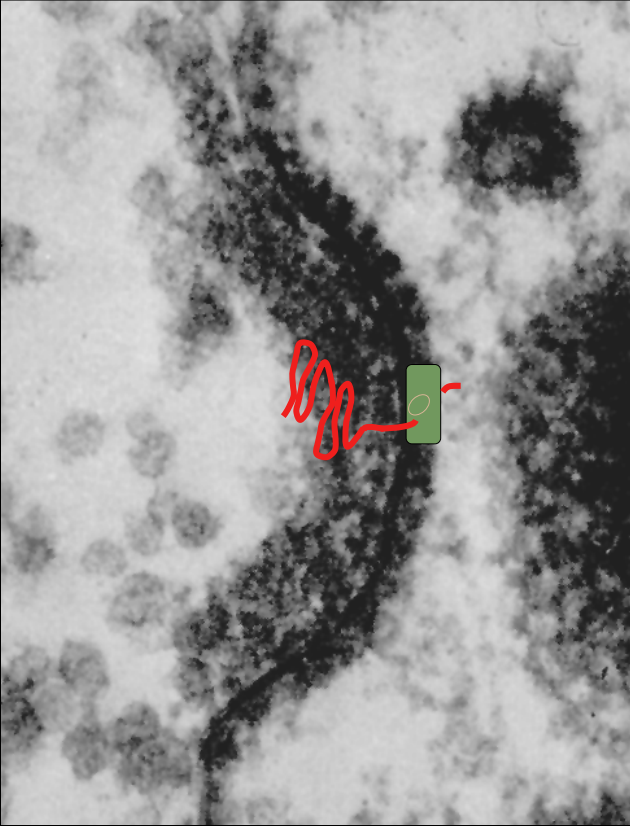
Protein Coat



Lumen

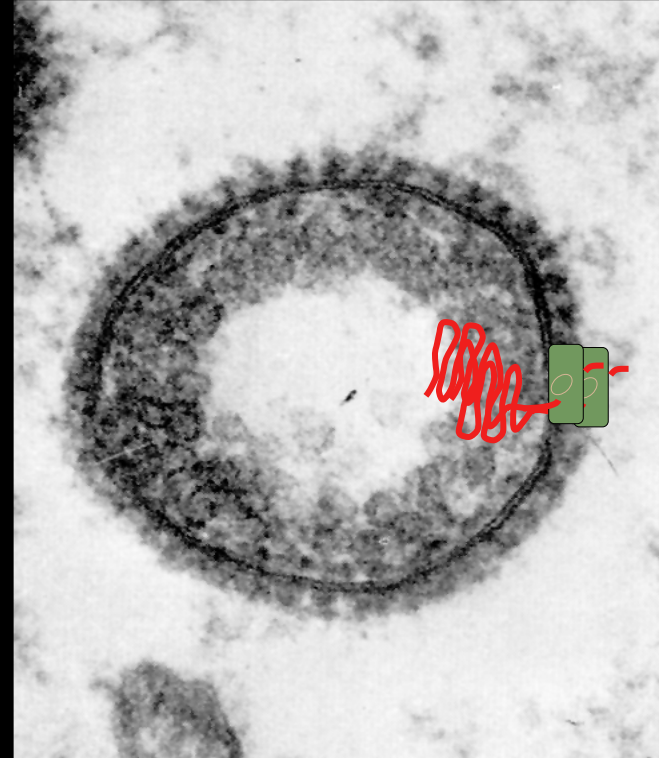
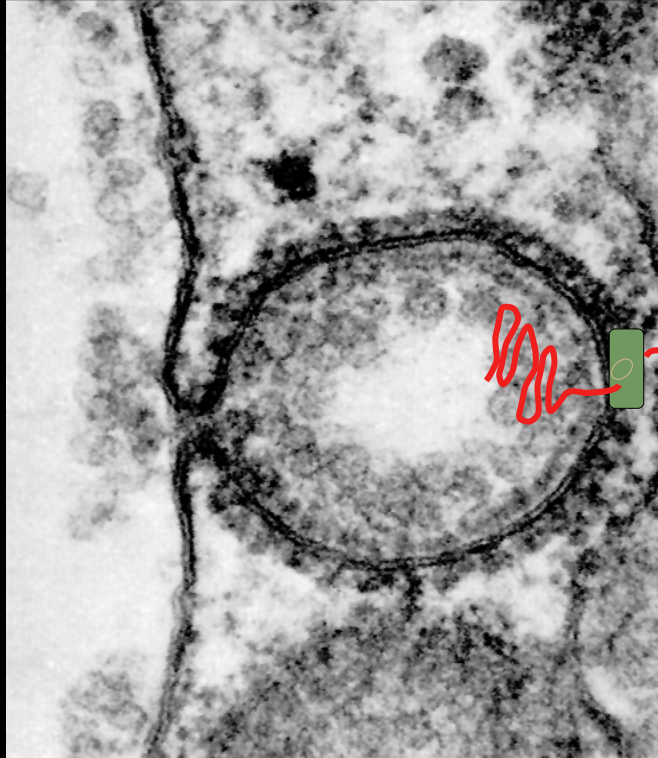
Cytosol

# Clathrin-coated Vesicles

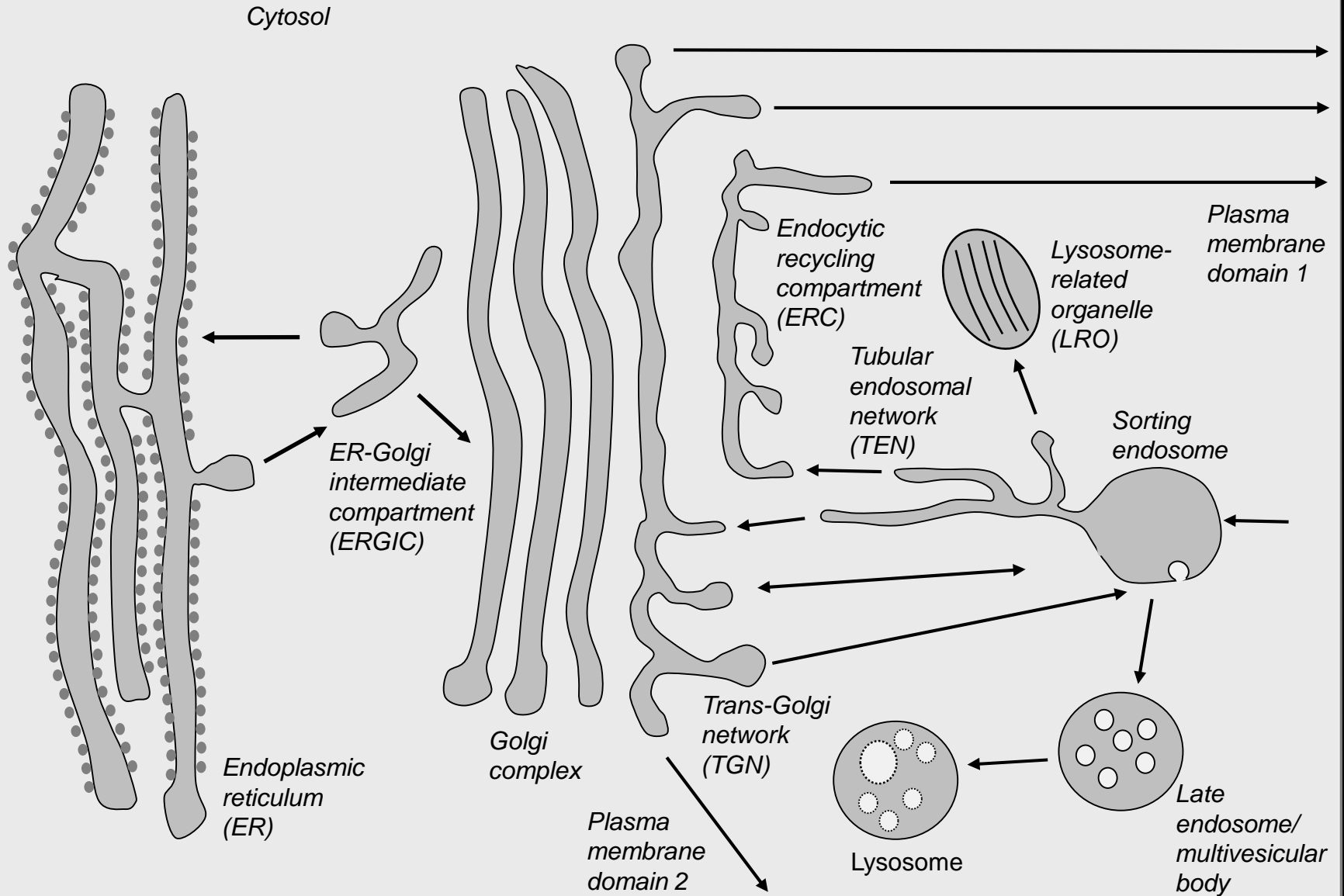




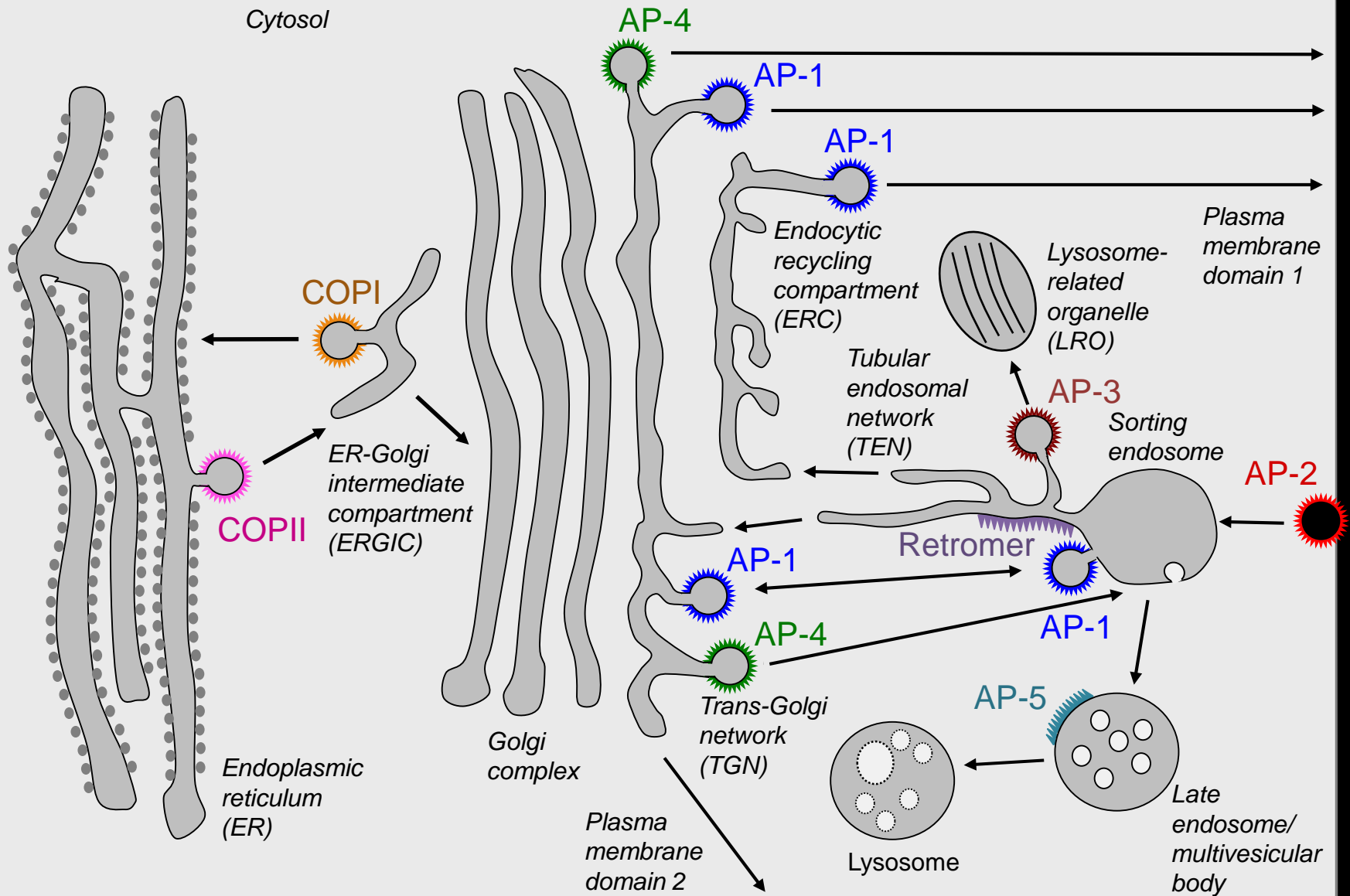
# Clathrin-coated Vesicles



# The Endomembrane System

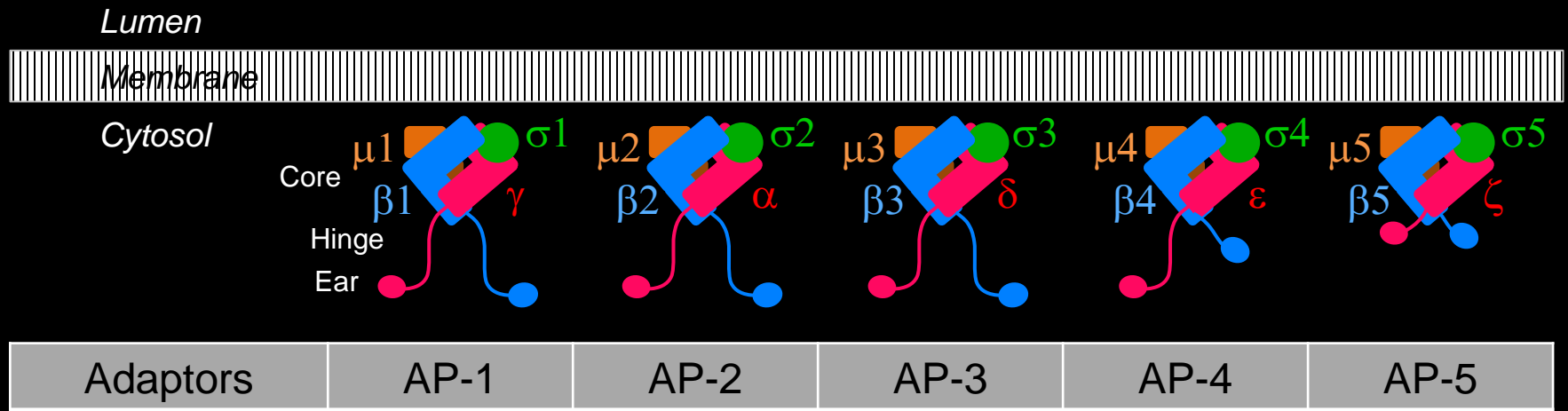


# Protein Coats in the Endomembrane System

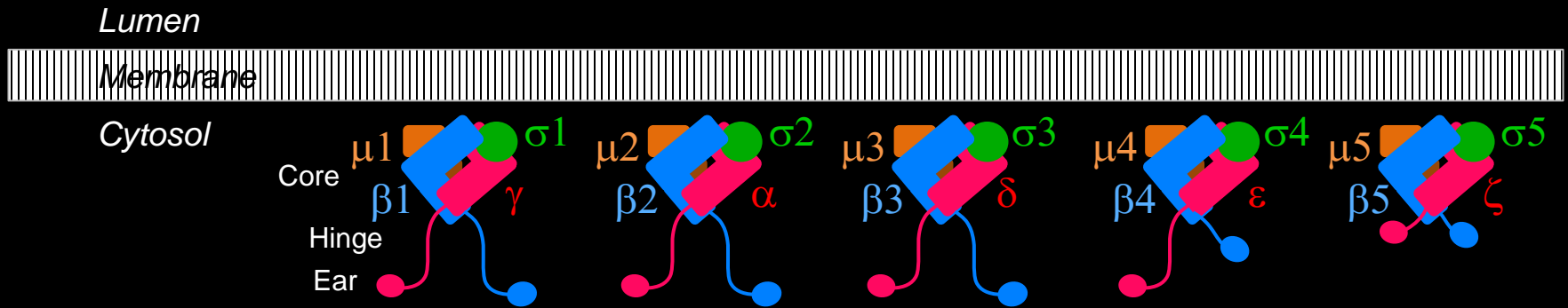




# Adaptor Protein (AP) Complexes

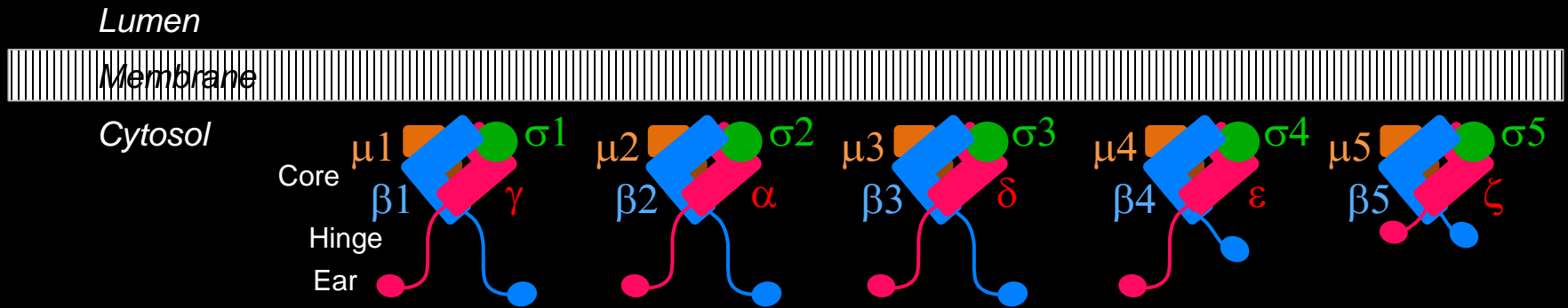


# Coatopathies



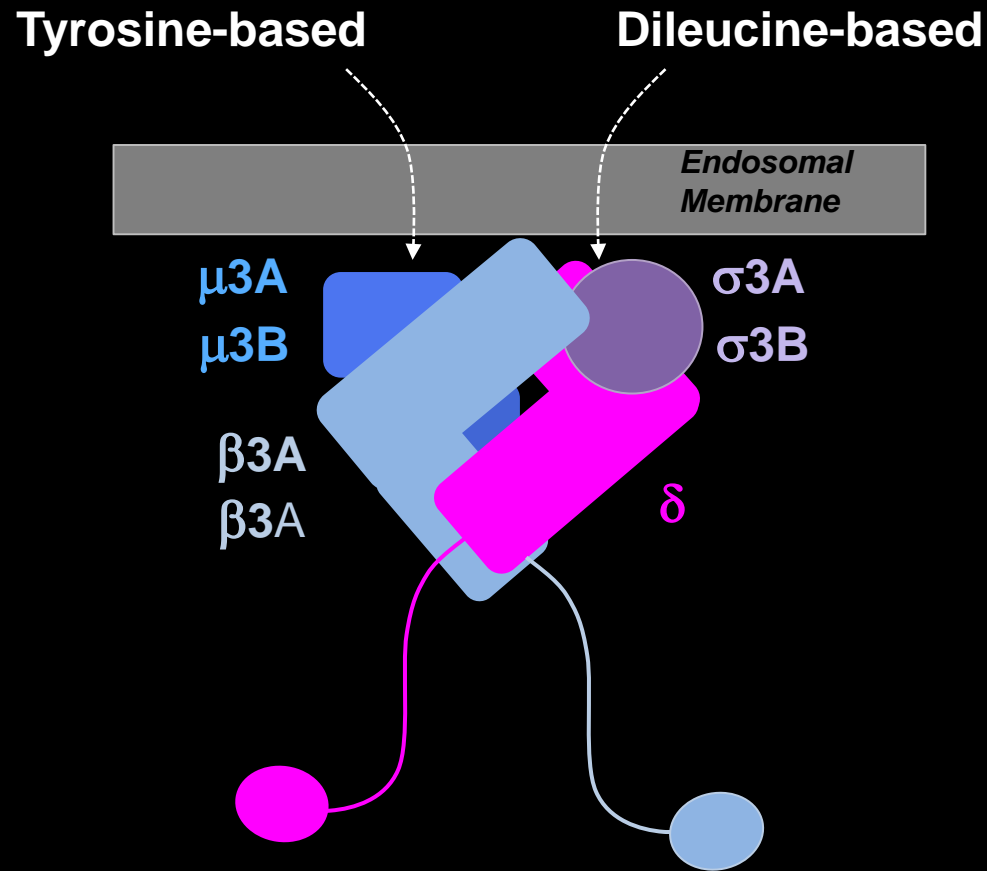
Adaptors	AP-1	AP-2	AP-3	AP-4	AP-5
Diseases (coatopathies)	<p>MEDNIK syndrome (<math>\sigma 1A</math>)</p> <p>Fried/Pettigrew syndrome (<math>\sigma 1B</math>)</p> <p>Pustular psoriasis 15 (<math>\sigma 1C</math>)</p>	<p>Hypocalciuric hypercalcemia type III (<math>\mu 2</math>)</p>	<p>Hermansky-Pudlak syndrome (HPS) type 2 (<math>\beta 3A</math>) and type 10 (<math>\delta</math>)</p> <p>Early onset epileptic encephalopathy (EOEE) type 48 (<math>\beta 3B</math>)</p>	<p>Hereditary spastic paraplegia (HSP) types 47 (<math>\beta 4</math>), 50 (<math>\mu 4</math>), 51 (<math>\epsilon</math>) and 52 (<math>\sigma 4</math>)</p>	<p>Hereditary spastic paraplegia type 48 (<math>\zeta</math>)</p>

# Coatopathies

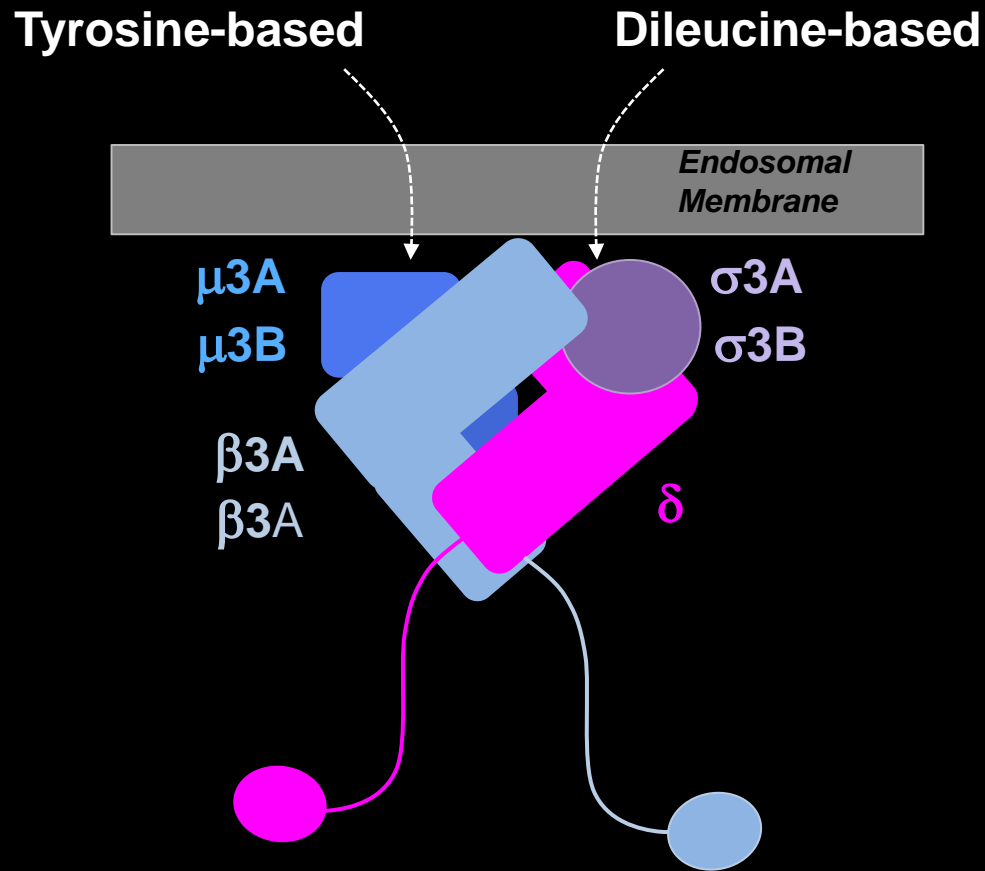


Adaptors	AP-1	AP-2	AP-3	AP-4	AP-5
Diseases (coatopathies)	<p>MEDNIK syndrome (<math>\sigma 1A</math>)</p> <p>Fried/Pettigrew syndrome (<math>\sigma 1B</math>)</p> <p>Pustular psoriasis 15 (<math>\sigma 1C</math>)</p>	<p>Hypocalciuric hypercalcemia type III (<math>\mu 2</math>)</p>	<p>Hermansky-Pudlak syndrome (HPS) type 2 (<math>\beta 3A</math>) and type 10 (<math>\delta</math>)</p> <p>Early onset epileptic encephalopathy (EOEE) type 48 (<math>\beta 3B</math>)</p>	<p>Hereditary spastic paraplegia (HSP) types 47 (<math>\beta 4</math>), 50 (<math>\mu 4</math>), 51 (<math>\epsilon</math>) and 52 (<math>\sigma 4</math>)</p>	<p>Hereditary spastic paraplegia type 48 (<math>\zeta</math>)</p>

# AP-3



# What is the Physiological Role of AP-3?



# BLAST Search Identifies Garnet As *Drosophila* AP-3 $\delta$

```
>gi|24641854|ref|NP_524785.2| garnet CG10986-PB [Drosophila melanogaster]
gi|22832217|gb|AAF48307.2| CG10986-PB [Drosophila melanogaster]
Length = 1034
```

Score = 912 bits (2358), Expect = 0.0

Identities = 497/848 (58%), Positives = 609/848 (71%), Gaps = 49/848 (5%)

```
Query: 1 MALKMVKGS-IDRMFDKNLQDLVRGIRNHKEDEAKYISQCIDEIKQELKQDNIAVKANAV 59
MALK VKG+ +RMFDKNL DLVRGIRN+K++EAKYIS CI+EIKQEL+QDNI+VK NAV
```

```
Sbjct: 1 MALKKVKGNFFERMFDKNLTDLVRGIRNNKDNEAKYISTCIEEIKQELRQDNISVKCNAV 60
```

```
Query: 60 CKLTYLQMLGYDISWAAFNIIEVMSASKFTFKRIGYLAASQSFHEGTDVIMLTTNQIRKD 119
KLT+Y+QMLGYDISWA FNIIIEVMS+S+FT KRIGYLAASQ FH ++++MLTTN IRKD
```

```
Sbjct: 61 AKLTYIQMLGYDISWAGFNIIIEVMSSSRFTCKRIGYLAASQCFHPDSELLMLTTNMIRKD 120
```

```
Query: 120 LSSPSQYDTGVALTGLSCFVTPDLARDLANDIMTLMSHTKPYIRKKAVLIMYKVFLKYPE 179
L+S +QYD GVAL+GLSCF++PDL+RDLANDIMTLM S TKPY+R KAVL+MYKVFL+YPE
```

```
Sbjct: 121 LNSQNQYDAGVALSGLSCFISPDLRDLANDIMTLMSSTKPYLRMKAVLMMYKVFLRYPE 180
```

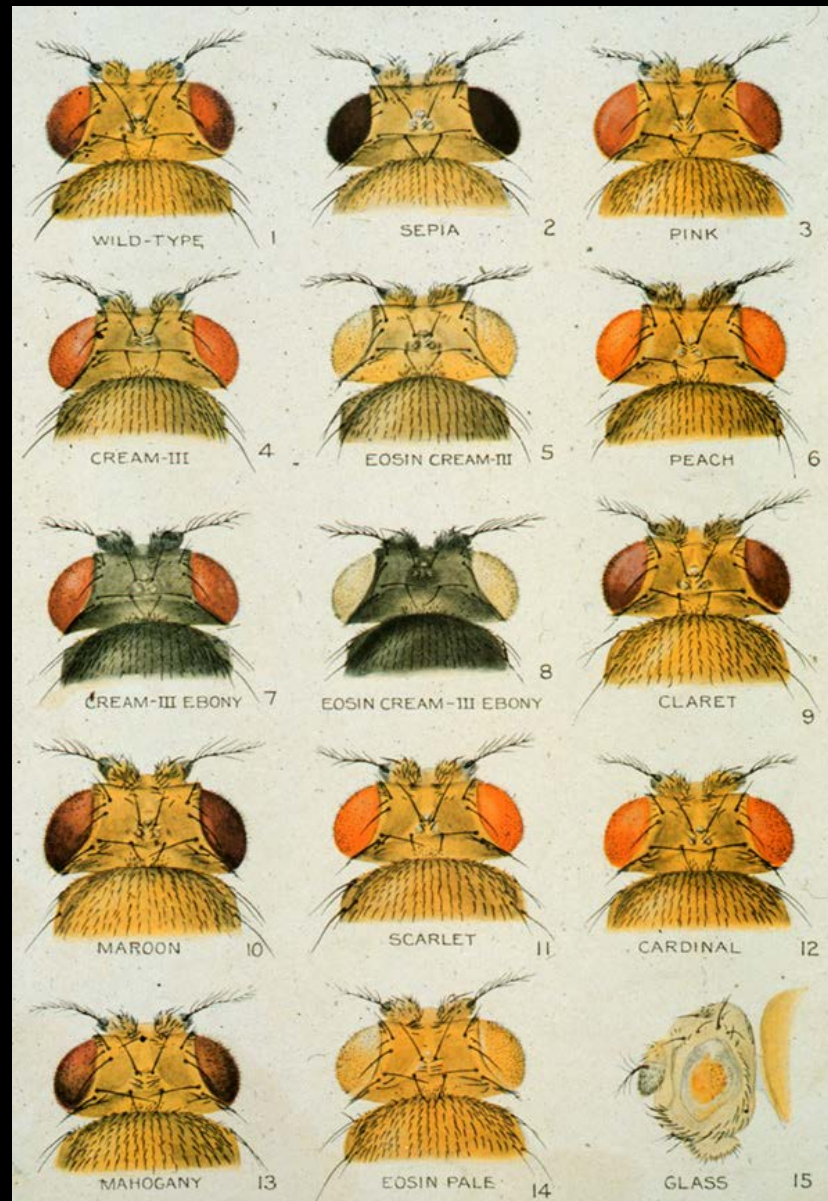
```
Query: 180 SLRPAFPRLKEKLEDPDPGVQSAAVNVICELARRNPKNYLSLAPLFFKLMTSSTNNWVLI 239
+LRPAFP+LKEKLEDPDPGVQSAAVNVICELAR+NPKNYL LAP+FFKLMT+STNNW+LI
```

```
Sbjct: 181 ALRPAFPKLKEKLEDPDPGVQSAAVNVICELARKNPKNYLP LAPIFFKLMTTSTNNWMLI 240
```

```
Query: 240 KIIKLFGALTPLEPRLGKKLIEPLTNLIHSTSAMSLLYECVNTVIAVLISLSSGMPNHSA 299
KIIKLFGALTPLEPRLGKKLIEPLTNLIHSTSAMSLLYEC+NTVIAVLIS+SSGMPNHSA
```

```
Sbjct: 241 KIIKLFGALTPLEPRLGKKLIEPLTNLIHSTSAMSLLYECINTVIAVLISISSGMPNHSA 300
```

# *Drosophila* Pigmentation Mutants



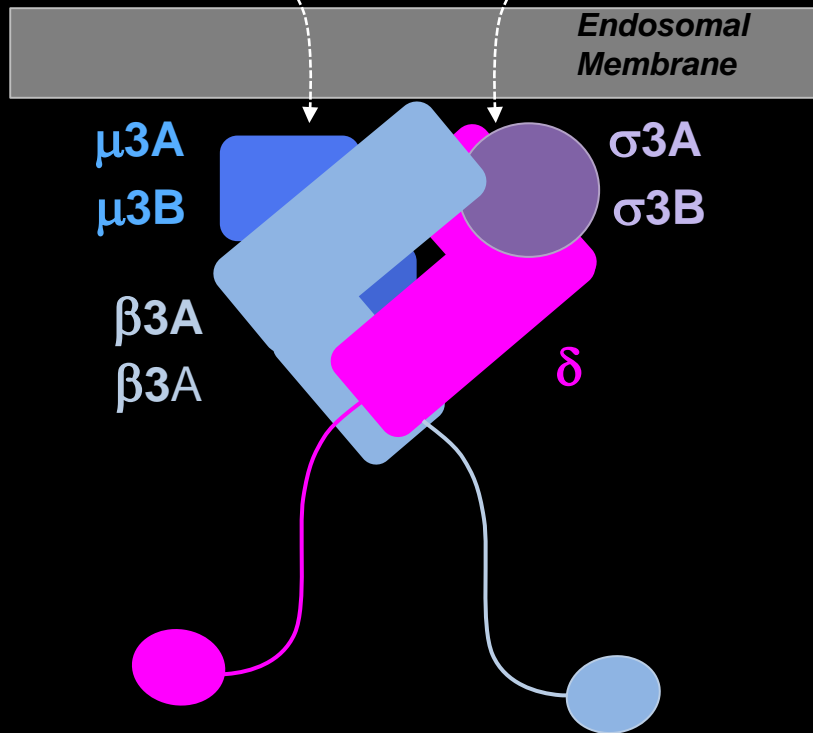
*Edith Wallace*



# AP-3 Defects in *Drosophila* Pigmentation Mutants

Tyrosine-based

Dileucine-based



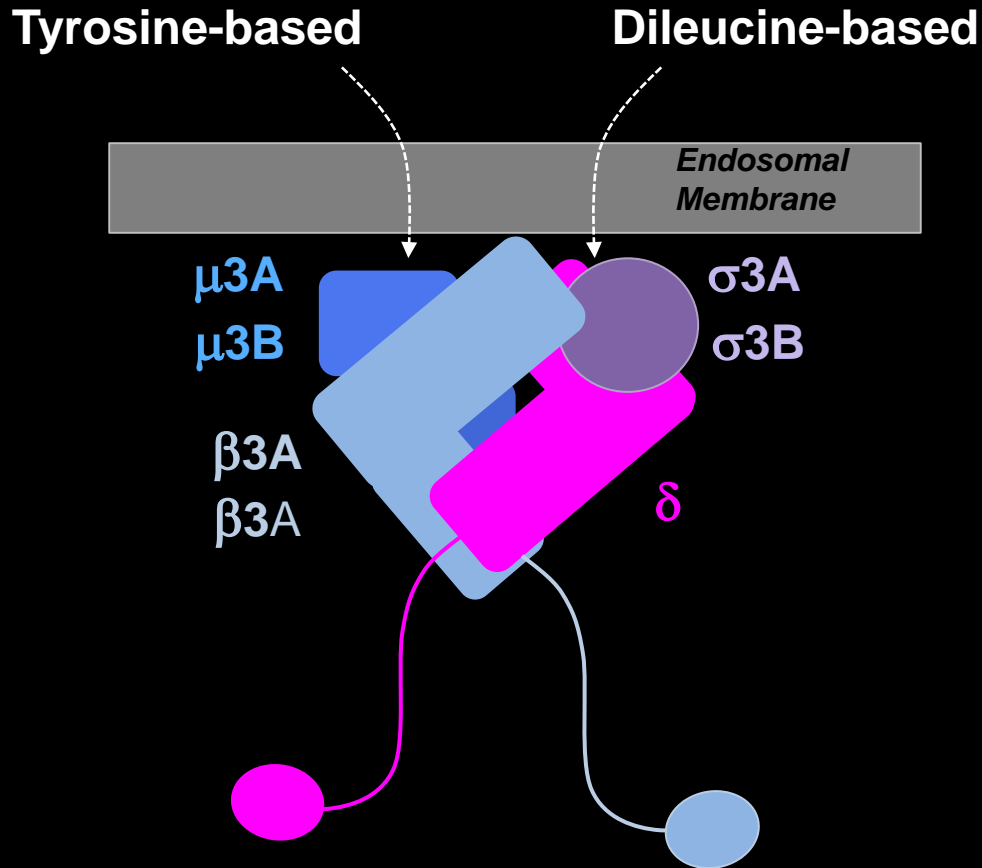
Subunit

$\delta$

Mutant

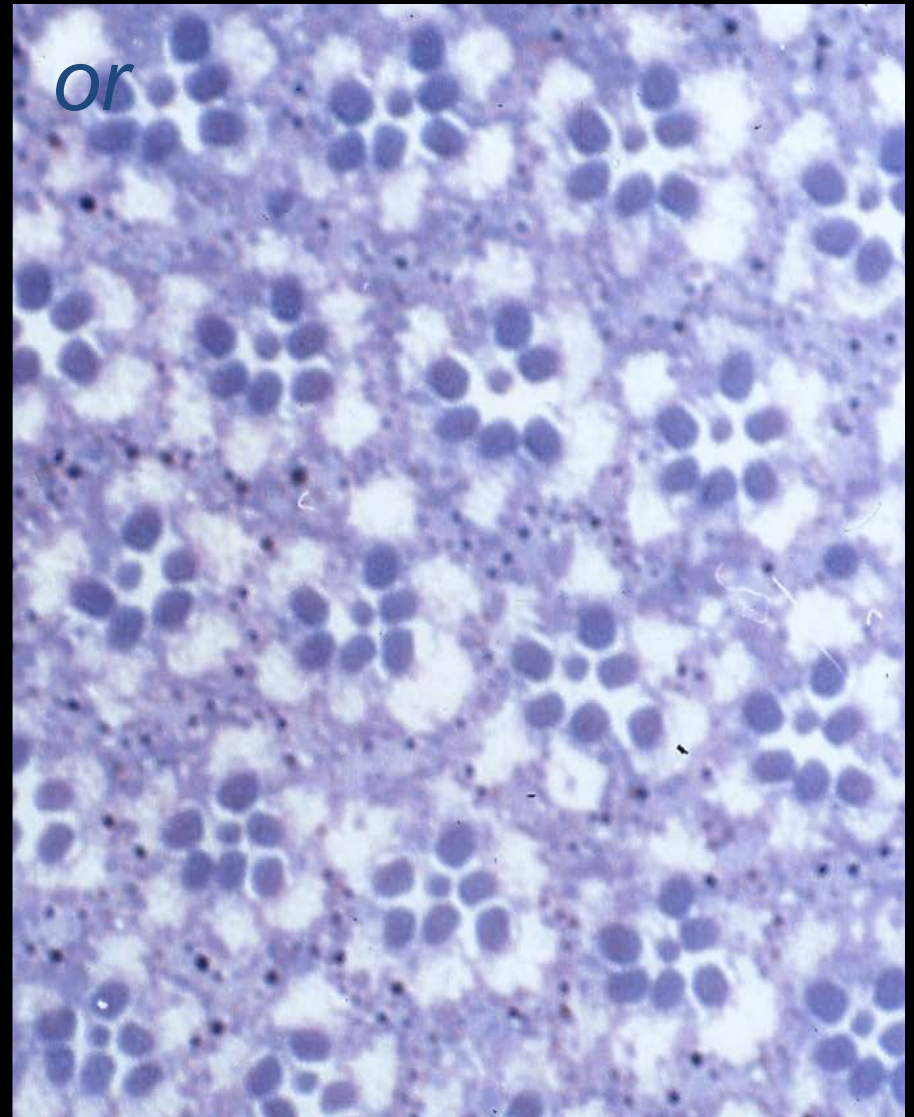
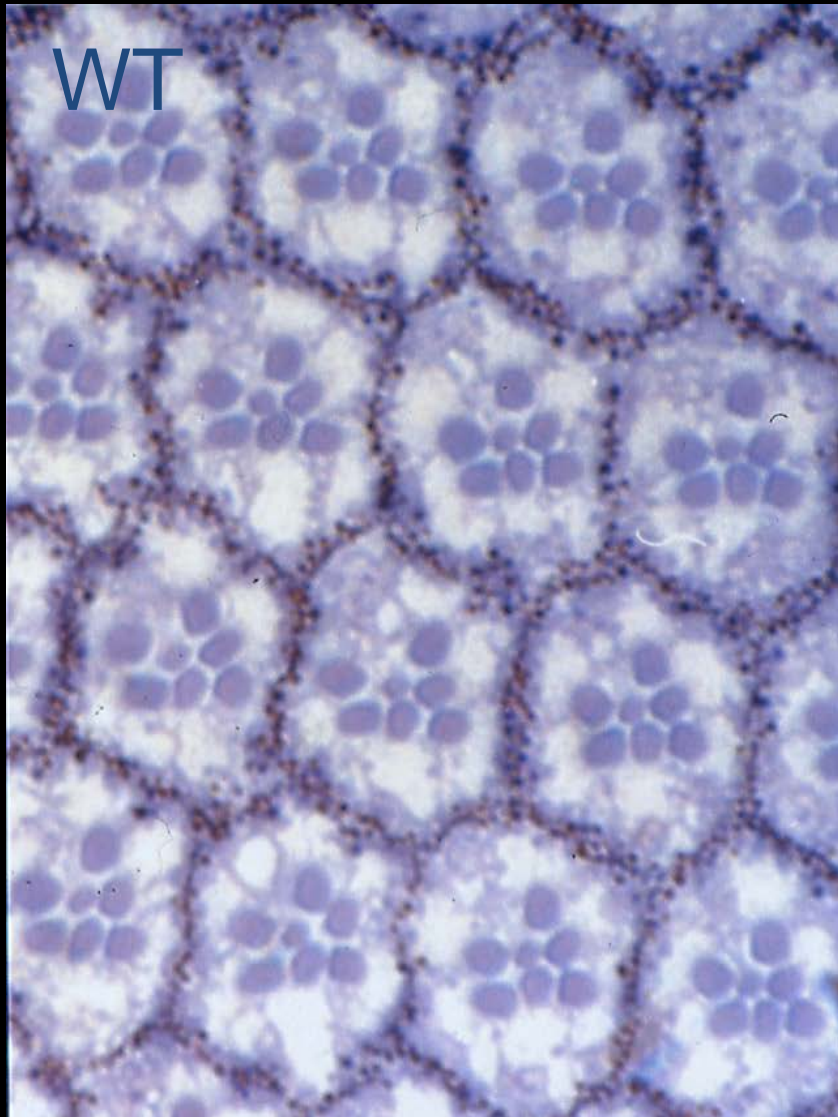
*garnet*

# AP-3 Defects in *Drosophila* Pigmentation Mutants

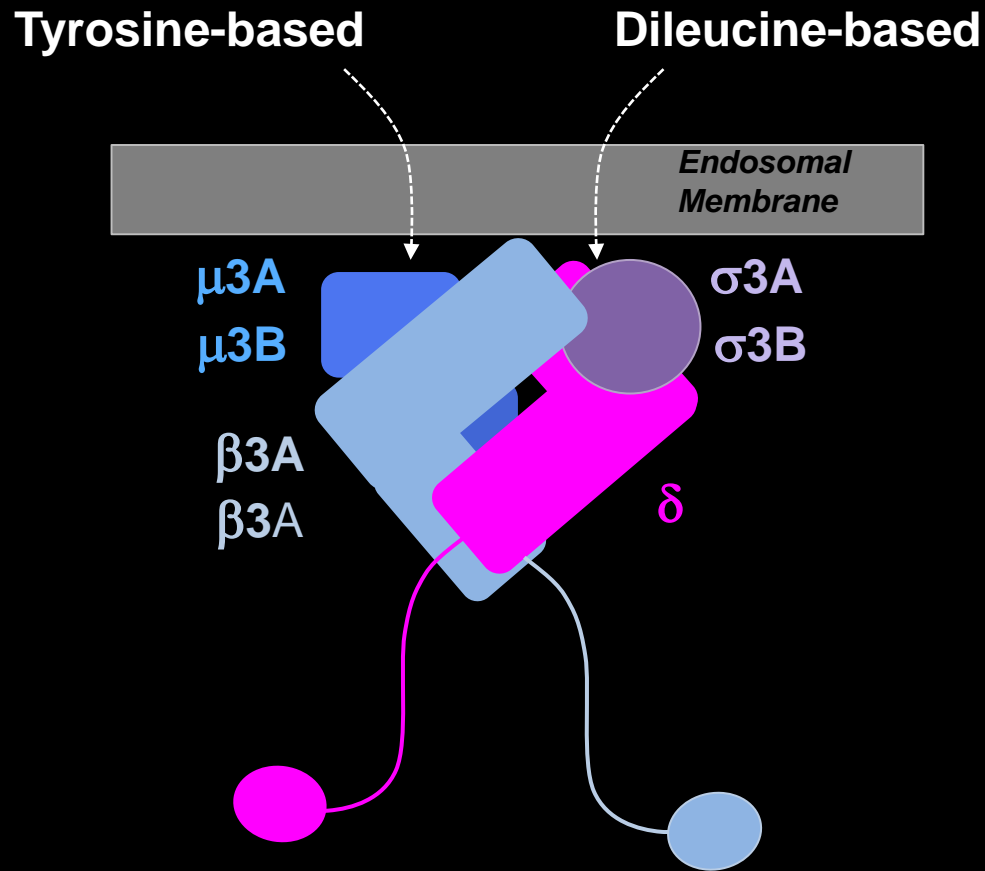


<u>Subunit</u>	<u>Mutant</u>
$\delta$	<i>garnet</i>
$\beta$ 3	<i>ruby</i>
$\mu$ 3	<i>carmine</i>
$\sigma$ 3	<i>orange</i>

# Sections of Wild-Type and Orange Eyes



# Could Mutations in AP-3 Cause Pigmentation Defects in Humans?



## Albinism

Richard A. King ■ Vincent J. Hearing  
Donnell J. Creel ■ William S. Oetting

1. Melanocytes represent cells, yet they are cells that account in the skin, hair, and number of critical steps processes either directly of these genes production, such as albinism
2. Melanocytes in the either directly, as in keratinocytes, or indirectly produced by other

### Hermansky Pudlak Syndrome Gene (*HPS1*)

Hermansky-Pudlak syndrome (HPS) (MIM 203300) is a very rare type of albinism, but two isolated populations in Puerto Rico and Switzerland, provided an opportunity to map one of the HPS loci (*HPS1*) to chromosome 10q23.<sup>60–62</sup> Both mapping strategies were based on a common founder for the mutated gene in these two isolates. The syntenic murine region contains the *pale ear* (*ep*) and *ruby eye* (*ru*), two candidate mouse models for HPS, and *ep* is the murine homologue of *HPS1*.<sup>63,64</sup> The gene consists of 20 exons and spans 30.5 kb.<sup>299</sup> In the initial isolation of the *HPS1* gene, a 3.6-kb transcript was identified that encodes a 700-amino acid protein with a size of 79.3 kDa.<sup>63,299</sup> Mutations in the 3' portion of this gene were associated with the HPS phenotype, indicating that this was the correct transcript responsible for the HPS phenotype.<sup>63</sup> A second 1.5-kb transcript of the *HPS1* gene that encodes a 324-amino acid protein was also identified.<sup>67</sup> The two cDNA transcripts are from the same gene and result from alternative splicing.<sup>67</sup> Both transcripts are polyadenylated, and contain transmembrane domains and a putative melanosomal localization signal. The 3.6-kb transcript codes for a transmembrane protein containing two transmembrane domains, where the 1.5-kb transcript is missing the transmembrane domain in the carboxy end of the protein. There are no homologies between the putative *HPS1* gene product and known proteins and its function are currently unknown. Initial analysis using confocal immunofluorescence has shown a cytoplasmic and membrane-associated distribution of the protein, suggesting a soluble and nonsoluble

and has been identified as a lysosomal trafficking regulator (LYST). This would explain the abnormal trafficking of melanosomal proteins found in the melanocytes. Several mutations in the mouse have been described.<sup>304</sup> The *bg<sup>11</sup>* is a 5-kb deletion at the 3' end of the gene that disrupts three exons, which would result in a truncated protein with probable splicing abnormalities as well. The *bg<sup>21</sup>* mutation has a drastically reduced level of transcription, due to a 116-bp insertion of a LINE1 sequence in the coding region producing a truncated protein.<sup>303</sup> The *bg<sup>81</sup>* allele has a nucleotide substitution of a C to T at bp 2027, producing a nonsense mutation that is predicted to lack 1442 amino acids.<sup>304</sup>

### Ocular Albinism (OA1) (MIM 300500)

The ocular albinism type 1 gene (OA1; Nettleship-Falls X-linked OA) maps to chromosome Xp22.<sup>305,306</sup> Both the human (*OAI*) and the murine (*Moa1*) genes have been isolated.<sup>307,308</sup> In humans, the *OAI* gene is divided into 9 exons within a 40-kb region.<sup>307</sup> The gene codes for a protein of 424 amino acids that contains several putative transmembrane regions.<sup>307,309,310</sup> The amino acid sequence does not share identity with any known proteins and its function is unknown. The gene is expressed almost exclusively in the retinal pigment epithelium<sup>307</sup> and cutaneous melanocytes,<sup>310</sup> and at a much lower level in the brain and adrenal tissues.<sup>307</sup> Although the clinical manifestations involve primarily the eye, the protein is a membrane glycoprotein localized to the melanosome in ocular and cutaneous melanocytes, indicating OA1 is really a type of OCA with changes in the eye and skin melanocytes.<sup>310</sup>

# *Hermansky-Pudlak Syndrome (HPS)*

---

# *Hermansky-Pudlak Syndrome (HPS)*

---

- **Autosomal recessive disorder**



# *Hermansky-Pudlak Syndrome (HPS)*

---

- **Autosomal recessive disorder**
- **Oculocutaneous albinism**

# *Hermansky-Pudlak Syndrome (HPS)*

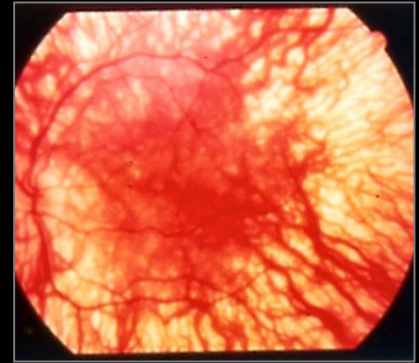
---



*Hair and skin*



*Iris*



*Retina*

# *Hermansky-Pudlak Syndrome (HPS)*

---

- **Autosomal recessive disorder**
- **Oculocutaneous albinism**

*Abnormal melanosomes*

# ***Hermansky-Pudlak Syndrome (HPS)***

---

- **Autosomal recessive disorder**
- **Oculocutaneous albinism**

***Abnormal melanosomes***

- **Prolonged bleeding**

***Absence of platelet dense granules***

# ***Hermansky-Pudlak Syndrome (HPS)***

---

- **Autosomal recessive disorder**
- **Oculocutaneous albinism**

***Abnormal melanosomes***

- **Prolonged bleeding**

***Absence of platelet dense granules***

- **Fibrosis of the lungs, inflammatory colitis**

# ***Hermansky-Pudlak Syndrome (HPS)***

---

- **Autosomal recessive disorder**
- **Oculocutaneous albinism**

***Abnormal melanosomes***

- **Prolonged bleeding**

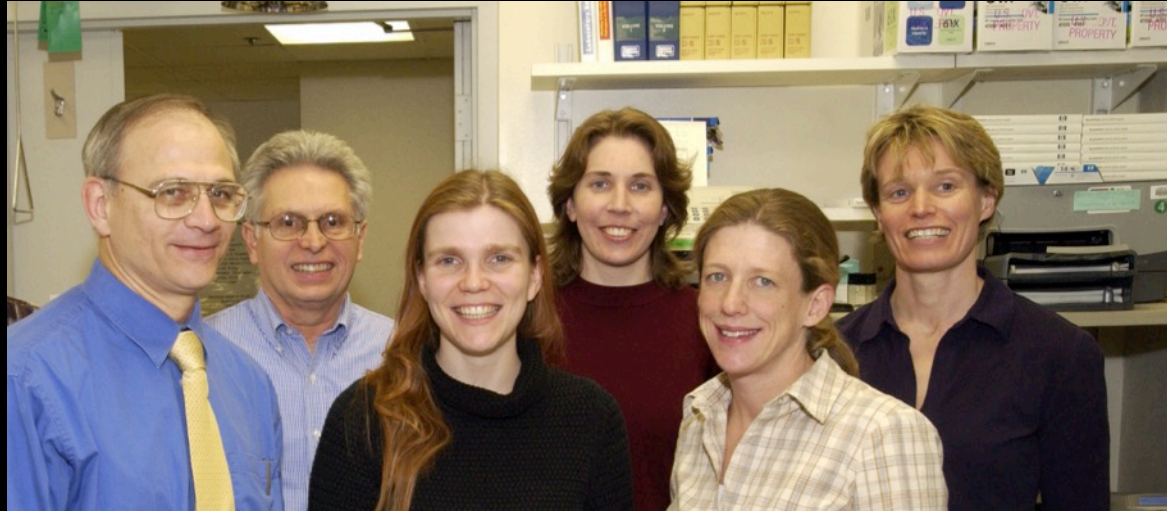
***Absence of platelet dense granules***

- **Fibrosis of the lungs, inflammatory colitis**

***Abnormal lung lamellar bodies, ceroid lipofuscin in macrophages***

# *William Gahl and Colleagues (NICHD/NHGRI)*

---





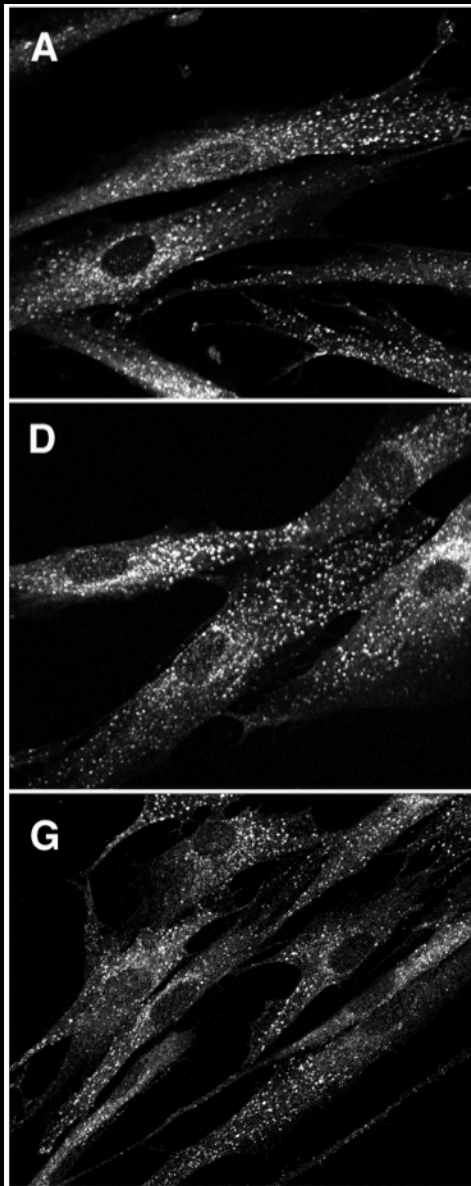
# HPS Patients



Photos Courtesy of Bill Gahl

# *AP-3 Defects in HPS Fibroblasts*

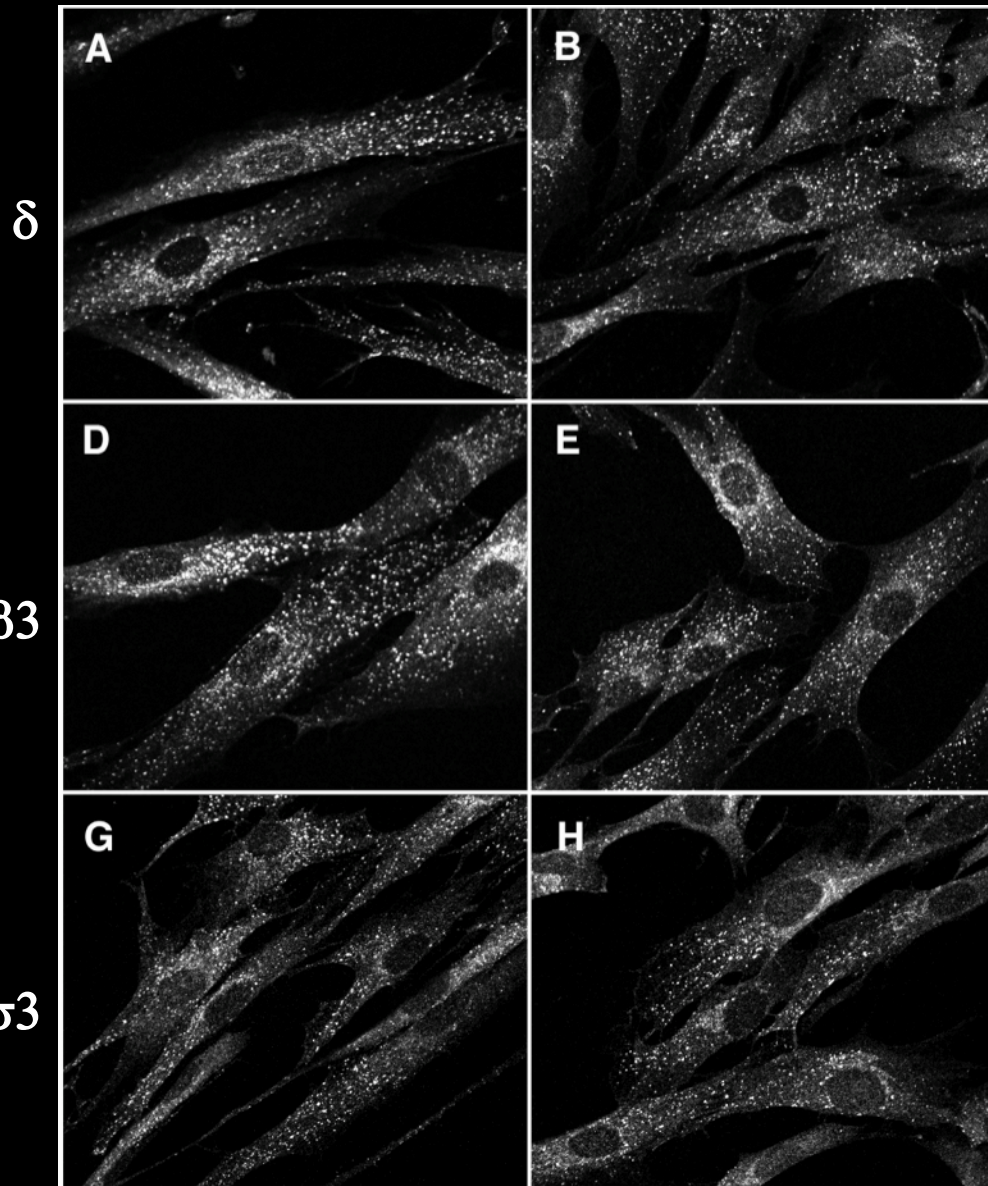
Normal



# AP-3 Defects in HPS Fibroblasts

Normal

Patient 8



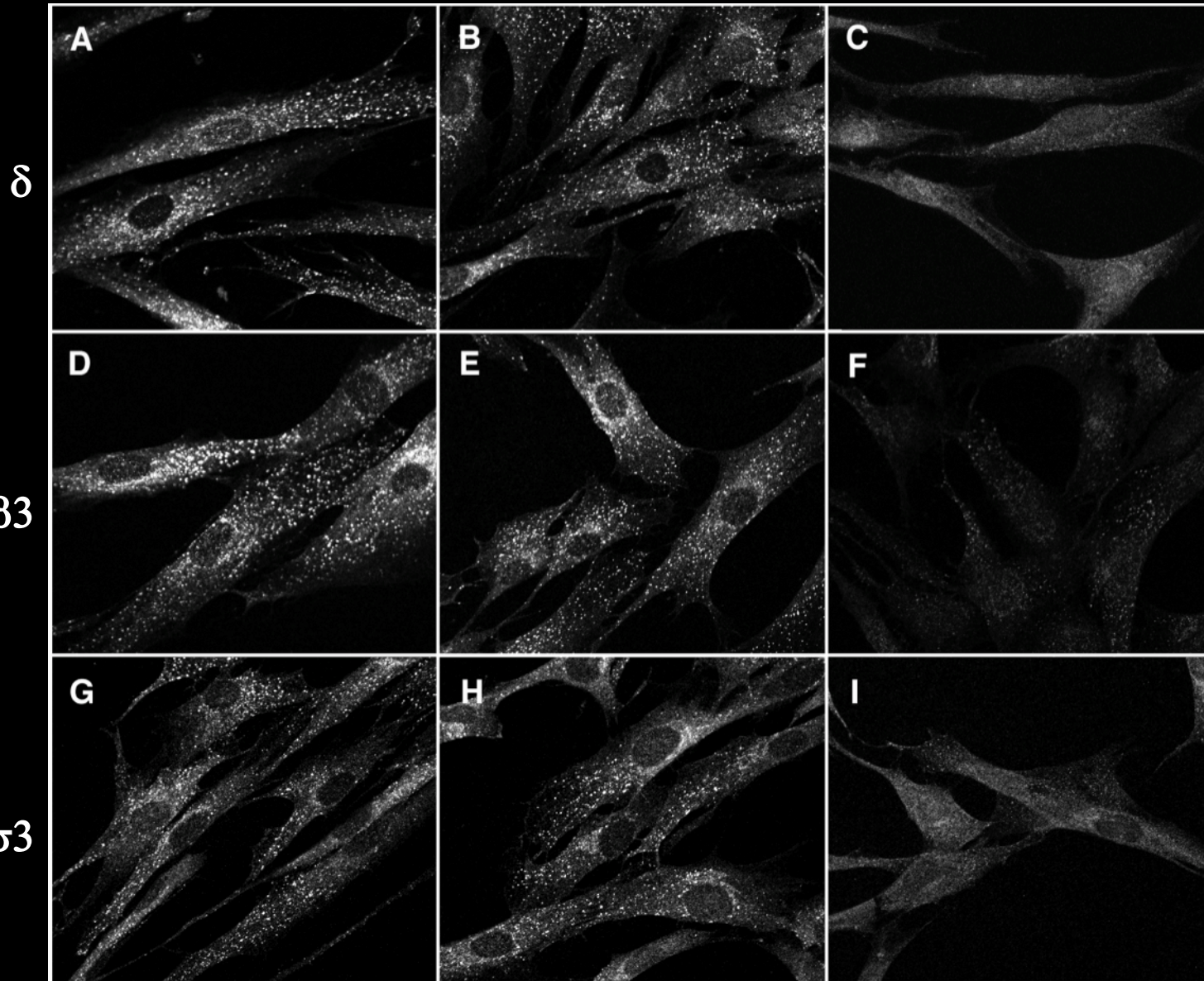


# AP-3 Defects in HPS Fibroblasts

Normal

Patient 8

Patient 40

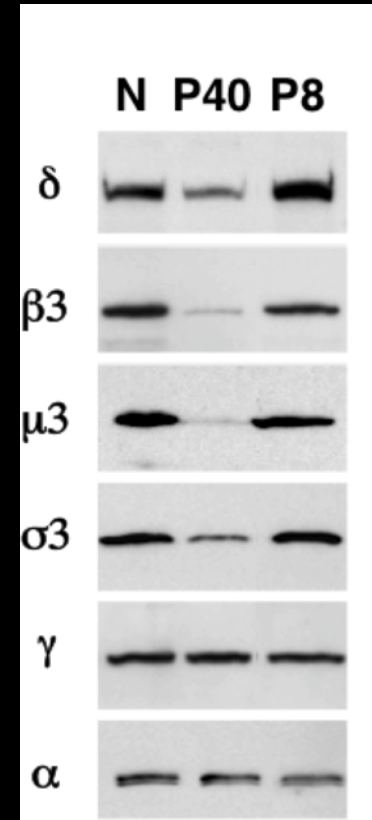
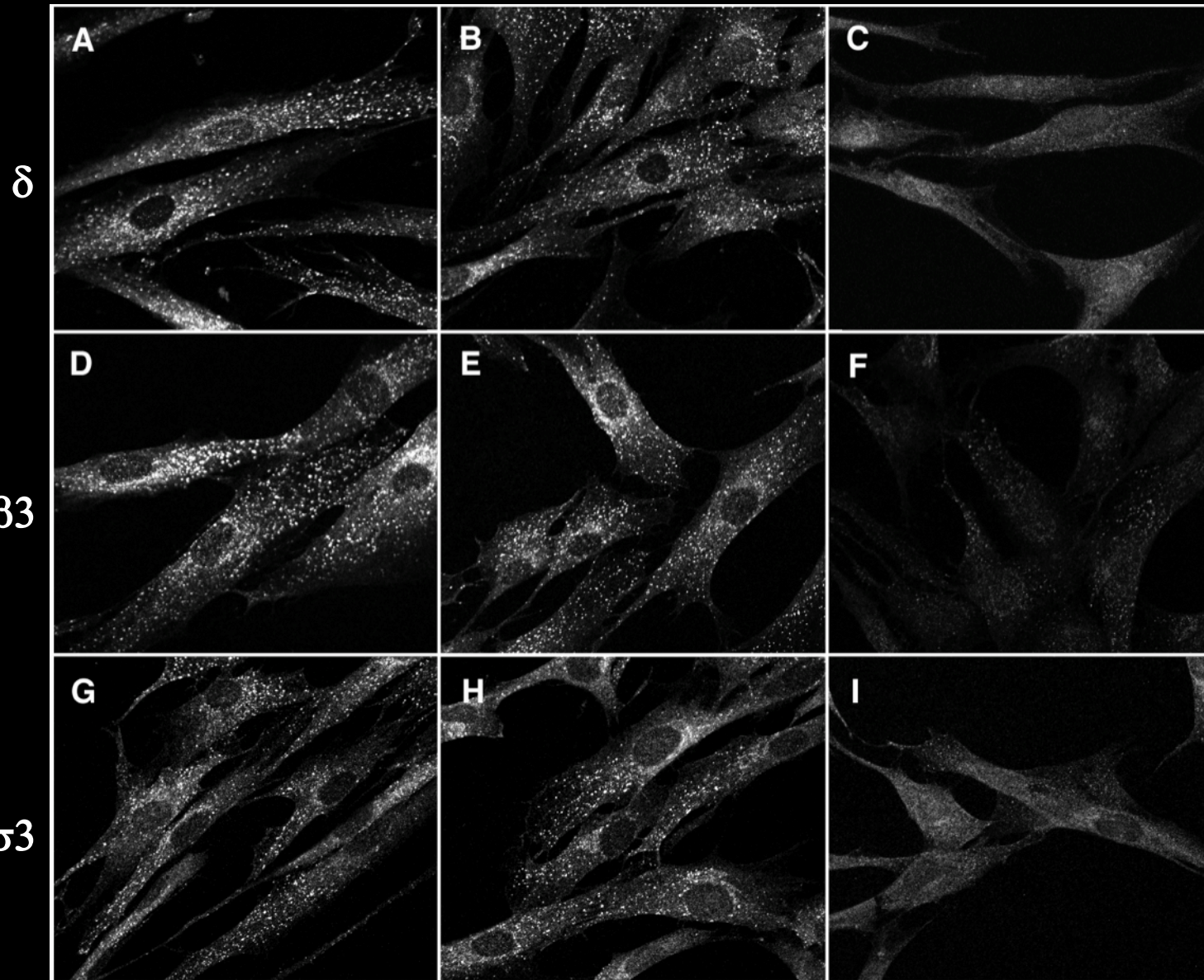


# AP-3 Defects in HPS Fibroblasts

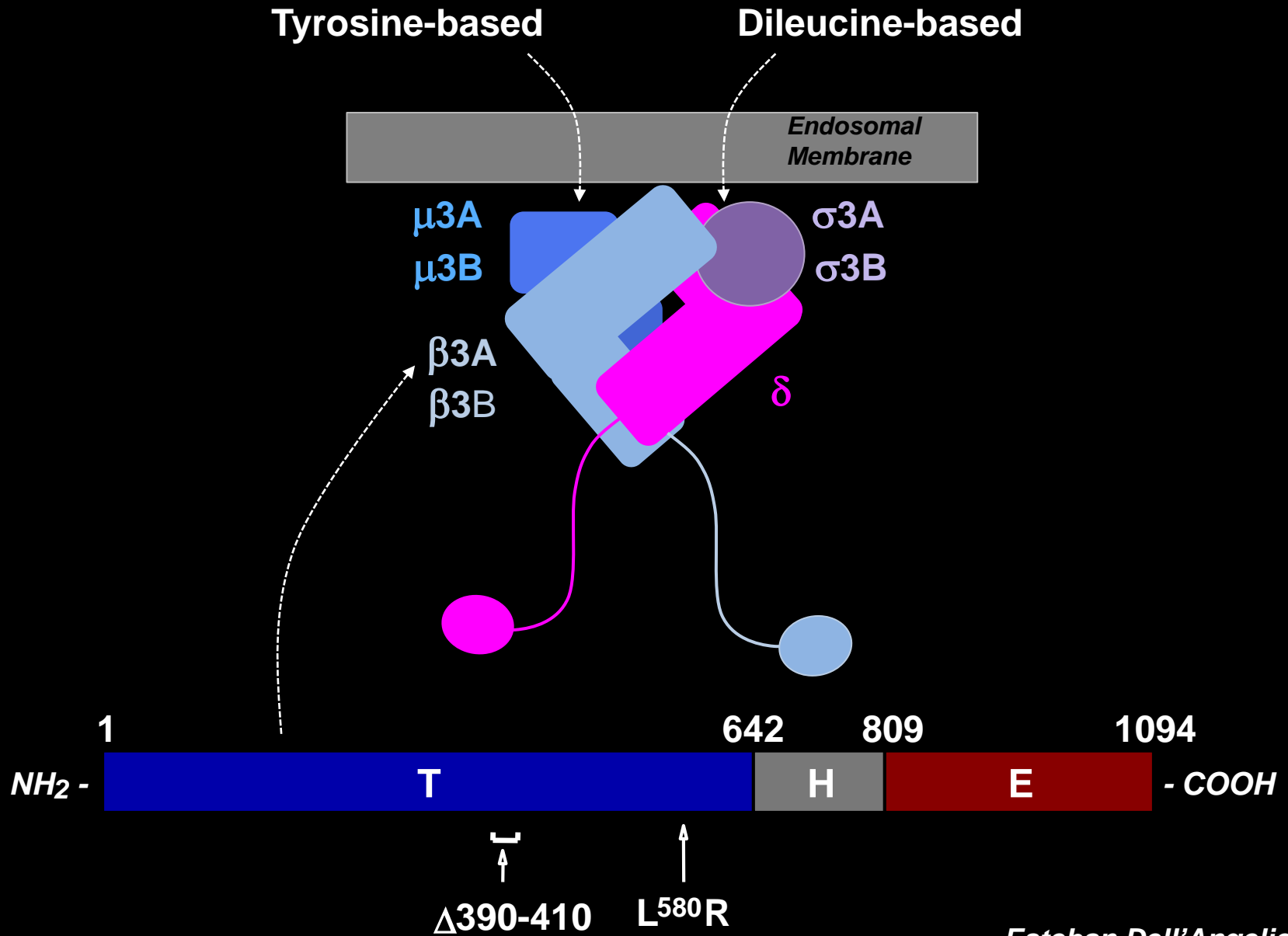
Normal

Patient 8

Patient 40

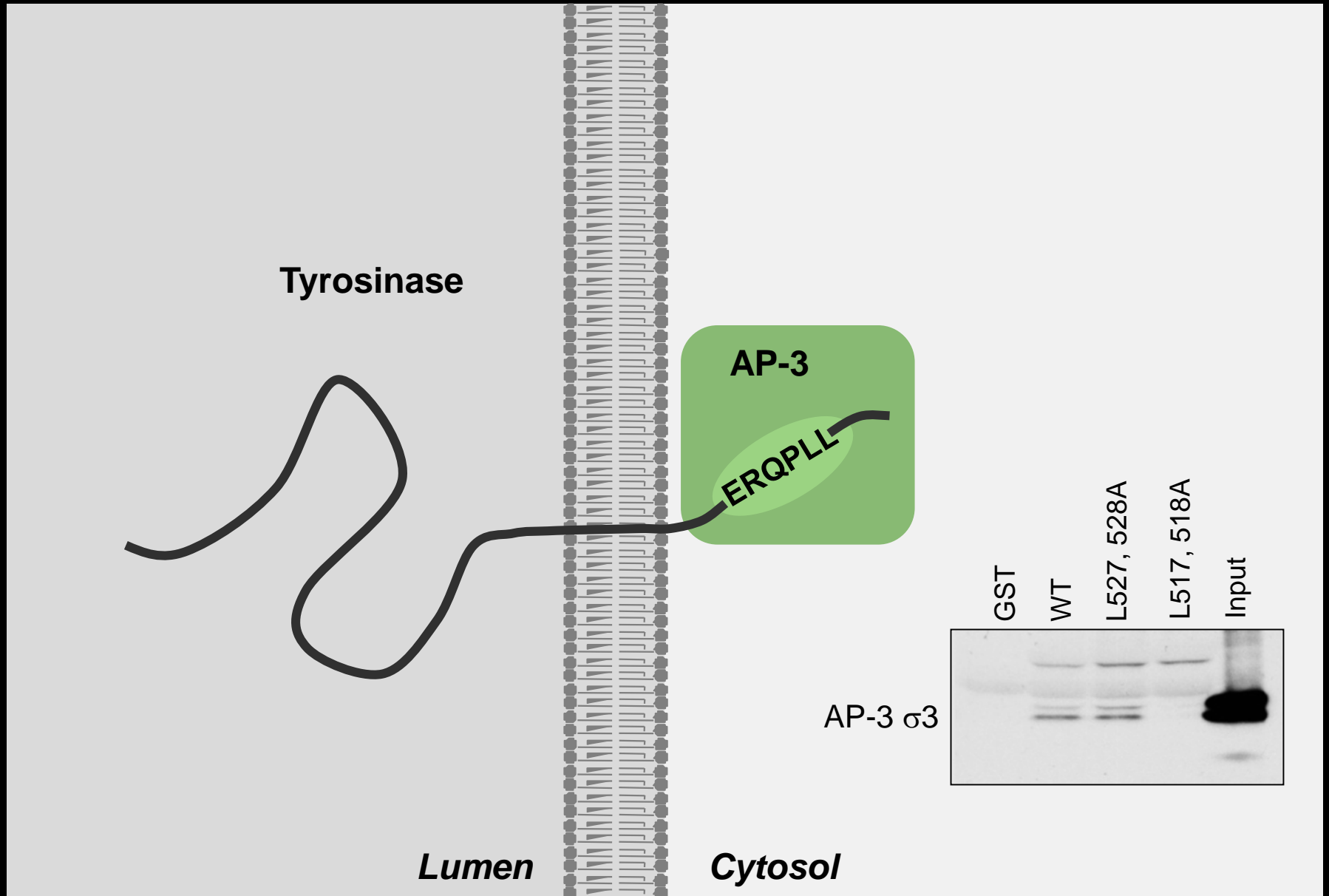


# Mutations in $\beta 3A$ in HPS-2

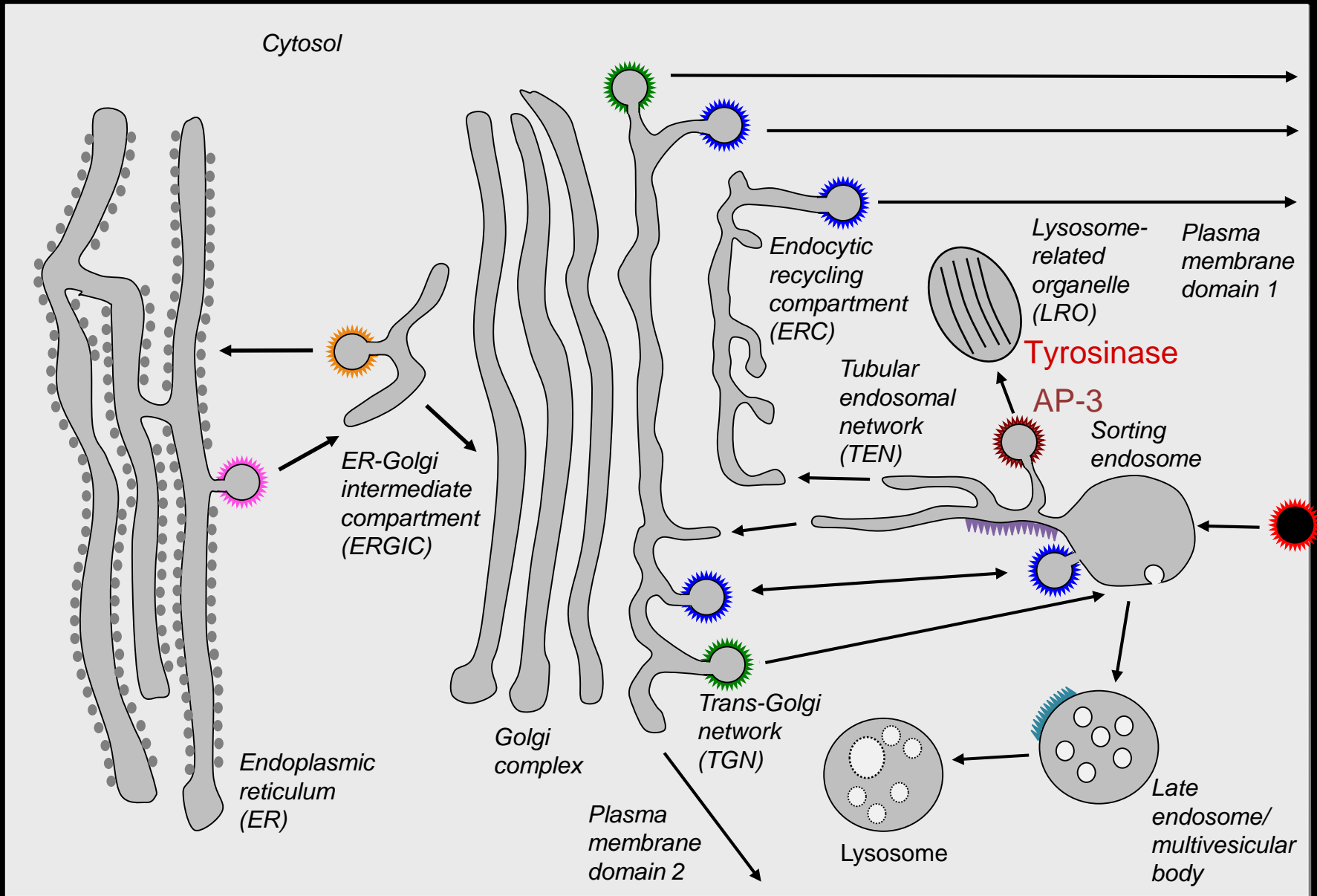




# Signal-Mediated Sorting of Tyrosinase by AP-3



# AP-3 Sorts Tyrosinase to Melanosomes





# Heterogeneity of HPS



# Human HPS Types

---



*Photo Courtesy of Bill Gahl*

**HPS1**

**HPS2**

**HPS3**

**HPS4**

**HPS5**

**HPS6**

**HPS7**

**HPS8**

**HPS9**

**HPS10**

# Human and Mouse HPS Types



*Photo Courtesy of Bill Gahl*

**HPS1**

**HPS2**

**HPS3**

**HPS4**

**HPS5**

**HPS6**

**HPS7**

**HPS8**

**HPS9**

**HPS10**

**Pale ear**

**Pearl**

**Cocoa**

**Light ear**

**Ruby eye-2**

**Ruby eye**

**Sandy**

**Reduced pigmentation**

**Muted**

**Mocha**

**Cappuccino**

**Pallid**

**Subtle gray**

**Gunmetal**



# Biogenesis of Lysosome-Related Organelles Complexes (BLOCs)

---

## BLOC-1

BLOS1

BLOS2

Snapin

Dysbindin

Pallidin

Muted

Cappuccino

BLOS3

## BLOC-2

HPS3

HPS5

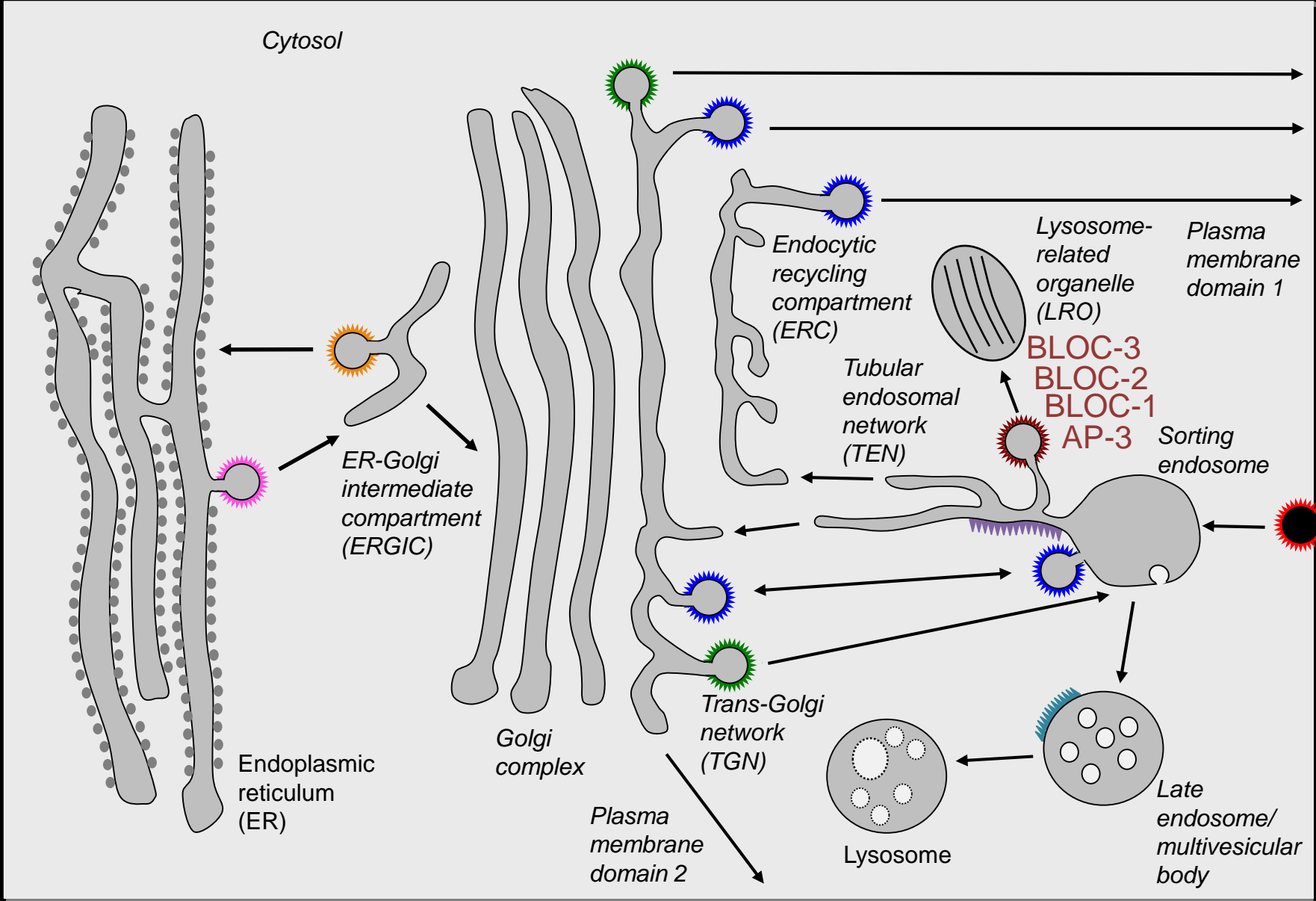
HPS6

## BLOC-3

HPS1

HPS4

# Biogenesis of Lysosome-Related Organelles Complexes (BLOCs)



# ***BLOC-One-Related Complex (BORC)***

---

## **BLOC-1**

**BLOS1**

**BLOS2**

**Snapin**

**Dysbindin**

**Pallidin**

**Muted**

**Cappuccino**

**BLOS3**

## **BLOC-2**

**HPS3**

**HPS5**

**HPS6**

## **BLOC-3**

**HPS1**

**HPS4**

## **BORC**

**BLOS1**

**BLOS2**

**Snapin**

**Myrlysin**

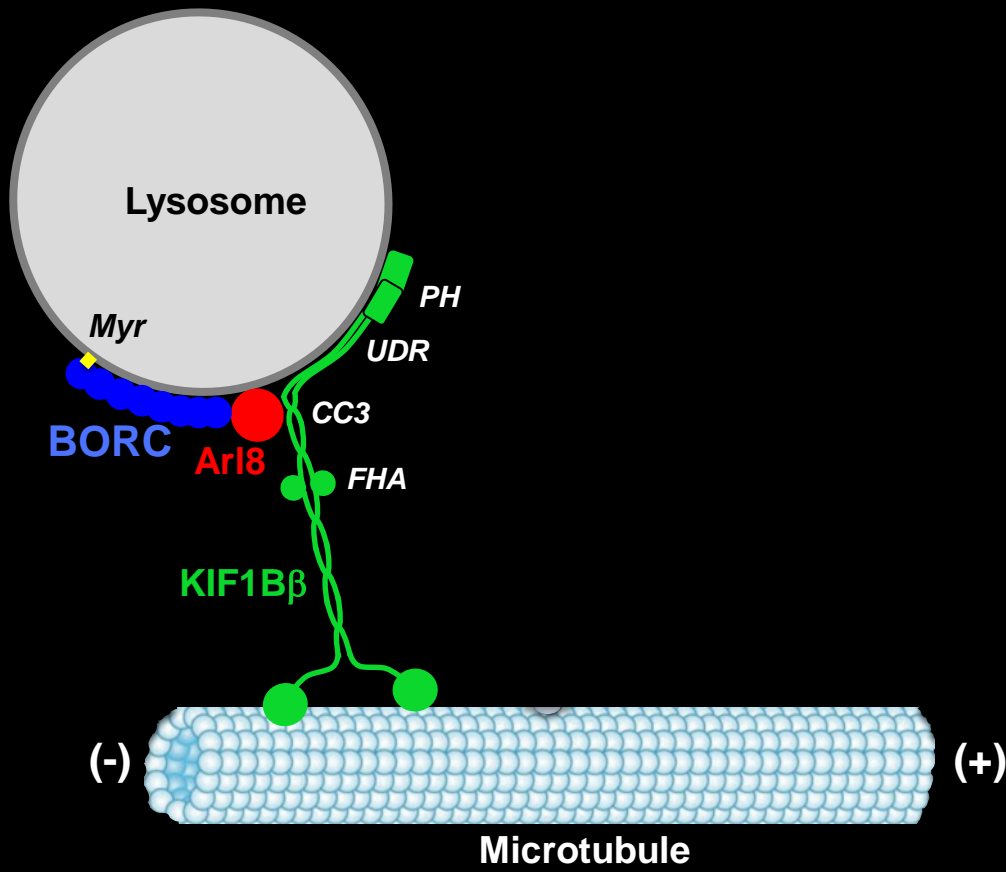
**Lyspersin**

**Diaskedin**

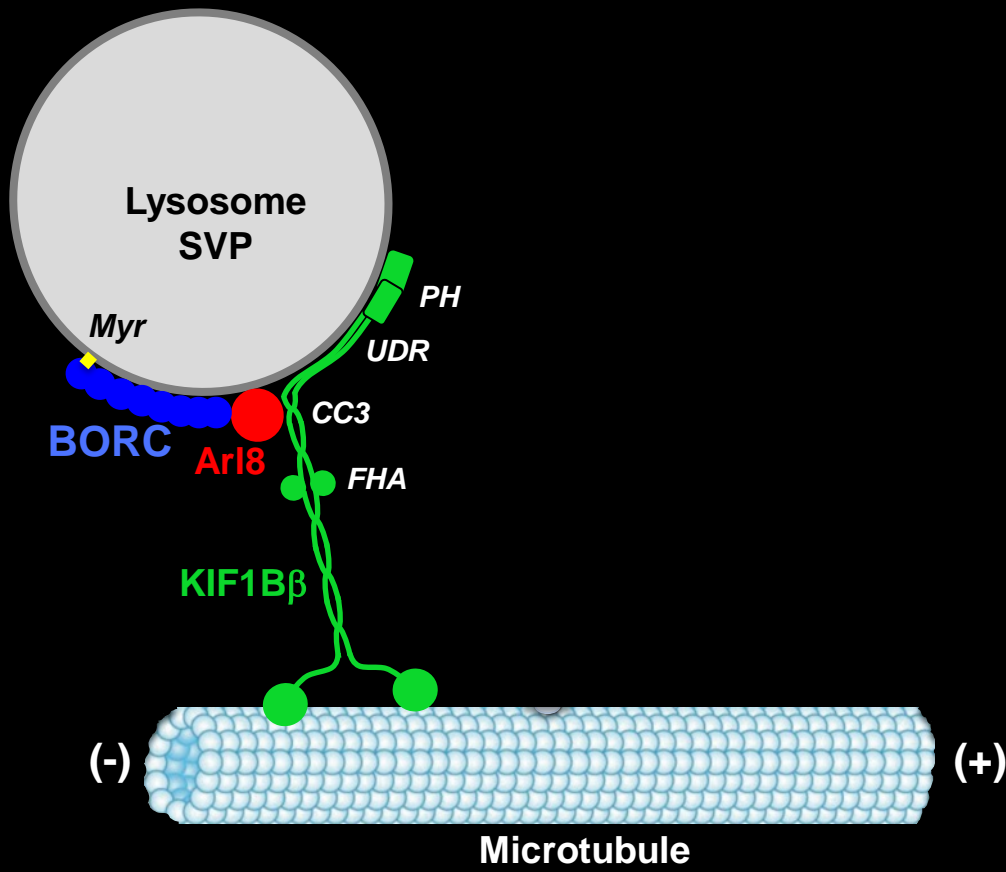
**MEF2BNB**

**KXD1**

# *BORC Couples Lysosomes and SVPs to Kinesins*

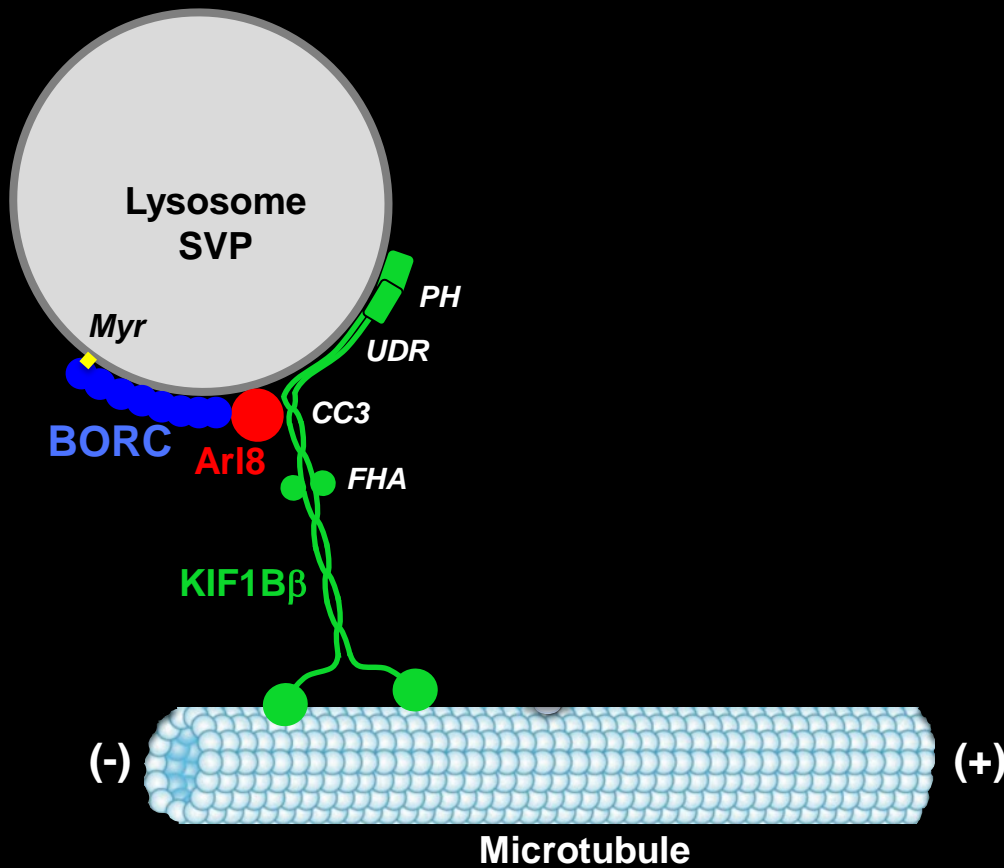


# ***BORC Couples Lysosomes and SVPs to Kinesins***



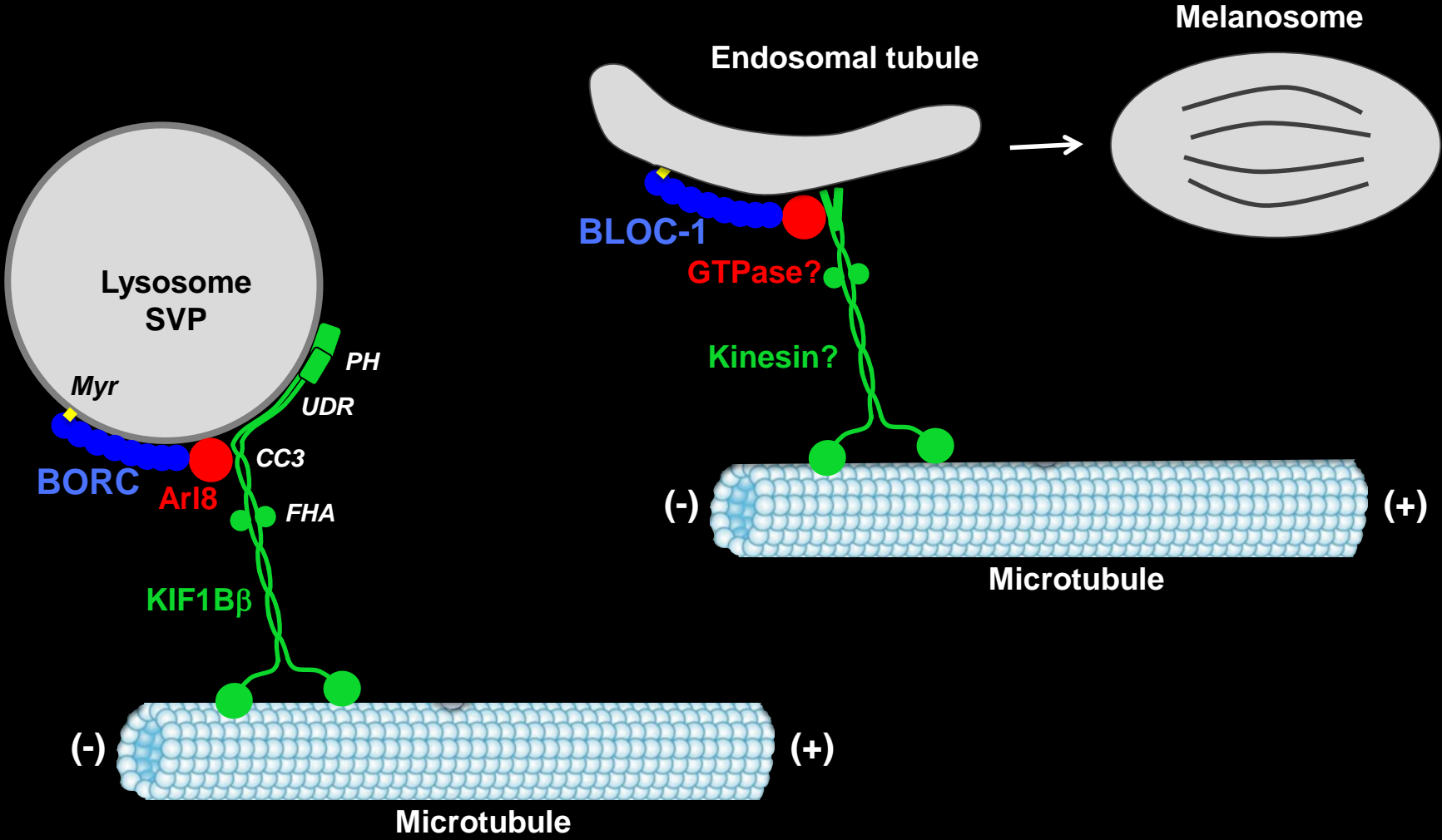


# ***BORC Couples Lysosomes and SVPs to Kinesins***



**BORCS7/Diaskedin: major schizophrenia susceptibility gene**

# Does BLOC-1 Couple Melanosomal Carriers to Kinesin?



# *Conclusions*

---

**AP-3 is involved in the biogenesis of lysosome-related organelles in flies, mice and humans**

# *Conclusions*

---

**AP-3 is involved in the biogenesis of lysosome-related organelles in flies, mice and humans**

**Mutations in AP-3 are the cause of Hermansky-Pudlak syndrome type 2**

# *Conclusions*

---

**AP-3 is involved in the biogenesis of lysosome-related organelles in flies, mice and humans**

**Mutations in AP-3 are the cause of Hermansky-Pudlak syndrome type 2**

**AP-3 mediates the sorting of tyrosinase from endosomes to melanosomes by recognition of a dileucine sorting signal**

# *Conclusions*

---

**AP-3 is involved in the biogenesis of lysosome-related organelles in flies, mice and humans**

**Mutations in AP-3 are the cause of Hermansky-Pudlak syndrome type 2**

**AP-3 mediates the sorting of tyrosinase from endosomes to melanosomes by recognition of a dileucine sorting signal**

**BLOC-1, BLOC-2 and BLOC-3 are novel components of a molecular machinery for the biogenesis of lysosome-related organelles**

# *Conclusions*

---

**BORC couples lysosomes and SVPs to kinesins**

**BLOC-1 may act in a similar manner to couple  
melanosome-bound carriers to kinesins**

# ***Thanks!***

---

***Esteban Dell' Angelica***

***Chean Eng Ooi***

***Chris Mullins***

***Kengo Moriyama***

***José Martina***

***Lisa Hartnell***

***Chris Schindler***

***Jing Pu***

***Charly Guardia***

***(CBMB, NICHD)***

***Bill Gahl***

***(NICHD/NHGRI)***

***Mickey Marks***

***(U. Pennsylvania)***

***Graça Raposo***

***(Curie Institute)***



***Eunice Kennedy Shriver National Institute  
of Child Health and Human Development***