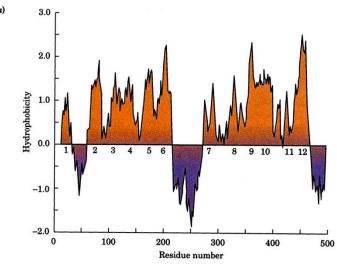
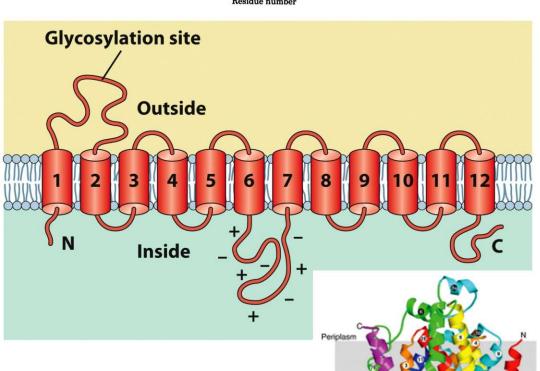
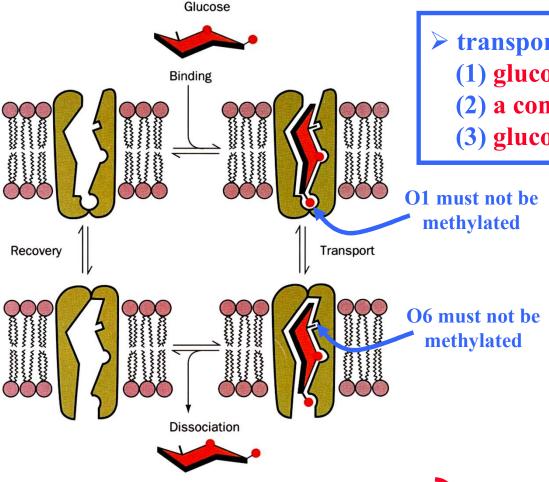
### A Closer Look at the Glucose Transporter



- integral membrane proteins have a directionality and do not flip-flop
- ⇒ they cannot work as an ionophore (mobile carrier)!
  - > the erythrocyte glucose transporter:
  - ♣ 55 kD glycoprotein; 492 aa
  - four major domains:
    - (1) 12 membrane-spanning  $\alpha$ -helices;
    - (2) large highly charged cytoplasmatic domain (between helices 6 & 7)
    - (3) smaller, carbohydrate bearing external domain (between helices 1 & 2)
    - (4) cytoplasmatic C-terminal domain
  - accounts for 2% of erythrocyte membrane proteins
  - > must be asymmetric:
  - galactose oxidase oxidizes oligosaccharide chain only on the outside;
  - \* trypsin disrupts transport only from the inside of an erythrocyte ghost



### Glucose Transporter: A Passive Gated Pore



> transport is passive:

How?

- (1) glucose is bound on the outside,
- (2) a conformational change occurs,
- (3) glucose is released on the inside
  - > transporter "only" equilibrates glucose concentration, but inside the cell glucose is rapidly consumed!

> But 15 min after insulin administration,  $J_{max}(glucose)$  increases by 6- to 12-fold, while  $K_M$  stays constant

> 20 min to 2 h after insulin withdrawal, glucose uptake returns to normal

Exocytosis

Endocytosis

Endocytosis

Endocytosis

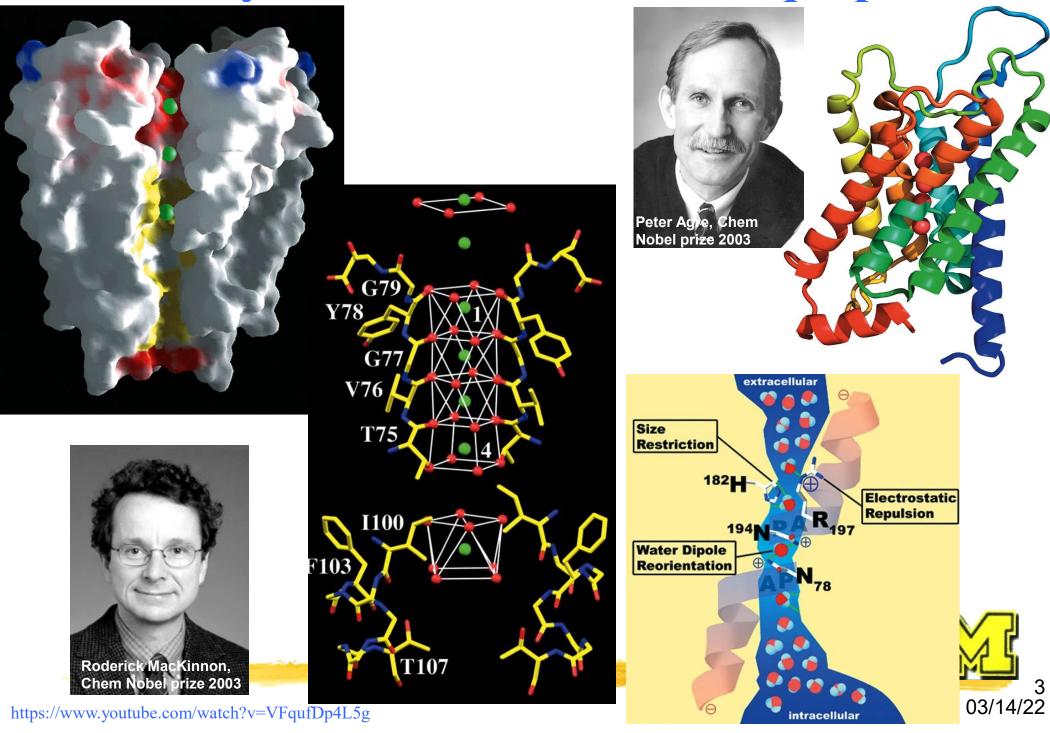
Membranous

vesicle

Plasma

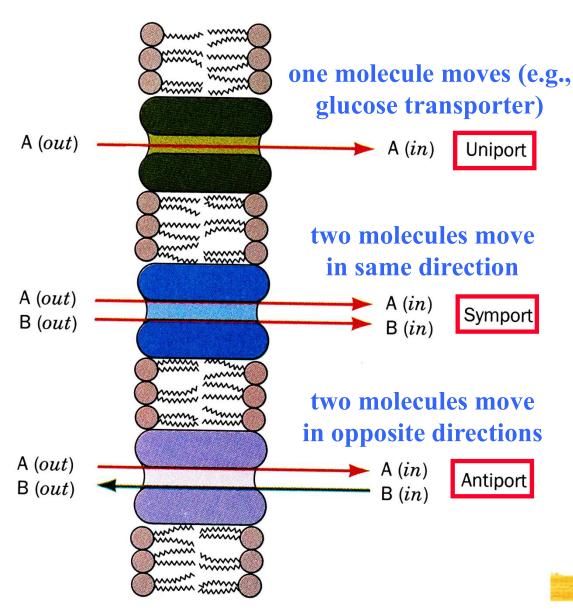
membrane

### Selectivity of K<sup>+</sup> Channels and Aquaporines



### **ATP-Driven Active Transport**

#### **Categories of mediated transport:**



Electroneutral: Simultaneous charge neutralization (symport of opposite charges or antiport of like charges)

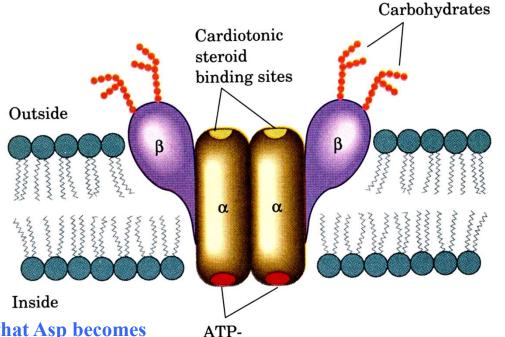
**Electrogenic:** Transport results in charge separation across membrane

Active transport against a gradient is endergonic ⇒ often coupled to ATP hydrolysis

- **Examples:**
- ♣ P-type ATPases; in plasma membrane, directly phosphorylated by ATP; inhibited by vanadate (VO<sub>4</sub><sup>3-</sup>) as P<sub>i</sub> analog
- **♣ F-type ATPases**; in mitochondrial membrane; important for oxidative phosphorylation
- **♣ V-type ATPases**; in plant vacuolar membranes; analogous to F-type ATPases

Nils Walter: Chem 451

### The (Na<sup>+</sup>-K<sup>+</sup>)-ATPase of Plasma Membranes

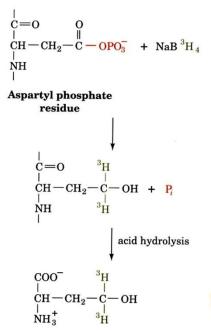


binding sites

- > The facts:
- $(\alpha\beta)_2$  subunit composition
- $\clubsuit$   $\alpha$  subunit: 110 kD, non-glycosylated; contains ATPase activity and ion-binding sites; 8 transmembrane  $\alpha$ -helices
- \* β subunit: 55 kD, glycosylated; 1 transmembrane α-helix
- electrogenic antiport
- ♣ also called (Na<sup>+</sup>-K<sup>+</sup>) "pump"
- \* sequential kinetic mechanism accounts for coupling of active transport with ATP hydrolysis

03/14/22





Homoserine

2 Conformational **ATP**  $Mg^{2+}$ states: E4 • ATP • 3Na+ 1. ATP binding 2. formation of  $3Na^{+}(in)$ "high-energy" aspartyl phosphate intermediate  $2K^+(in) \leftarrow$ Inside 6. K+ transport and 3. Na\* transport Nat binding Relaxation Outside → 3Na<sup>+</sup>(out) 5. phosphate 4. K+ binding hydrolysis  $H_2O$ 2K (out)

# Chapter 20: What have we learned about transport?

- > Thermodynamics, kinetics, nomenclature (symport, antiport...)
- > Carrier and Channel-Forming Ionophores as models

	Non-mediated	Mediated	
		<u>Passive</u>	<u>Active</u>
Carrier	No	Yes	Yes
Transport direction	[High]→[Low]	[High]→[Low]	[Low]→[High]
Energy used?	No	No	Yes
Examples	$O_2$ , $H_2O$	Glucose	(Na <sup>+</sup> -K <sup>+</sup> )-ATPase, Ca <sup>2+</sup> -ATPase, Translocation systems

#### **Glycolysis** Glucose Glucose-6-phosphate 2Glyceraldehyde-3-phosphate glyceraldehyde-- 2NAD+ 3-phosphate dehydrogenase 2NADH 2 1,3-Bisphosphoglycerate 2Pyruvate 2NAD+ pyruvate dehydrogenase 2NADH 2Acetyl-CoA 2NADH 20xaloacetate 2NAD+ 2Citrate 2Malate dehydrogenase 2lsocitrate Citric acid 2Fumarate cycle isocitrate 2NAD+ 2FADH<sub>2</sub> succinate dehydrogenase dehydrogenase 2NADH 2FAD 2Succinate α-ketoglutarate 2α-Ketoglutarate dehydrogenase 2Succinvl-CoA 2NAD+

# **Electron Transport and Oxidative Phosphorylation**

Voet & Voet, Chapter 22

- ⇒ Complete glucose oxidation:  $C_6H_{12}O_6 + 6O_2 \rightarrow 6CO_2 + 6H_2O;$   $\Delta G^{0'} = -2,823 \text{ kJ/mol}$ 
  - > Oxidation = loss of electrons:

Two half-reactions: 
$$C_6H_{12}O_6 + 6H_2O \rightarrow 6CO_2 + 24H^+ + 24e^-$$
  
 $6O_2 + 24H^+ + 24e^- \rightarrow 12H_2O$ 

➤ How can the energy of these electrons be transferred to ATP?

24 e<sup>-</sup> are carried in 10 NADH and 2 FADH<sub>2</sub>

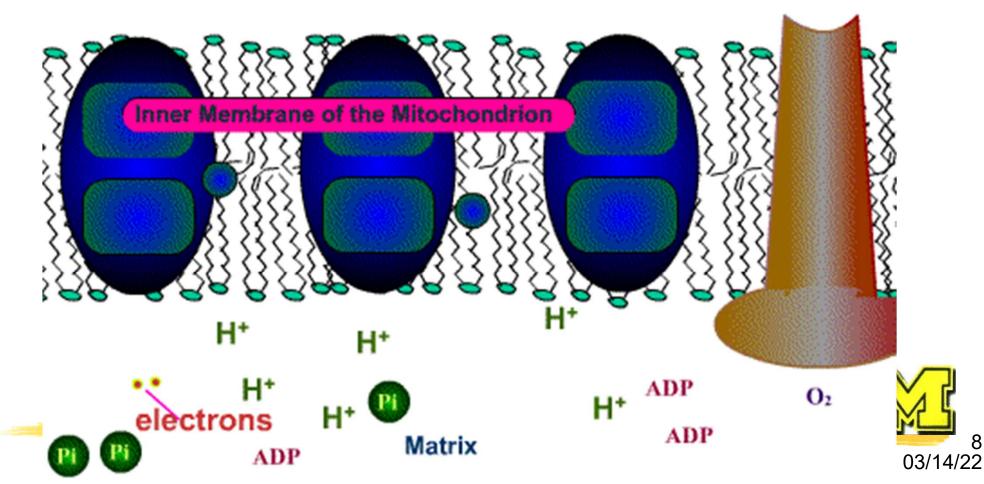
Pass into electron-transport chain (participate in reduction-oxidation of >10 redox centers)

Reduce O<sub>2</sub> to H<sub>2</sub>O, expel H<sup>+</sup> from mitochondrion

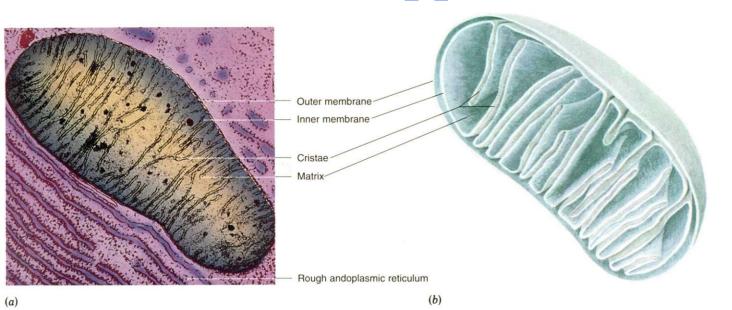
H<sup>+</sup> gradient drives ATP production

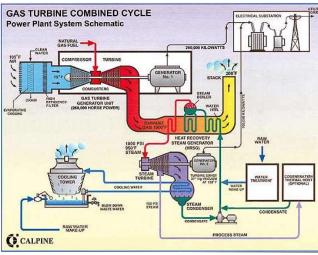
# The Movie of Electron Transport and Oxidative Phosphorylation





### Where it All Happens: The Mitochondrion





Outer membrane:
Outer face  $(2806 \text{ particles} \bullet \mu\text{m}^{-2})$ Inner face  $(770 \text{ particles} \bullet \mu\text{m}^{-2})$ Outer face  $(2120 \text{ particles} \bullet \mu\text{m}^{-2})$ Inner face  $(4208 \text{ particles} \bullet \mu\text{m}^{-2})$ 

#### The mitochondrion contains:

- > Pyruvate dehydrogenase
- > Citric acid cycle enzymes
- **Enzymes that catalyze fatty acid oxidation**
- Enzymes and redox proteins for electron transport and oxidative phosphorylation

⇒ The cellular "power plant"