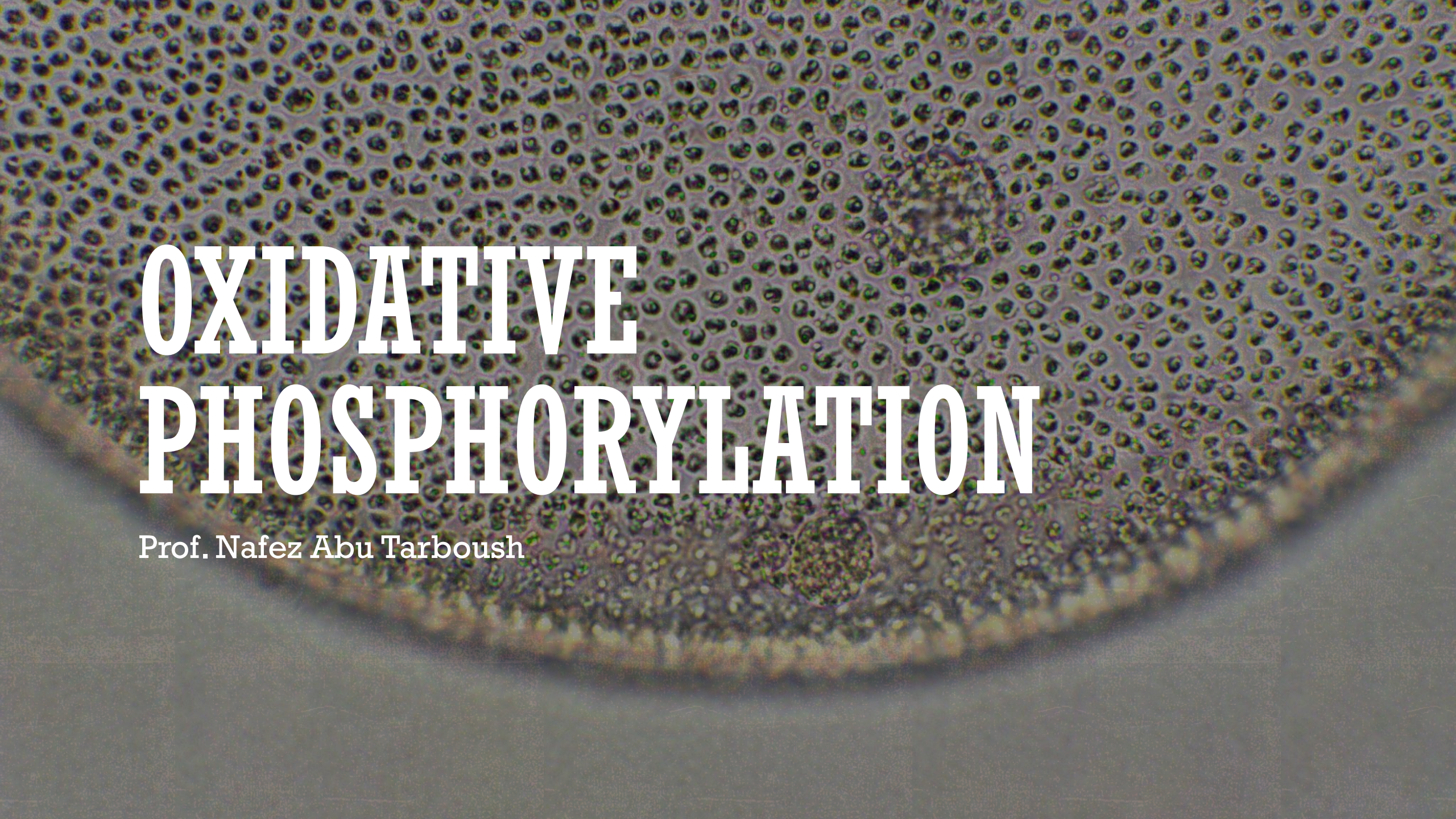
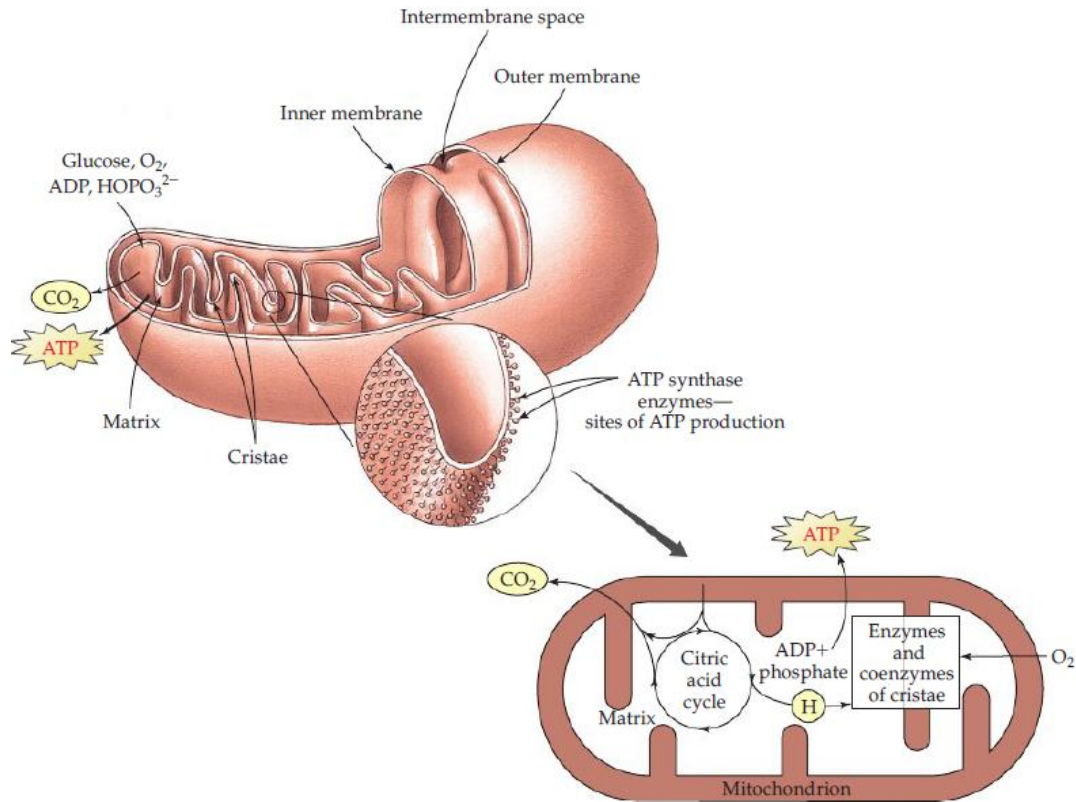




OXIDATIVE PHOSPHORYLATION

Prof. Nafez Abu Tarboush

MITOCHONDRIAL ARCHITECTURE



- OMM: permeable (MW<5,000); (transmembrane channels)
- IMM: impermeable (H^+); specific transporters
- IMM: components of respiratory chain and ATP synthase
- Matrix: pyruvate dehydrogenase complex; TCA cycle enzymes, fatty acid β -oxidation pathway, and the pathways of amino acid oxidation
- Matrix contains all pathways of fuel oxidation except glycolysis

TABLE 20.3: Location of enzymes in mitochondria

Mitochondria, outer membrane:

Monoamino oxidase

Acyl CoA synthetase

Phospholipase A2

In between outer and inner membrane:

Adenylate kinase

Creatine kinase

Inner membrane, outer surface:

Glycerol-3-phosphate dehydrogenase

Inner membrane, inner surface:

Succinate dehydrogenase

Enzymes of respiratory chain

Soluble matrix:

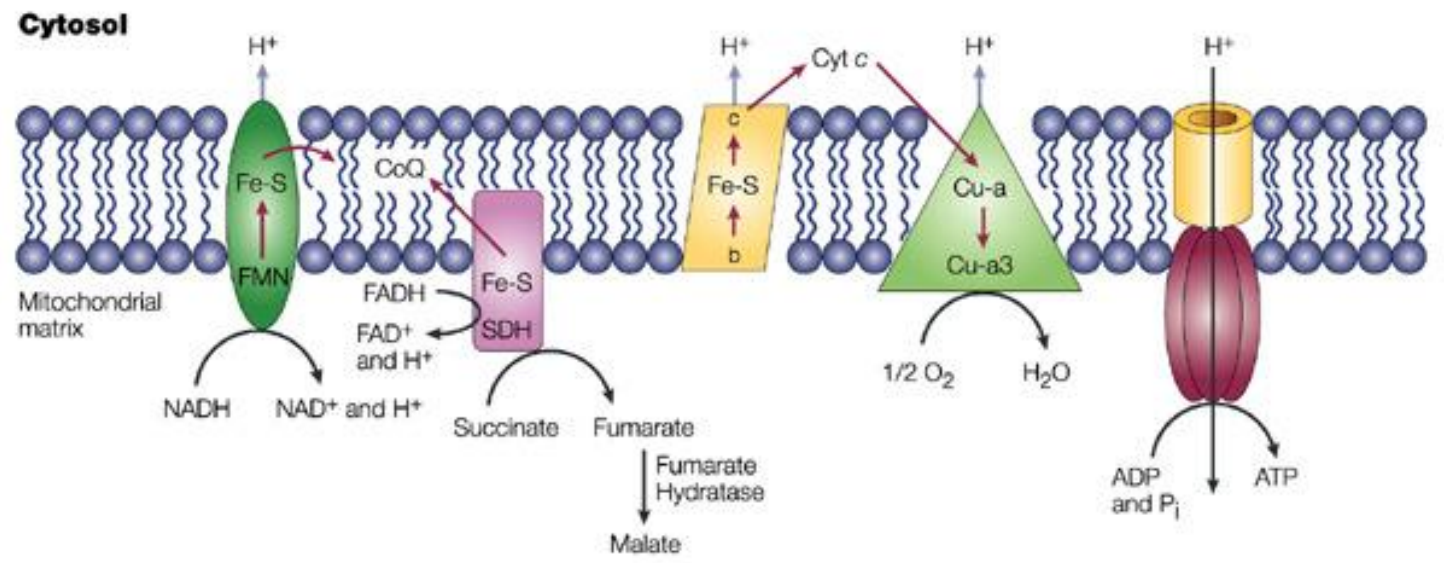
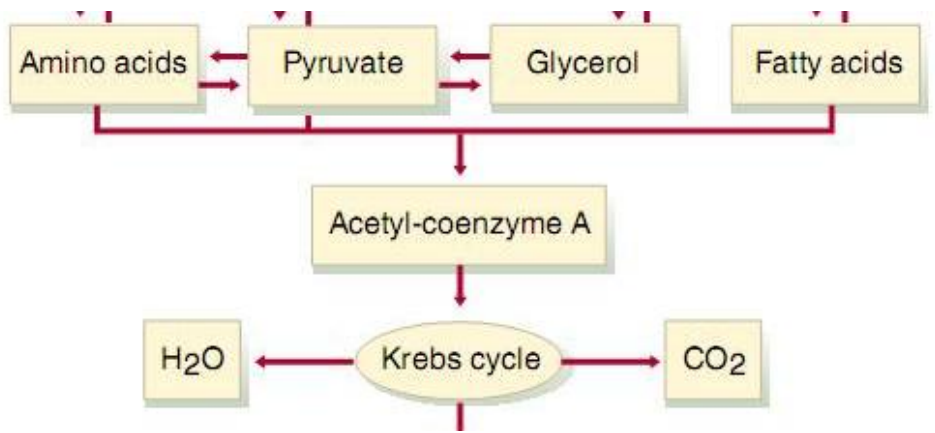
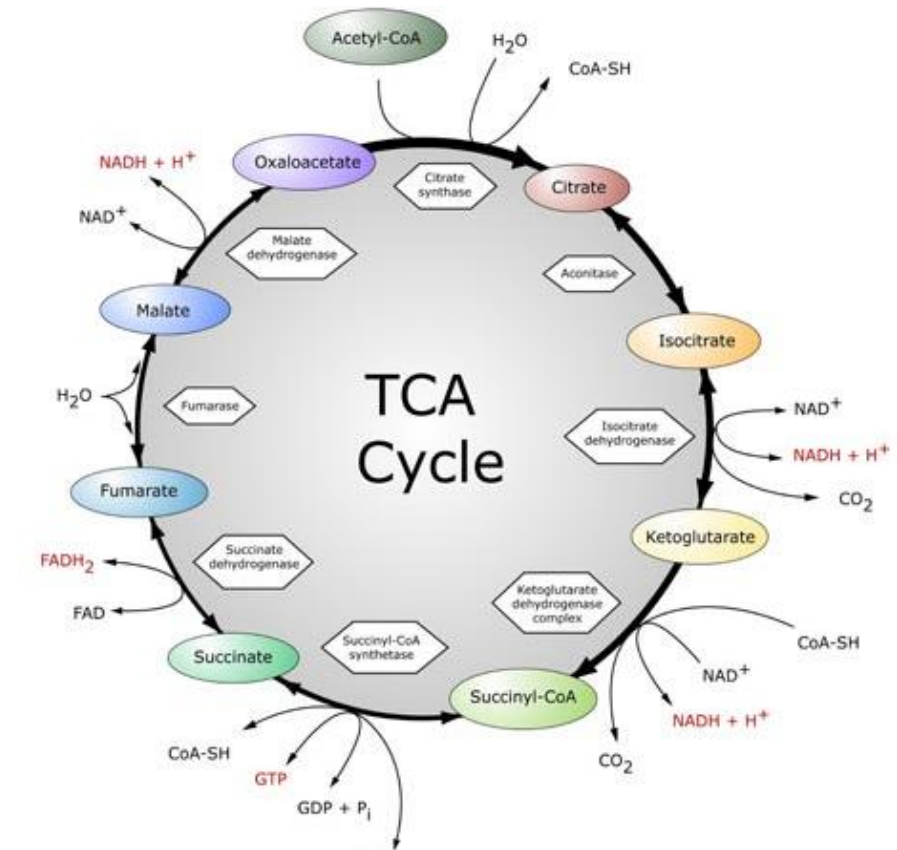
Enzymes of citric acid cycle

Enzymes of beta oxidation of fatty acid

MITOCHONDRIAL MEMBRANES

- **Inner membrane:**
 - 22% cardiolipin
 - No cholesterol
- **Outer membrane:**
 - Similar to cell membrane
 - Less than 3% cardiolipin
 - 45% cholesterol



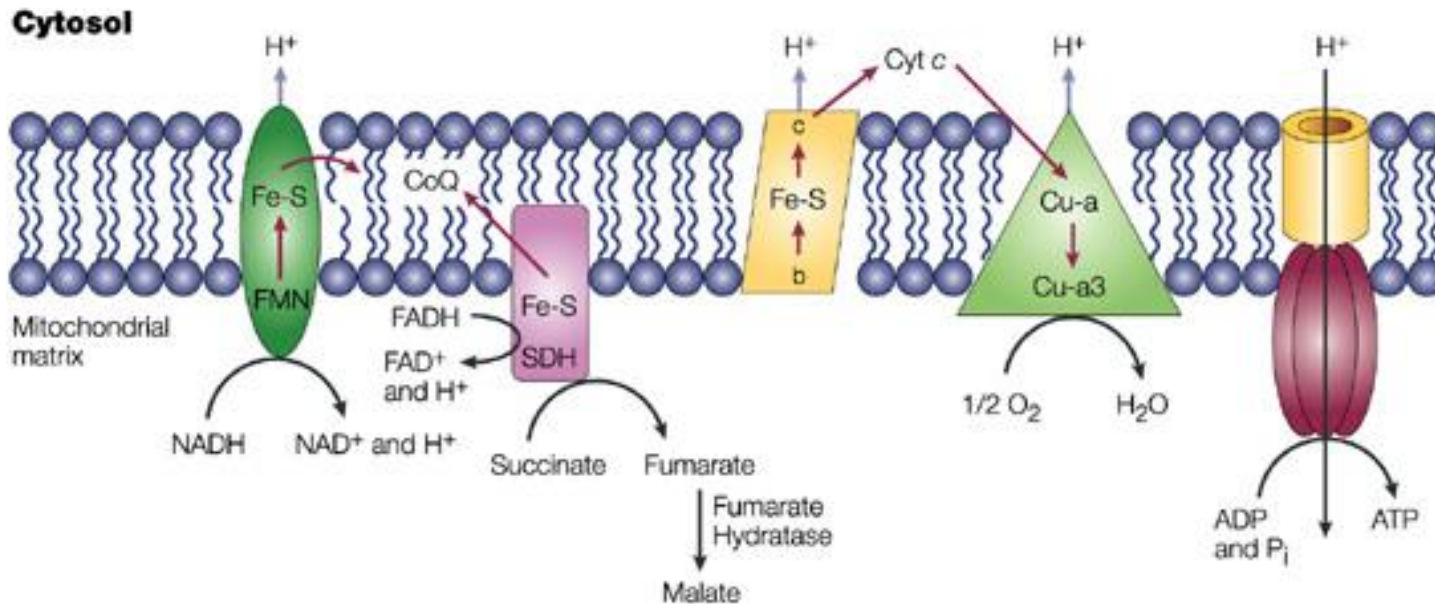


Complex	I	II	III	IV	V
Nuclear gene mutations	<i>NDU FS1, 2, 4, 7,8</i> <i>NDU FV1</i>	<i>SDHA, B, C, D</i>	<i>BCS1L</i>	<i>SURF1, SCO1, SCO2, COX10</i>	
Phenotype	Leigh syndrome Leukodystrophy	Leigh syndrome Paraganglioma	Hepatopathy Encephalopathy	Leigh syndrome Cardioencephalo-	Cardioencephalo- myopathy

THE OXIDATIVE PHOSPHORYLATION, WHERE ARE WE?

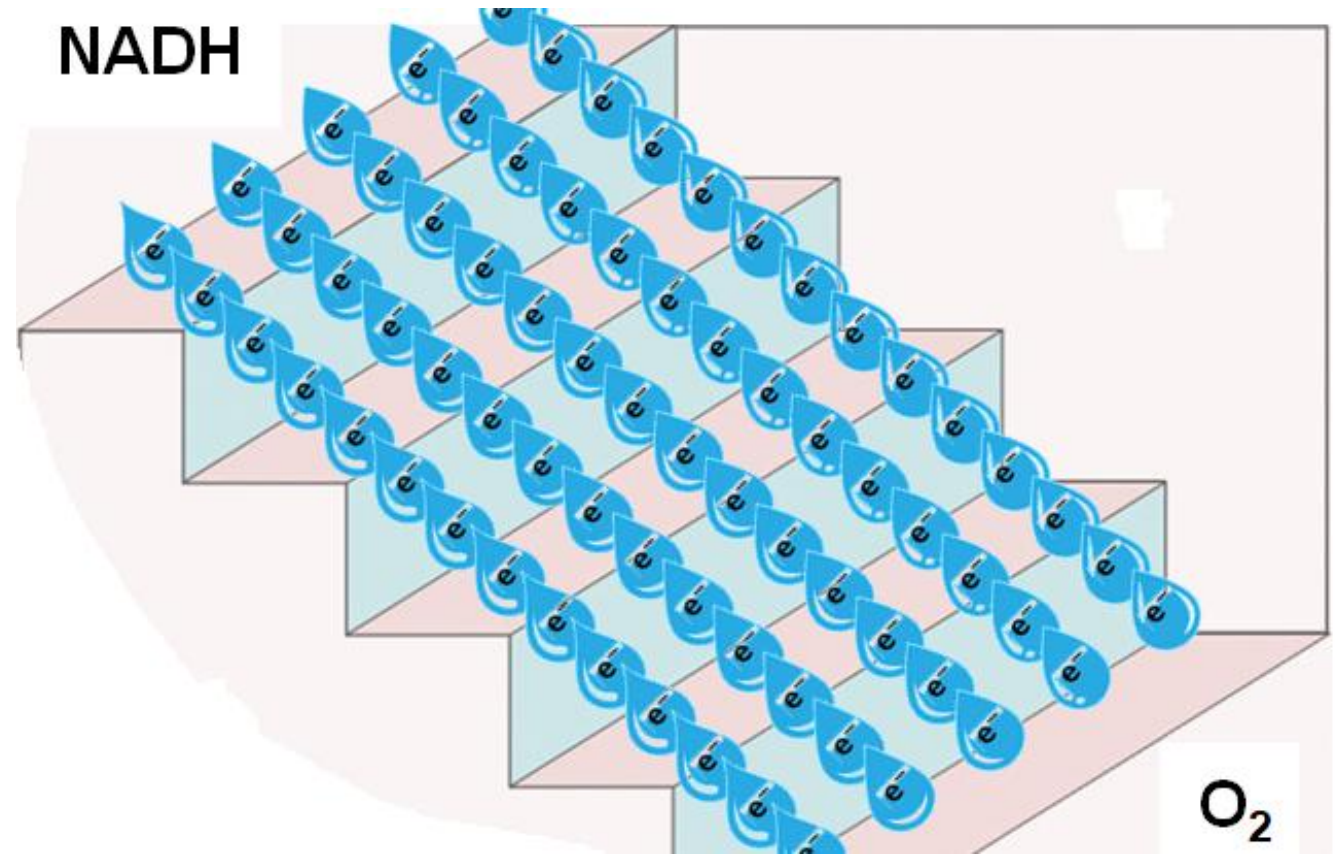
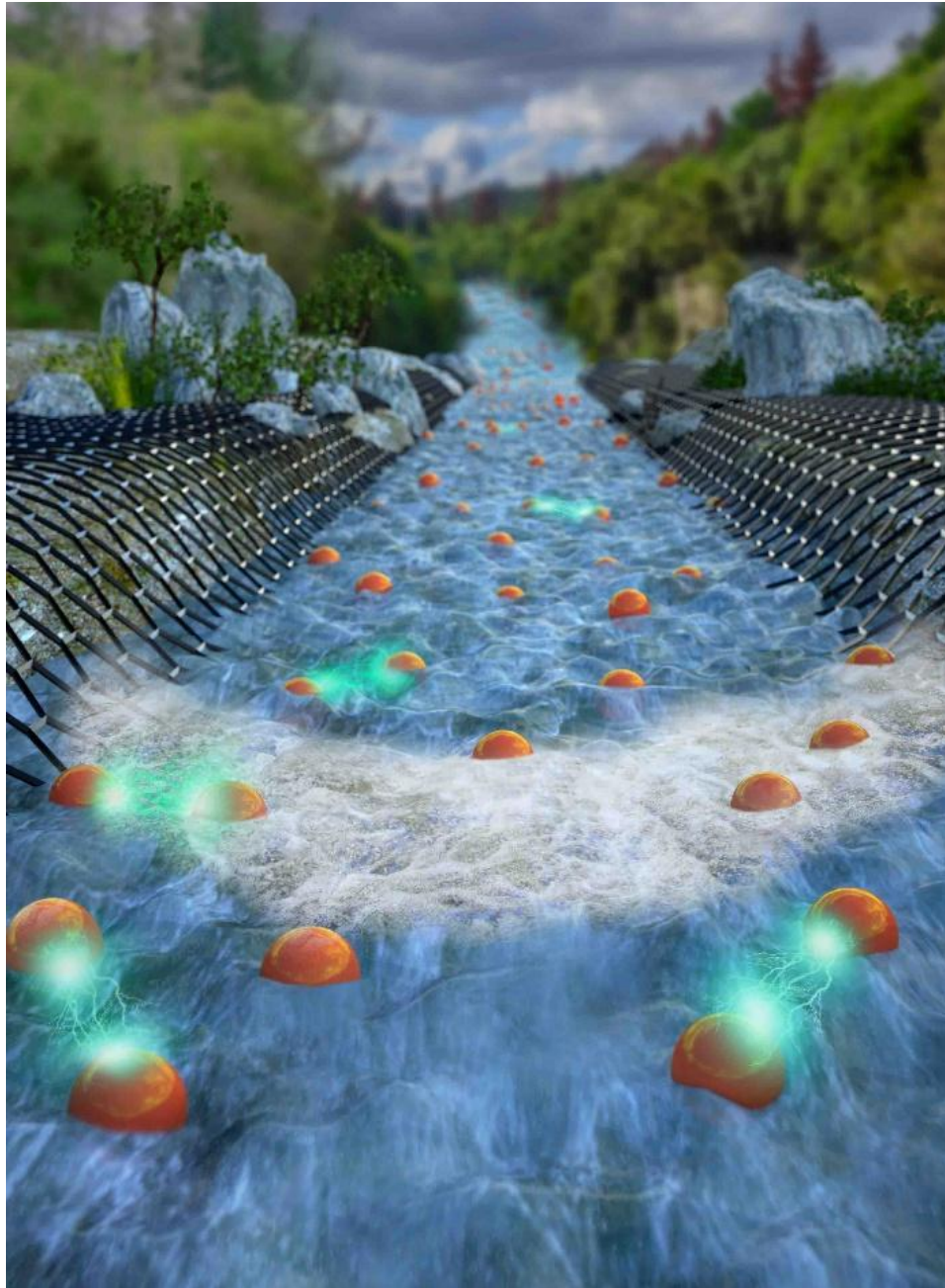
Stages: Digestion; Acetyl-CoA; TCA; OxPhos

(OXPHOS) OXIDATIVE PHOSPHORYLATION

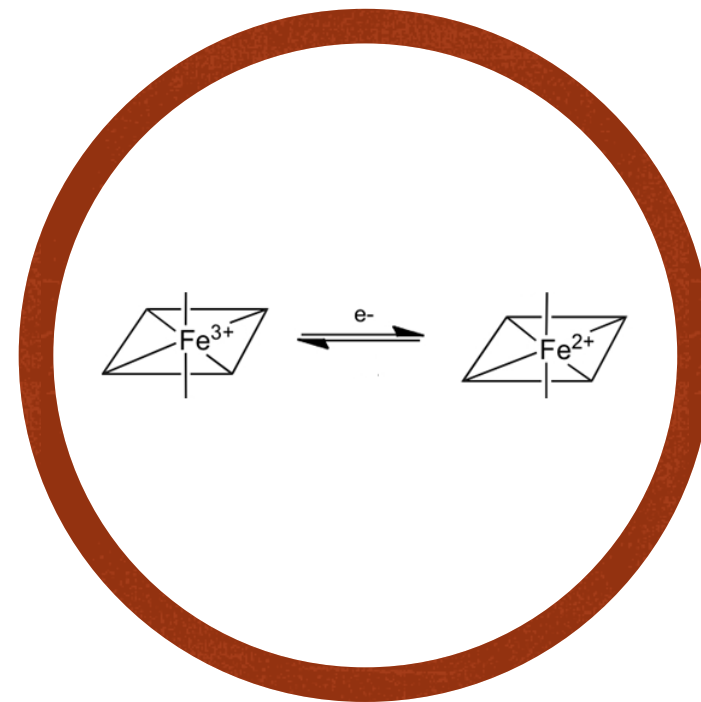
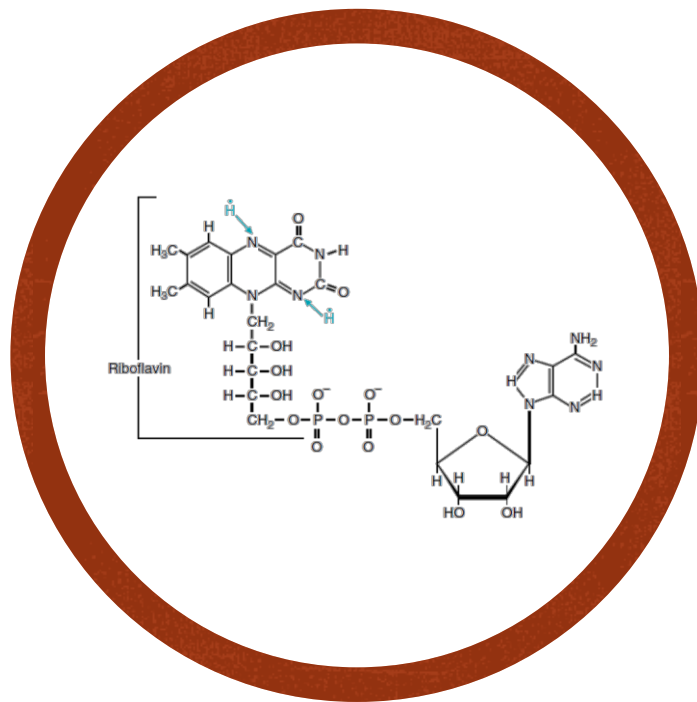
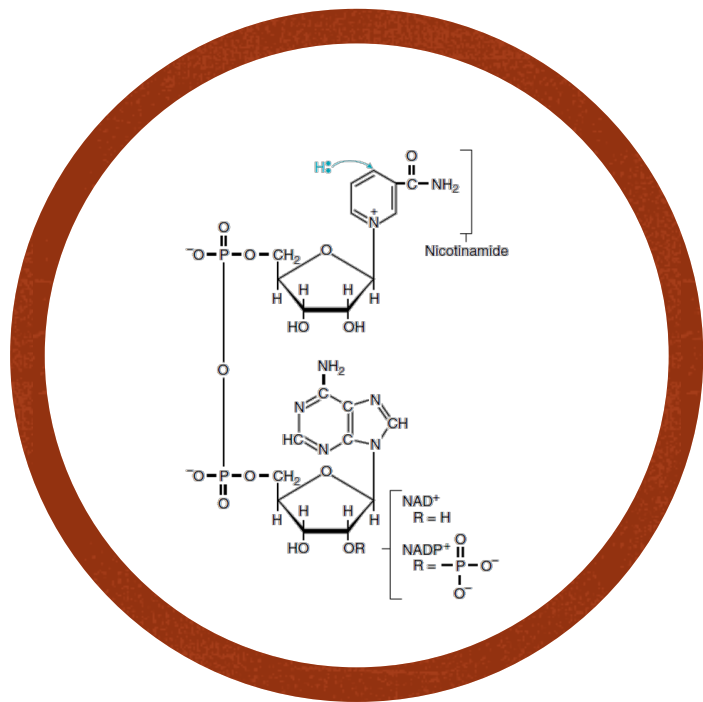


- Generation of ATP aided by O₂ reduction
- Peter Mitchell (1961): the chemiosmotic theory
- 3 major aspects:
 - ✓ (1) Flow of electrons
 - ✓ (2) The free energy available (exergonic) is coupled to transport of protons
 - ✓ (3) Flow of protons provides the free energy for synthesis of ATP (ATP synthase)



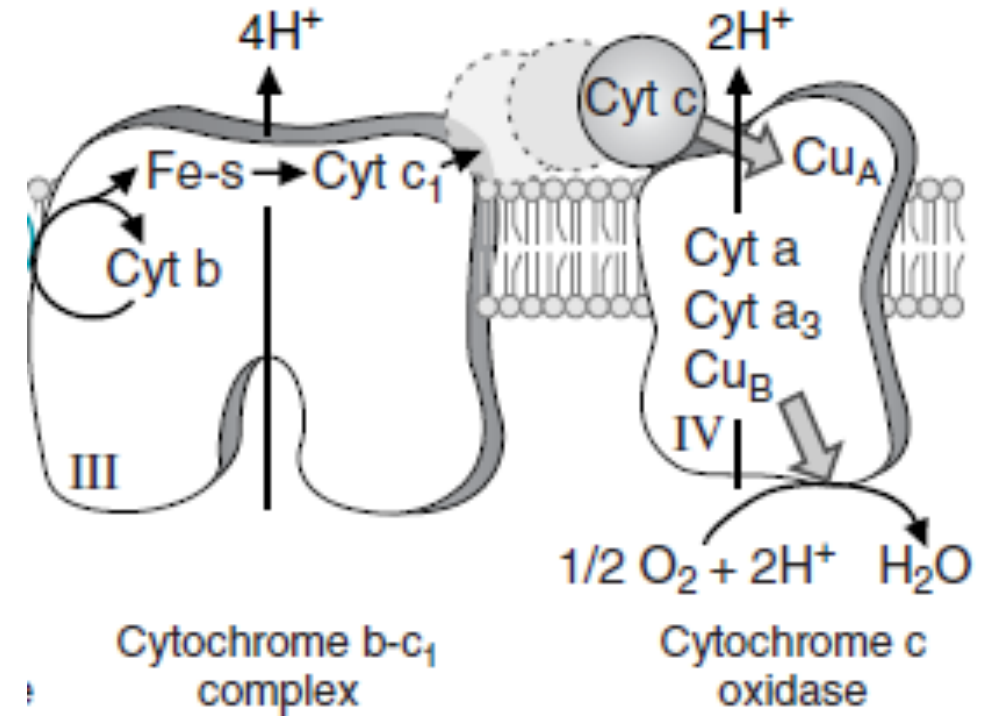
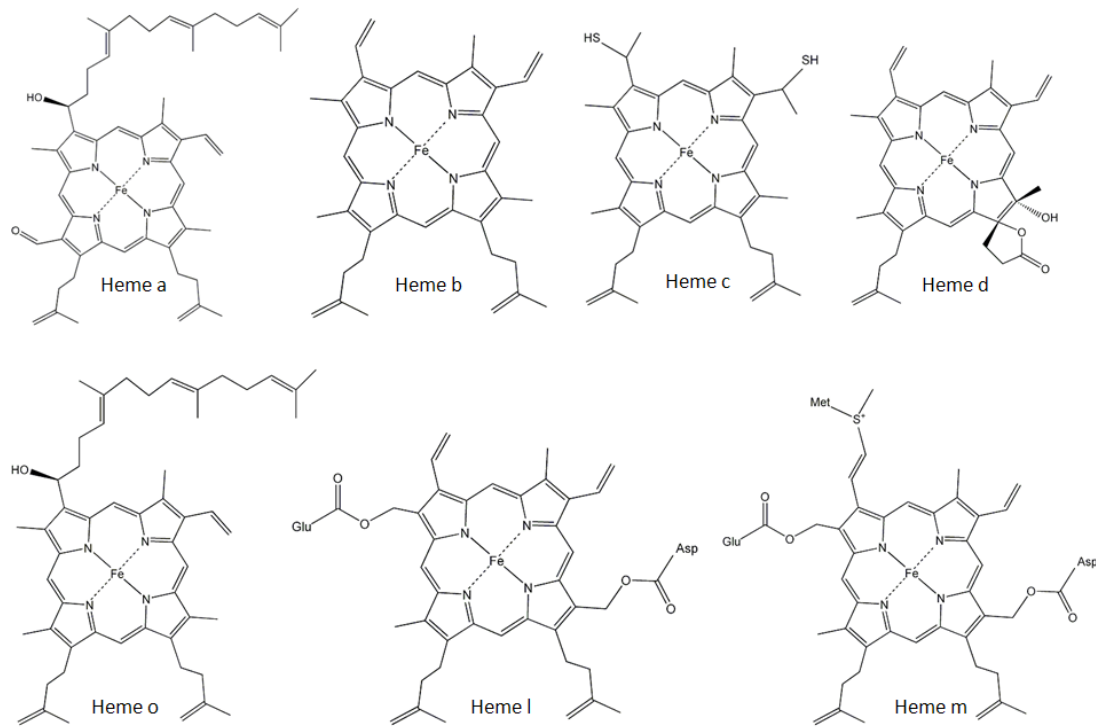


THE ELECTRON TRANSPORT CHAIN: THE RIVER OF ELECTRONS THAT PROVIDES A STEPWISE ENERGY RELEASE



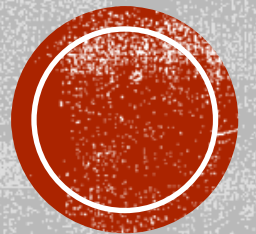
THE CAST OF CHARACTERS: ELECTRON CARRIERS

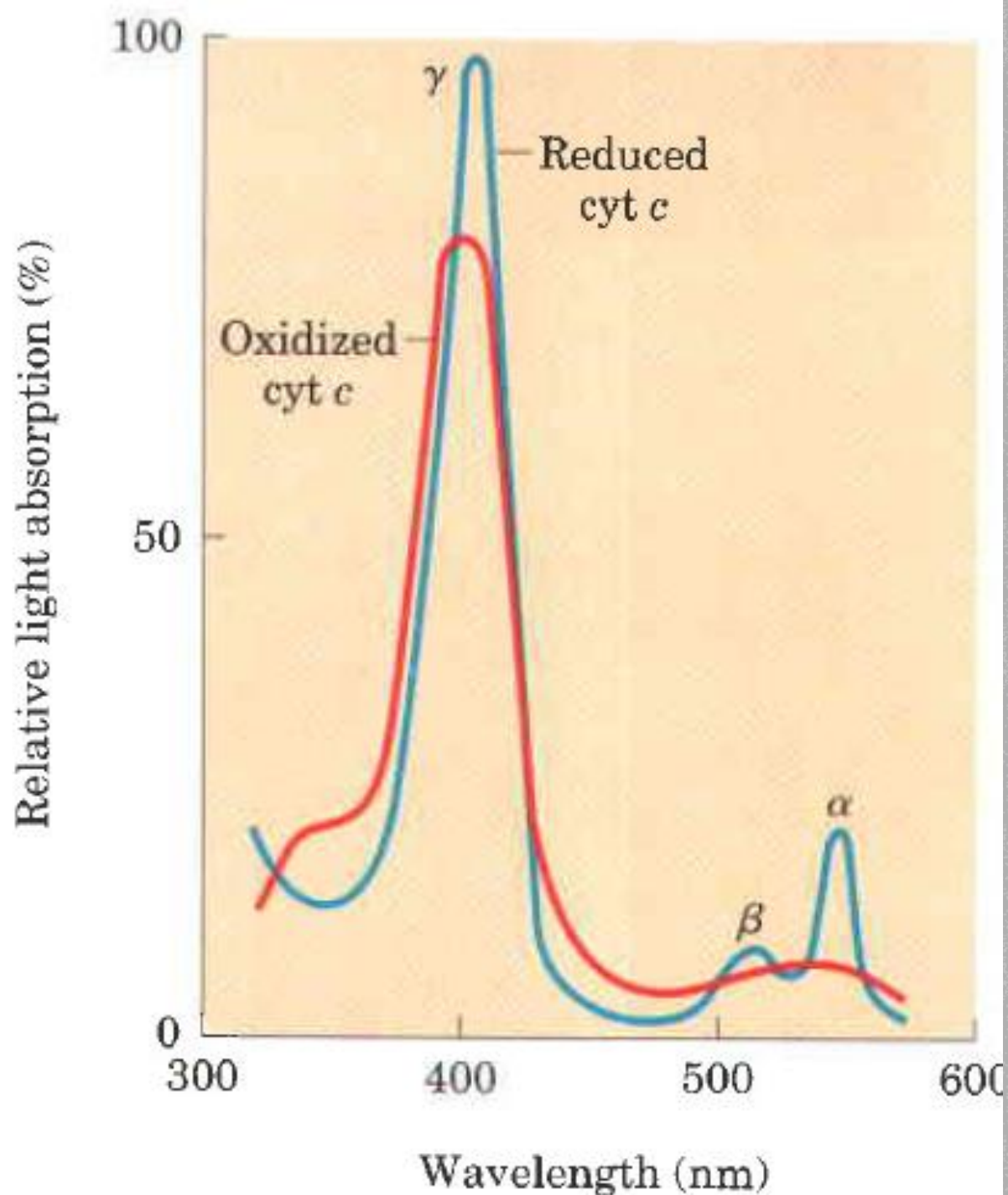




OTHER ELECTRON-CARRYING MOLECULES “CYTOCHROMES”

Mitochondria contain three classes of cytochromes (a, b, & c)

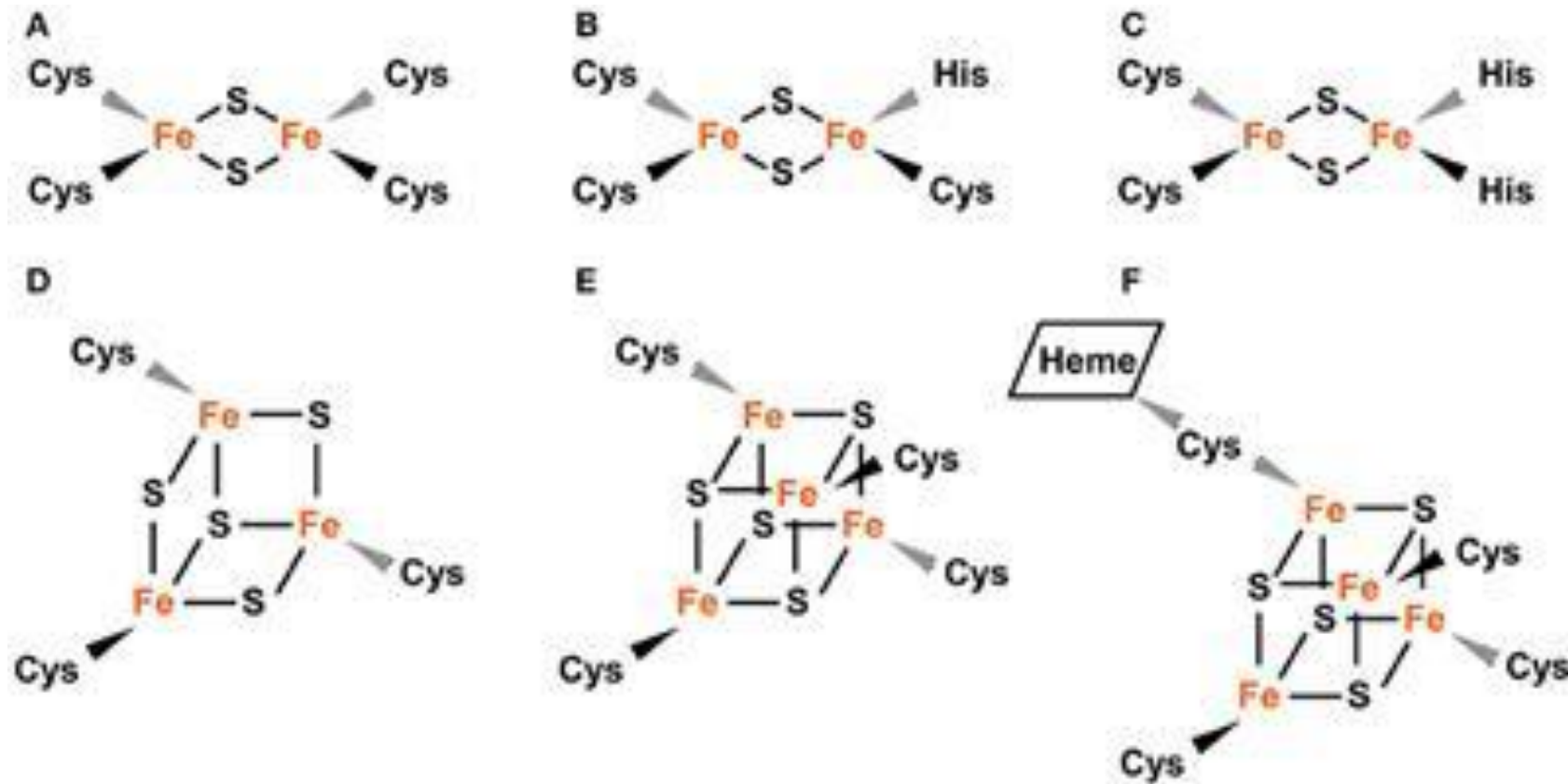




OTHER ELECTRON-CARRYING MOLECULES: "CYTOCHROMES"

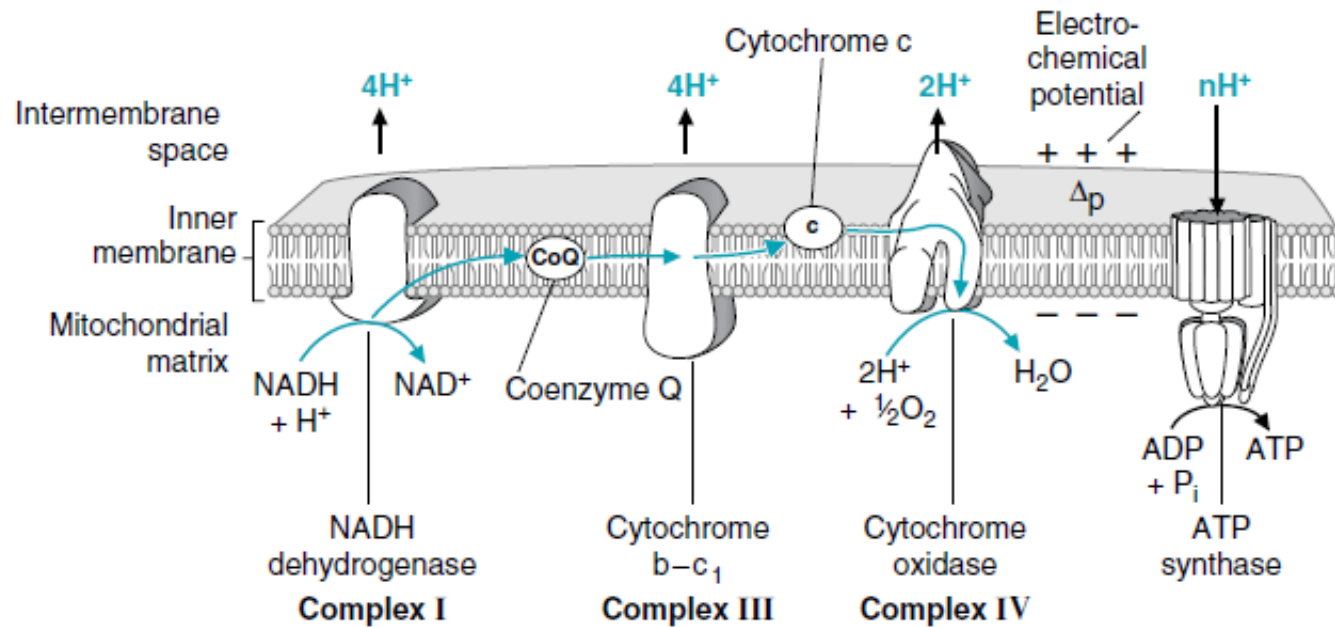
- Light absorption: α band (~ 600 nm in type a; ~ 560 nm in type b; ~ 550 nm in type c)
- Some cytochromes are named by the exact α band wavelength:
 - Cytochrome b_{562} ; Cytochrome c_{550} ; Cytochrome c_{551}
- Heme can carry one electron
- ΔE° depends on the protein





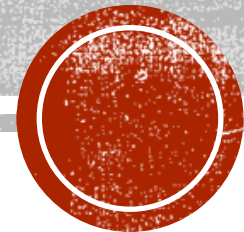
OTHER ELECTRON-CARRYING MOLECULES: IRON-SULFUR CLUSTERS

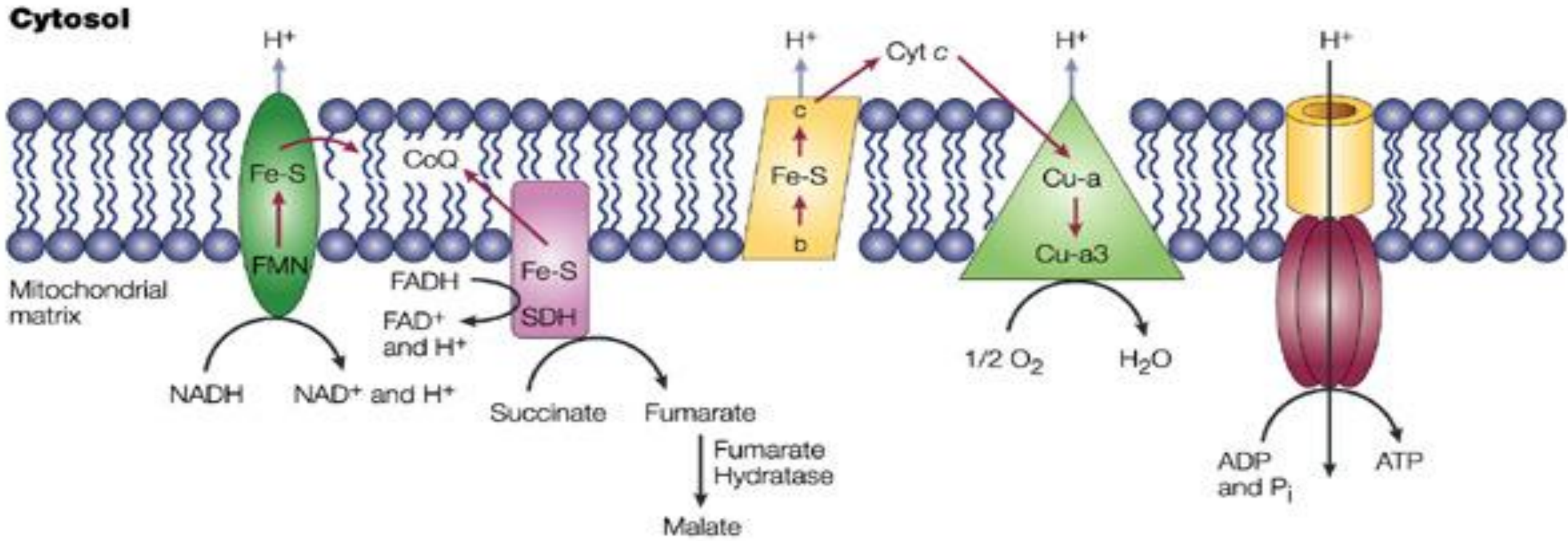




REQUIREMENTS OF OXPHOS

- Electrons' donor (NADH or FADH_2) & acceptor (O_2)
- ETC of proteins
- ATP synthase
- An intact IMM



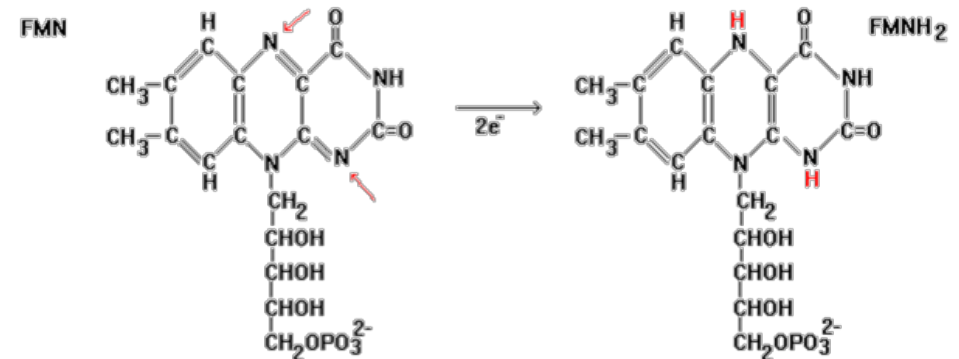
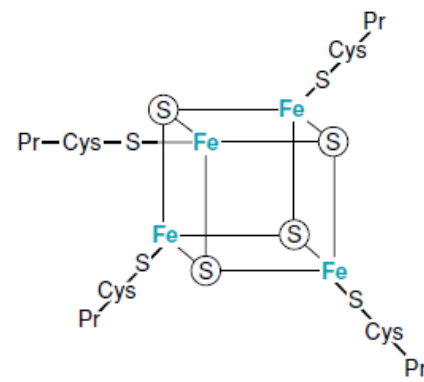
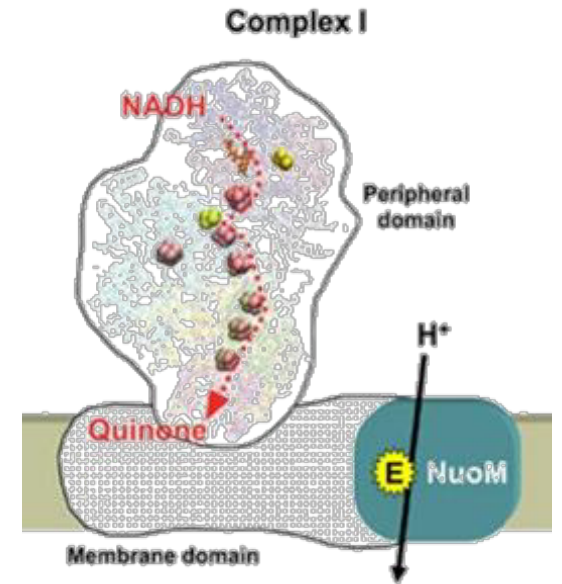
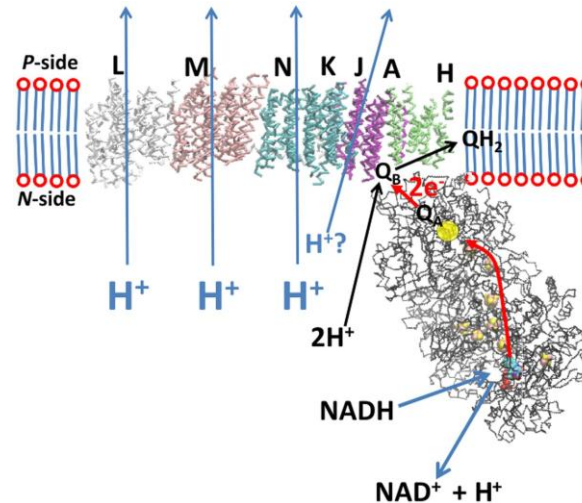
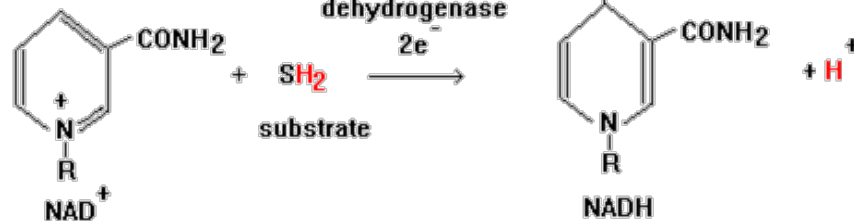


OXIDATIVE PHOSPHORYLATION (OXPHOS)



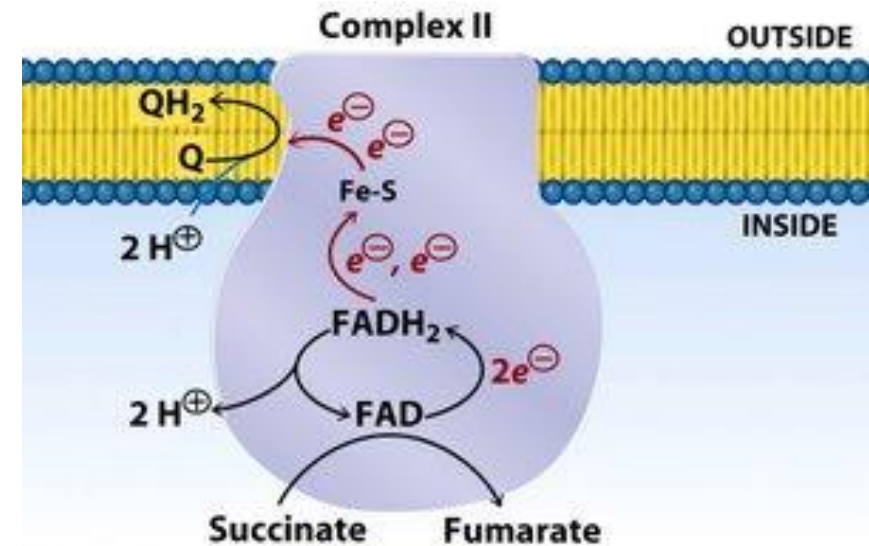
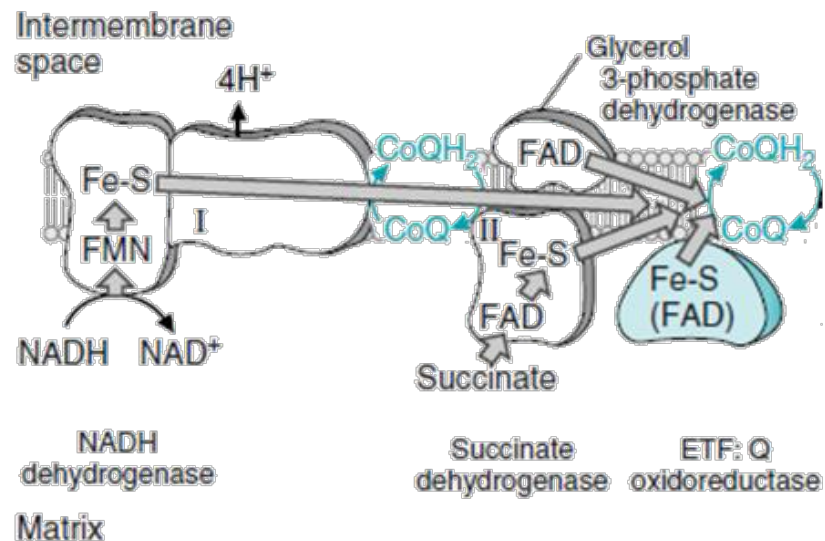
OXI-RED COMPONENTS OF THE ETC: THE NADH DEHYDROGENASE GIANT - COMPLEX I

- NADH-Q oxidoreductase
- More than 25 polypeptide chain
- A huge flavoprotein membrane-spanning complex
- FMN is tightly bound
- 7-9 Fe-S centers of at least two different types
- Drop in energy ≈ 13 to 14 kcal
- Binds NADH & CoQ
- 4 H^+

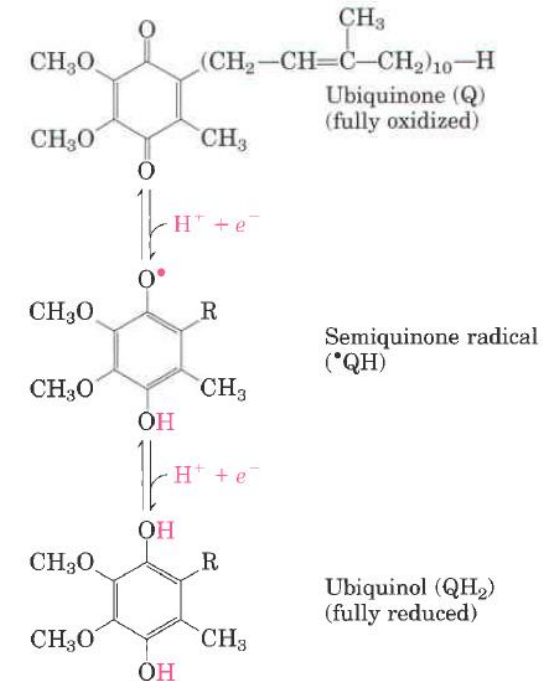
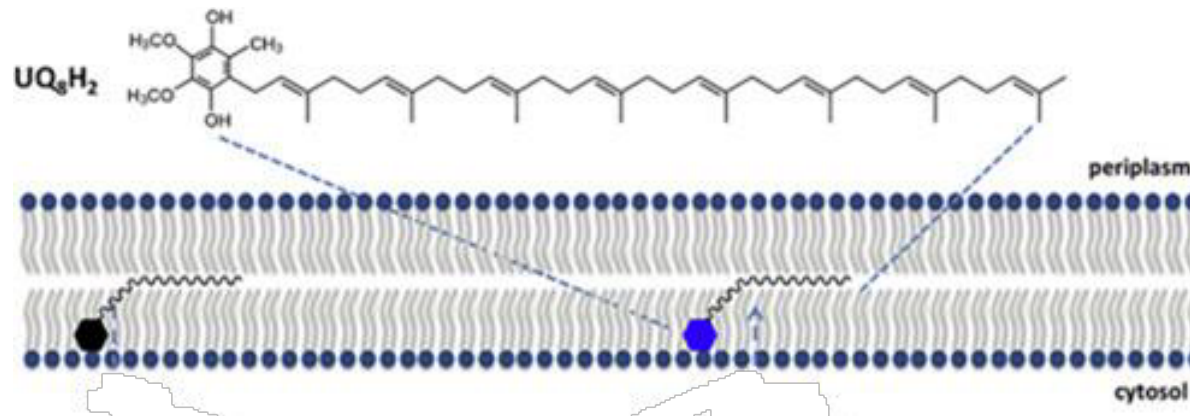


OXI-RED COMPONENTS OF THE ETC: “SUCCINATE DEHYDROGENASE” – COMPLEX II; THE CITRIC ACID CYCLE LINK

- Succinate Dehydrogenase & other flavoproteins
- TCA cycle
 - ✓ ETF-CoQ oxidoreductase (ex. fatty acid oxidation); Fatty acyl CoA dehydrogenase; Mitochondrial glycerol phosphate dehydrogenase
 - ≈ 0 Kcal, H^+ ?

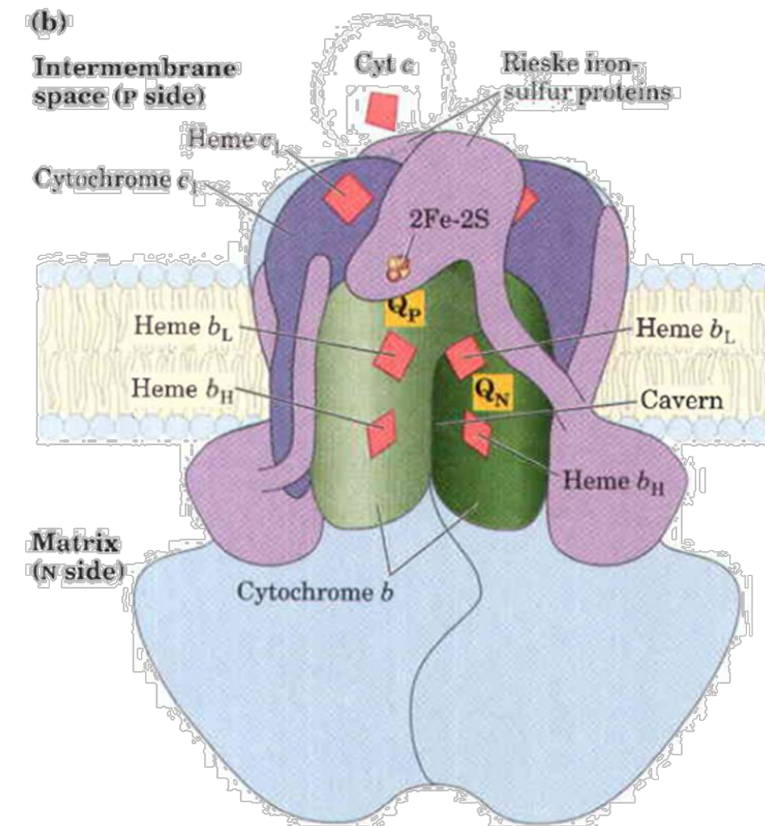


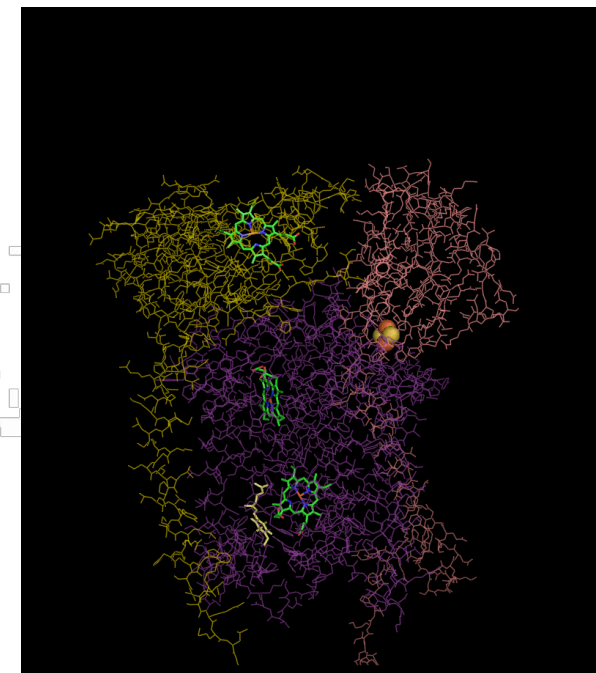
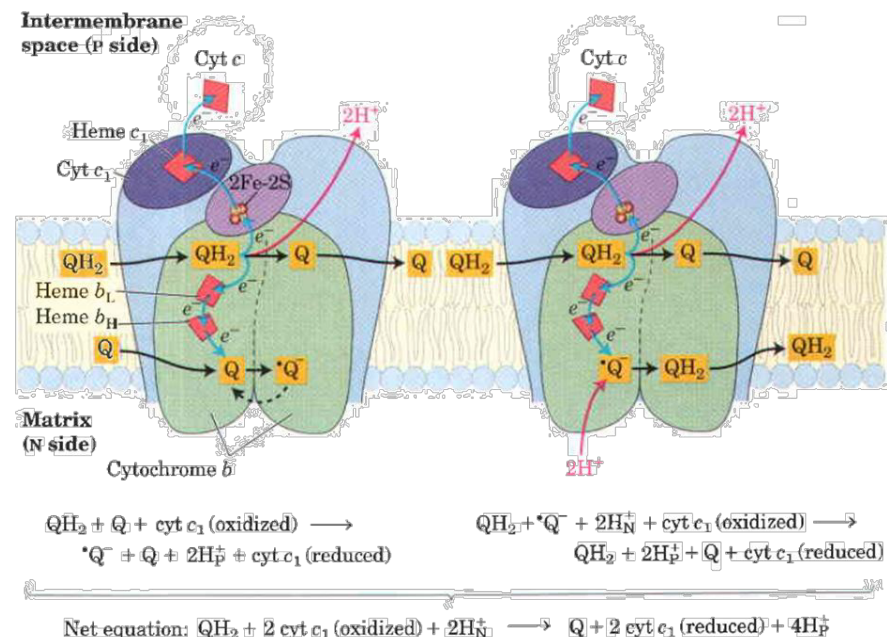
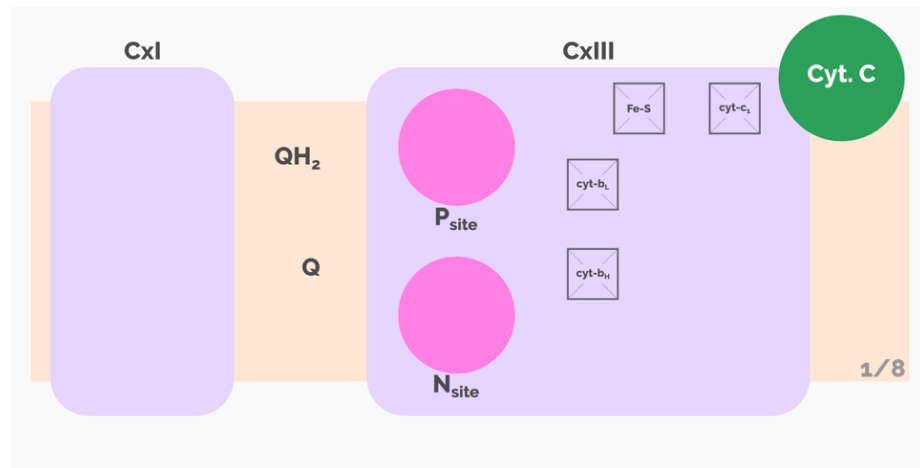
- COc1cc(C(=O)c2cc(C)c([C@@H](C2)C/C=C/C)cc1OC)cc(=O)c3cc(C)cc(C)cc3



OXI-RED COMPONENTS OF THE ETC: “CYTOCHROME BC₁” – COMPLEX III

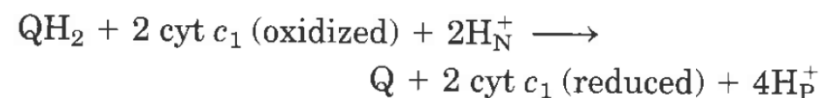
- Q-cytochrome c Oxidoreductase
- 11 subunits including two cytochrome subunits
- Contains iron sulfur center
- Contain three heme groups in two cytochrome subunits
- b_L and b_H in cytochrome b; c type in cytochrome c₁
- Contain two CoQ binding sites
- 4H⁺

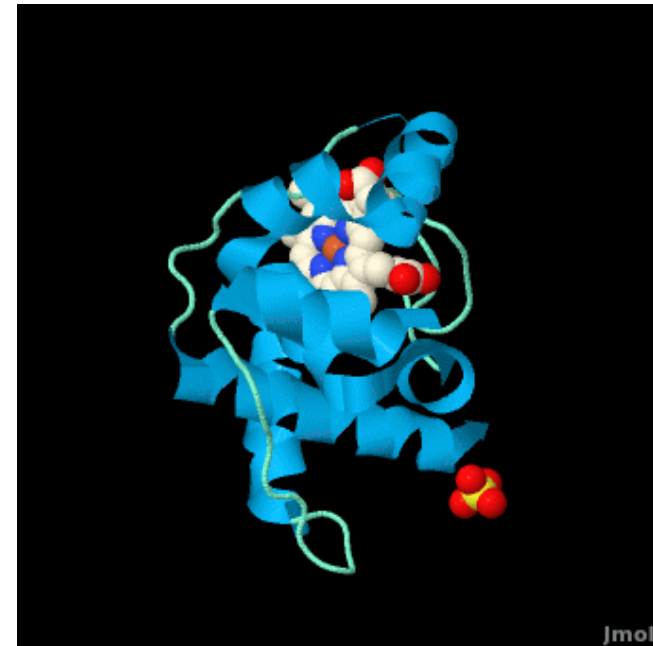
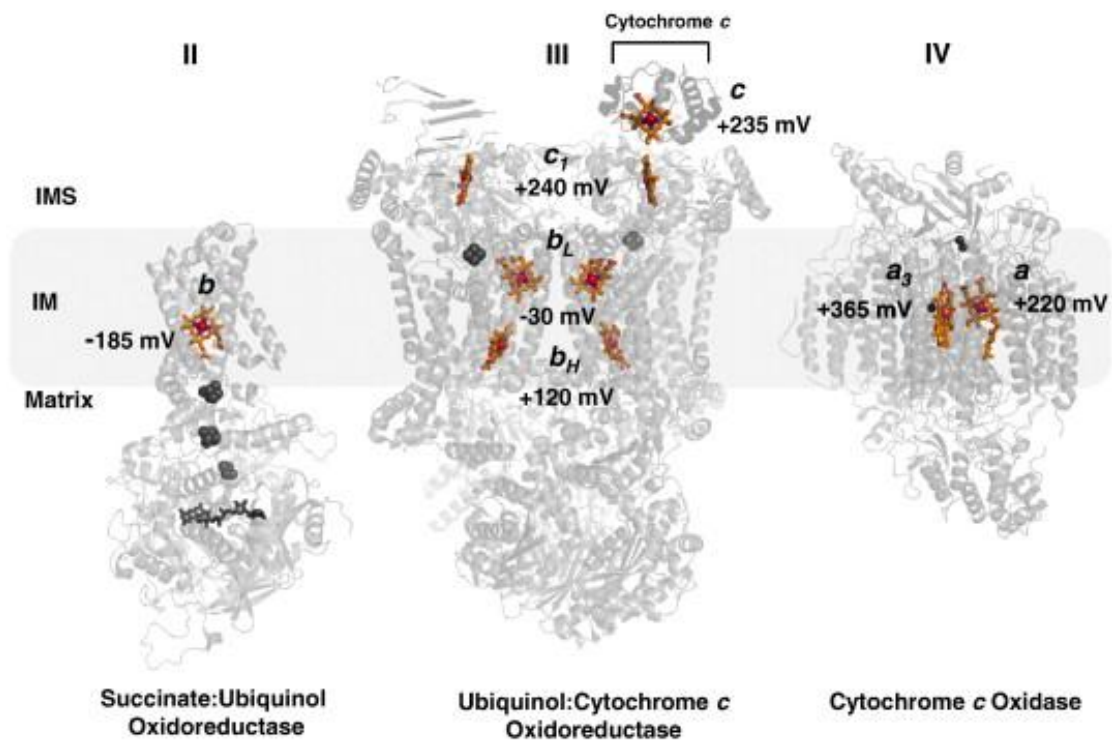




THE Q-CYCLE

- Partial reduction is hazardous
- Accommodates the switch between 2e⁻/1e⁻
- Explains the measured stoichiometry of 4 H⁺/2e⁻



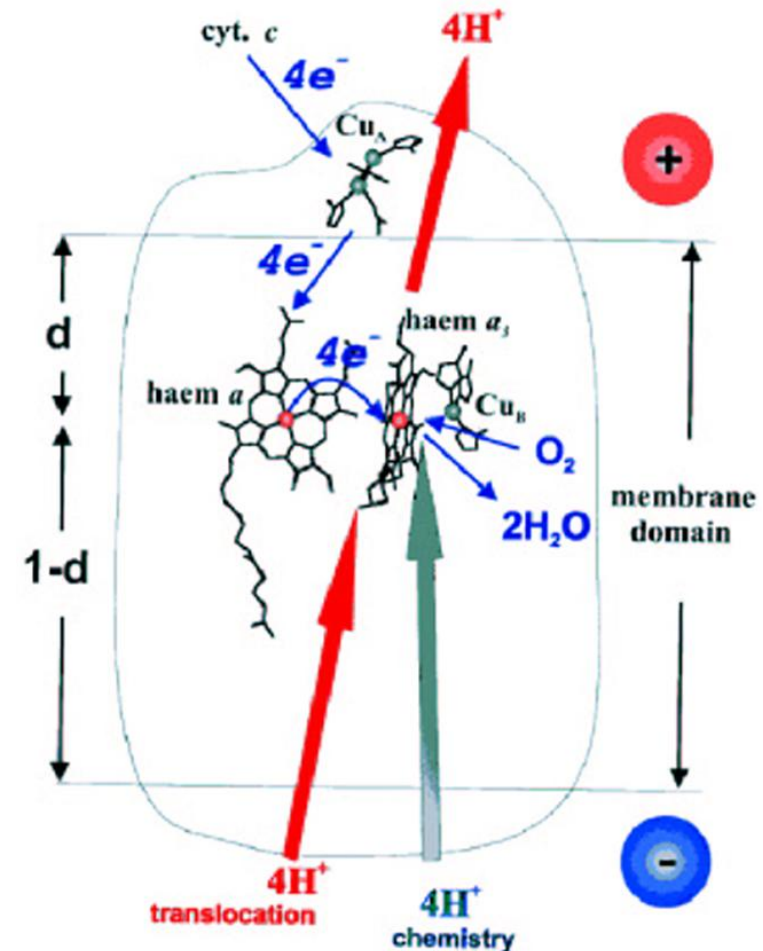


CYTOCHROME C: THE SURFACE MESSENGER



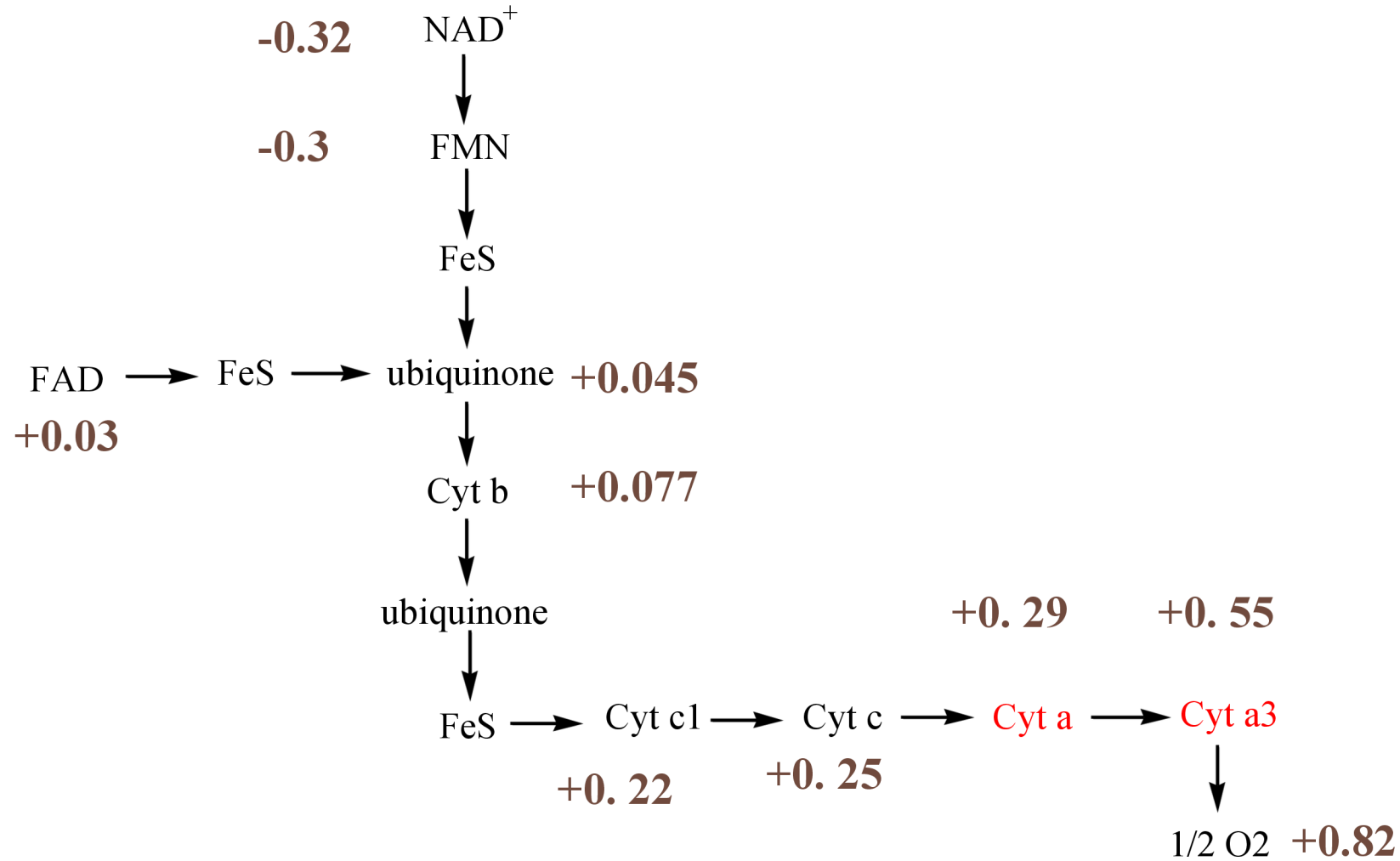
OXI-RED COMPONENTS OF THE ETC: “CYTOCHROME C OXIDASE” – COMPLEX IV

- Contains cytochrome a and a₃
- Contains two copper sites
- Contains oxygen binding sites
- O₂ must accept 4 electrons to be reduced to 2 H₂O (2H⁺/2e⁻)
- Cytochrome c is one electron carrier
 - $\text{Cyt } c_{\text{red}} + 4\text{H}^+ + \text{O}_2 \rightarrow \text{Cyt } c_{\text{ox}} + 2\text{H}_2\text{O}$
- Cytochrome oxidase has a much lower K_m for O₂ than myoglobin (hemoglobin, myoglobin, complex IV)
- Partial reduction of O₂ is hazardous



THE RIGHT ARRANGEMENT: HOW CAN WE PROVE IT?

1. MEASURING THE STANDARD REDUCTION POTENTIALS (REAL?)



THE RIGHT ARRANGEMENT: HOW CAN WE PROVE IT?

1. MEASURING THE STANDARD REDUCTION POTENTIALS (REAL?)

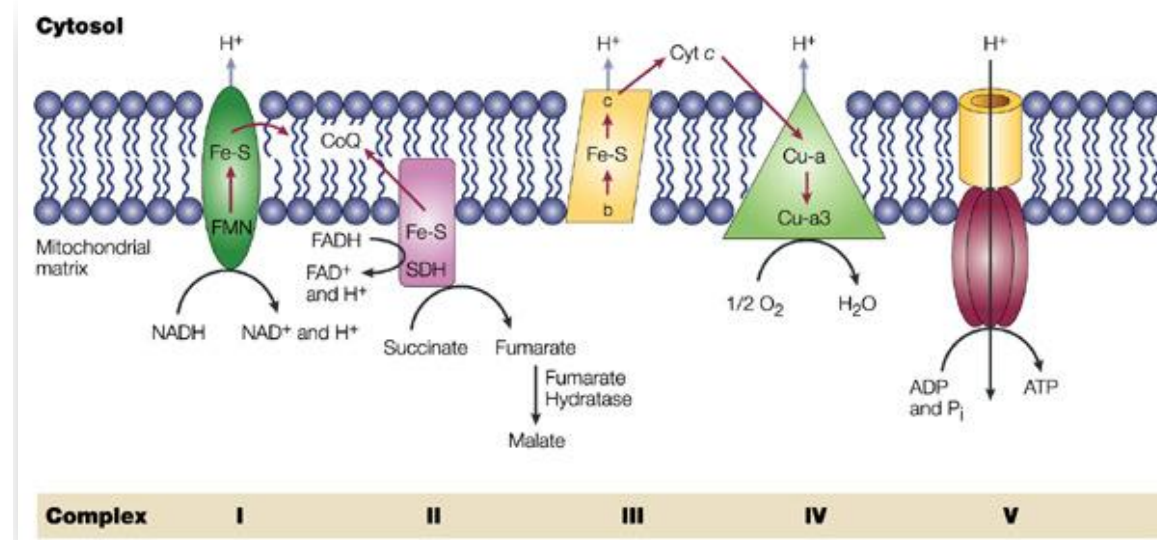
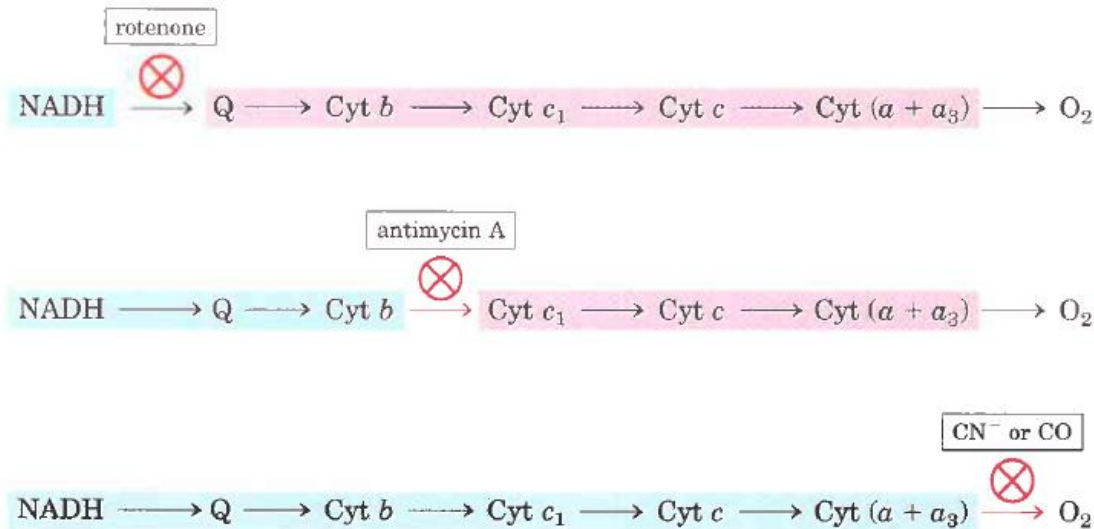
Redox reaction (half-reaction)	E'° (V)
$2\text{H}^{+} + 2e^{-} \longrightarrow \text{H}_2$	-0.414
$\text{NAD}^{+} + \text{H}^{+} + 2e^{-} \longrightarrow \text{NADH}$	-0.320
$\text{NADP}^{+} + \text{H}^{+} + 2e^{-} \longrightarrow \text{NADPH}$	-0.324
$\text{NADH dehydrogenase (FMN)} + 2\text{H}^{+} + 2e^{-} \longrightarrow \text{NADH dehydrogenase (FMNH}_2\text{)}$	-0.30
$\text{Ubiquinone} + 2\text{H}^{+} + 2e^{-} \longrightarrow \text{ubiquinol}$	0.045
$\text{Cytochrome } b (\text{Fe}^{3+}) + e^{-} \longrightarrow \text{cytochrome } b (\text{Fe}^{2+})$	0.077
$\text{Cytochrome } c_t (\text{Fe}^{3+}) + e^{-} \longrightarrow \text{cytochrome } c_t (\text{Fe}^{2+})$	0.22
$\text{Cytochrome } c (\text{Fe}^{3+}) + e^{-} \longrightarrow \text{cytochrome } c (\text{Fe}^{2+})$	0.254
$\text{Cytochrome } a (\text{Fe}^{3+}) + e^{-} \longrightarrow \text{cytochrome } a (\text{Fe}^{2+})$	0.29
$\text{Cytochrome } a_3 (\text{Fe}^{3+}) + e^{-} \longrightarrow \text{cytochrome } a_3 (\text{Fe}^{2+})$	0.35
$\frac{1}{2}\text{O}_2 + 2\text{H}^{+} + 2e^{-} \longrightarrow \text{H}_2\text{O}$	0.8166

NADH $\rightarrow$ Q $\rightarrow$ cytochrome b $\rightarrow$ cytochrome c1 $\rightarrow$ cytochrome c $\rightarrow$ cytochrome a $\rightarrow$ cytochrome a3 $\rightarrow$ O2

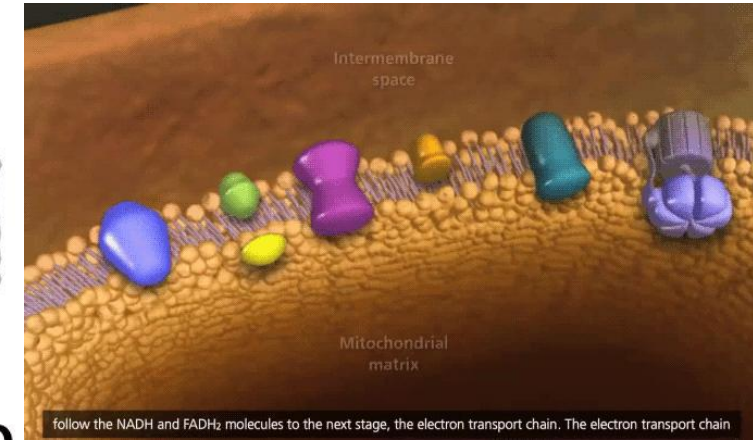
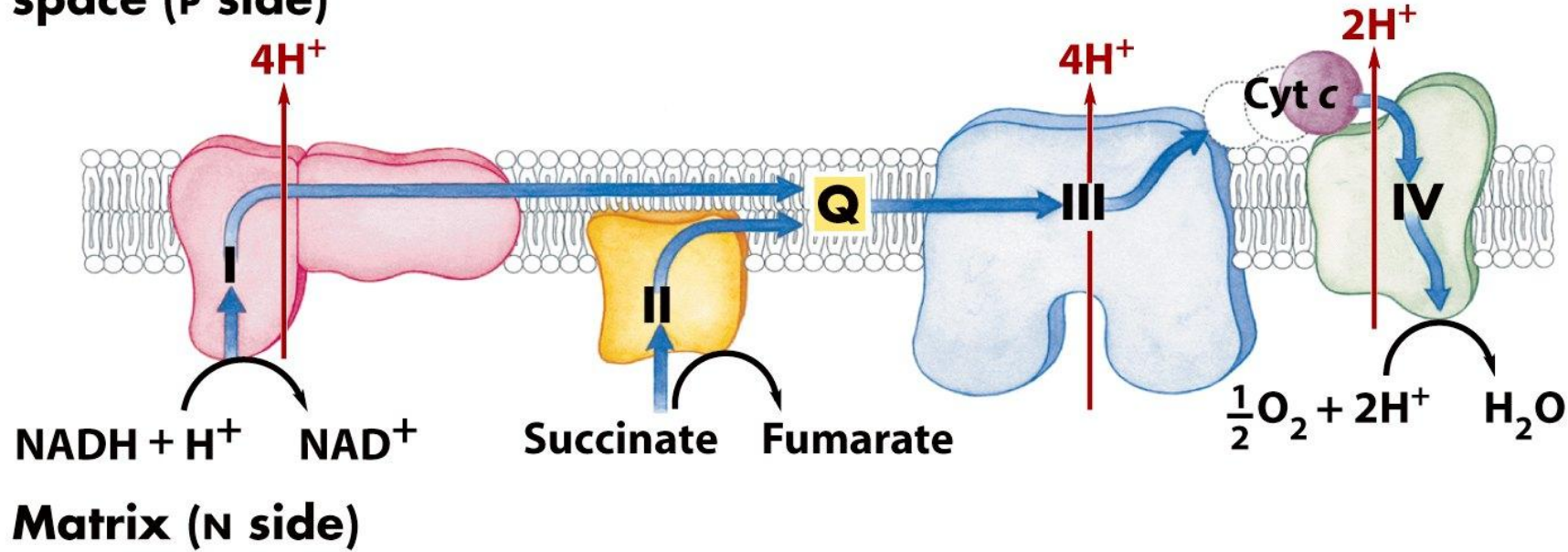
THE RIGHT ARRANGEMENT: HOW CAN WE PROVE IT?

NADH → Q → cytochrome b → cytochrome c₁ → cytochrome c → cytochrome a → cytochrome a₃ → O₂

- 2. Reduction of the entire ETC with no O₂
- 3. Addition of inhibitors



**Intermembrane
space (P side)**



PUMPING OF PROTONS

- For every 2 electrons passing:
- 4H^+ (complex I); 0H^+ (complex II); 4H^+ (complex III), 2H^+ (complex IV)

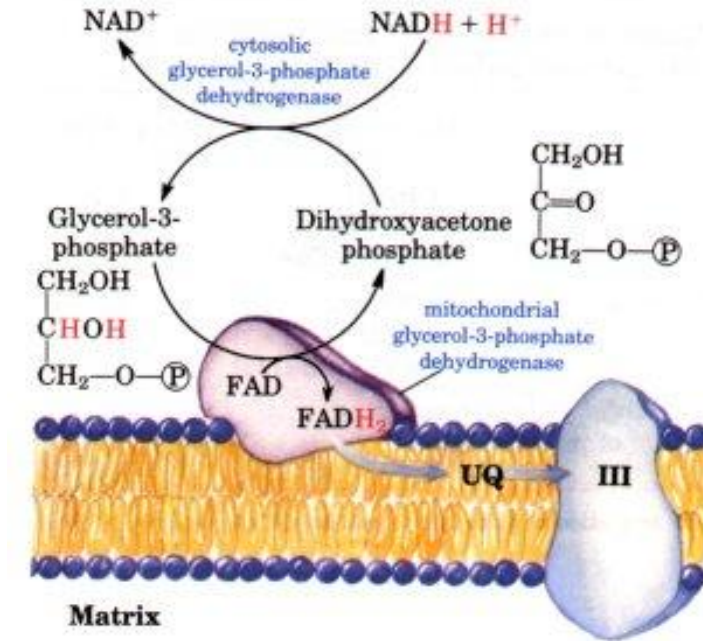
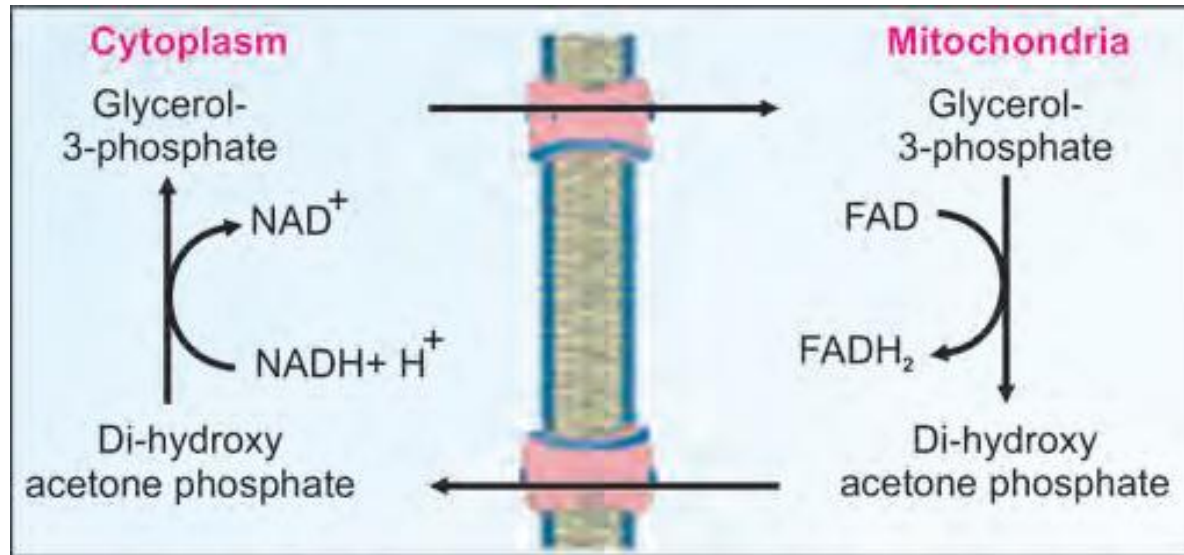


Box 37.3: NAD⁺ dependent enzymes

1. Lactate dehydrogenase (lactate → pyruvate) (see Fig. 9.14)
2. Glyceraldehyde-3-phosphate dehydrogenase (glyceraldehyde-3-phosphate → 1,3-bisphosphoglycerate) (see Fig.9.10)
3. Pyruvate dehydrogenase (pyruvate → acetyl CoA) (see Fig.9.22)
4. Alpha ketoglutarate dehydrogenase (alpha ketoglutarate → succinyl CoA) (see Fig.19.2)
5. Beta hydroxyacyl CoA dehydrogenase (beta hydroxyacyl CoA → beta ketoacyl CoA (see Step 3, Fig.12.9)
6. Glutamate dehydrogenase (Glutamate → alpha ketoglutarate (see Fig.15.9)

**SOME NADH
PRODUCING
ENZYMES**





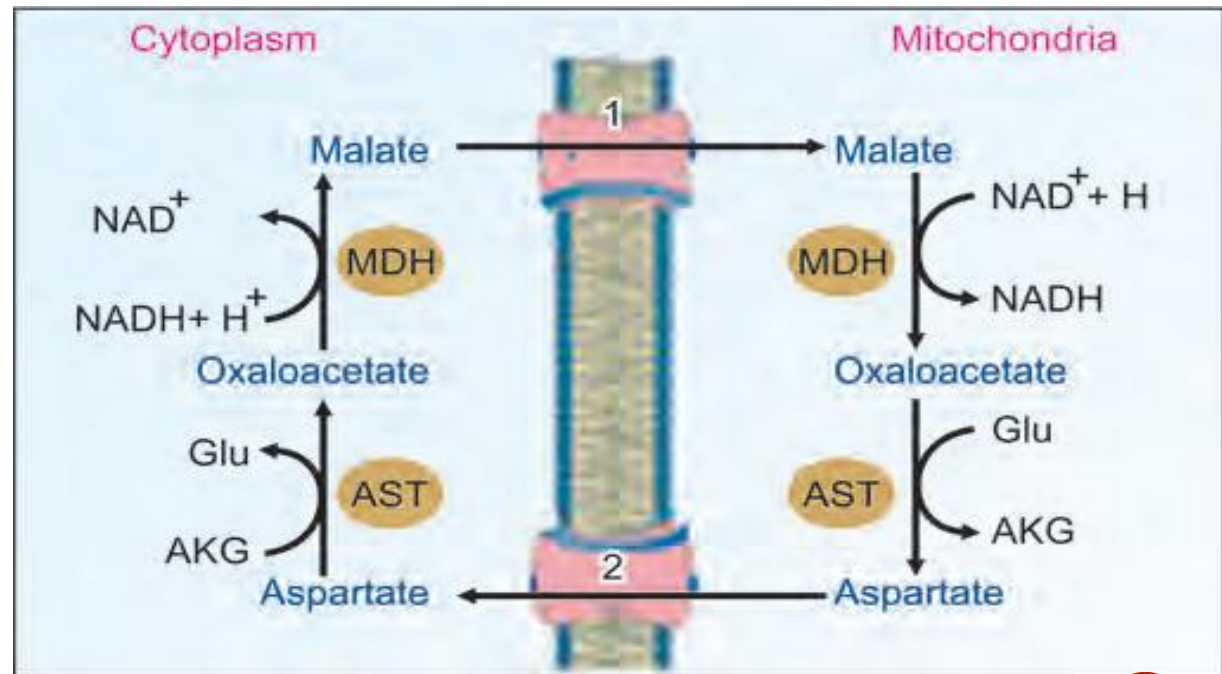
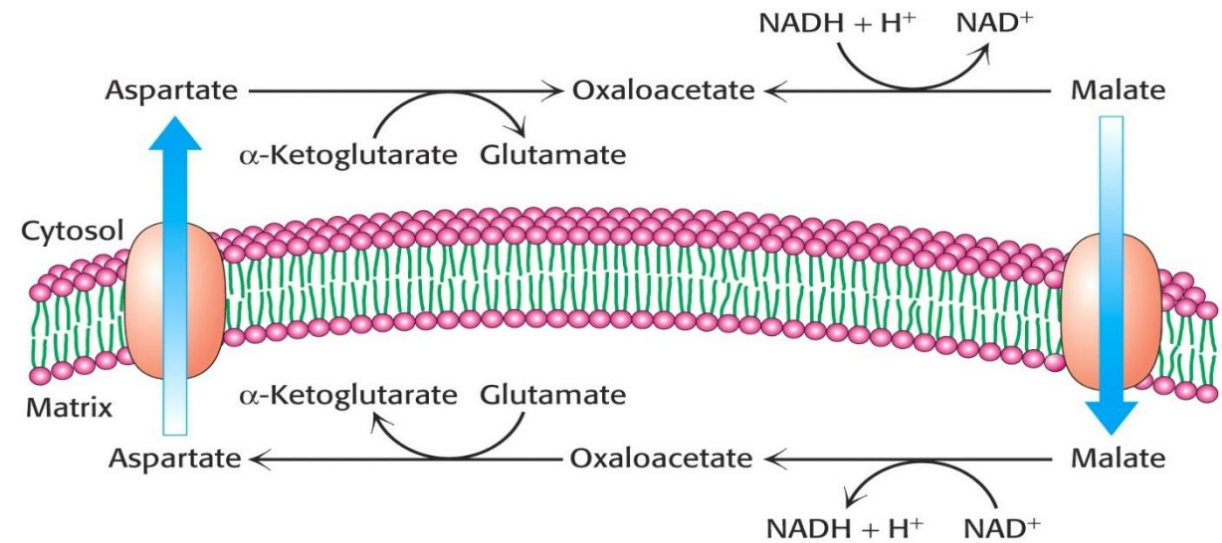
MITOCHONDRIAL SHUTTLING SYSTEMS “CYTOSOLIC NADH”

- Glycerol 3-phosphate shuttle
- In skeletal muscle and brain
- Glycolytic pathway as an example
- How NADH passes?
- ATP yield?



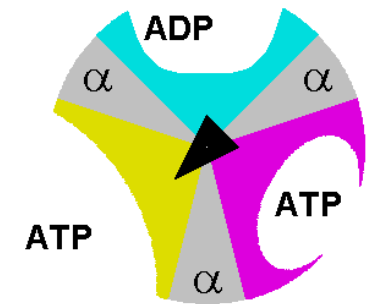
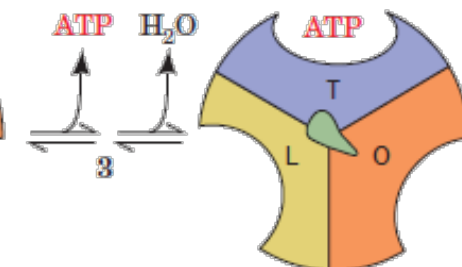
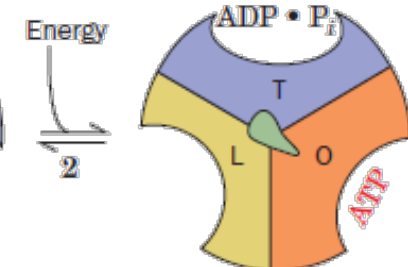
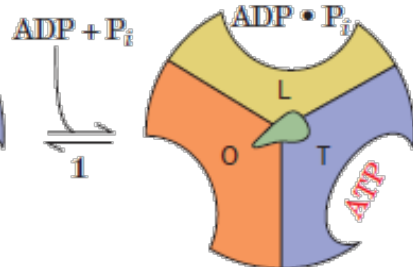
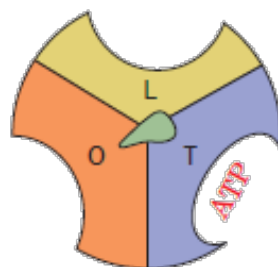
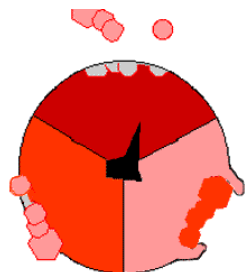
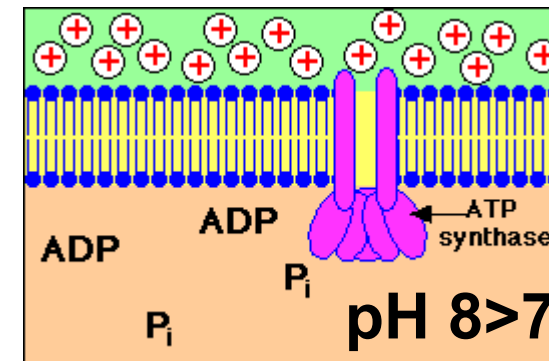
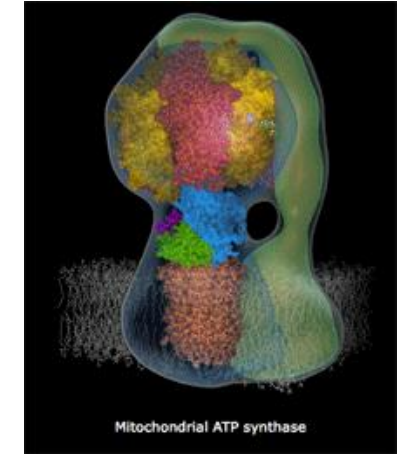
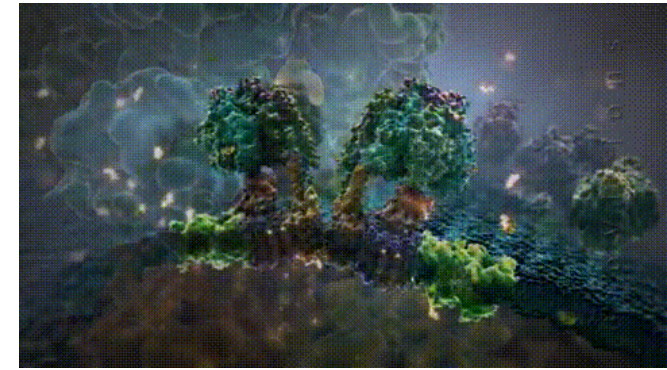
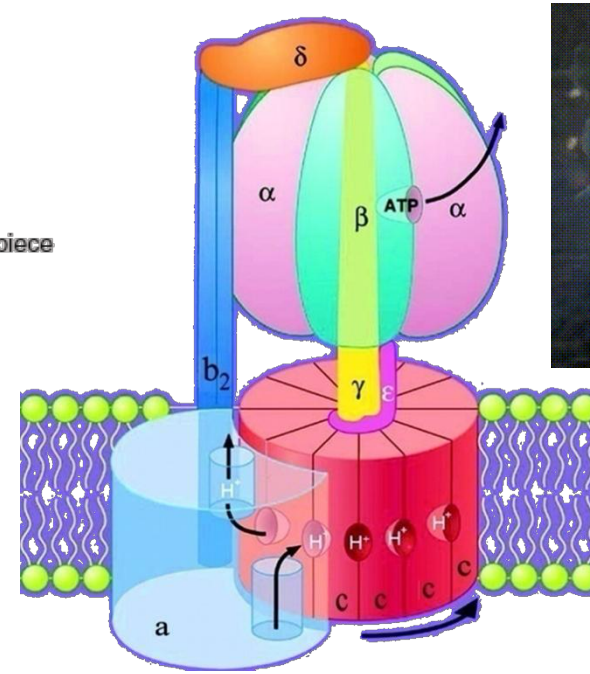
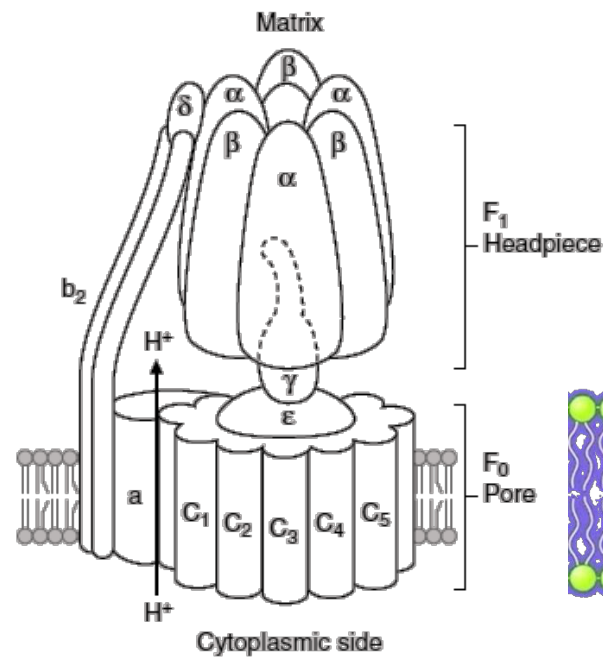
MITOCHONDRIAL SHUTTLING SYSTEMS - "CYTOSOLIC NADH"

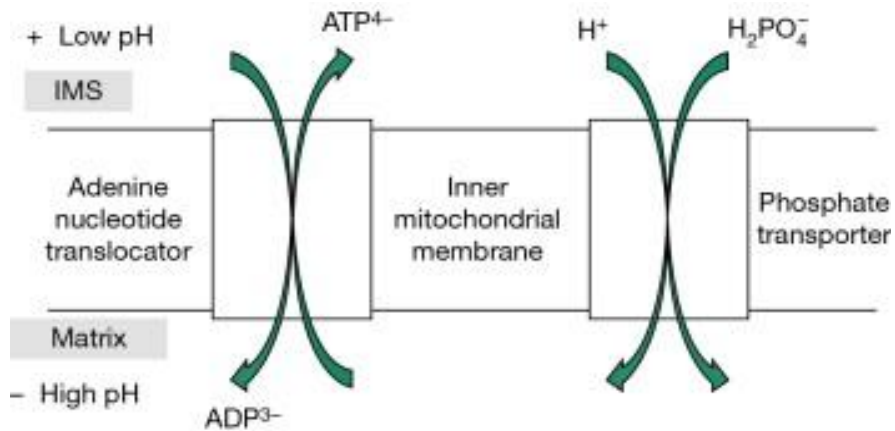
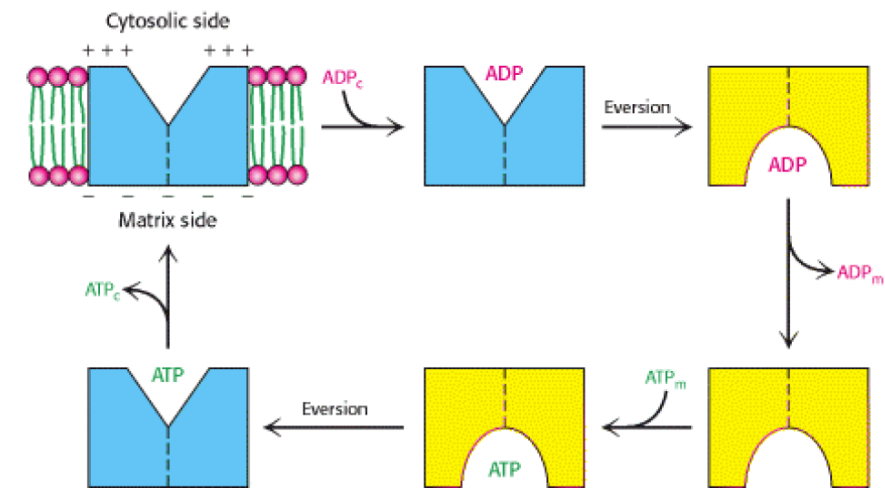
- Malate-Aspartate shuttle
- operates mainly in liver, kidney
- 2 membrane carriers & 4 enzymes
- Readily reversible (vs. Glycerol 3-phosphate shuttle)
- NADH can be transferred only if the NADH/NAD⁺ ratio is higher in the cytosol than in the mitochondrial matrix
- Exchange of key intermediates between mitochondria & cytosol



ATP SYNTHASE

- F₁:
 - “γ” subunit: rotates
 - “β” subunit: binds
 - “α” subunit: structural
 - 3 conformations: tight (T), loose (L), open (O)
- F₀:
 - “a” subunit: point of entry & exit
 - “c” subunit rotates
 - 4H⁺/ATP
- Can run backwards

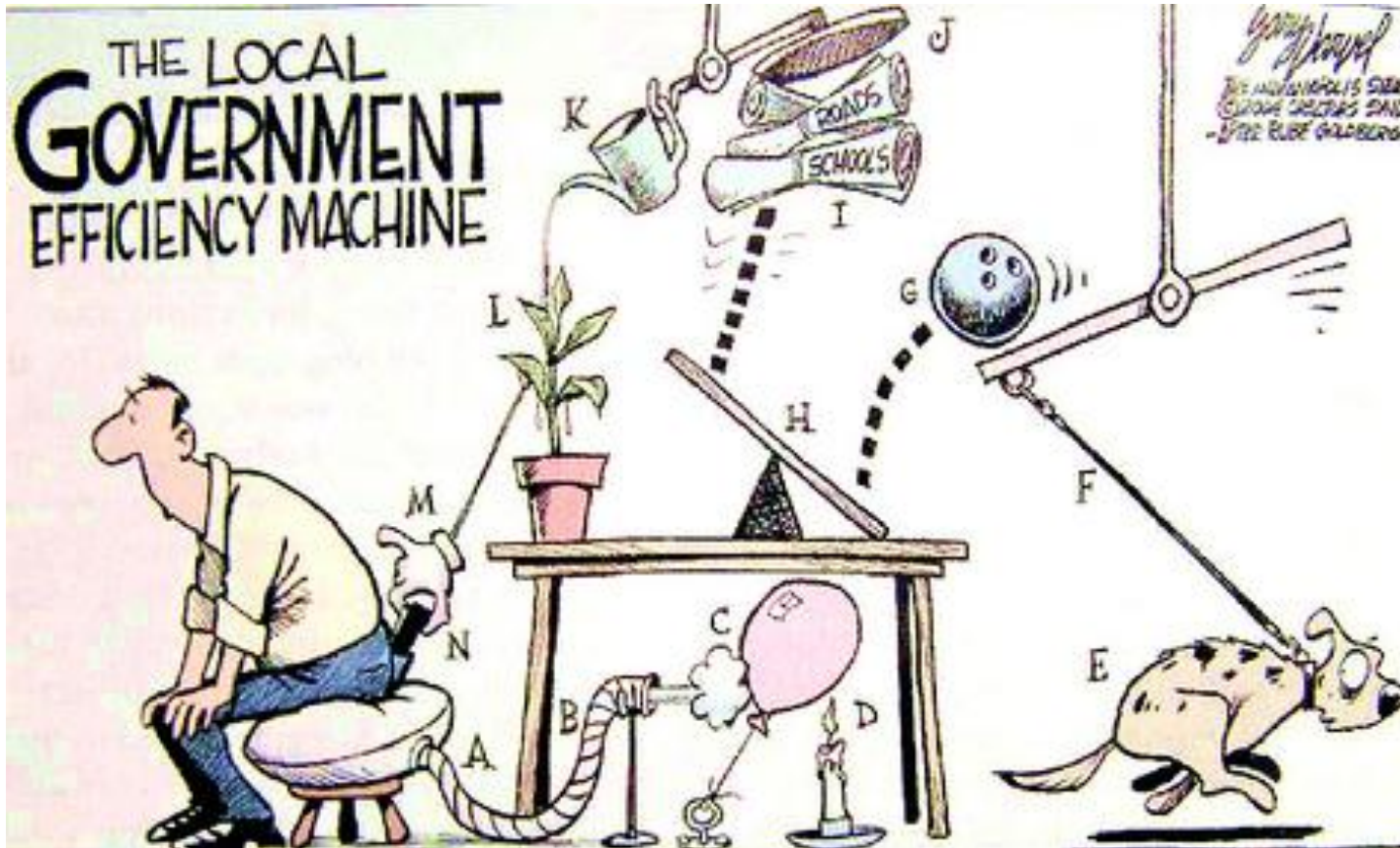




MITOCHONDRIAL SHUTTLING SYSTEMS “ATP/ADP”

- ATP-ADP Translocase (also called adenine nucleotide translocase or ANT)
- The flows of ATP and ADP are coupled (ADP enters only if ATP exits, and vice versa)
- Highly abundant (14% of IMM proteins)
- Contains a single nucleotide-binding site (alternates)
- Similar affinity to ATP and ADP
- Endergonic (25% of ETC)
- Inhibition leads to subsequent inhibition of cellular respiration





ENERGY YIELD FROM THE ETC

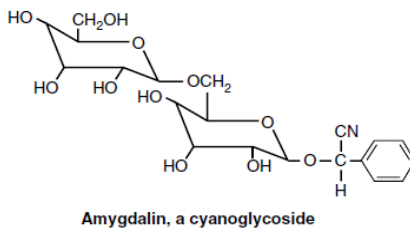
- NADH, -53 kcal, ATP?
- FADH₂, -41 kcal, ATP?
- ΔG° is so negative, never reversible
- ATP machine efficiency, (anions, Ca^{+2} , heat, phosphate, substrates)



REGULATION – THE NEED FOR ATP

- In skeletal muscles, 20% drop in ATP concentration
- In the heart, Ca^{+2} activates TCA enzymes for extra push (NADH, ATP), no drop
- ET is tightly coupled to phosphorylation (simultaneously)
- ADP is the most important factor in determining the rate (respiratory control)





Anit-cancerous drug

الصفحة الرئيسية > مخطبات

أشهر جرائم القتل العائلية في المملكة

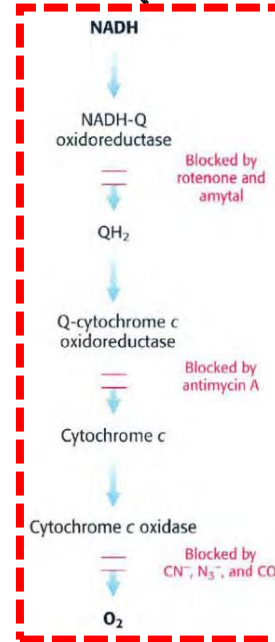
جراسا نيوز -

جراسا -تعرض فيما يلي قائمة بأشهر جرائم القتل العائلية التي حدثت في الاردن خلال السنوات الماضية ، والتي كان لكل منها وقع الصدمة حين وقوعها لما تمثله من فعل غريب على المجتمع وأعرافه ، فضلا عن مخالفتها الشرائع السماوية والقوانين النافذة والطبيعة الإنسانية بعامه.

قضية السيانيذ

أول جريمة من نوعها يرتكبها أب ضد ولديه ، إذ قام الاب بوضع مادة السيانيذ في كأس الحليب وطلب من طفليه ان يشربا منه ، حيث فارقا الحياة بعد 10 دقائق من مغادرة الام المنزل لتعود وتجددهما جثتين هامدتين.

وقد ادين الاب بعقوبة الاعدام شنقا الا ان والده اسقط الحق الشخصي كونه وليا عن الطفلين وحكم عليه بالاشغال المؤبدة.



REGULATION — INHIBITION (COUPLING)

➤ Specific inhibitors:

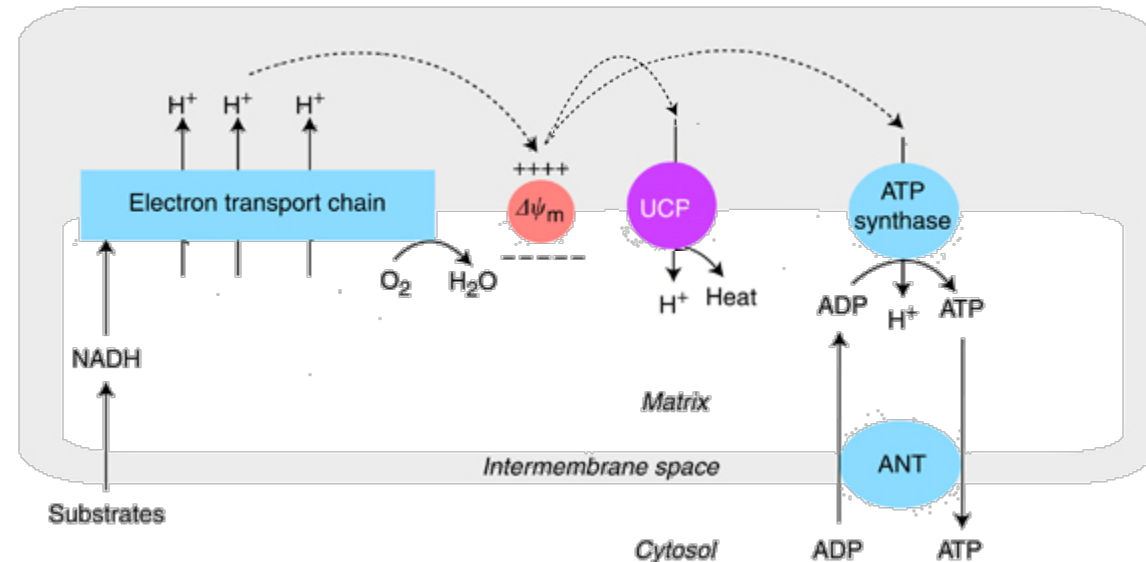
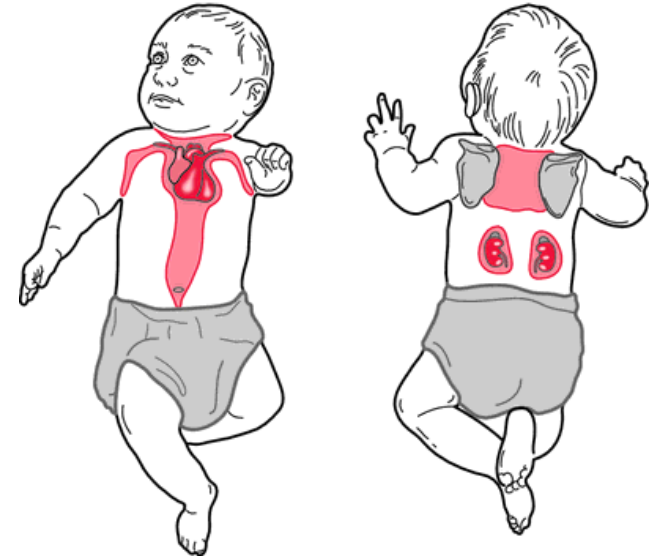
- ✓ Cyanoglycosides such as amygdalin are present in edible plant pits

Specific inhibitor	Target
Rotenone (insecticide) & Amytal (sedative)	NADH-Q oxidoreductase
Antimycin A (antibiotic)	Q-cytochrome c oxidoreductase
Cyanide (CN ⁻), Azide (N ₃ ⁻), & (CO)	Cytochrome c oxidase
Oligomycin (antibiotic)	ATP synthase



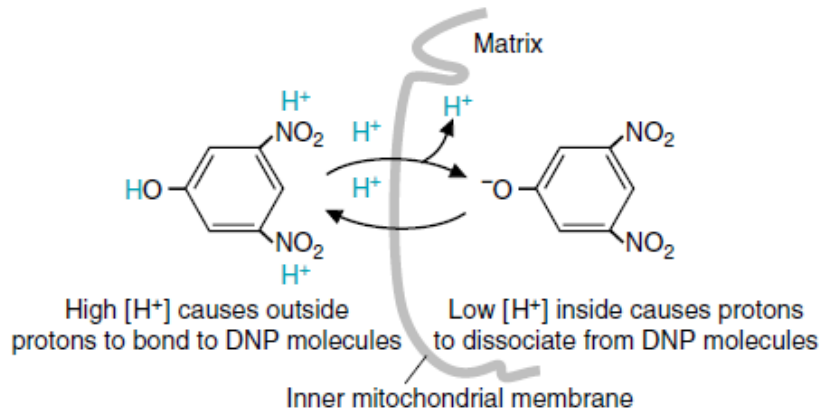
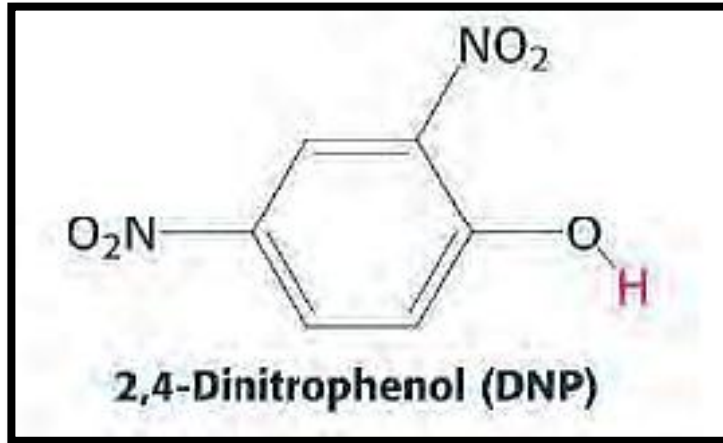
REGULATION – UNCOUPLING REGULATED – UNCOUPLING PROTEINS (UCPS)

- Short-circuiting ATP synthase
- UCP1 (thermogenin):
 - ✓ Brown adipose tissue, non-shivering thermogenesis
 - ✓ Infants: neck, breast, around kidneys
 - ✓ Fatty acids directly activates UCP1
- UCP2 (most cells); UCP3 (skeletal muscle); {UCP4, UCP5} (brain)
- Obesity tendency in some populations



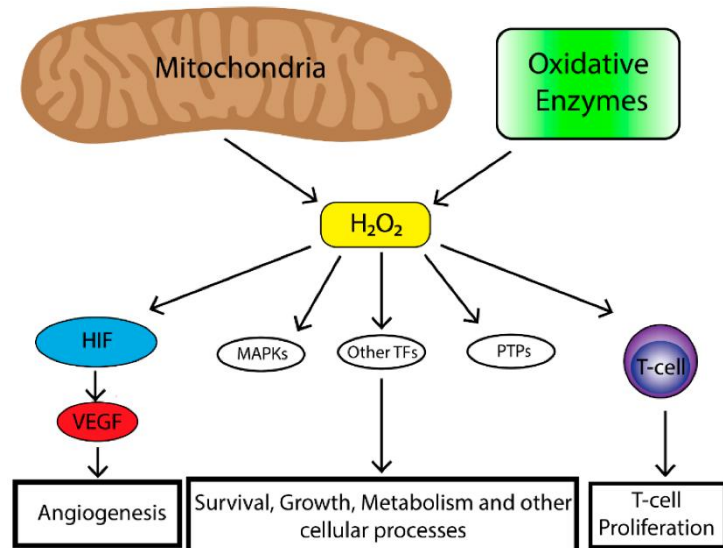
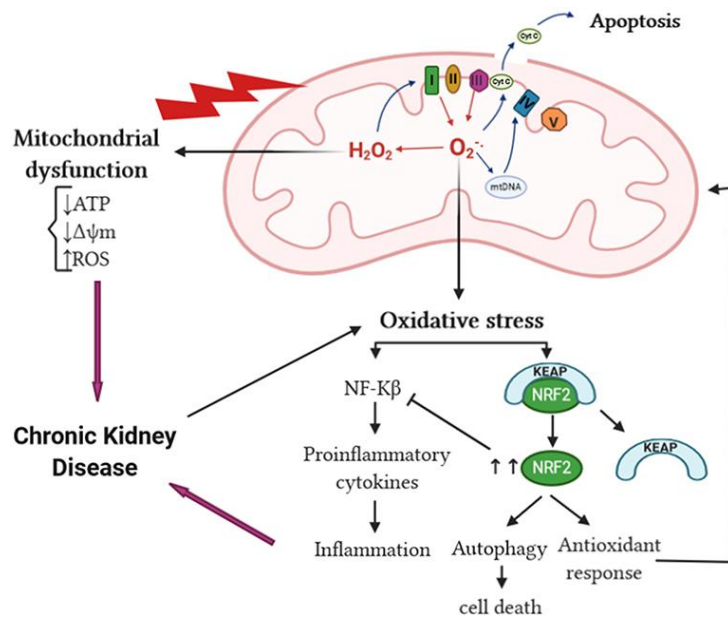
REGULATION – UNCOUPLING

UNREGULATED – CHEMICAL UNCOUPLERS



- What is uncoupling?
- How does it occur? Dissipation of PMF
- What is the result?
- Is it physiological or not?
- 2,4-dinitrophenol (DNP) & other acidic aromatic compounds
- What changes happen? ↑ O₂ consumption, ↑ NADH oxidation
- Soviet soldiers were given DNP, FDA banned DNP (1938)



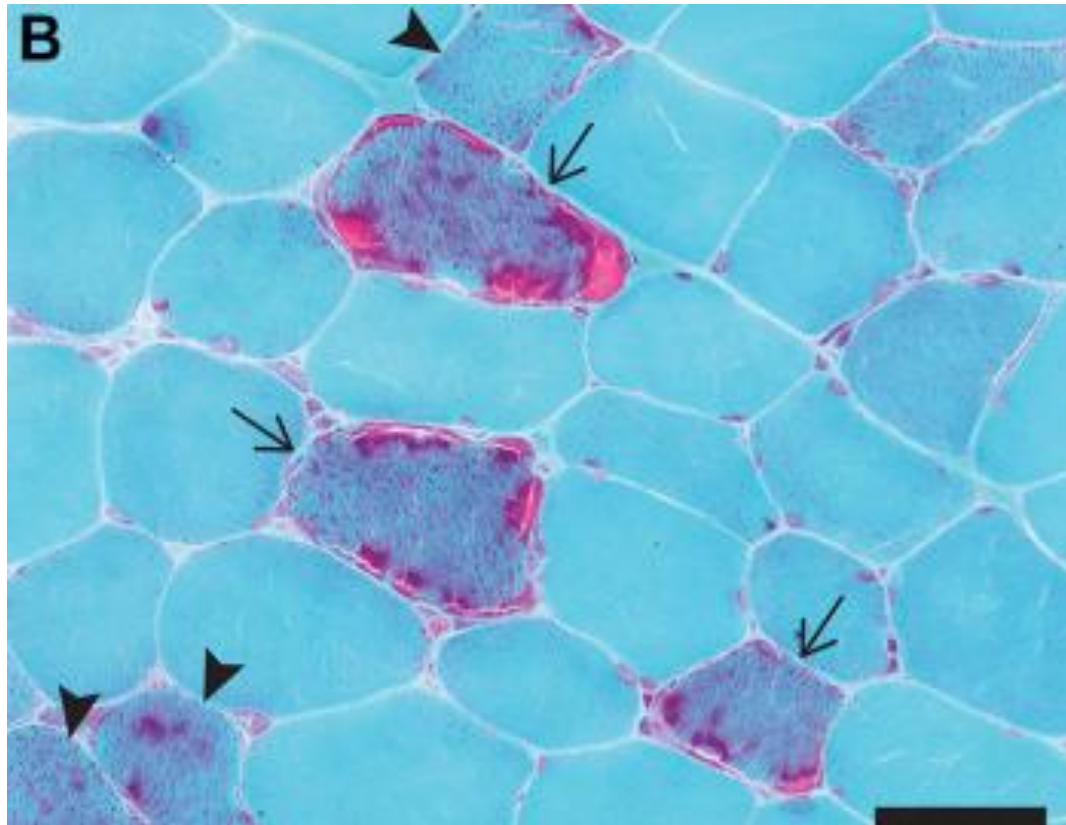


THE DARK SIDE: REACTIVE OXYGEN SPECIES (ROS)

- E.g., FMN, semiquinone in Q cycle
- Single electrons can leak and reduce O_2 to superoxide ($O_2^{\bullet-}$)
- **Pathophysiology:**
 - **Primary ROS:** Superoxide ($O_2^{\bullet-}$). Converted by Superoxide Dismutase (SOD) to H_2O_2
 - **Damage:** lipids (peroxidation), proteins (carbonylation), and DNA (strand breaks)
 - **Role in Disease:** Contributes to aging, neurodegenerative diseases (Parkinson's, Alzheimer's), ischemia-reperfusion injury, and inflammatory damage

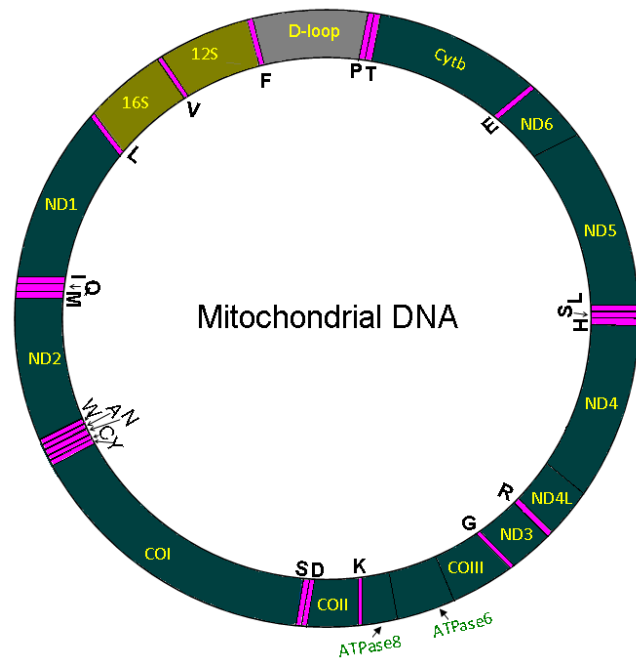
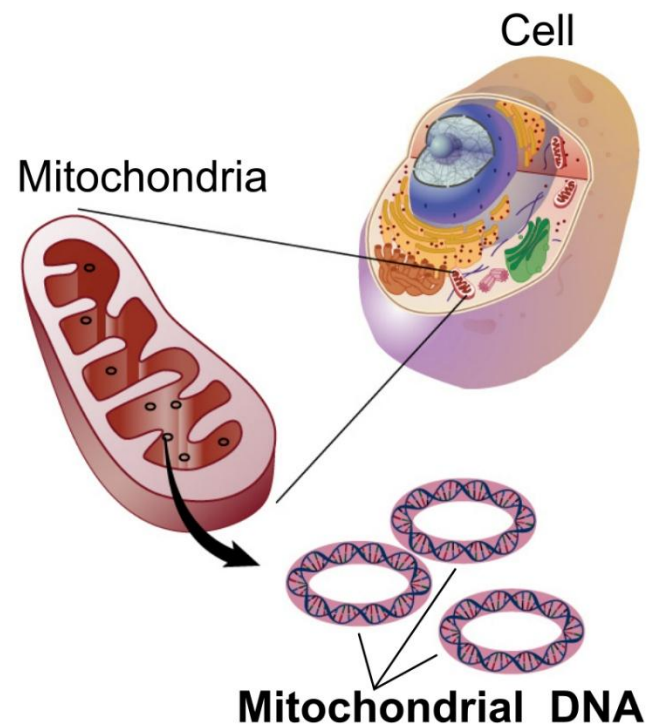


MITOCHONDRIAL DISEASES: GENETICS AND PRINCIPLES



- A muscle biopsy with "ragged red fibers" (Gomori trichrome stain)
- Etiology:
 - nDNA or mtDNA
 - mtDNA Inheritance: Maternal. High mutation rate, no histones, poor repair.
 - Heteroplasmy: A mixture of wild-type and mutant mtDNA within a cell.
 - Symptoms appear when the mutant load exceeds a threshold.
- Tissues Affected: High-energy demand tissues: CNS, muscle, heart, liver, kidney.





OXPHOS DISEASES – 1 (GENETIC)

- Mitochondrial DNA and OXPHOS Diseases
 - Small (16,569) base pair, double-stranded, circular DNA
 - Encodes 13 subunits: 7 (I) , 1 (III), 3 (IV), 2 (F0)
 - Also encodes necessary components for translation of its own mRNA: a large and small rRNA and tRNAs
 - Maternal inheritance, replicative segregation and heteroplasmy
 - Accumulation of somatic mutations with age
 - Highest ATP demands: CNS, heart, skeletal muscle, and kidney, liver

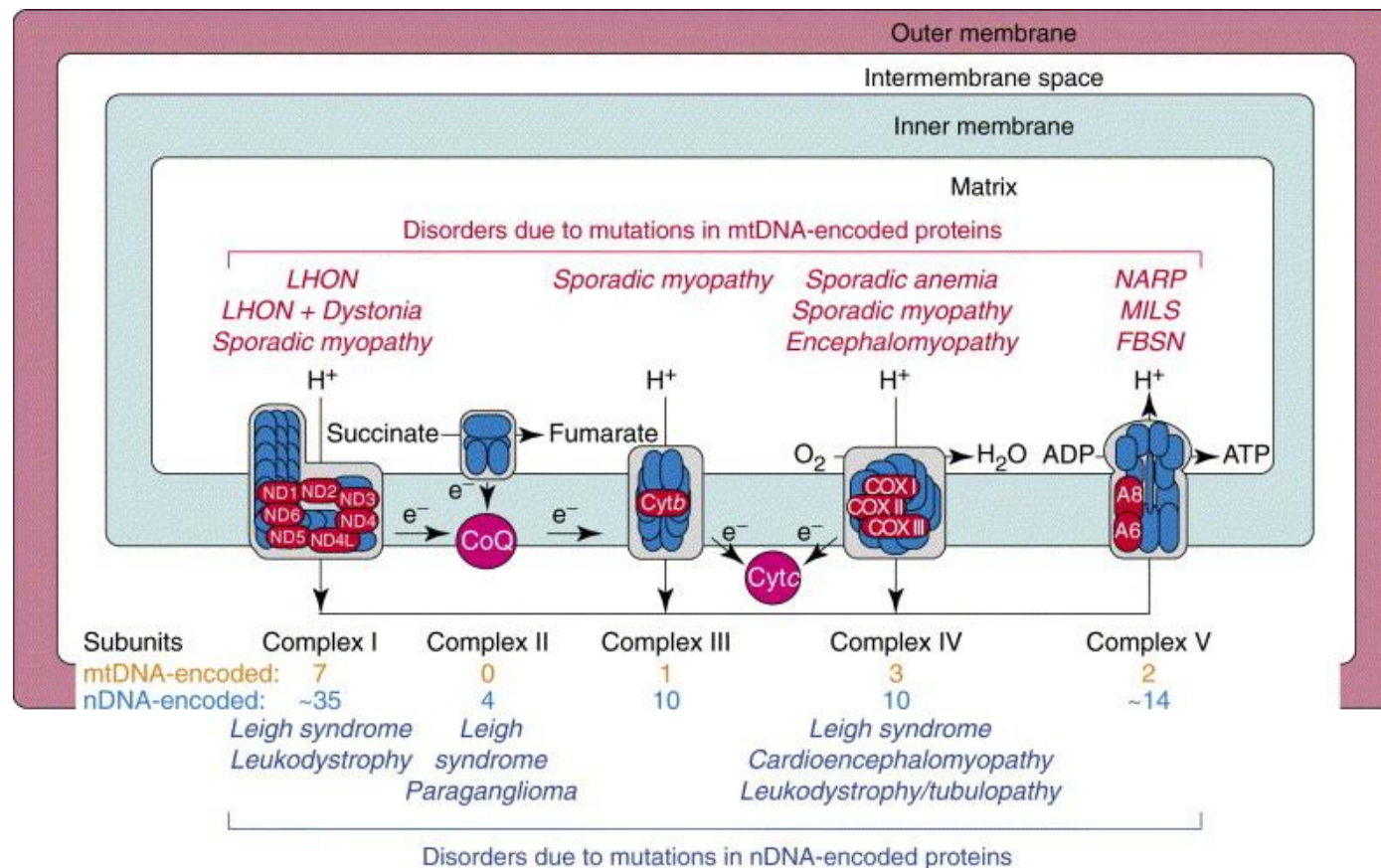


OXPPOS DISEASES

— 2 (GENETIC)

- Nuclear Genetic Disorders of Oxidative Phosphorylation
 - 1,000 proteins
 - Usually autosomal recessive
 - Expressed in all tissues
 - Phenotypic expression with high ATP demand

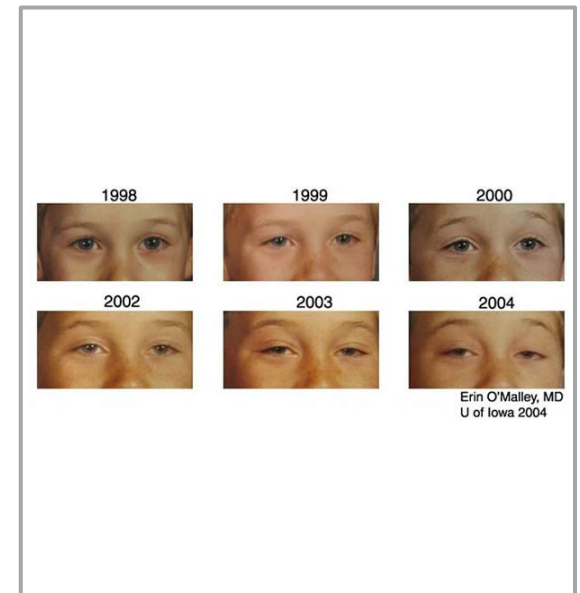
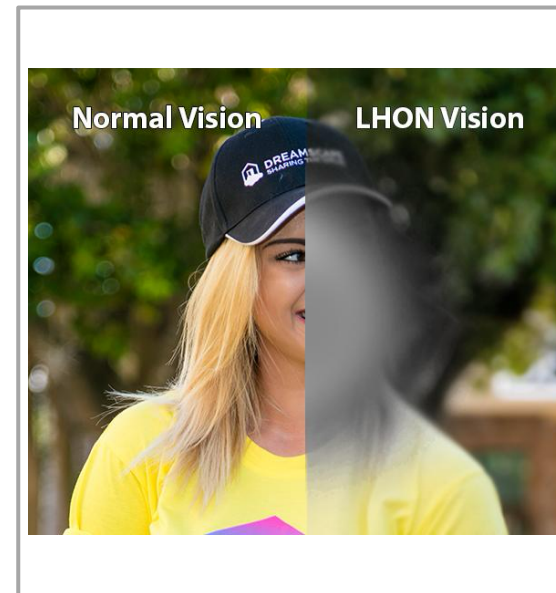
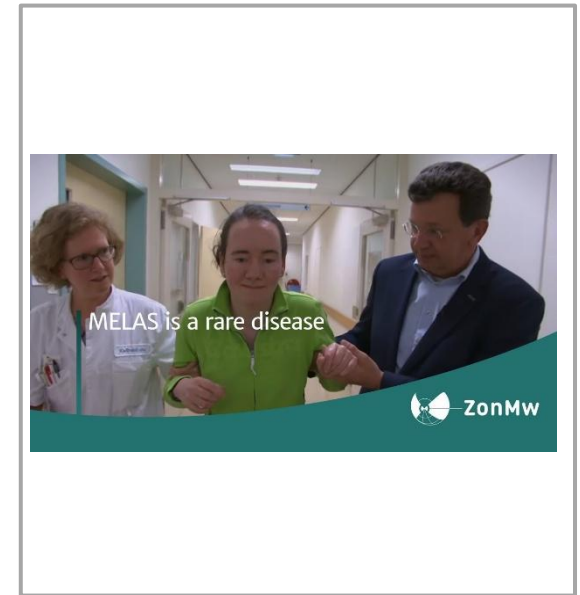
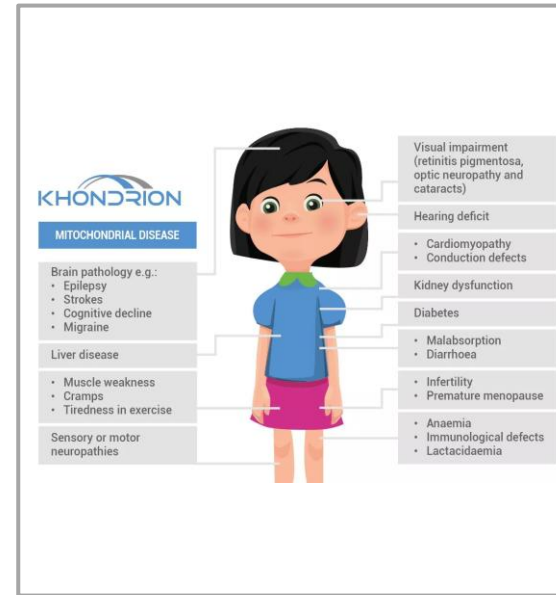
OXPHOS DISEASES (GENETIC)



MITOCHONDRIAL DISEASES 2: CLINICAL EXAMPLES

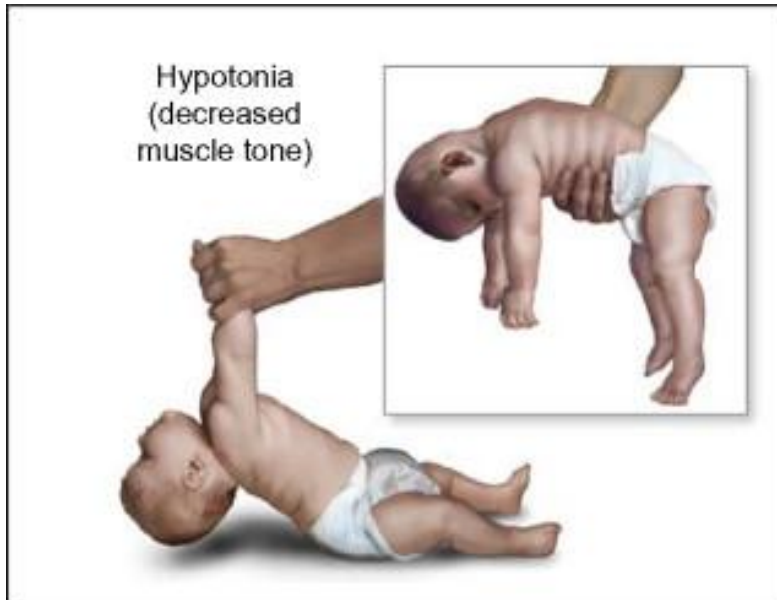
■ Examples:

- **Leber's Hereditary Optic Neuropathy (LHON):** mutation in Complex I. Causes sudden, painless bilateral vision loss in young adults.
- **MELAS:** Mitochondrial Encephalomyopathy, Lactic Acidosis, and Stroke-like episodes. Caused by mtDNA point mutation.
- **Chronic Progressive External Ophthalmoplegia (CPEO):** ptosis and inability to move the eyes. Often associated with large-scale mtDNA deletions.



INTERACTIVE CASE STUDY

CLINICAL CORRELATION: A DIAGNOSTIC PUZZLE



■ **Presentation:**

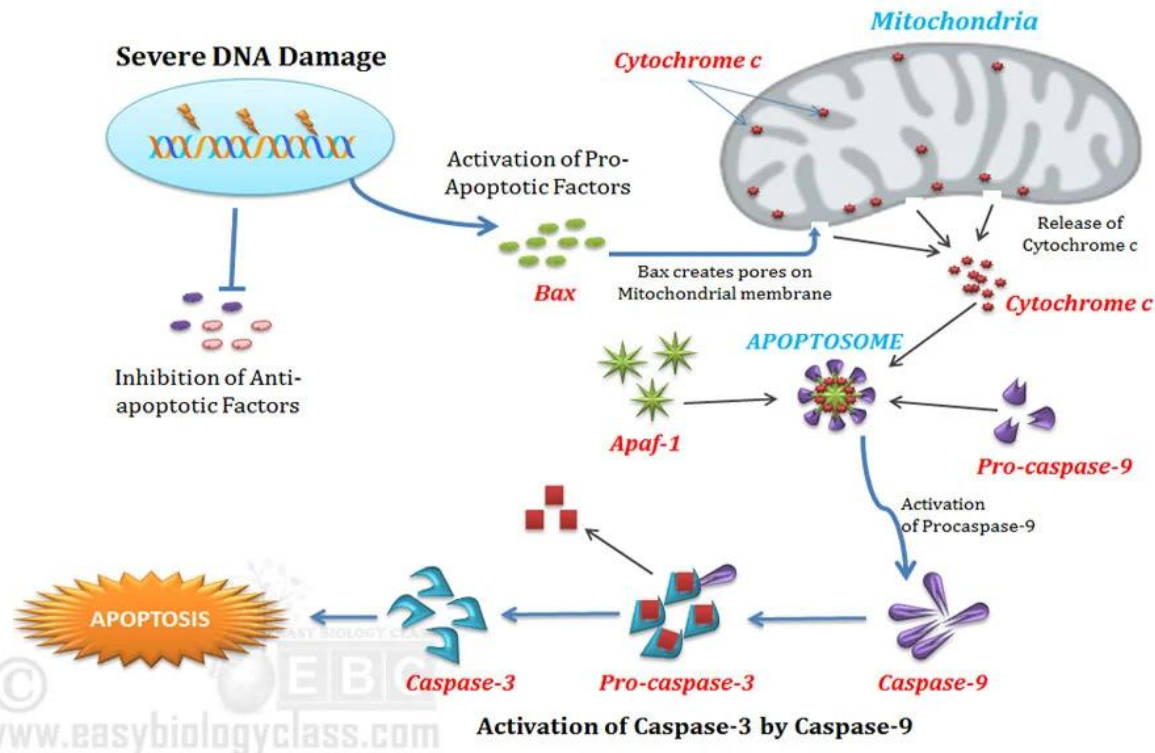
- A 4-year-old boy presents with progressive muscle weakness, exercise intolerance, and developmental delay.
- Lab tests: profound **lactic acidosis**.
- Muscle biopsy: **ragged red fibers**.

■ **Questions:**

- What is the most likely category of disease?
 - **(Mitochondrial Myopathy)**
- What is the biochemical basis for lactic acidosis?
 - **(Impaired OXPHOS → anaerobic glycolysis)**
- What molecular tests could be ordered?
 - **(mtDNA sequencing, nuclear gene panel for OXPHOS proteins)**



INTRINSIC PATHWAY OF APOPTOSIS (Mitochondria Mediated Programmed Cell Death Pathway)



BEYOND ENERGY: MITOCHONDRIA IN APOPTOSIS (CELL DEATH) AND AGING

- **Apoptosis:** In response to severe stress, mitochondria release Cytochrome c and other proteins, which activate caspases and trigger programmed cell death.
- **The Mitochondrial Theory of Aging:** Accumulation of mtDNA mutations over a lifetime → progressive decline in OXPHOS capacity → increased ROS production → further damage → tissue dysfunction and aging.

