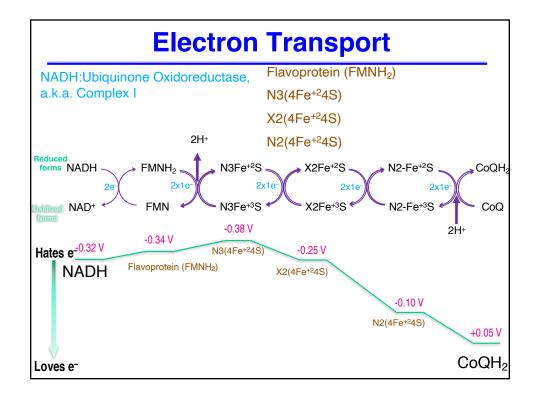
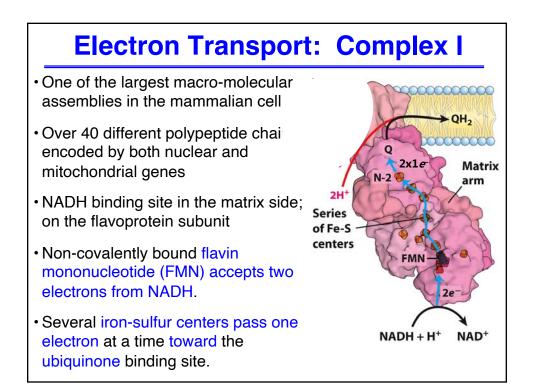
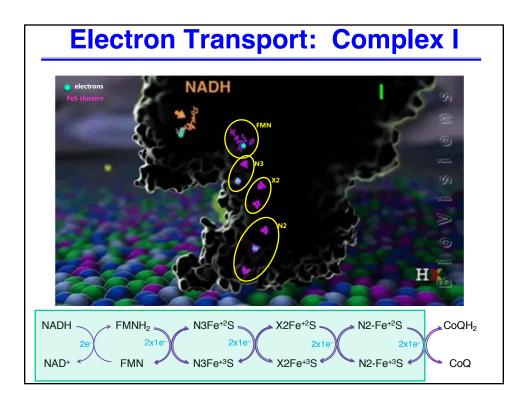
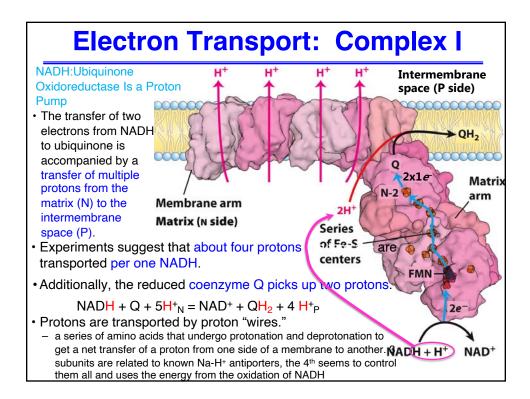
| BI/CH 422/622  |                   |
|--|-------------------|
| OUTLINE:<br>Introduction and review<br>Transport   |                   |
|  |                   |
| Pasteur: Anaerobic vs Aerobic  | Exam-1 material   |
| Fermentations  | Exam-2 material   |
| Pyruvate   |                   |
| pyruvate dehydrogenase (ox-decarbox; S-ester)<br>Krebs' Cycle  |                   |
| How did he figure it out?<br>Overview<br>8 Steps<br>Citrate Synthase (C-C)<br>Aconitase (=, -OH)<br>Isocitrate dehydrogenase (ox-decarbox; =0)<br>Succinyl-CoA synthetase (sub-level phos)<br>Succinate dehydrogenase (=)<br>Fumarase (-OH)<br>Malate dehydrogenase (=O)<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary<br>Suffmary | arrow Window      |
| Oxidative Phosphorylation<br>Energetics (-0.16 V needed for making ATP)  | <u>mow window</u> |
| Mitochondria<br>Transport (2.4 kcal/mol needed to transport H+ out)  |                   |
| Electron transport<br>Discovery<br>Four Complexes<br>Complex I: NADH → CoQH2   |                   |
| Complex II: Succinate $\rightarrow$ CoQH <sub>2</sub>  |                   |
| Complex III: CoQH₂ → Cytochrome C (Fe²+)<br>Complex IV: Cytochrome C (Fe²+) → H₂O  |                   |

|  |               | oonents of t<br>spiratory Ch |  |   |                                 |  |
|--|---------------|------------------------------|--|---|---------------------------------|--|
| Enzyme<br>complex/protein  | Mass<br>(kDa) | Number of<br>subunitsª       | Prosthetic<br>group(s)   | Reduction<br>potential<br>( <i>E</i> ₀́V) | Binding sites<br>for:           | Inhibted by:                             |
| NADH dehydrogenase   | 850           | 45 (14)                      | FMN, Fe-S  | -0.36                                     | NADH, CoQ                       | amytal, rotenone                         |
| l Succinate<br>Jehydrogenase   | 140           | 4                            | FAD-E, Fe-S  | 0.09<br>(Co                               | 5 Succinate,<br>Q) CoQ          | malonate                                 |
| II Ubiquinone:<br>cytochrome c<br>oxidoreductase <sup>b</sup>                                  | 250           | 11                           | Hemes b, c <sub>1</sub> ,<br>Fe-S                              | 0.17                                      | CoQ,<br>Cytochrome c            | antimycin a                              |
| Cytochrome c <sup>c</sup>  | 13            | 1                            | Heme   | 0.25                                      | i (Cyt c)                       |  |
| V Cytochrome oxidase <sup>b</sup>  | 204           | 13 (3–4)                     | Hemes a, a <sub>3</sub> ;<br>Cu <sub>A</sub> , Cu <sub>B</sub> | 0.57                                      | Cytochrome c,<br>O <sub>2</sub> | Cyanide, azide,<br>CO                    |
| Number of subunits in the b<br>Mass and subunit data are<br>Cytochrome <i>c</i> is not part of | for the mone  | omeric form.                 |  | 0.8<br>(Oz<br>lexes III and IV            | 2)                              | otein.                                   |
|  |               |                              |  |   | Solubilization v                | ith detergent<br>exchange chromatography |

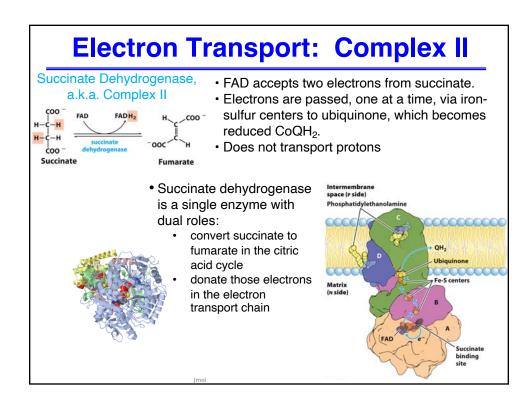




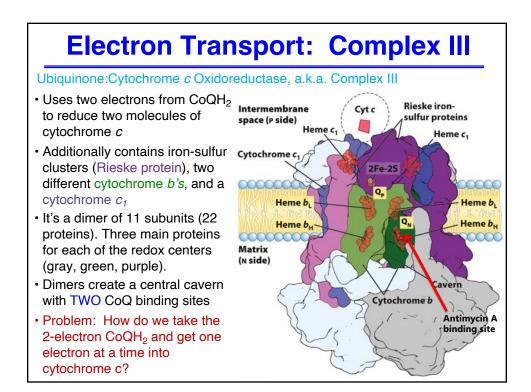


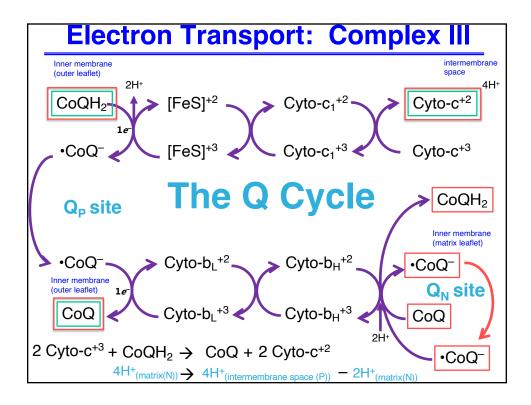


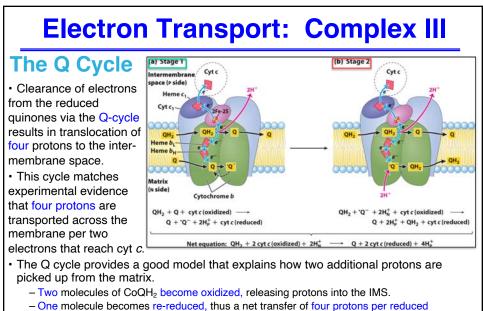
| TABLE The Protein Components of the   19-3 Mitochondrial Respiratory Chain   |               |                        |  |   |                                 |                       |  |
|--|---------------|------------------------|--|---|---------------------------------|-----------------------|--|
| Enzyme<br>complex/protein  | Mass<br>(kDa) | Number of<br>subunitsª | Prosthetic<br>group(s)   | Reduction<br>potential<br>( <i>E</i> 。´V) | Binding sites<br>for:           | Inhibted by:          |  |
| I NADH dehydrogenase   | 850           | 45 (14)                | FMN, Fe-S  | -0.36                                     | NADH, CoQ                       | amytal, rotenone      |  |
| II Succinate<br>dehydrogenase  | 140           | 4                      | FAD-E, Fe-S  | 0.09                                      | Succinate,<br>CoQ               |                       |  |
| III Ubiquinone:<br>cytochrome c<br>oxidoreductase <sup>b</sup>   | 250           | 11                     | Hemes b, c <sub>1</sub> ,<br>Fe-S                              | 0.17                                      | CoQ,<br>Cytochrome c            | antimycin a           |  |
| Cytochrome c <sup>c</sup>  | 13            | 1                      | Heme   |   |                                 |                       |  |
| IV Cytochrome oxidase <sup>b</sup>   | 204           | 13 (3–4)               | Hemes a, a <sub>3</sub> ;<br>Cu <sub>A</sub> , Cu <sub>B</sub> | 0.57                                      | Cytochrome c,<br>O <sub>2</sub> | Cyanide, azide,<br>CO |  |
| <sup>a</sup> Number of subunits in the b <sup>b</sup> Mass and subunit data are fi<br><sup>c</sup> Cytochrome c is not part of a | or the mone   | omeric form.           |  | lexes III and IV                          | as a freely soluble pr          | otein.                |  |

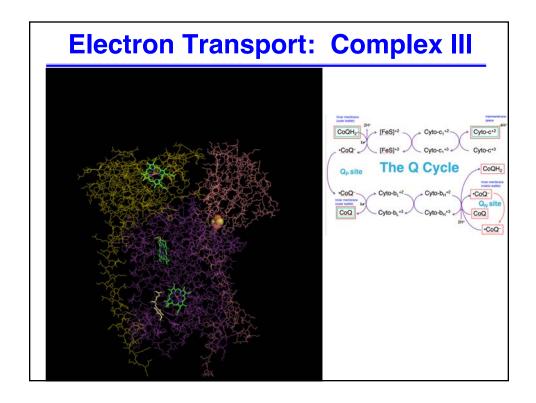


| TABLE The Protein Components of the   19-3 Mitochondrial Respiratory Chain                     |                           |  |  |  |                                 |                       |  |
|--|---------------------------|--|--|--|---------------------------------|-----------------------|--|
| Enzyme<br>complex/protein  | Mass<br>(kDa)             | Number of<br>subunitsª                     | Prosthetic<br>group(s)   | Reduction<br>potential<br>( <i>E</i> ₀´ V) | Binding sites<br>for:           | Inhibted by:          |  |
| NADH dehydrogenase   | 850                       | 45 (14)                                    | FMN, Fe-S  | -0.36                                      | NADH, CoQ                       | amytal, rotenone      |  |
| II Succinate<br>dehydrogenase  | 140                       | 4  | FAD-E, Fe-S  | 0.09                                       | Succinate,<br>CoQ               |                       |  |
| III Ubiquinone:<br>cytochrome <i>c</i><br>oxidoreductase <sup>b</sup>                          | 250                       | 11   | Hemes b, c <sub>1</sub> ,<br>Fe-S                              | 0.17                                       | CoQ,<br>Cytochrome c            | antimycin a           |  |
| Cytochrome c <sup>c</sup>  | 13                        | 1  | Heme   |  |                                 |                       |  |
| V Cytochrome oxidase <sup>b</sup>  | 204                       | 13 (3–4)                                   | Hemes a, a <sub>3</sub> ;<br>Cu <sub>A</sub> , Cu <sub>B</sub> | 0.57                                       | Cytochrome c,<br>O <sub>2</sub> | Cyanide, azide,<br>CO |  |
| Number of subunits in the base and subunit data are ficture c is not part of a Hates e +0.05 \ | for the mono<br>an enzyme | omeric form.<br>complex; it move<br>.077 V |  | ,  | as a freely soluble pr          | otein.                |  |
|  | Cyte                      | chrome b                                   | +0.20 V  |  | +0.22 V                         | +0.254                |  |



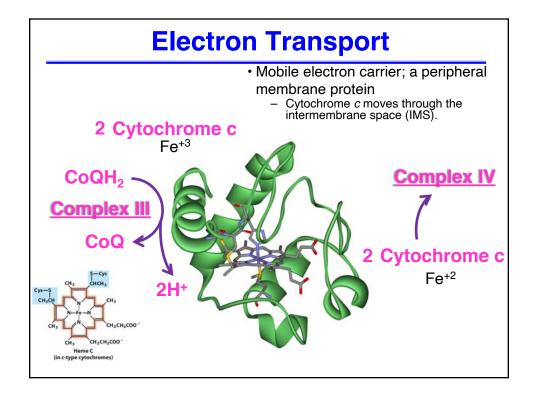




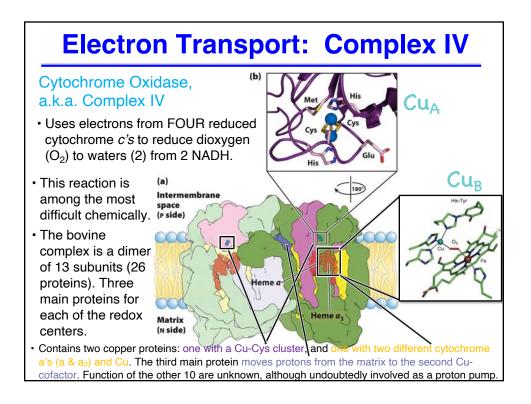


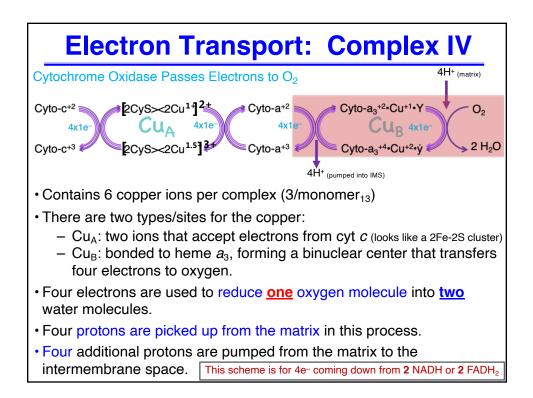
### **Electron Transport**

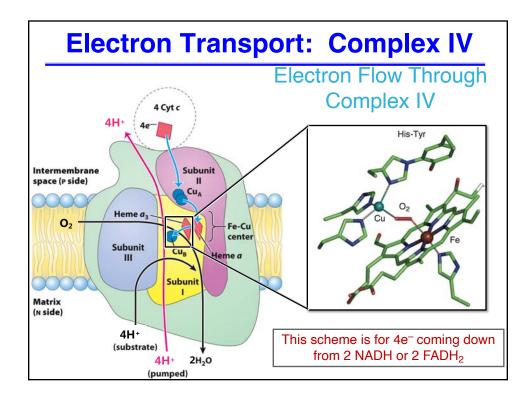
### TABLE The Protein Components of the Mitochondrial Respiratory Chain 19-3 Reduction Enzyme Number of Prosthetic **Binding sites** Mass potential (*E*°´V) Inhibted by: complex/protein (kDa) subunits<sup>a</sup> group(s) for: I NADH dehydrogenase 850 45 (14) FMN, Fe-S -0.36 NADH, CoQ amytal, rotenone II Succinate Succinate, 140 4 FAD-E, Fe-S 0.09 dehydrogenase CoQ III Ubiquinone: Hemes b, $c_1$ , CoQ, 0.17 250 11 antimycin a ${\it cytochrome} \ c$ Fe-S Cytochrome c oxidoreductaseb Cytochrome $c^{c}$ 13 1 Heme Hemes a, a<sub>3</sub>; Cytochrome c, Cyanide, azide, 204 0.57 IV Cytochrome oxidase<sup>b</sup> 13 (3-4) Cu<sub>A</sub>, Cu<sub>B</sub> 02 со <sup>a</sup>Number of subunits in the bacterial complexes in parentheses. <sup>b</sup>Mass and subunit data are for the monomeric form. <sup>c</sup>Cytochrome c is not part of an enzyme complex; it moves between Complexes III and IV as a freely soluble protein.

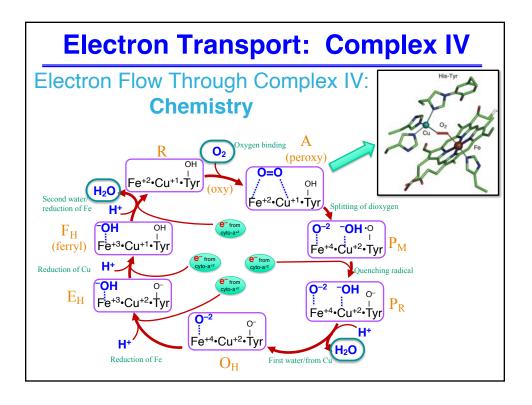


| Enzyme<br>complex/protein   | Mass<br>(kDa) | Number of subunitsª | Prosthetic<br>group(s)   | Reduction<br>potential<br>( <i>E</i> ₀́V) | Binding sites<br>for:           | Inhibted by:          |
|---|---------------|---------------------|--|---|---------------------------------|-----------------------|
| I NADH dehydrogenase  | 850           | 45 (14)             | FMN, Fe-S  | -0.36                                     | NADH, CoQ                       | amytal, rotenone      |
| II Succinate<br>dehydrogenase   | 140           | 4                   | FAD-E, Fe-S  | 0.09                                      | Succinate,<br>CoQ               |                       |
| III Ubiquinone:<br>cytochrome <i>c</i><br>oxidoreductase <sup>b</sup> | 250           | 11                  | Hemes b, c <sub>1</sub> ,<br>Fe-S                              | 0.17                                      | CoQ,<br>Cytochrome c            | antimycin a           |
| Cytochrome c <sup>c</sup>   | 13            | 1                   | Heme   |   |                                 |                       |
| IV Cytochrome oxidase <sup>b</sup>                                    | 204           | 13 (3–4)            | Hemes a, a <sub>3</sub> ;<br>Cu <sub>A</sub> , Cu <sub>B</sub> | 0.57                                      | Cytochrome c,<br>O <sub>2</sub> | Cyanide, azide,<br>CO |





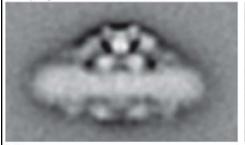


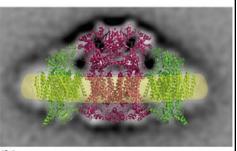


# **Electron Transport**

Multiple Complexes Associate Together to Form a "Respirasome"

Courtesy of Egbert Boekema



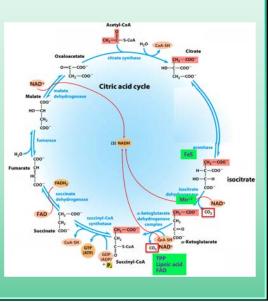


**Complex III and Complex IV** 



### **Rare Fumarase Deficiency**

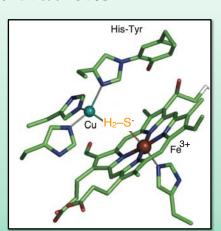
- Loss of activity in TCA enzymes is rare as it is lethal to the cell. However, a few very rare cases have been described.
- One such case was a severe deficiency of fumarase.
- Urine and blood have high levels of fumarate, succinate, α-ketoglutarate, citrate, and malate.
- Humans with this disease have neurological impairment, encephalomyopathy (brain/muscle malady), dystonia (muscle teaching)



# **Clinical Correlations**

### **Cyanide poisoning**

- CN gas or KCN ingestion causes rapid loss of mitochondrial function and death.
- Cyanide works by inhibition of cytochrome oxidase by binding tightly to the Fe<sup>3+</sup> of heme- $a_3$ . Mitochondrial respiration, and energy production ceases, and cell death rapidly follows.
- Cyanide is one of the most potent and rapidly acting poisons known. Other poisons do the same thing: CO,  $H_2S$ ,  $N_3^-$
- If detected early enough, the antidote is to offer the CN more Fe<sup>3+</sup> sites to bind and titrate it off of the heme-*a*<sub>3</sub>.



 Creation of "metHb" by oxidation of Hb using various nitrates (Fe<sup>2+</sup> → Fe<sup>3+</sup>) can work due to vast amounts of Hb.