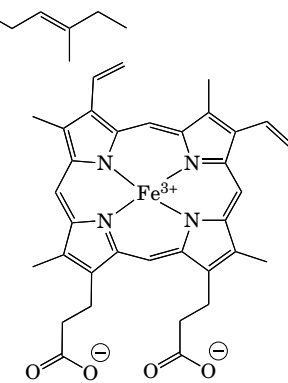
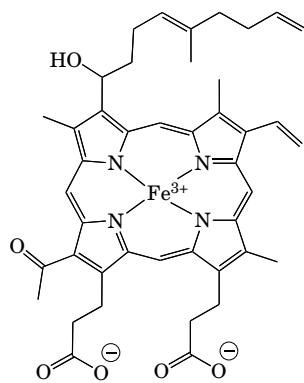
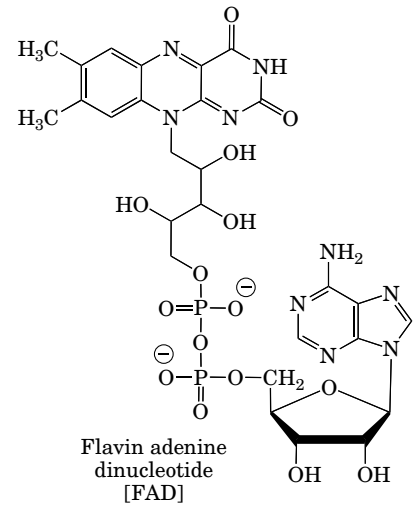
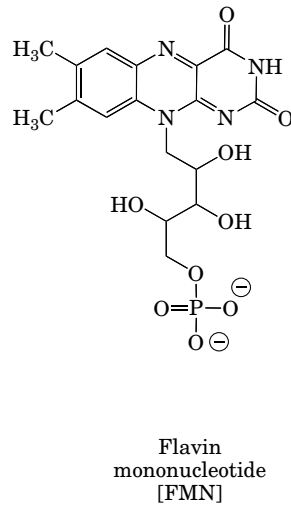
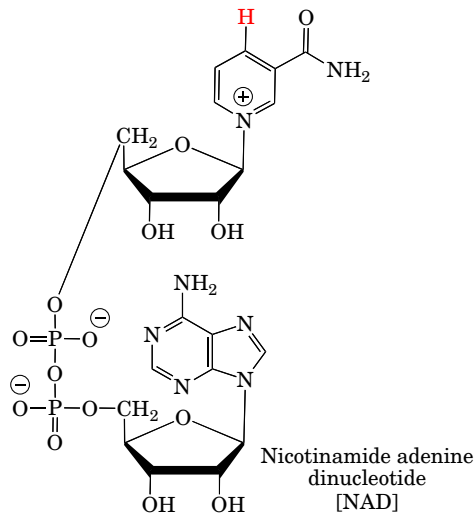
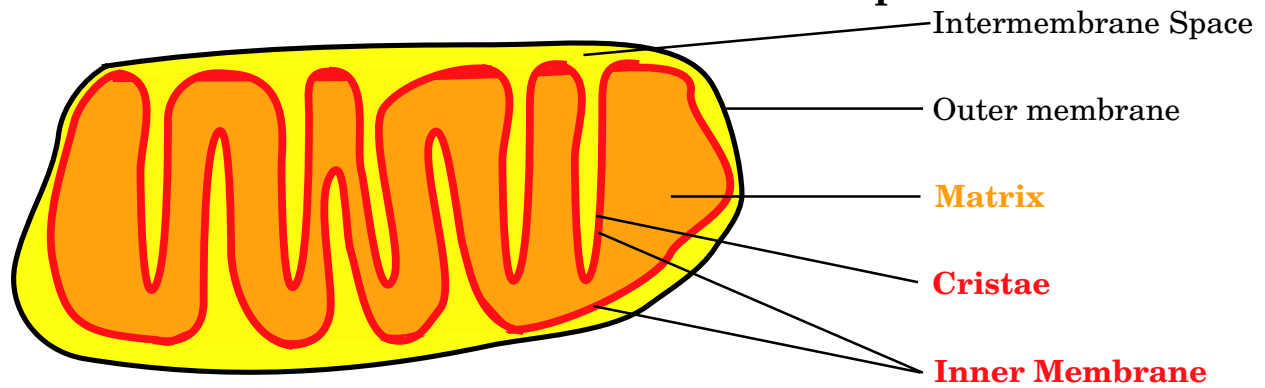
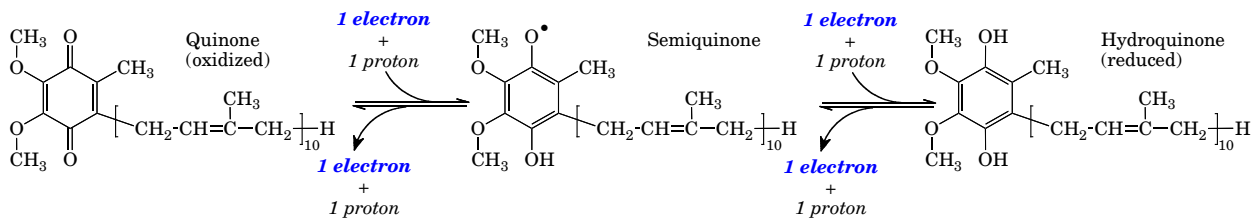
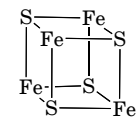
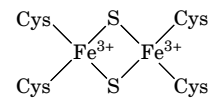
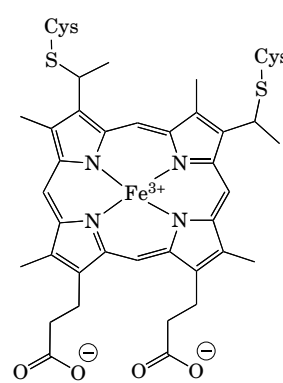
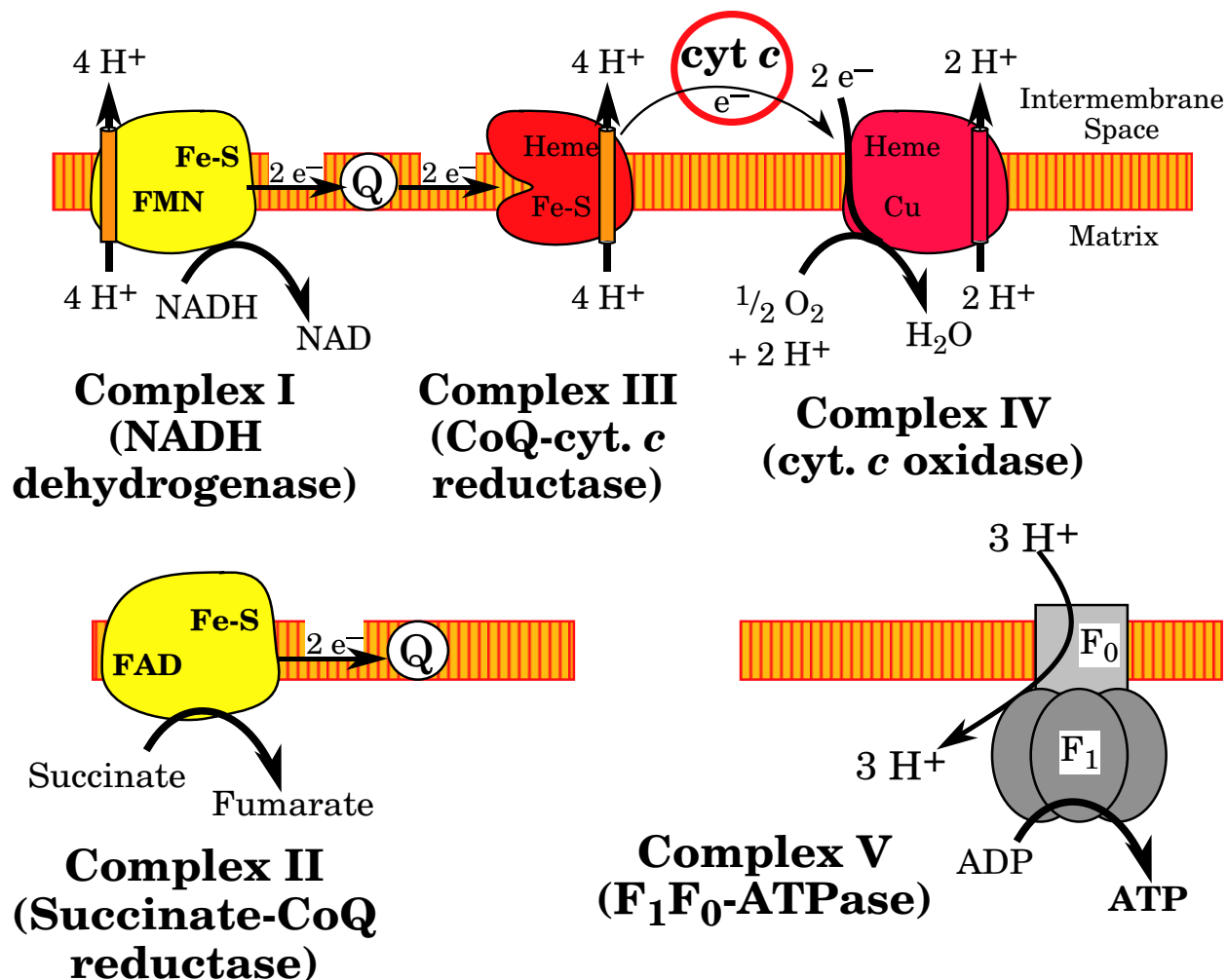


Mitochondrial Structure and Electron Transport



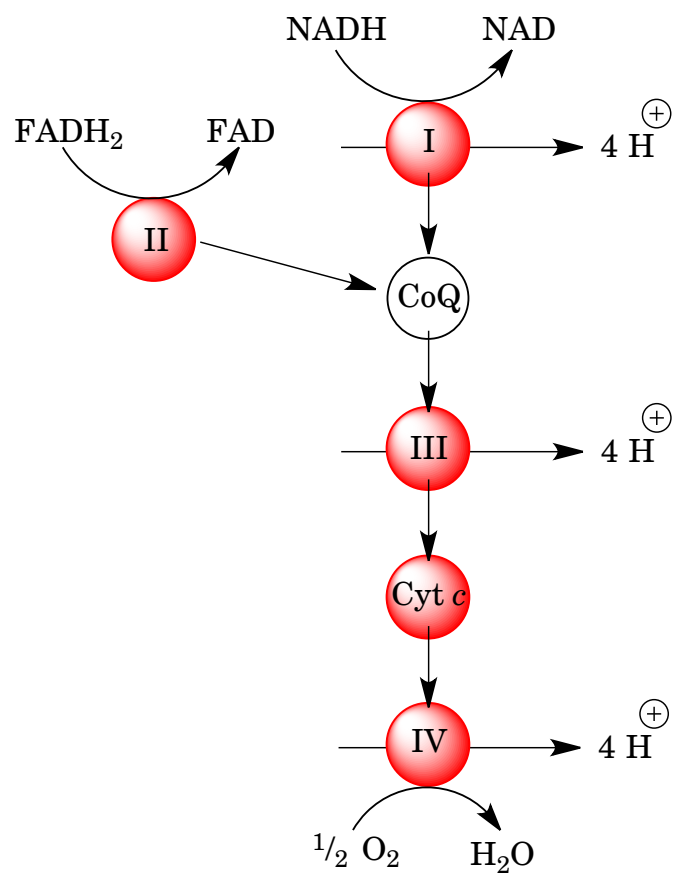
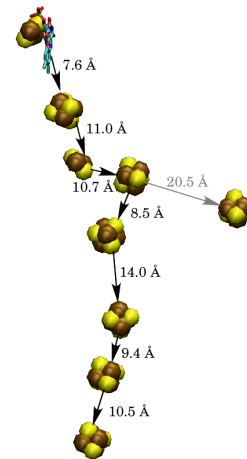
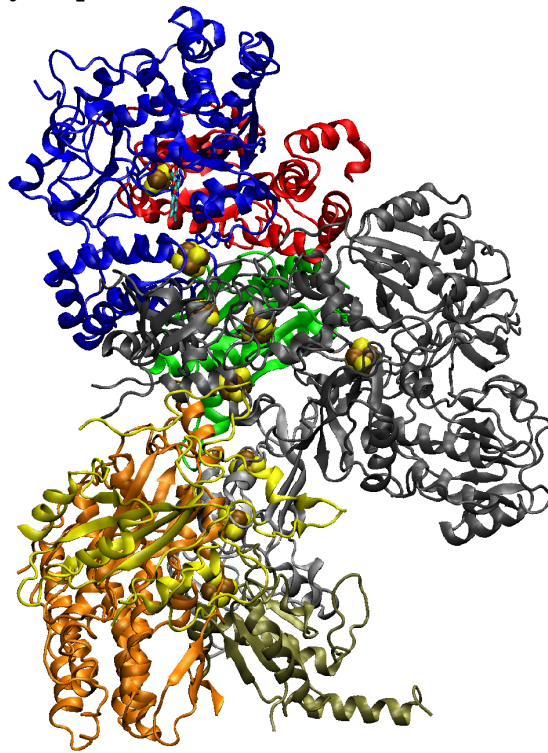
Iron-Protoporphyrin IX



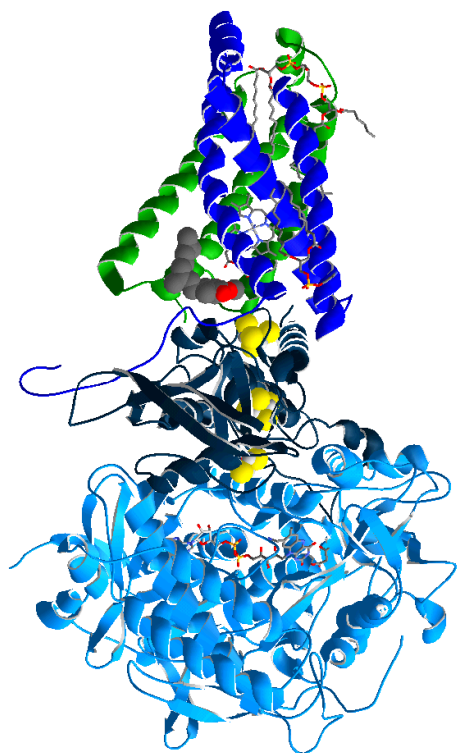


Compound	Reduction potential (V)
NADH	-0.315
Complex I Fe-S	-0.38 to -0.27
Succinate	+0.031
Complex II FAD	-0.04
Complex II Fe-S	-0.245 to -0.03
Complex II heme	-0.08
Coenzyme Q	+0.045
Complex III heme <i>b</i>	-0.03 to +0.03
Complex III Fe-S	+0.28
Complex III heme <i>c</i> ₁	+0.215
Cytochrome <i>c</i>	+0.235
Complex IV heme	+0.21 to +0.385
Complex IV Cu	+0.245 to +0.34
Oxygen	+0.815

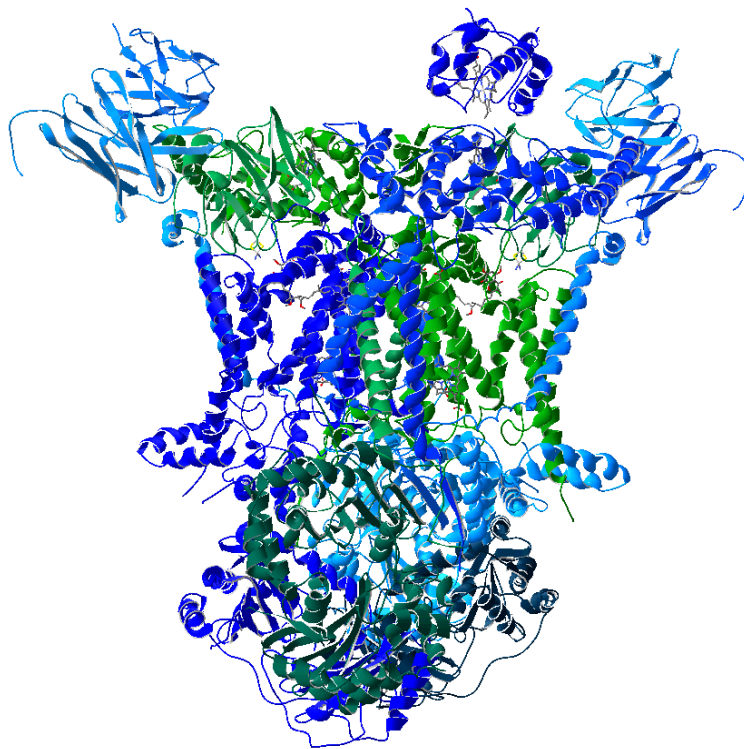
Complex I Hydrophilic Domain from *Thermus thermophilus* (pdb ID 2FUG)



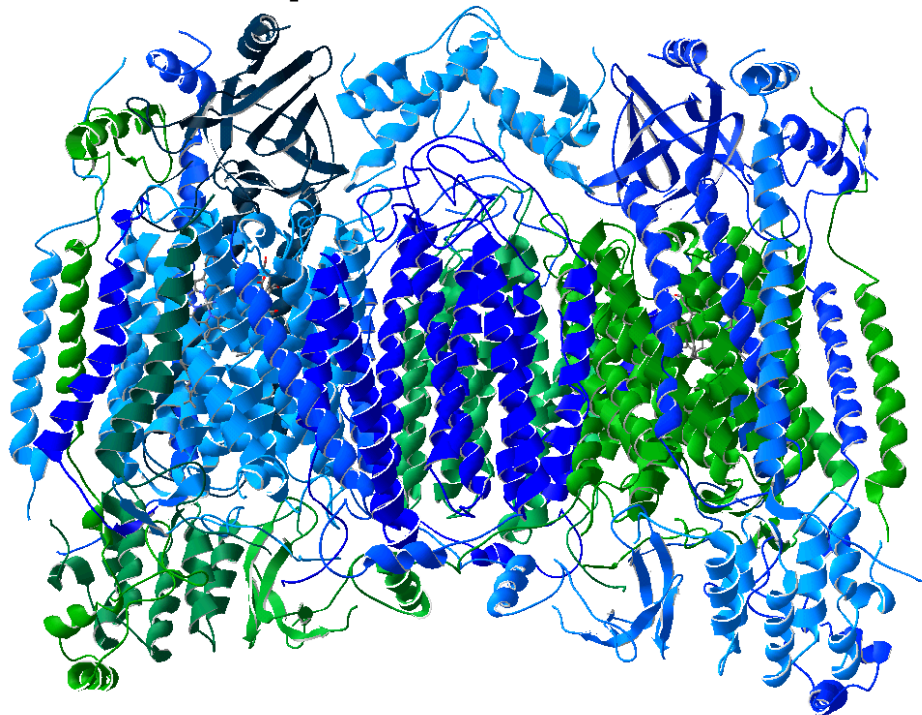
Complex II from *Escherichia coli* (pdb ID 1NEK)



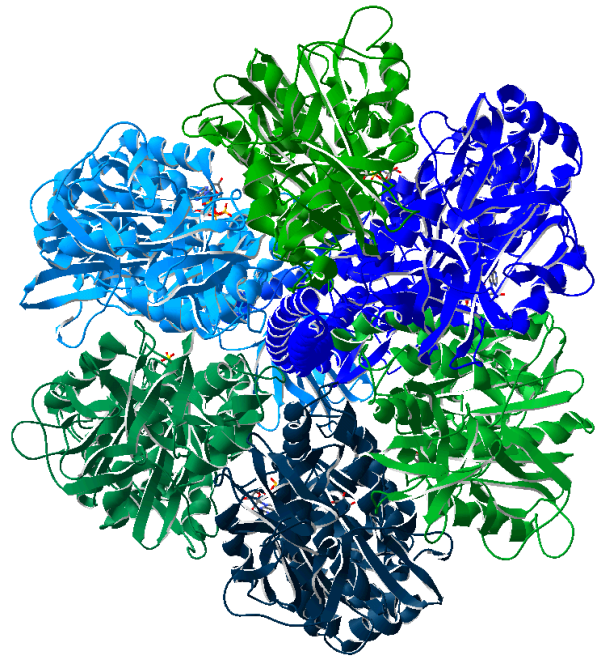
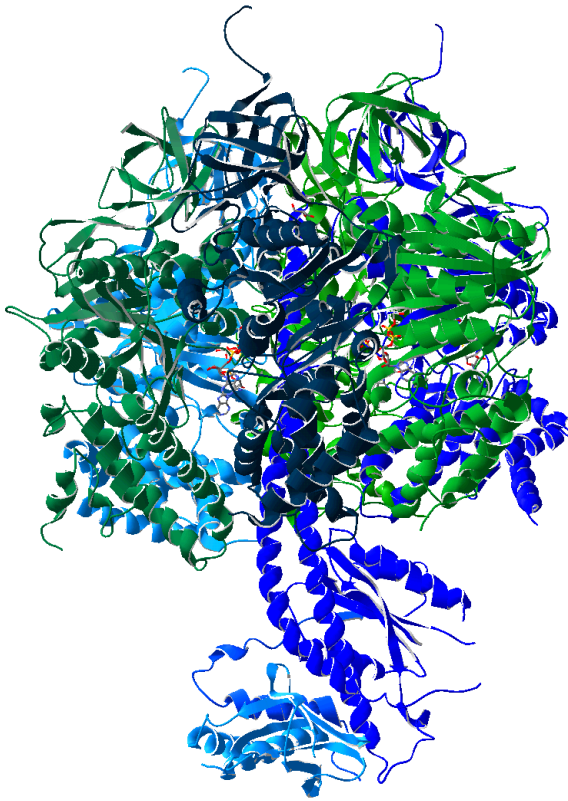
Complex III and cytochrome *c* from *Saccharomyces cerevisiae* (pdb ID 1KYO)



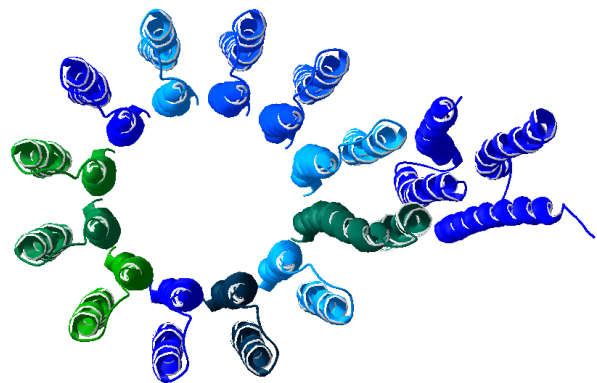
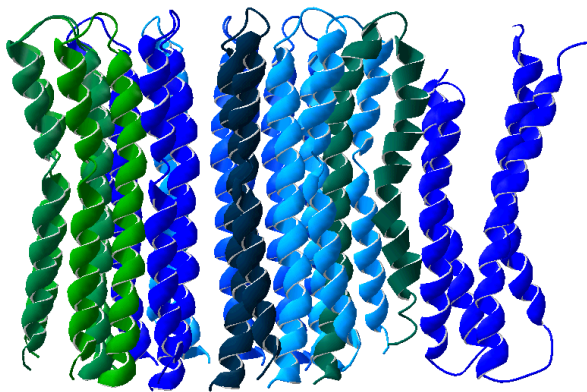
Complex IV from *Bos taurus* (pdb ID 2OCC)



F_0 - F_1 ATPase (ATP Synthase)

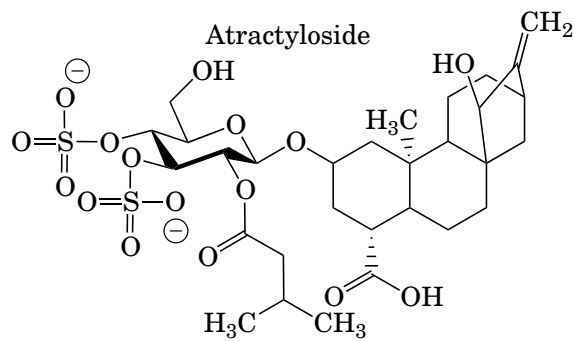
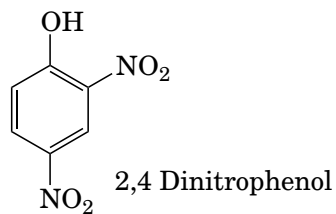
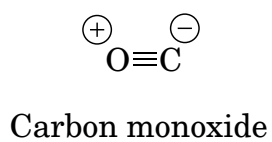
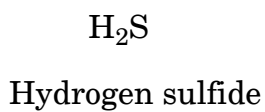
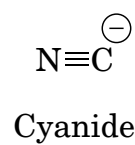
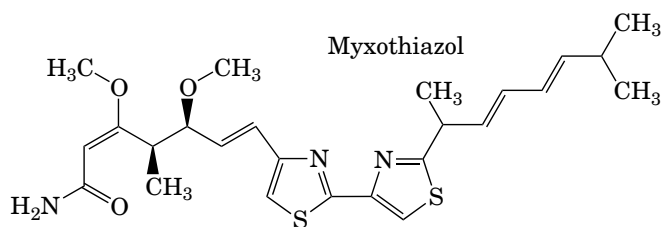
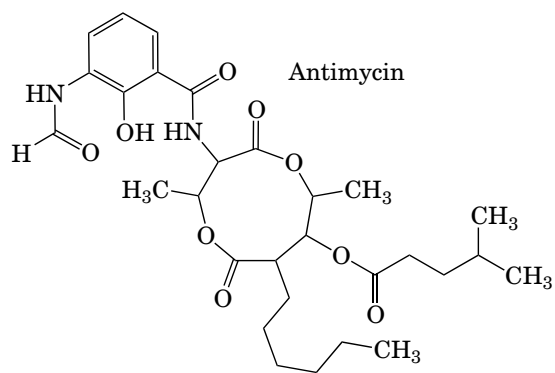
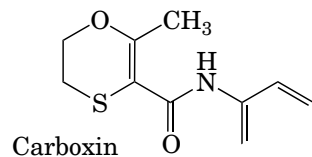
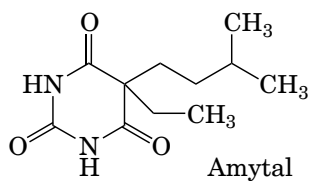
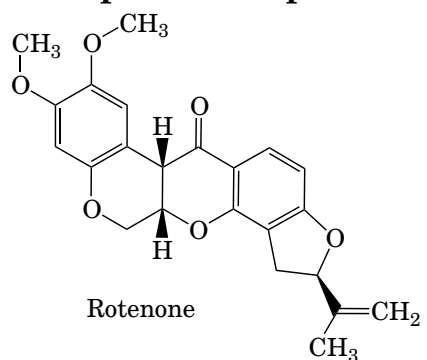


pdb ID 1E79



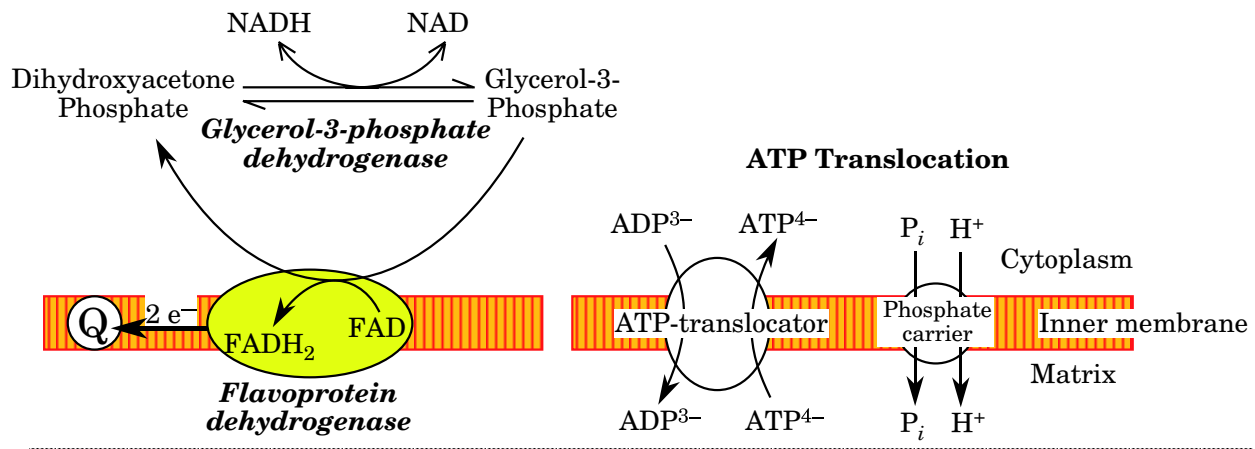
pdb ID 1C17

Examples of compounds that affect mitochondrial ATP synthesis

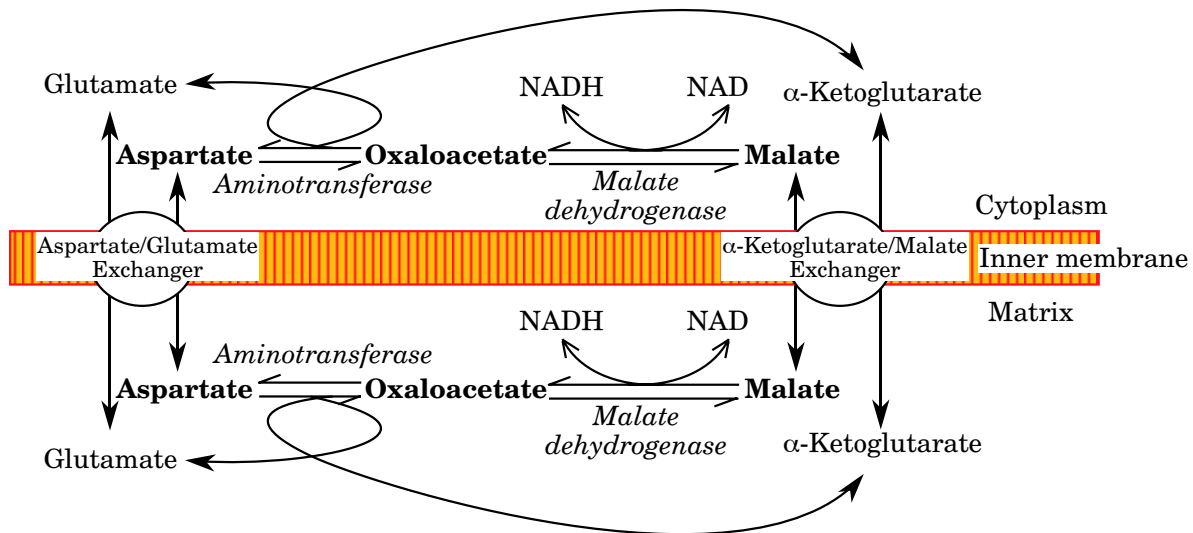


Examples of Mitochondrial Shuttle Pathways

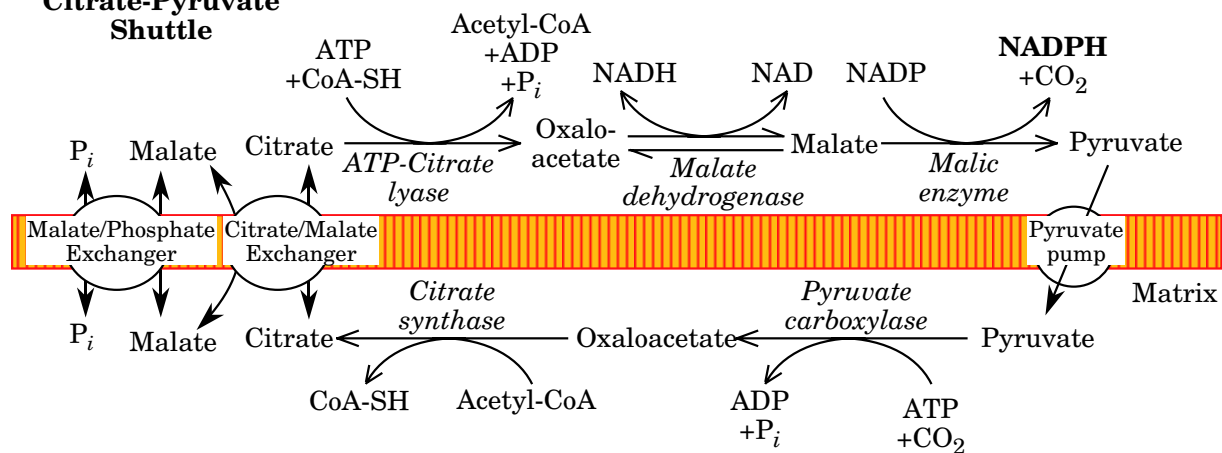
Glycerophosphate Shuttle



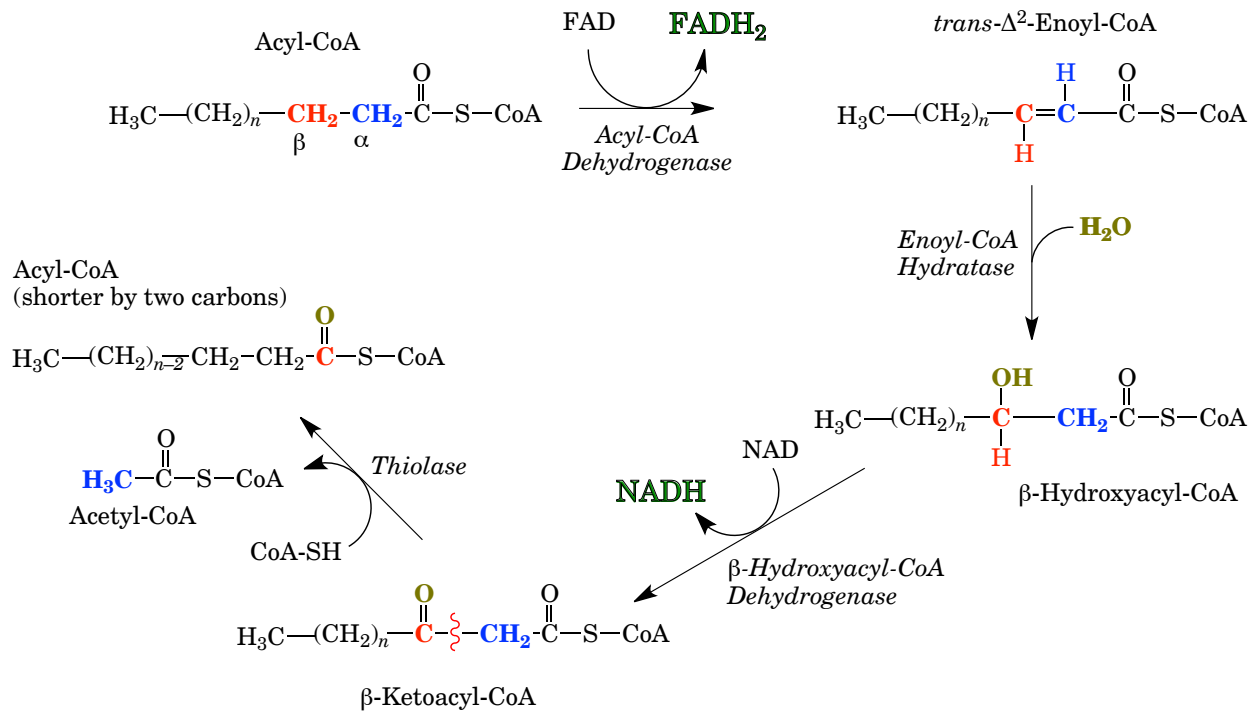
Malate-Aspartate Shuttle



Citrate-Pyruvate Shuttle



Fatty Acid β -oxidation



Comparison of Energetics of Metabolism for Glucose and Stearic Acid

Energetic molecule	Glucose	Stearate ↓ Acetyl-CoA	9 Acetyl-CoA ↓ CO ₂	Stearate (total)
Products				
ATP	4 → 4 ATP	-2	9	7 → 7 ATP
NADH	10 → 30 ATP	8	27	35 → 105 ATP
FADH ₂	2 → 4 ATP	8	9	17 → 34 ATP
Total	38 ATP			146 ATP