

Subunit-1.3.

Mitochondrial Electron Transport chain
and Oxidative Phosphorylation
(Part-II)

By

Dr. Sumita Kumari Shaha

A.P. & Head

P.G. Dept. of Zoology

Maharaja College

Asst. Prof. Complexes

Characteristics of Respiratory chain
and Transfer of Electrons :-

Characteristics of proteins subunits constituting different respiratory chain complexes differ remarkably from each other. The interactions of red chains of proteins with cofactors, hemes and Fe-S centers determine redox potential of redox centers or complexes.

Complex-I :- The Complex I of respiratory chain is also named NADH

dehydrogenase or NADH : ubiquinone Oxidoreductase. It is a metalloprotein, formed of 41 polypeptide chains, one

flavoprotein with flavin mononucleotide (FMN) as coenzyme and six iron-sulfur centers, all linked covalently to

other proteins. This complex is L-shaped with larger axis placed

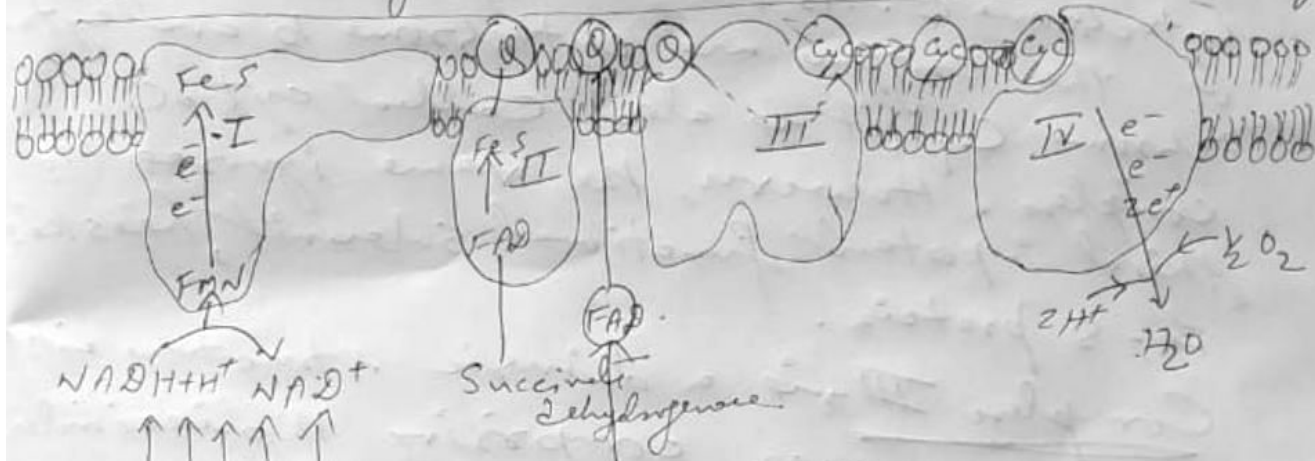
along the length of IM and smaller

atom projected in the matrix. The Complex I receives electrons pair as hydride ion from reduced NAD and a proton from matrix. The FMN receives hydrogen atoms or electrons and proton in single forming semiquinone (FMN^{•-}) and dihydro form (FMNH₂). From FMNH₂ electrons move in single to Fe-S centers and protons to matrix. Flow of electrons from Fe-S center to Coenzyme Q and proton from matrix bring about reduction of coenzyme Q to CoQH₂. This flow of electrons and protons from reduced NAD to CoQ generates energy (exergonic process), which pumps four protons (for every pair of electrons) to the intermembrane space from matrix. Exact process of this protein pumping is speculative.

Complex II - The Complex II of res. chain is basically iron-bound enzyme succinate dehydrogenase of Krebs's cycle. The enzyme comprises four subunits. Two of these subunits C & D are transmembrane in nature and mainly helical, whereas subunits A and B are globular, protruding into matrix. The subunit A possesses binding sites for succinate and coenzyme FAD whereas subunit B possesses three Fe-S centers. The two transmembrane subunits (C & D) possess a heme named heme

b sandwiched between two subunits, A binding site for CO₂ is present mainly in subunit C.

Coenzyme Q (Ubiquinone) - also called Ubiquinone, is a small hydrophobic component of respiratory chain. Being small and hydrophobic, it is easily



1. Pyruvate dehydrogenase
2. α-Ketoglutarate dehydrogenase
3. Malate dehydrogenase
4. Isocitrate dehydrogenase
5. Glyceraldehyde-3-P dehydrogenase

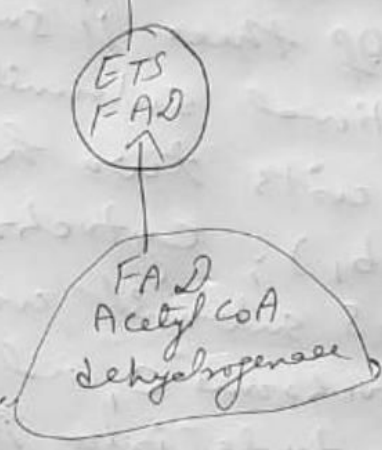


Diagram showing four ETC Complexes (I, II, III and IV) of respiratory chain with two mobile components (cyt c) and cytochrome c (c) along with sources of reduced NAD and reduced FAD feeding the respiratory chain with electron pairs.

- Diffusible into lipid bilayer of IM. It acts as a mobile electron carrier between immobilized components of res. chain. Complexes I and III as well as Complexes II and IV.

Complex III : The Complex III (cytochrome b_6 complex) also called ubiquinone, cytochrome c oxidoreductase is so called because it acts as an enzyme in the transfer of electrons from Coenzyme Q to cytochrome c. As the main functional subunit of the enzyme are cytochrome b and cytochrome c_1 , it is also called cyt b_6 complex. However, the name ignores the third important functional subunit, Rieske's Fe-S protein.

Complex III is a dimeric structure with each monomer comprising 11 dissimilar structures. Of these 11 subunits, 3 are main functional subunits. They are cytochrome b with two hemes (heme b_2 and heme b_1); cytochrome c_1 and Rieske's Fe-S subunit with 2 Fe-2S center. Part of cytochrome c_1 and Rieske's Fe-S subunit protrude into intermembrane space and thus provide ideal site for interaction of cytochrome c. The cytochrome c is not a component of complex III but is a mobile and soluble component present in interaction with outer surface of IM. It acts as an electron transferring shuttle between complex III and IV. Two molecules of CO, participate in single is the transfer of electrons through heme centers present in all the three subunits. The electrons ($2e^-$) from each CO & 1/2

(Pg 5)

are transferred in single through two different electron transfer circuits. Loss of electrons from O_2H_2 is coupled to loss of protons to intermembrane space. One electron transfer circuit comprises O_2H_2 forming a reduced cyt c. The other electron transfer circuit comprises O_2H_2 with four reduced electrons. Two electrons go to cyt c forming two reduced cyt c and two other electrons go to CO_2 picking up two protons from matrix to generate O_2H_2 . The second CO_2 is left in oxidised form (CO_2).

Cytochrome c :- Like other cytochromes,

cytochrome c is also a hemoprotein, but unlike them is a soluble protein and is present not as integral protein of IM but is associated with outer surface of IM by electrostatic interactions. This makes it a mobile component of respiratory chain, hence it acts as a shuttle carrying electrons between Complex III and IV. Quite unlike other cytochromes where heme is non-covalently bound to protein, heme is covalently bound to protein in cyt c through S of cysteinyl residues. Cytochrome c receives electron singly from cyt c, of complex III and shuttles to complex IV. Donating electron in single to binuclear copper center of complex IV.

- Control part IV