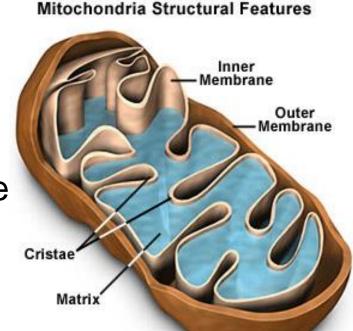


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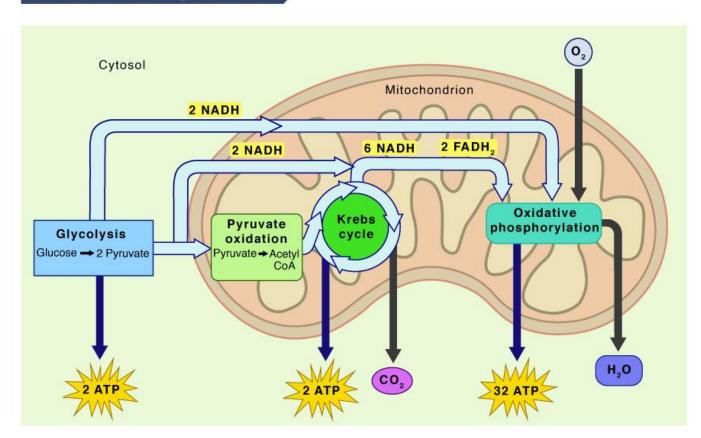
- Citric acid, Tricarboxylic acid cycle (TCA) or Krebs cycle is a central pathway used by all aerobic organisms to generate energy through the oxidation of acetate (in the form of acetyl CoA) into CO₂ and ATP. Also it releases the energy-rich molecules: NADH and FADH2
- It occurs in mitochondrial matrix except reaction 6 in which succinate dehydrogenase enzyme is found in inner mitochondrial membrane (it is the only transmembrane protein in Krebs cycle)



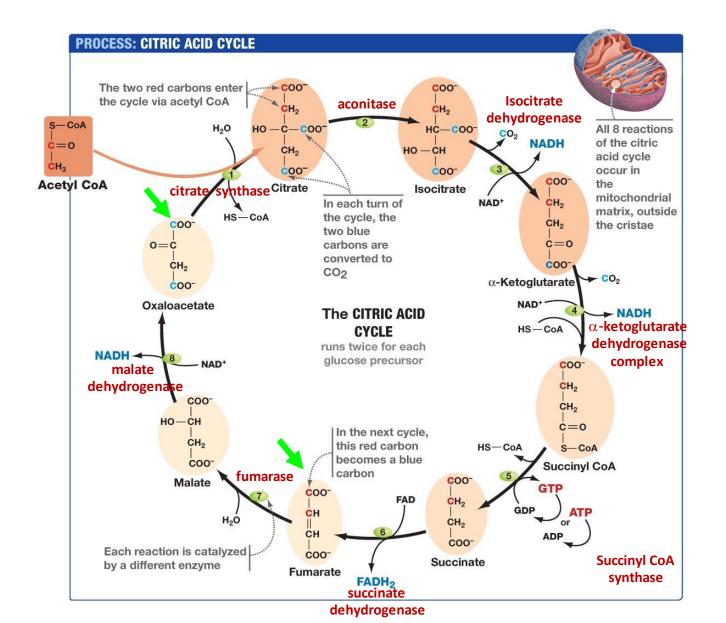
- حونه
- To proceed in Krebs cycle we need the presence of:
 - 1. Mitochondria
 - 2. 02

Aerobic Respiration

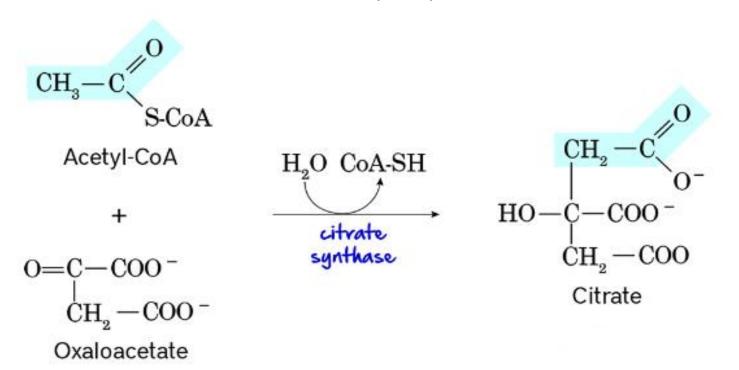








- Krebs cycle is a series of 8 reactions run twice / glucose molecule:
- Step 1: The irreversible condensation of acetyl CoA (2C) and oxaloacetate (4C) via citrate synthase to form citrate (6C)

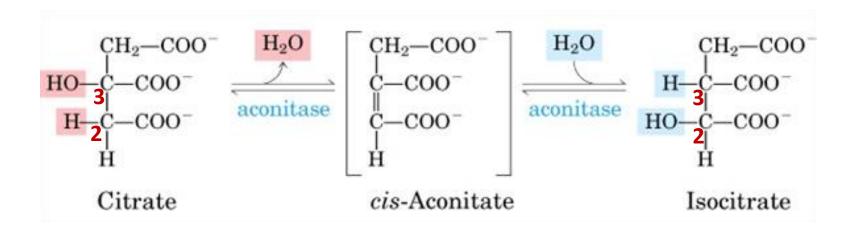


 Oxaloacetate is already found in matrix. It can be produced in several ways in nature. For example, it is generated from an ATP-dependent carboxylation of pyruvate catalyzed by pyruvate carboxylase. This reaction occurs in the matrix

 Step 2: Aconitase enzyme catalyzes the reversible isomerization of citrate to isocitrate (isomers differ in the position of OH group from C3 to C2)

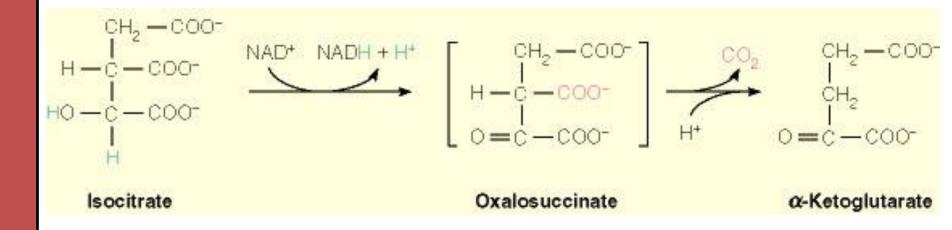


 This isomerization reaction is pre-required step to prepare substrates for decarboxylation reaction

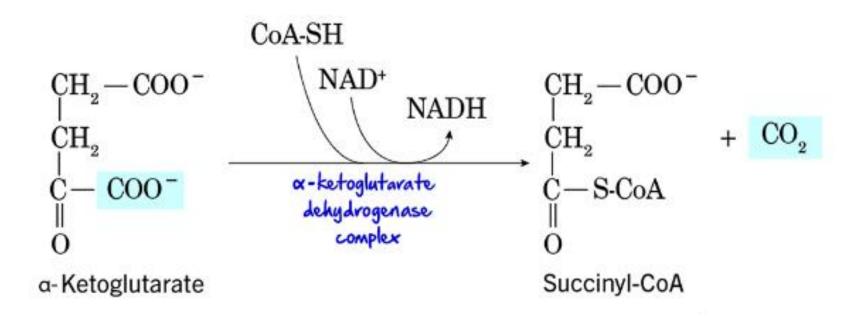


It involves successive dehydration and hydration reactions

- Step 3: Isocitrate dehydrogenase catalyzes the first oxidative decarboxylation of isocitrate (6C) to α -ketoglutarate (5C) resulting in the release of first CO₂ and the formation of first NADH molecule
- It involves successive oxidation and decarboxylation reactions

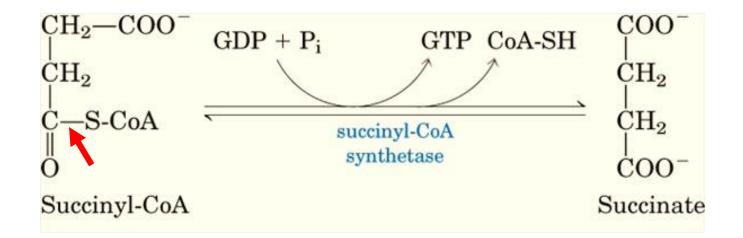


Step 4: α-ketoglutarate dehydrogenase complex catalyzes the oxidative decarboxylation of α-ketoglutarate (5C) to succinyl CoA (4C) releasing the second CO₂ and producing the second NADH molecule



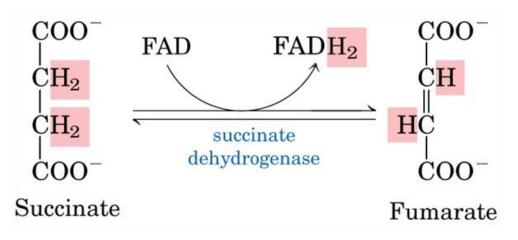


 Step 5: Succinyl CoA synthetase (also known as succinate thiokinase) generates the first ATP (e.g. brain & heart tissues) or GTP (e.g. liver tissues). The thioester bond of succinyl-CoA is energy-rich and can drive the phosphorylation of ADP or GDP (substrate-level phosphorylation)





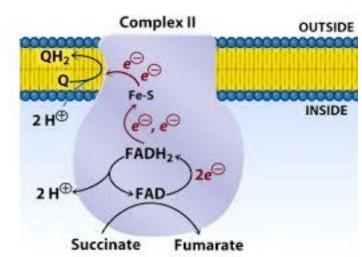
 Step 6: Succinate dehydrogenase catalyzes the oxidation of succinate to fumarate and consequently, the reduction of prosthetic group FAD into FADH₂

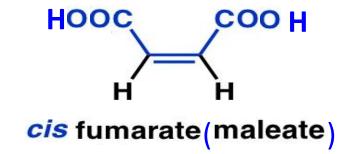


$$H_3C$$
 H_3C
 H_3C



- Succinate dehydrogenase is the only enzyme found in the inner membrane of mitochondria
- FAD is more powerful oxidizing agent than NAD+



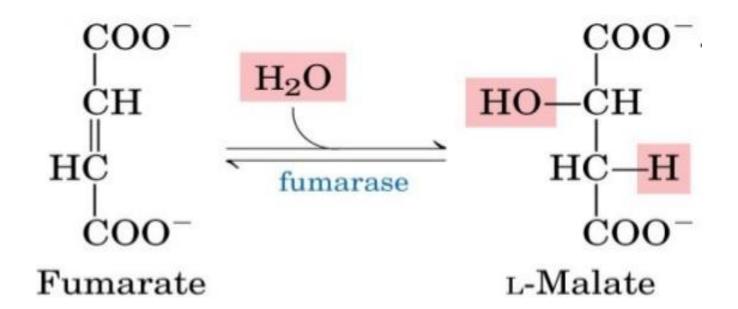


COOH

trans fumarate



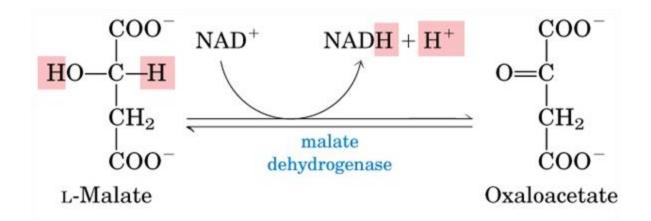
 Step 7: Fumarate is converted to L-malate in a hydration reaction catalyzed by fumarase (reversible reaction)



Fumarase is another stereospecific enzyme



 Step 8: L-malate is oxidized to regenerate oxaloacetate via malate dehydrogenase enzyme thus generating the third NADH (reversible)



 At the end of krebs cycle, the products of oxidation of one glucose via glycolysis and TCA are:

ATP Yield per one Glucose



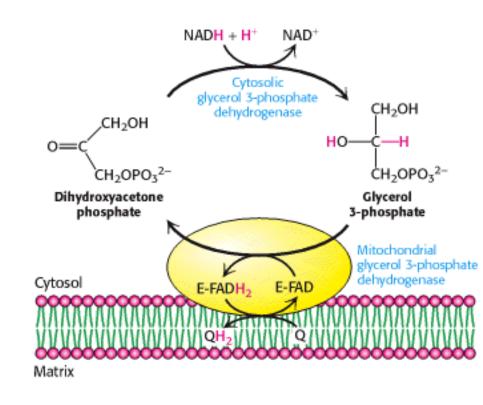
Stage	ATP produced by substrate-level phosphorylation	
Glycolysis	2 ATP	
Acetyl CoA production	none	
Krebs Cycle	2 ATP	
Total/glucose 4 ATP molecules		

Stage	Electron-carrier	Total H ⁺	ATP synthase
	molecule	pumped	4H⁺── 1 ATP
Glycolysis	2 NADH	12-20	3-5 ATP
Acetyl CoA production	2 NADH	20	5 ATP
Krebs Cycle	6 NADH	60	15 ATP
	2 FADH ₂	12	3 ATP

Total/glucose 26-28 ATP produced by oxidative phosphorylation

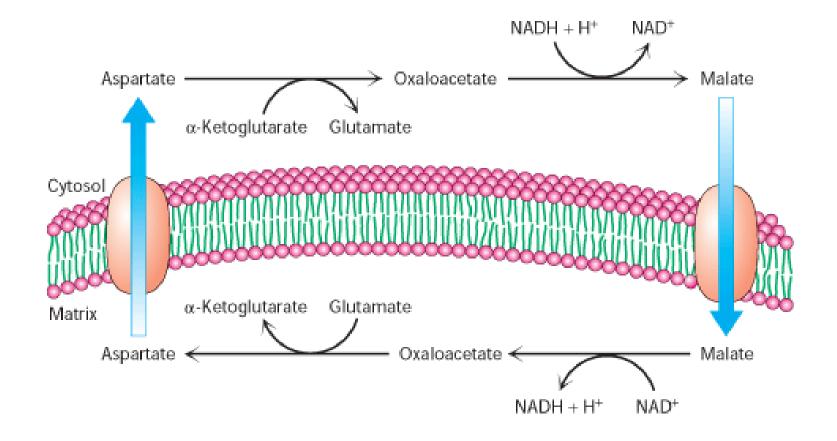
Cytosolic NADH Shuttling

- The electrons carried by cytosolic NADH (i.e. NADH generated by glycolysis) will be shuttled to the matrix by one of two mechanisms:
- 1. DHAP/G3P shuttle: it is active in brain and skeletal muscle. This pathway delivers the 2e from cytosolic NADH to mitochondrial FAD



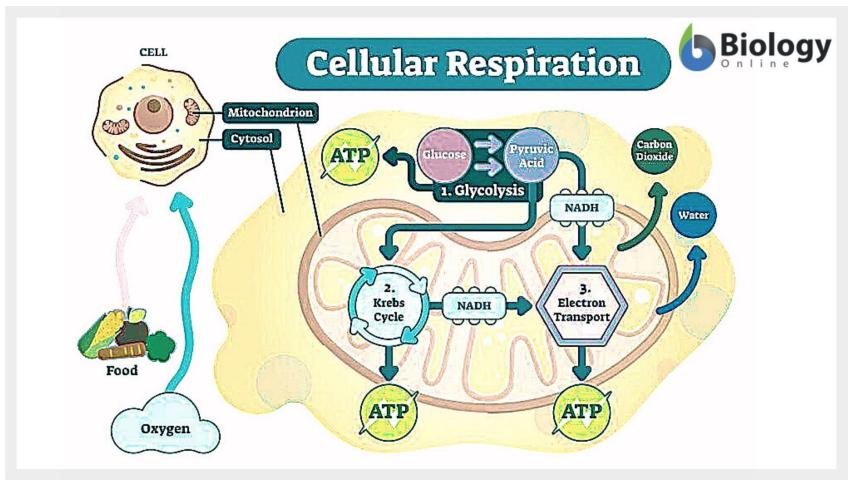
NADH Shuttling

 Aspartate/malate shuttle: it is active in liver and heart. This pathway delivers the 2e from cytosolic NADH to mitochondrial NAD+ (found in the matrix)



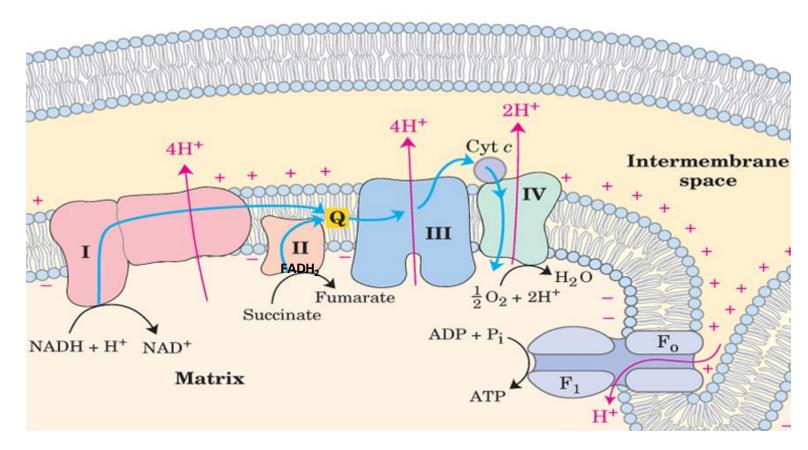
Aerobic Respiration





Oxidative Phosphorylation

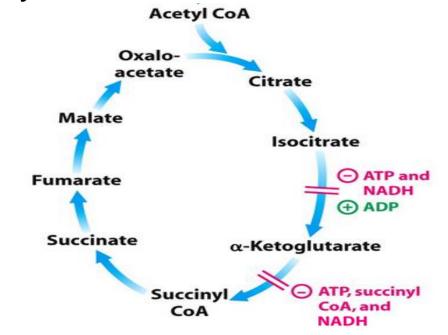




 TCA is considered as a part of aerobic metabolism although it does not use O₂ in any of its reaction ??

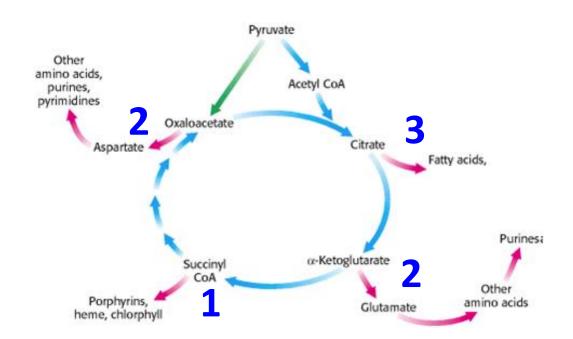
TCA Cycle Regulation

- Reactions 3 and 4 are key sites for allosteric regulation:
- Isocitrate dehydrogenase is activated by ADP and directly inhibited by NADH and ATP
- α-ketoglutarate dehydrogenase activity is inhibited by NADH, ATP and succinyl-CoA



Biosynthetic Role of TCA Intermediates

 In addition to its role in catabolism and energy generation, TCA intermediates have anabolic role in biosynthesis of other molecules:



Biosynthetic Role of TCA Intermediates



- Succinyl CoA is used in synthesis of heme and other porphyrins
- Oxaloacetate and α-ketoglutarate are converted by transamination to the corresponding amino acids aspartate and glutamate, respectively
- Citrate in some tissues is transported to cytosol where it is converted back to acetyl CoA for fatty acids biosynthesis

Anaplerotic Pathway

 Anaplerotic pathways (from Greek word meaning filling up) are the processes that replenish TCA intermediates so that the flow of carbon out of the cycle is balanced by these reactions

- Reactions that replenish oxaloacetate:
- Pyruvate carboxylase (PC) catalyzes the irreversible ATP-dependent carboxylation of pyruvate to generate oxaloacetate (occurs in the matrix)

Anaplerotic Pathway

Transamination replenish TCA intermediates:

Transamination are reversible reaction in which an amino acid loses an amino group thereby converted itself to a keto acid

$$R_1$$
— C — $COO^- + R_2$ — C — $COO^- + R_2$ — C — $COO^- + R_2$ — C — COO^-

 For example glutamate and aspartate undergo transamination to generate the TCA intermediates α-ketoglutarate and oxaloacetate, respectively

