



# **METABOLISM**

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## ❖ TCA cycle

- It is called **Krebs cycle**, **Citric acid cycle** and **Tricarboxylic acid cycle (TCA)**
  - It represents the oxidation of acetyl group of acetyl CoA to  $\text{CO}_2$
  - It involves the release of energy in the form of GTP and stored in NADH and  $\text{FADH}_2$
  - No  $\text{O}_2$  is required (*anaerobic* pathway)
- All of the major nutrients can be converted to acetyl CoA in the second stage of metabolism
- TCA cycle occurs in the *mitochondrial matrix* except succinate dehydrogenase (**step 6**) occurs in the *inner mitochondrial membrane*
- Oxaloacetate (4C) is the final intermediate of the TCA and it has no net synthesis or degradation
- **Each turn** of the TCA cycle involves:
  - Release of **2  $\text{CO}_2$** 
    - ✓ The source of carbons released is Acetyl CoA
  - Transfer of 3 pairs of electrons in the form of hydride ions to  $\text{NAD}^+$  to form **3 NADH**
  - Transfer of 1 electron in the form of hydrogen atom to reduce FAD to  **$\text{FADH}_2$**
  - Substrate level phosphorylation which results in the formation of **GTP** from GDP and  $\text{P}_i$

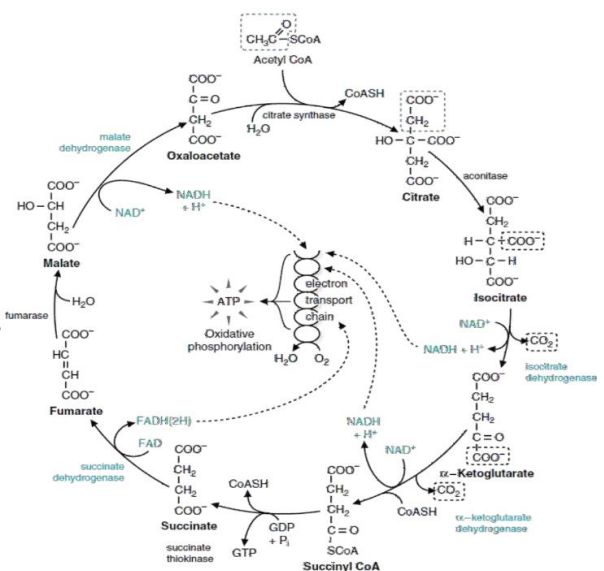
### ◆ Steps and Enzymes of TCA cycle

#### 1) Citrate synthase

- It produces **citrate** by the condensation of acetyl with oxaloacetate (OAA)
- This step is inhibited by **citrate**, **ATP** and **succinyl CoA** and activated by **substrates**
- Citrate activates acetyl CoA carboxylase (Fatty acid synthesis), and also citrate represents a source of Acetyl CoA which is used in the synthesis of FAs
- Citrate inhibits phosphofructokinase (glycolysis)
- Citrate Stimulates fructose-1,6-bisphosphatase (gluconeogenesis)
- $\Delta G^\circ = -9 \text{ kcal/mol}$

#### 2) Aconitase

- It isomerizes citrate (3° alcohol) into **isocitrate** (2° alcohol) by forming an intermediate called aconitate
- Dehydration followed by hydration
- **Fluoroacetate** (rat poison, a non-competitive inhibitor), **succinyl CoA** and **NADH** inhibits aconitase
- Activated by **ADP** and  **$\text{Ca}^{+2}$**
- $\Delta G^\circ = +1.5 \text{ kcal/mol}$



### 3) Isocitrate dehydrogenase complex

- It undergoes oxidative decarboxylation of isocitrate into  ***$\alpha$ -ketoglutarate***
- Release  ***$CO_2$***  and reduces  $NAD^+$  into  ***$NADH + H^+$***
- It is considered as the **rate limiting step**
- Activated by ***ADP*** and  $Ca^{+2}$ 
  - ✓ ADP lowers  $K_m$  10 folds which increases the affinity of the enzyme to its substrate
- Inhibited by ***ATP*** and ***NADH***
- $\Delta G^\circ = -5 \text{ kcal/mol}$

### 4) $\alpha$ -Ketoglutarate dehydrogenase complex

- It undergoes oxidative decarboxylation of  $\alpha$ -Ketoglutarate into ***Succinyl CoA***
- Release  ***$CO_2$***  and reduces  $NAD^+$  into  ***$NADH + H^+$***
- This complex is a multimolecular aggregate of 3 enzymes
  - ✓ E1 (decarboxylase) requires thiamine pyrophosphate (***TPP***) as a cofactor
  - ✓ E2 (dihydrolipoyl transacylase) requires ***lipoic*** acid and ***CoA*** as cofactors
  - ✓ E3 (dihydrolipoyl Dehydrogenase) requires ***FAD*** and  $NAD^+$  as cofactors
- This complex is one of  $\alpha$ -Ketoacid dehydrogenase complexes which act in  $\alpha$ -Ketoglutarate, pyruvate and branched chain  $\alpha$ -Ketoacids
- Inhibited by ***ATP***, ***GTP***, ***NADH***, and ***succinyl CoA***, and activated by ***AMP*** and  $Ca^{+2}$
- Inhibited by ***Arsenite*** (non-competitive inhibitor)
- $\Delta G^\circ = -8 \text{ kcal/mol}$

TPP is a **Vitamin B1** derivative

Lipoic acid and FAD are covalently attached to E2 and E3

### 5) Succinyl thiokinase

- Succinyl CoA is converted into ***Succinate***, by the cleavage of ***thioester bond*** (high energy bond)
- It is coupled to phosphorylation of GDP to ***GTP*** (substrate level phosphorylation)
  - ✓ GTP and ATP are energetically interconvertible by the nucleoside diphosphate kinase reaction
- $\Delta G^\circ = -8 \text{ kcal/mol}$

### 6) Succinate dehydrogenase

- Oxidation of succinate into ***fumarate***
- The only dehydrogenation in TCA cycle that is not NAD-linked, but FAD to form  ***$FADH_2$*** 
  - ✓ Uses FAD instead of  $NAD^+$ , because succinate reducing power is not sufficient to reduce  $NAD^+$
- Placed in the **inner mitochondrial membrane** not the matrix
- Succinate dehydrogenase functions as Complex II of the electron transport chain
- ***Malonate*** is a competitive inhibitor
- $\Delta G^\circ = 0 \text{ kcal/mol}$

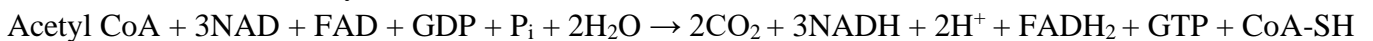
7) Fumarase (fumarate hydratase)

- **Reversible** hydration of fumarate to **L-malate**
- $\Delta G^\circ = 0.9$  kcal/mol

8) Malate dehydrogenase

- Completes the cycle by regenerating **OAA** (a regenerating substrate)
- It is the final of three reactions in which **NADH + H<sup>+</sup>** is produced
- $\Delta G^\circ = +7.1$  kcal/mol (highly positive), but it is driven by the very low concentration of OAA due to the highly exergonic citrate synthase reaction.

- The net reaction of TCA cycle is:



- The Overall net  $\Delta G$  of TCA cycle is negative (*favorable*, -228 Kcal/mole)
  - It produces GTP, 3 NADH, FADH<sub>2</sub> which energetically equals 10 ATP molecules
  - Most steps are *irreversible* steps
- Efficient burning depends mainly on the presence of carbohydrates to provide OAA
- *Fats can't be converted into glucose*, because pyruvate dehydrogenase reaction is an absolutely **irreversible** step
  - Pyruvate dehydrogenase is the enzyme that oxidizes pyruvate into Acetyl CoA
- It is a central pathway because it is **amphibolic** (contributes in *catabolism* and *anabolism*)
  - Citrate is important in the synthesis of FAs (in the liver)
  - Malate is important in the gluconeogenesis during fasting (in the liver)
  - Succinyl CoA is important in the **heme** biosynthesis in the bone marrow
  - $\alpha$ -Ketoglutarate is important in the production of GABA (an inhibitory neurotransmitter in the brain) and can be converted into glutamine which is an amino acid used in protein synthesis
  - OAA is important in the production of asparagine (protein synthesis)
- TCA is **Anapleiotropic** pathway where its intermediates must be maintained
  - OAA is replenished by pyruvate carboxylase, which:
    - ✓ Found in the liver, kidney, brain, adipocytes and fibroblasts (mainly liver and kidney)
    - ✓ Activated by **Acetyl CoA**
    - ✓ It requires **biotin (vitamin B7)** as a cofactor

## Past Papers

- All of the following enzymatic reactions happen during TCA cycle EXCEPT:
  - Phosphoryl transfer
  - Intramolecular phosphoryl transfer
  - Oxidation reduction
  - Dehydration
  - Decarboxylation
- How many high-energy phosphate molecules are produced in the conversion of Citrate to Succinate?
  - 0
  - 3
  - 6
  - 7
- What is the maximal amount of ATP produced from the oxidation of isocitrate to alpha-ketoglutarate?
  - 0
  - 2
  - 3
  - 1
- One of these is not an intermediate in Krebs cycle:
  - Citrate
  - Alpha ketoglutarate
  - Acetyl CoA
  - Fumarate
  - Oxaloacetate
- Which of the following is the coenzyme for dihydrolipoyl transacetylase:
  - NAD + FAD
  - CoA + NAD
  - Lipoic acid
  - TPP
  - Lipoic acid + CoA
- The reaction which results in the reduction of FAD into FADH<sub>2</sub> is:
  - Citrate to isocitrate
  - Malate to oxaloacetate
  - Succinate into fumarate
  - Fumarate to malate
- Which of the following is not a coenzyme of alpha ketoglutarate dehydrogenase?
  - NAD
  - FAD
  - Lipoic acid
  - ATP



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