

BI/CH 422/622

OUTLINE:

Introduction and review

Transport

Glycogenolysis

Glycolysis

Introduction & overview; 2 phases

Phase I

Phase II

Summary: logic, energetics, labeling studies

Other sugars

Pasteur: Anaerobic vs Aerobic

Fermentations

Lactate-lactate dehydrogenase

Exam-1 material

Acetoacetate decarboxylase

Ethanol-pyruvate decarboxylase & alcohol dehydrogenase

Exam-2 material

Krebs' Cycle

How did he figure it out?

Overview

8 Steps

Citrate Synthase

Aconitase

Isocitrate dehydrogenase

Ketoglutarate dehydrogenase

Succinyl-CoA synthetase

Succinate dehydrogenase

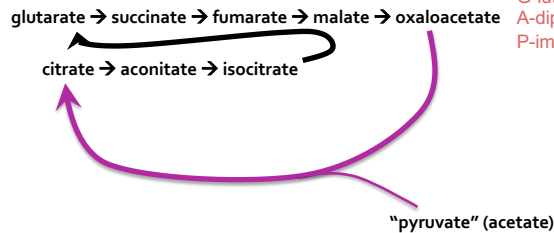
Fumarase

Malate dehydrogenase

Time B.C. (Before the Cycle)



Hans Krebs
1900-1981



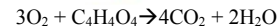
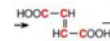
O-leic
M-alonic
S-uccinic
G-lutaric
A-dipic
P-imelic

Krebs confirmed that the pathway was consistent with succinate, fumarate, and malate proved to be useful because all these molecules increased oxygen consumption in the pigeon breast muscle.

Dr. Kornberg: Lecture 02.08.17
(19:54-20:39)

(1 min)

The first clue came from an experiment with fumarate. Krebs did careful measurements using the Warburg manometer. Fumarate gave greater than expected oxygen consumption in the pigeon breast muscle.



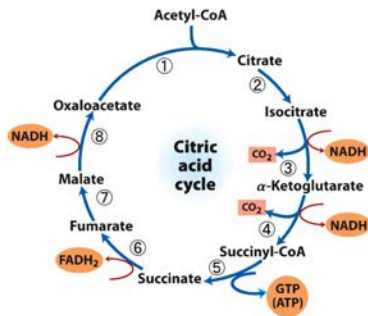
∴ 1 μmole fumarate would consume 3 μmole O₂

- 1) Malonic acid inhibition of the succinate → fumarate step prevented this increase.... BUT, succinate **accumulated!**
- 2) How can fumarate give rise to succinate? There must be a cycle
- 3) Tested by showing that using succinate or fumarate you could detect the formation of citrate.

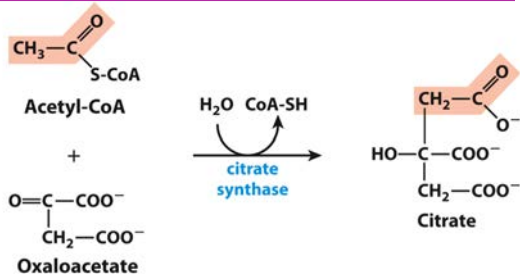
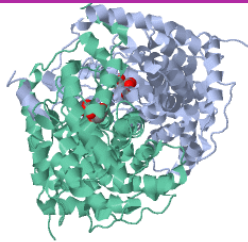
Later in 1937, he proposed that pyruvate would combine with oxaloacetate to make citrate in a cycle he called the Citric Acid Cycle. Later, Fritz Lipmann showed that it was acetyl-CoA and not pyruvate.

The Citric Acid Cycle

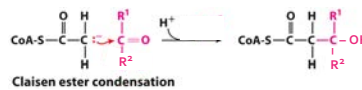
- Step 1: C-C bond formation between acetate (2C) and oxaloacetate (4C) to make citrate (6C)
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The Citric Acid Cycle: Citrate Synthase



- Joining of acetyl-CoA and oxaloacetate with C-C bond formation
- Highly thermodynamically favorable/irreversible ($\Delta G^\circ = -7.7$ kcal/mol)
 - regulated by substrate availability and product inhibition
- Activity largely depends on [oxaloacetate].
- Rate-limiting step of CAC
- Uses acid/base catalysis
 - Carbonyl of oxaloacetate is a good electrophile.
 - Methyl of acetyl-CoA is not a good nucleophile...
 - ...unless activated by deprotonation to form a carbanion.



The Citric Acid Cycle: Citrate Synthase

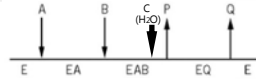
Mechanism

- Conformational change occurs upon binding oxaloacetate.
- Avoids unnecessary hydrolysis of thioester in acetyl-CoA

a) Open conformation:

Free enzyme does not have a binding site for acetyl-CoA. Ordered binding.

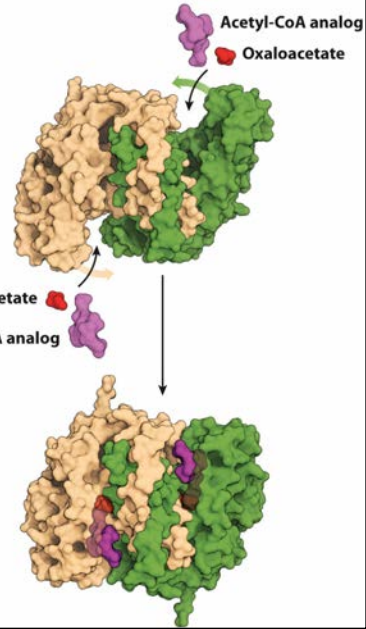
Sequential ordered ter bi



b) Closed conformation:

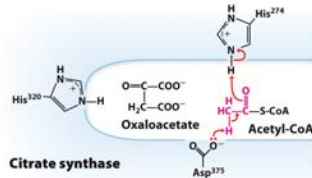
Binding of OAA creates binding for acetyl-CoA.
Reactive carbanion is protected.

(a)

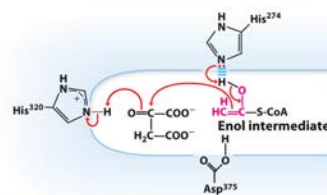
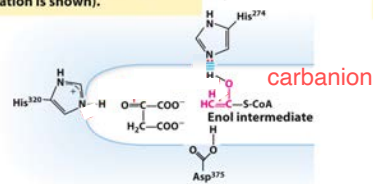
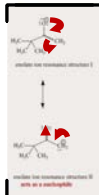


The Citric Acid Cycle: Citrate Synthase

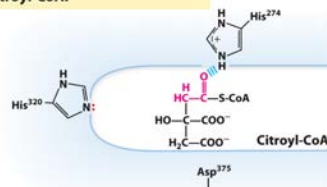
Mechanism



The thioester linkage in acetyl-CoA activates the methyl hydrogens. Asp³⁷⁵ abstracts a proton from the methyl group, forming an enolate intermediate. The intermediate is stabilized by hydrogen bonding to and/or protonation by His²⁷⁴ (full protonation is shown).



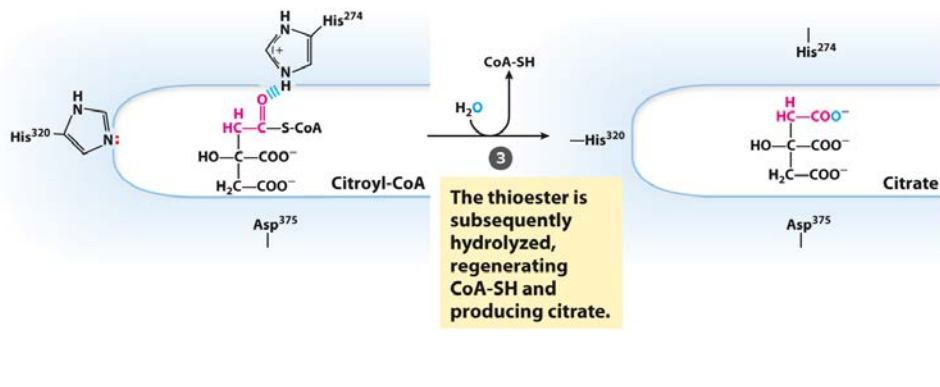
The enol(ate) rearranges to attack the carbonyl carbon of oxaloacetate, with His²⁷⁴ positioned to abstract the proton it had previously donated. His³²⁰ acts as a general acid. The result generates citryl-CoA.



The Citric Acid Cycle: Citrate Synthase

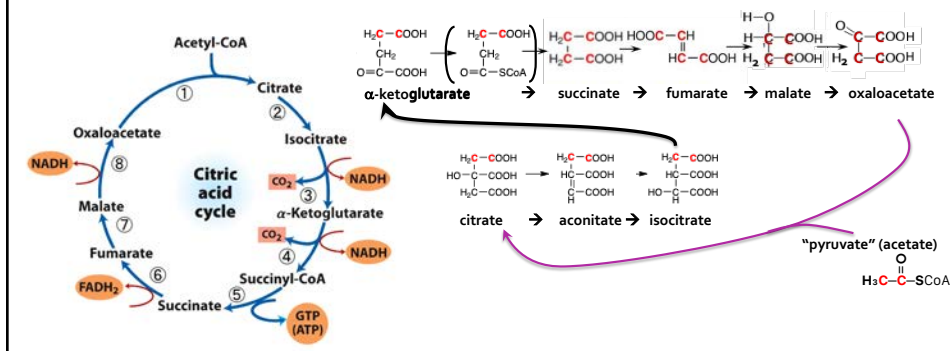
Mechanism

Hydrolysis of Thioester; citroyl-CoA



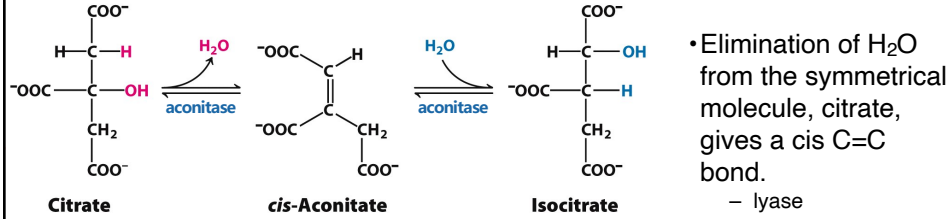
The Citric Acid Cycle

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- Step 1: C-C bond formation between acetate (2C) and oxaloacetate (4C) to make citrate (6C)
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 - Step 8: Dehydrogenation to give NADH



The Citric Acid Cycle: Aconitase

Isomerization by Dehydration/Rehydration



• Rationale:

- Citrate, a tertiary alcohol, is a poor substrate for oxidation.
- Isocitrate, a secondary alcohol, is a good substrate for oxidation.

• Thermodynamically **unfavorable/reversible** ($\Delta G^\circ = +3.2 \text{ kcal/mol}$)

- product concentration kept low to pull forward; citrate tends to “pool” with higher conc.

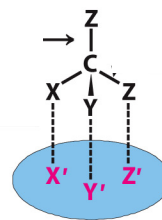
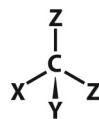
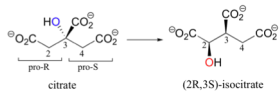
• Dehydration & Addition of H₂O to cis-aconitate is stereospecific.

- This was initially very confusing to bio/organic chemists
- Only R-isocitrate is produced by aconitase.
- A biochemist names A.G. Ogston clarified the situation by realizing that the enzyme spatially templates this symmetrical molecule by binding in only one way (e.g., clockwise or counter clockwise, not both)
- Distinguished by three-point attachment to the active site

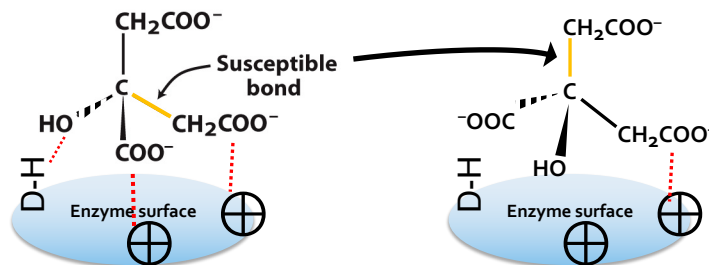
The Citric Acid Cycle: Aconitase

3-point attachment; prochirality

This binding protects the nascent acetate from chemistry

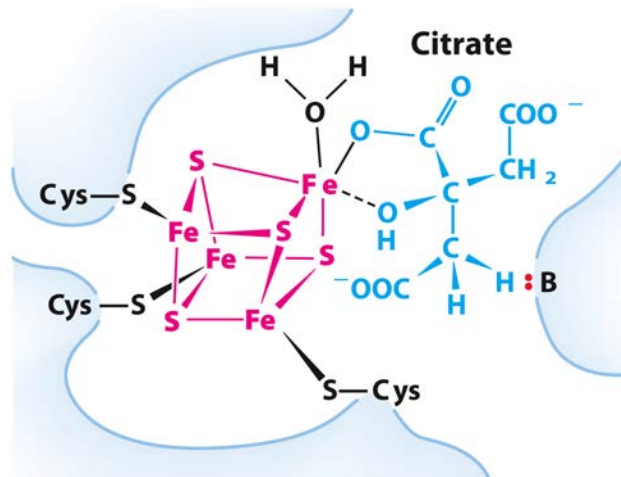


Active site has complementary binding points.



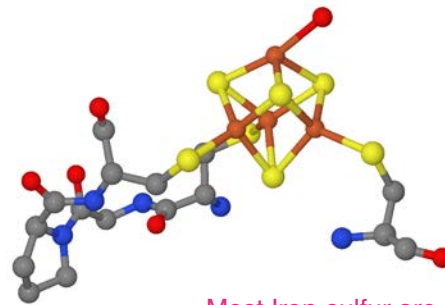
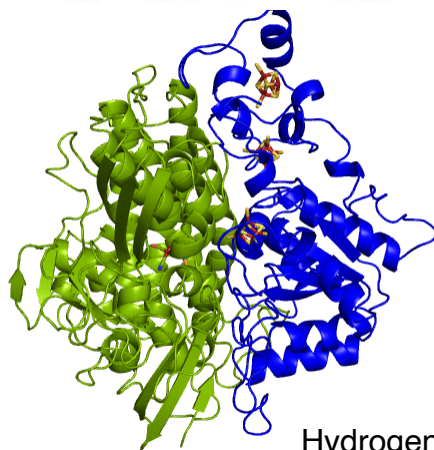
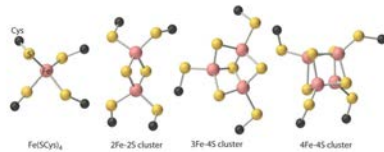
The Citric Acid Cycle: Aconitase

Iron-Sulfur Center in Aconitase

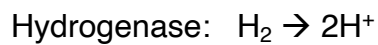


The Citric Acid Cycle: Aconitase

Iron-Sulfur Centers



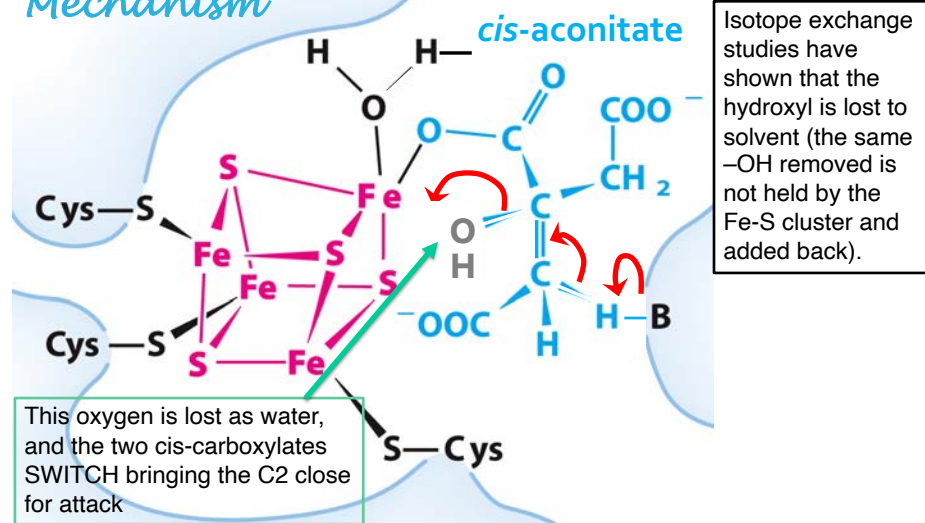
Most Iron-sulfur are involved in re-dox reactions (without proton transfers). In aconitase, the role is very different.



The Citric Acid Cycle: Aconitase

Water removal from *citrate* and subsequent addition to *cis*-aconitate are catalyzed by the **iron-sulfur center**: sensitive to oxidative stress.

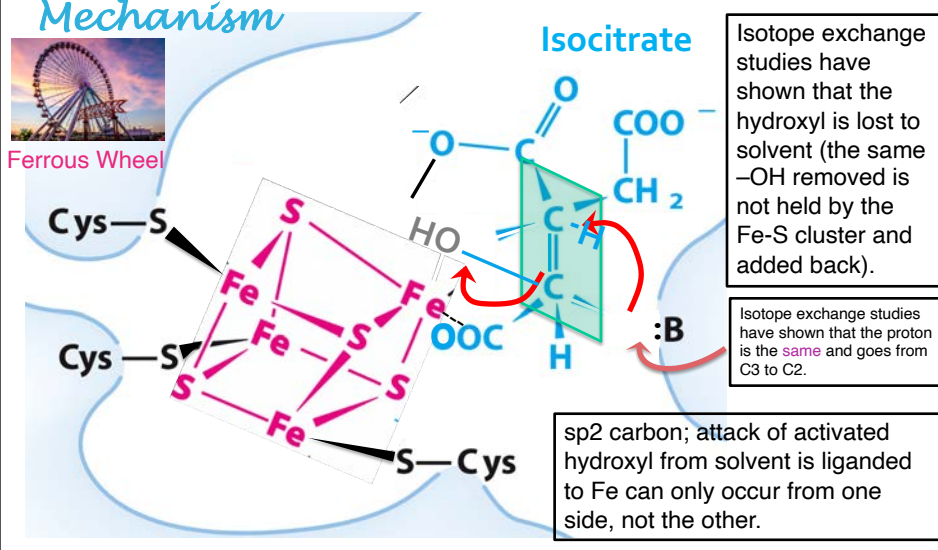
Mechanism



The Citric Acid Cycle: Aconitase

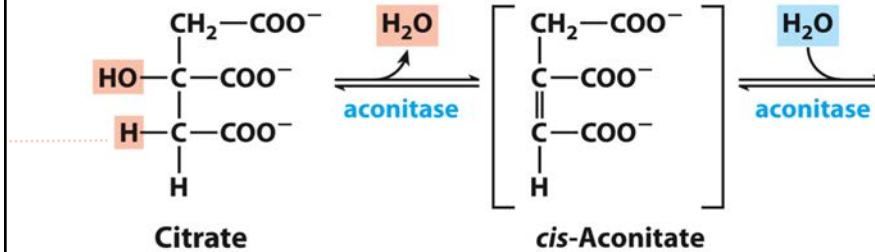
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Mechanism

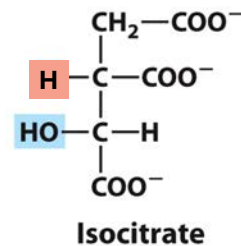


The Citric Acid Cycle: Aconitase

Isomerization by Dehydration/Rehydration



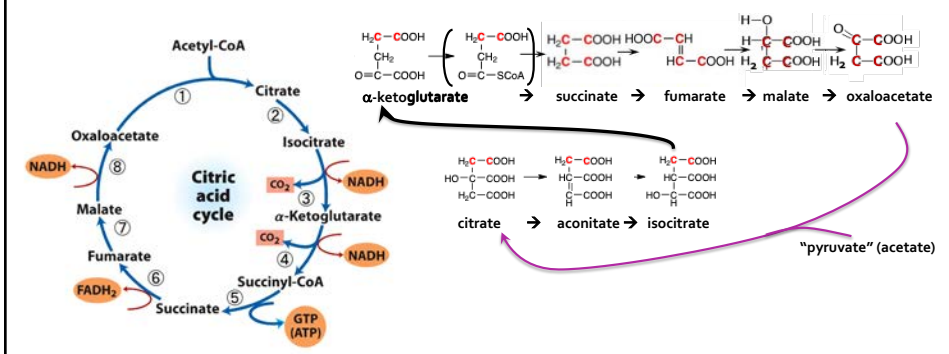
- Dehydration of H₂O from the symmetrical molecule, citrate, gives a *cis* C=C bond.
- Rehydration of H₂O to *cis*-aconitate is stereospecific.



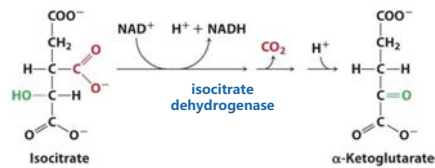
$$\Delta G'^{\circ} = 13.3 \text{ kJ/mol}$$

The Citric Acid Cycle

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- Aconitase** ✓
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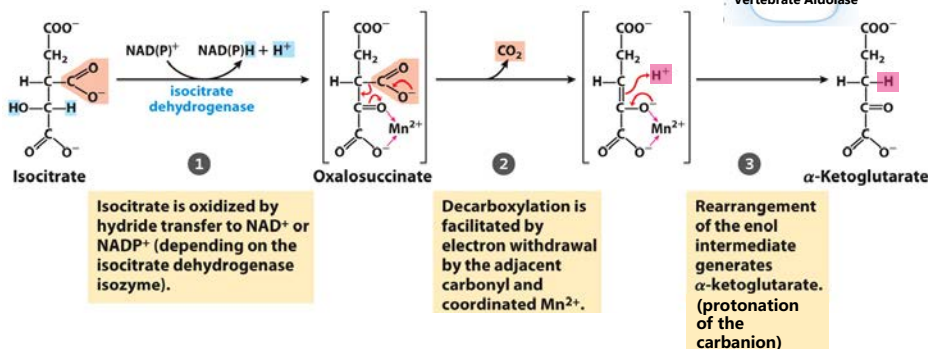
The Citric Acid Cycle: Isocitrate dehydrogenase



- Converting the C2 hydroxyl to a keto destabilizes the C-C bond to the carboxylate at C3 (C-C bond β -to carbonyl).
- This requires a 2-step process:
 - First perform an alcohol-to-keto dehydrogenation at C2 using NAD^+
 - Second, using carbonyl, allow for decarboxylation (the oxidation of the carboxylate to CO_2 , with the reduction of C3).
 - C2 is oxidized, C3 is reduced, Carboxylate is oxidized: Net oxidation is $2e^-$
- Isozymes are specific for NADP^+ (cytosolic) or NAD^+ (mitochondrial).
- Favorable but irreversible due to loss of CO_2 ($\Delta G^\circ = -2.0 \text{ kcal/mol}$)
- Regulated by [ATP] (OMSGAP)

The Citric Acid Cycle: Isocitrate dehydrogenase

Mechanism



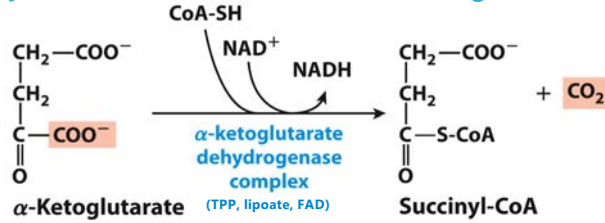
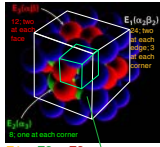
This mechanism is just like that of LDH or other dehydrogenases (Base abstracts $-\text{OH}$ proton, carbonyl forms, elimination of $:\text{H}-$)

The Mn^{++} cofactor stabilizes the α -keto acid, which destabilizes the middle carboxylate (electron sink)

This is an oxidative decarboxylation at carboxylate β -to carbonyl, so it does not need TPP!! Uses Mn & α -carbonyl

The Citric Acid Cycle: The α -Keto-Glutarate Dehydrogenase Complex

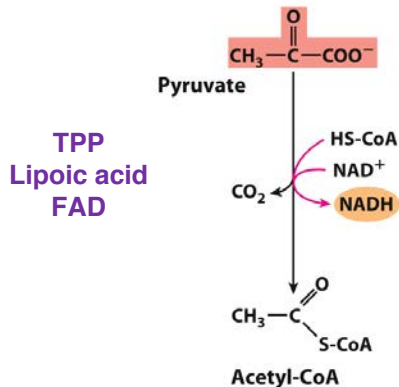
Oxidative Decarboxylation of an α -keto acid: α -Ketoglutarate Dehydrogenase



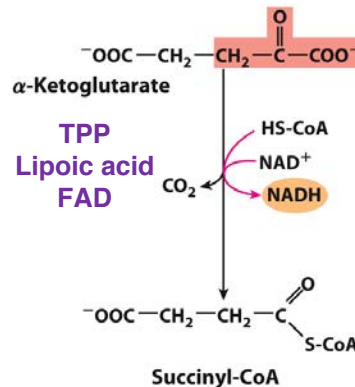
- Last oxidative decarboxylation in TCA cycle
 - full oxidation of all carbons of glucose:
 - Takes two turns of the cycle
 - The carbons oxidized are not directly from glucose because the carbons came from oxaloacetate, not acetyl-CoA
- Requires TPP, FAD, Lipoic acid cofactors Where have we seen this before?
- Succinyl-CoA is another higher-energy thioester bond.
- Highly thermodynamically **favorable/irreversible** ($\Delta G^\circ = -8.0 \text{ kcal/mol}$)
 - regulated by product inhibition

The Citric Acid Cycle: The α -Keto-Glutarate Dehydrogenase Complex

Pyruvate dehydrogenase complex

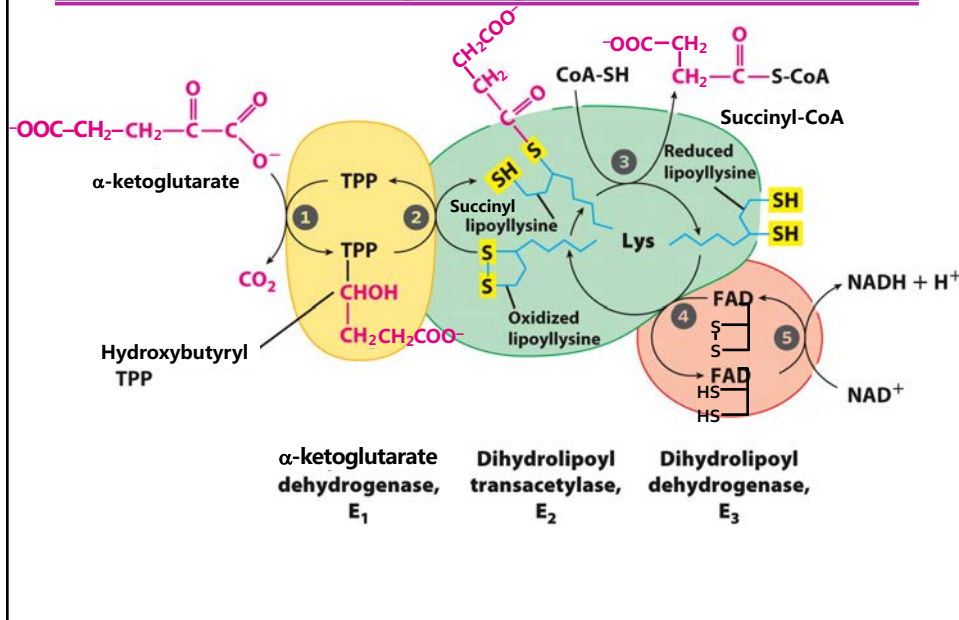


Citric acid cycle (α -KGDH)



- Complex similar to pyruvate dehydrogenase
 - same coenzymes, identical mechanisms, E2 & E3 are identical
 - active site of E1 different to accommodate different-sized substrates

The Citric Acid Cycle: The α -Keto-Glutarate Dehydrogenase Complex



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- Isocitrate dehydrogenase** ✓ • Steps 3–4: Oxidative decarboxylations to give 2 $NADH$
- α -ketoglutarate dehydrogenase complex** • Step 5: Substrate-level phosphorylation to give GTP
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- Step 8: Dehydrogenation to give $NADH$

