

Oxidative Pyruvate Decarboxylation and the Citric Acid Cycle

1

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Fates of pyruvate

- Aerobic: oxidize to CO₂
- Anaerobic: fermentation

2

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Core concept mastery check.....

3

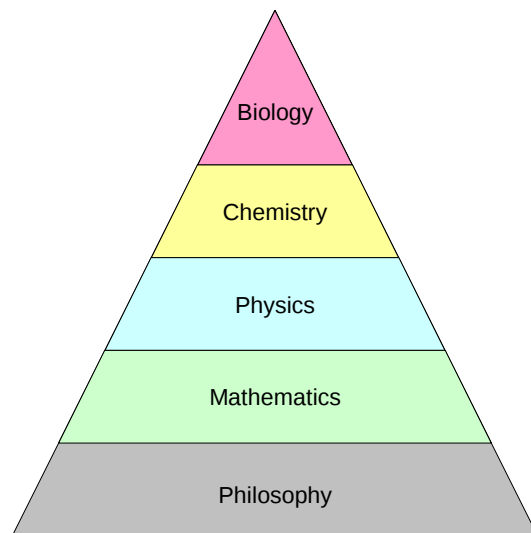
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If you want to really understand oxidative metabolism, you have to be able to keep track of electrons. To keep track of electrons, you have to know how to determine oxidation numbers!!

4

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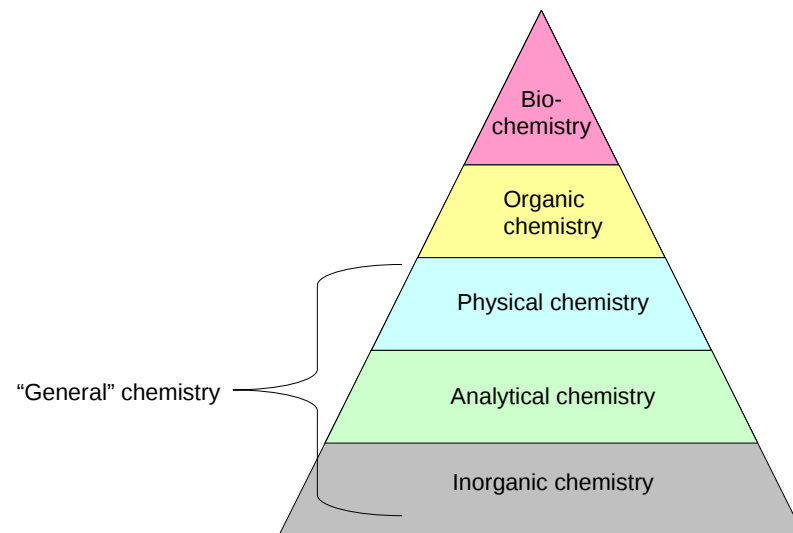
The hierarchy of scientific learning....



5

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The hierarchy of biochemical learning..



6

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Rules for assigning oxidation numbers

1. For an atom in its elemental form, oxidation # = 0
2. For a monatomic ion, oxidation # = ion charge
3. For binary compounds of nonmetals: Assign the element with greater electronegativity the "ion charge" it would have if it was in an ionic compound
4. The sum of oxidation #s is 0 in a neutral compound, while the sum is the overall charge for polyatomic ions

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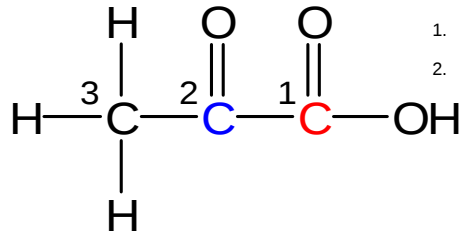
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Oxidation numbers for bioorganic compounds

1. Select the carbon atom you wish to assign an oxidation number
2. Assign oxidation numbers to the atoms bonded to the carbon atom of interest based on their electronegativities relative to the C you are focusing on:
 1. H = +1
 2. O = -2 (except for peroxides, where O = -1)
 3. N = -3
 4. S = -2
 5. C = 0
 6. Phosphorus-carbon bonds are rare in biology; in phosphate compounds, P = +5
3. If the atom bonded to the C of interest is in turn bonded to additional groups, this must be accounted for in assigning an oxidation number for the entire "group" relative to the C. For example, the "OH group" has oxidation number -1 since O is -2 and H is +1 (same for SH). The "NH₂ group" also has oxidation number -1 since N is -3 and each H is +1. For an ether O or S, the -2 charge on the central O/S must be split between the two carbons to which it is bonded, so O/S contributes -1 to each C (see examples following pages).
4. When focusing on another C atom in the same compound, remember that any C atoms bonded to it are assigned oxidation numbers of "0" for the purpose of calculating oxidation numbers, even if they have nonzero oxidation numbers relative to the overall molecule
5. The sum of oxidation #s for all atoms is 0 in a neutral compound, while the sum of oxidation #s is the overall charge for an ionic species. For organic compounds that have ionizable groups, I recommend drawing the structure of the compound in the nonionized form so you don't have to keep track of the net charge.
6. When assigned properly, oxidation numbers will allow you to keep track of which specific C atom(s) gain or lose electrons during an oxidation/reduction reaction

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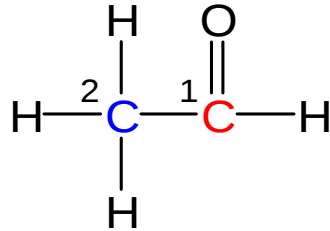
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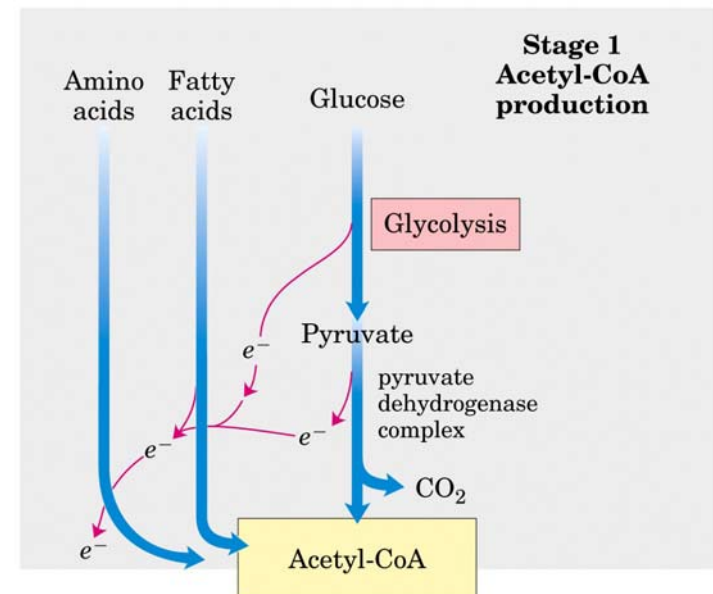
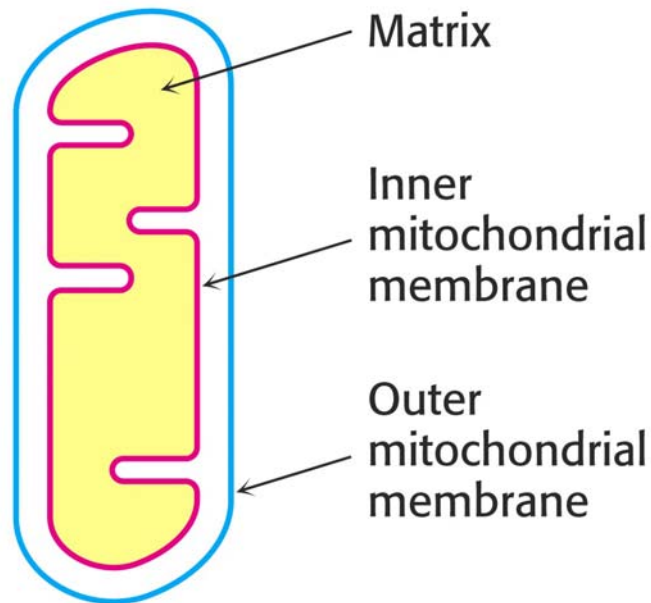
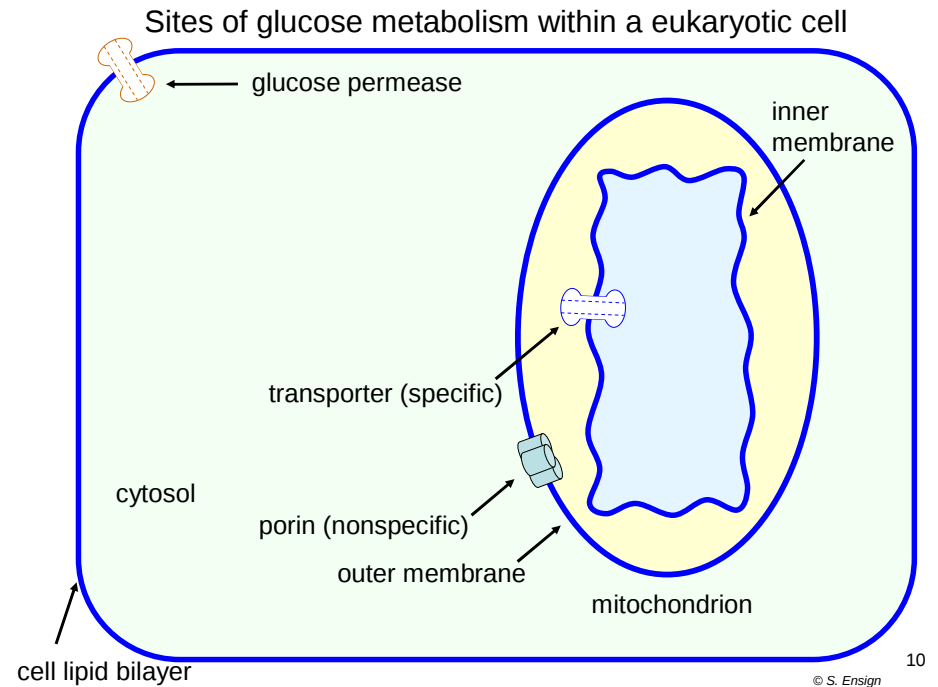
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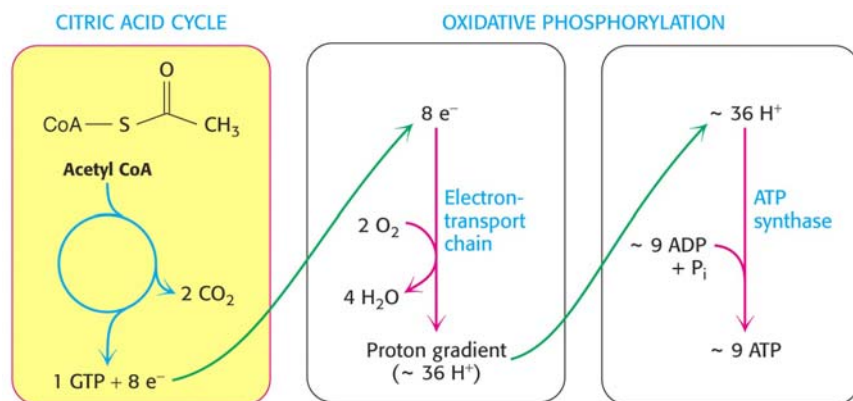
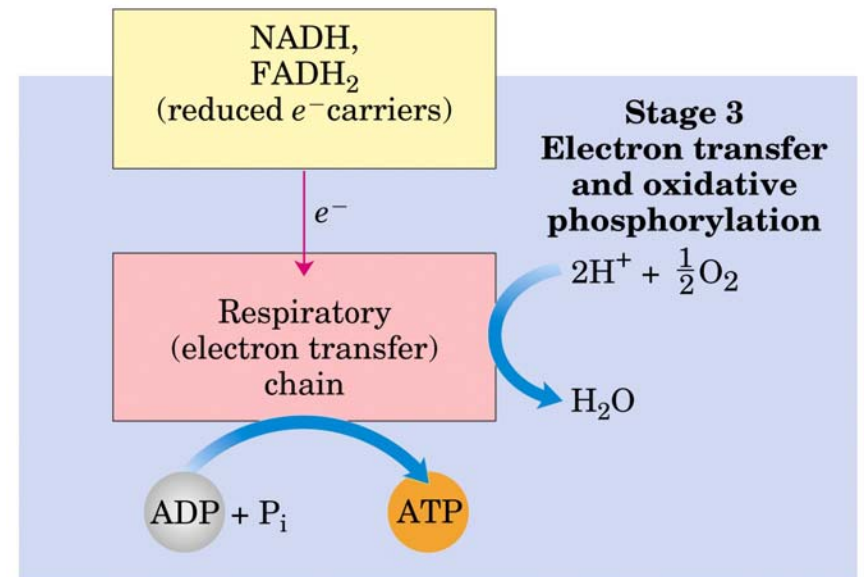
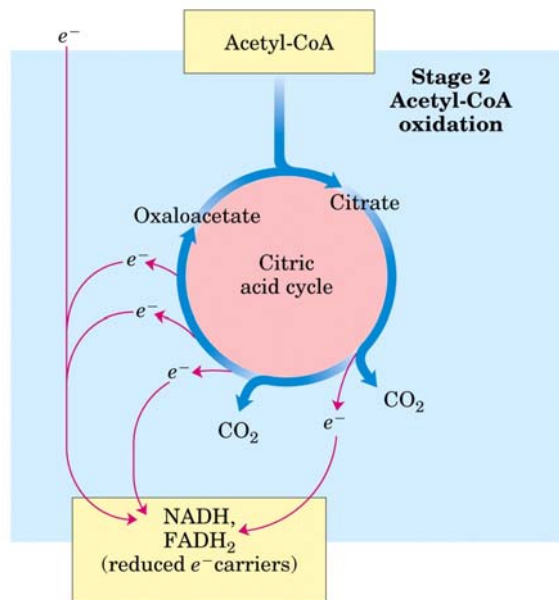
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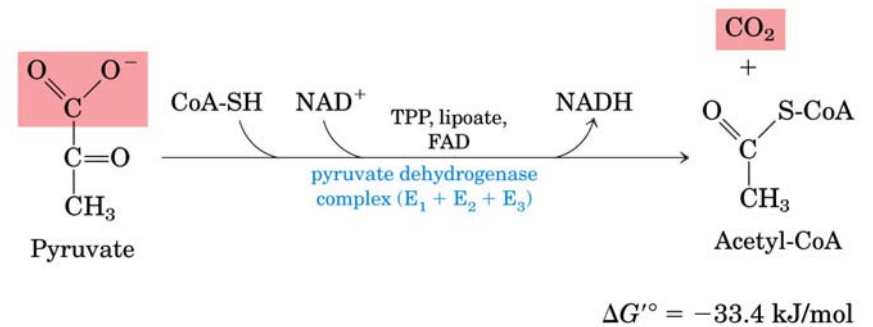


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5. The sum of oxidation #s for all atoms is 0 in a neutral compound, while the sum of oxidation #s is the overall charge for an ionic species.





Oxidative pyruvate decarboxylation



Pyruvate dehydrogenase

- A multienzyme complex consisting of three enzymes that work in unison
- E_1 : pyruvate dehydrogenase
- E_2 : dihydrolipoyl transacetylase
- E_3 : dihydrolipoyl dehydrogenase

Pyruvate dehydrogenase cofactors

Cofactor	Vitamin precursor	Function	Bound or dissociable?
Thiamine pyrophosphate	Thiamin	Decarboxylation	Bound, E_1
Lipoic acid (lipoamide)		e^- and acyl carrier	Bound, E_2
FAD	Riboflavin	e^- carrier	Bound, E_3
NAD ⁺	Niacin	e^- transfer	Diss. E_3
Coenzyme A	pantothenate	Acyl carrier	Diss. E_2

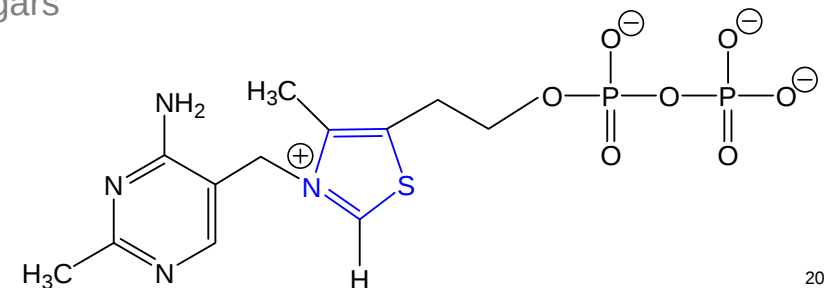
Introduction to the cofactors of the pyruvate dehydrogenase complex

- Three you have seen already
- Two which are new
- Tightly bound (catalytic):
 - TPP (noncovalent)
 - Lipoamide (covalent)
 - FAD (covalent in some enzymes, noncovalent in others)
- Substrates in the reaction (get transformed):
 - NAD⁺
 - Coenzyme A

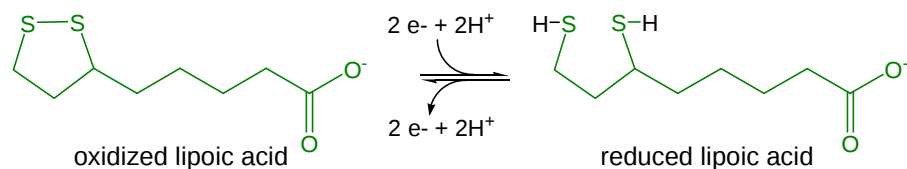
Review..

Thiamine pyrophosphate

- Pyruvate decarboxylase: “nonoxidative” decarboxylation- production of acetaldehyde
- Pyruvate dehydrogenase: “oxidative” decarboxylation
- Transketolase: intermolecular group transfer between sugars



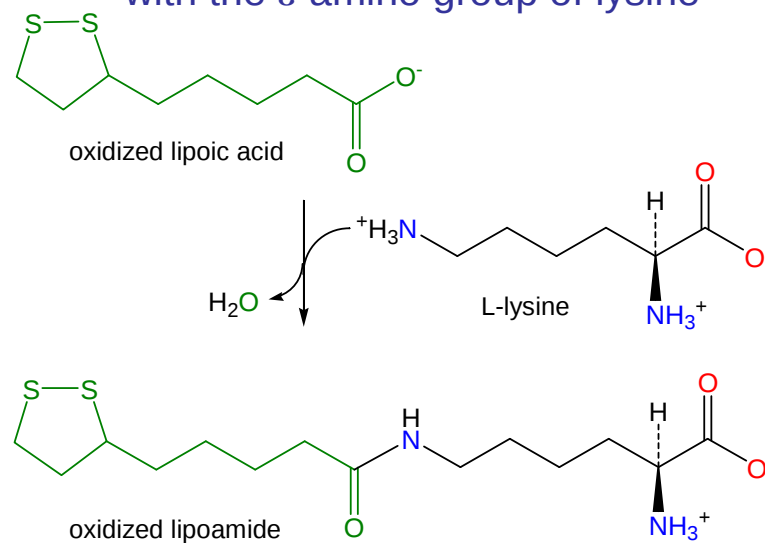
Lipoic acid



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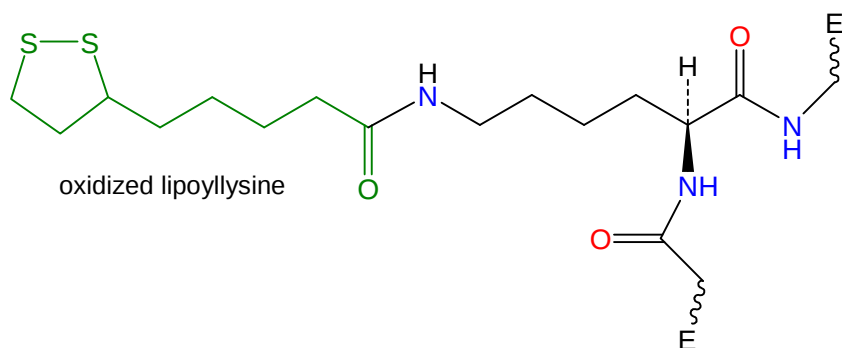
Lipoamide formation by reaction with the ϵ -amino group of lysine



22

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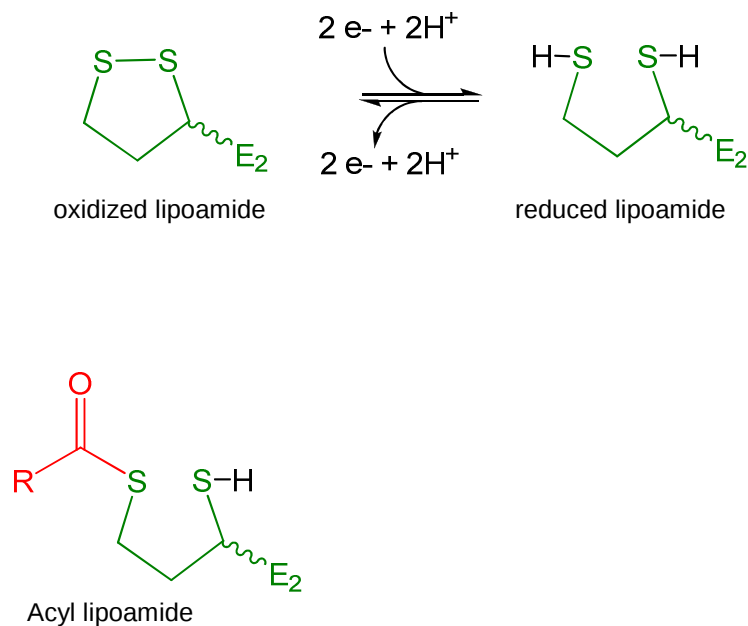
Lipoamide in a polypeptide



23

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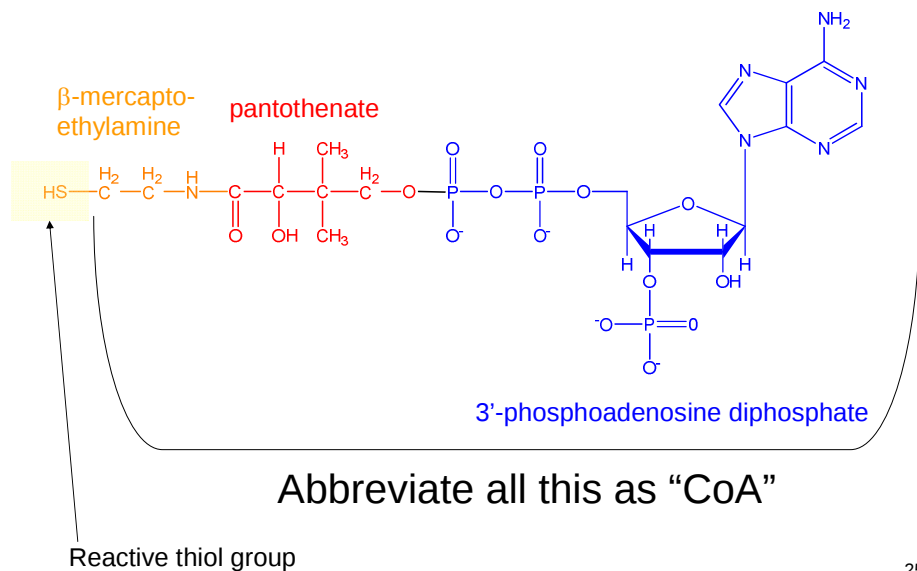
Biologically important forms of lipoamide



24

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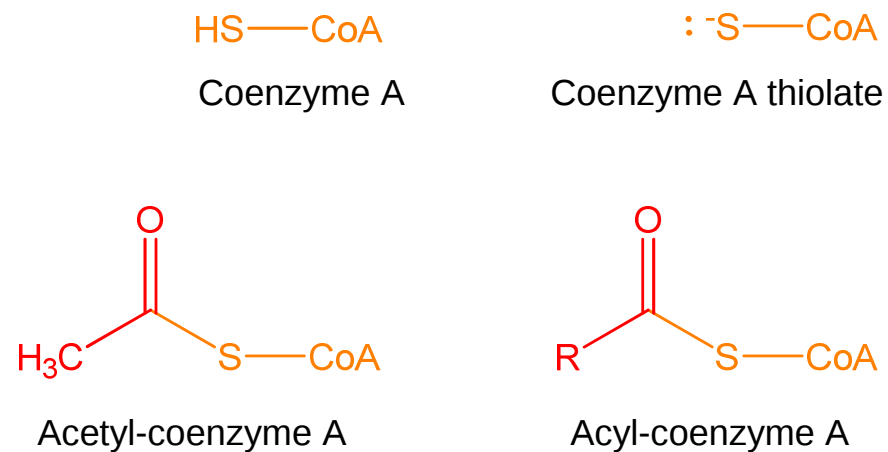
Coenzyme A, an acyl group carrier



25

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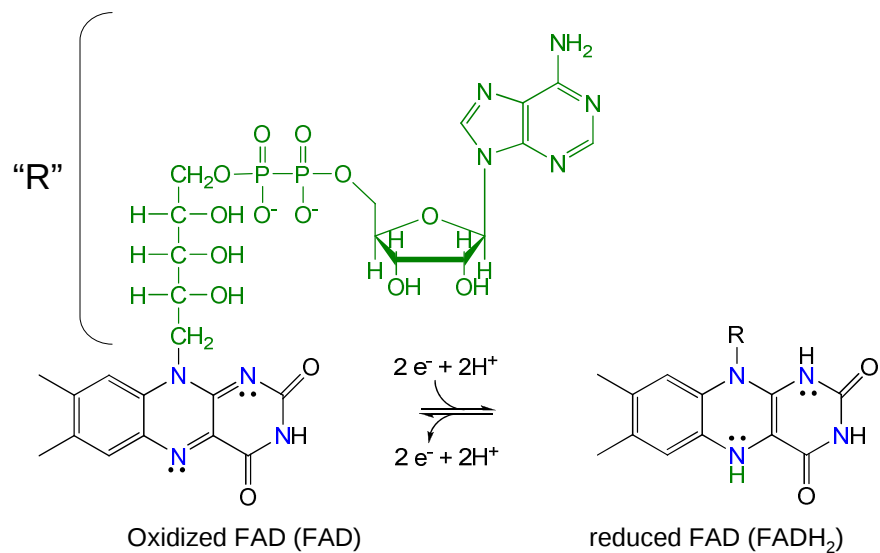
Coenzyme A, an acyl group carrier



26

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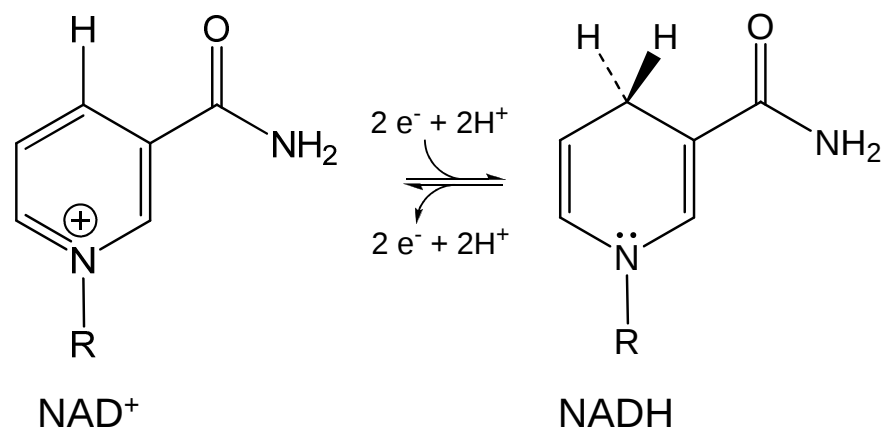
Flavin adenine dinucleotide



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Nicotinamide adenine dinucleotide



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Pyruvate dehydrogenase mechanism

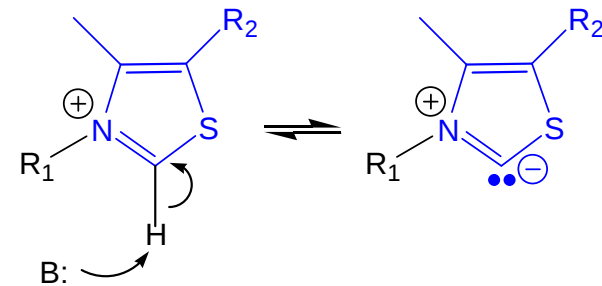
- The three enzyme components work in unison
- E_1 (pyruvate dehydrogenase) decarboxylates pyruvate as seen for pyruvate decarboxylase
- E_2 (dihydrolipoyl transacetylase) accepts the acetyl derived from pyruvate, and then transfers it to Coenzyme A. In the process lipoamide is reduced by two electrons
- E_3 (dihydrolipoyl dehydrogenase) reoxidizes lipoamide using FAD as the initial e^- acceptor and NAD^+ as the terminal e^- acceptor
- The lipoate group “pivots” on E_2 , translocating between E_1 and E_3 during catalysis

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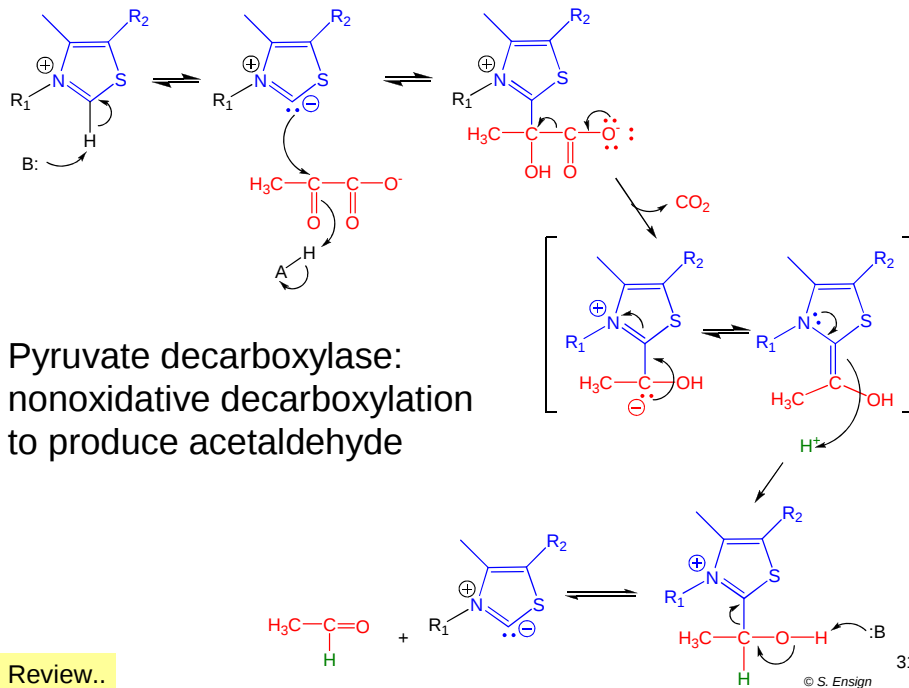
Review..

Recall mechanism of pyruvate decarboxylase



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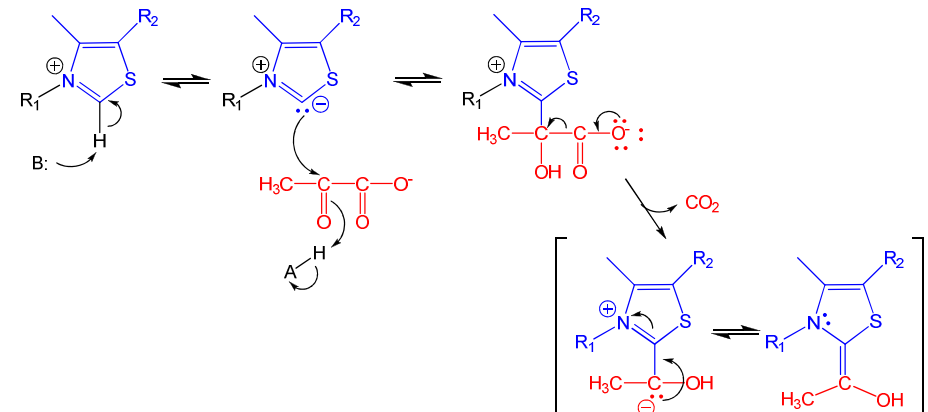
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Review..

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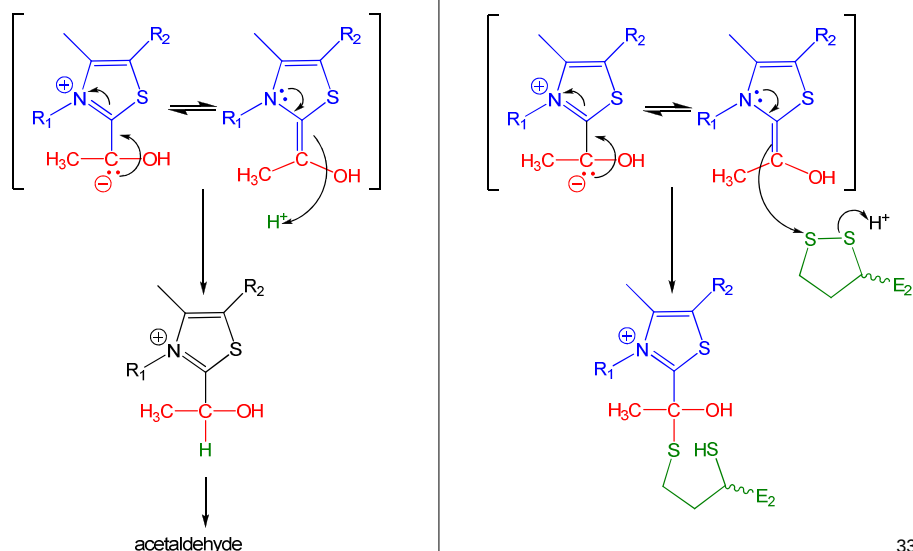


The mechanisms of pyruvate carboxylase and pyruvate dehydrogenase diverge at the level of the hydroxyethyl-TPP carbanion

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In oxidative pyruvate decarboxylation, the hydroxyethyl-TPP anion is transferred to lipoamide rather than H⁺



33

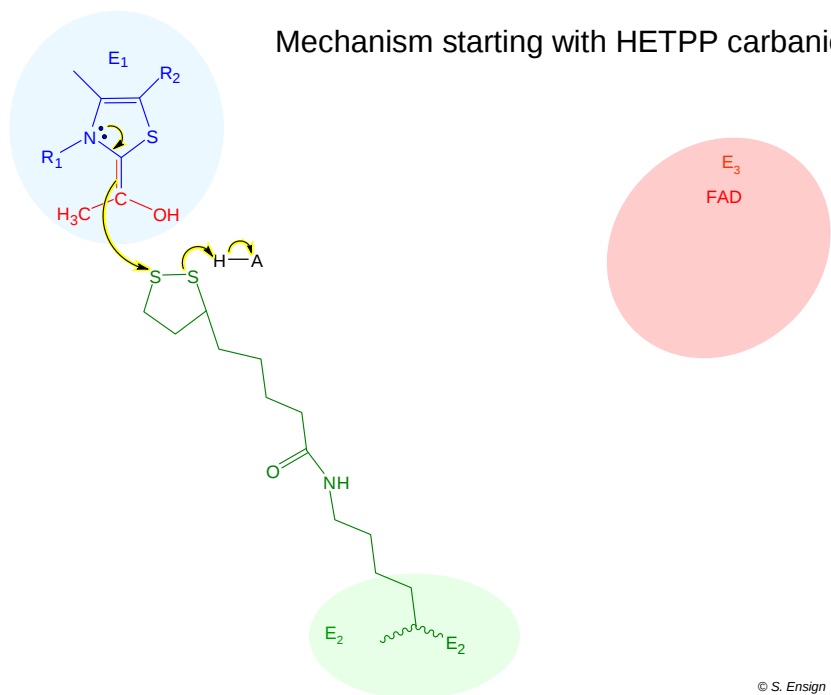
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Watch how the interplay of the activities of the three subunits is mediated by the movement of the flexible linker arm connecting the “business end” of lipoamide

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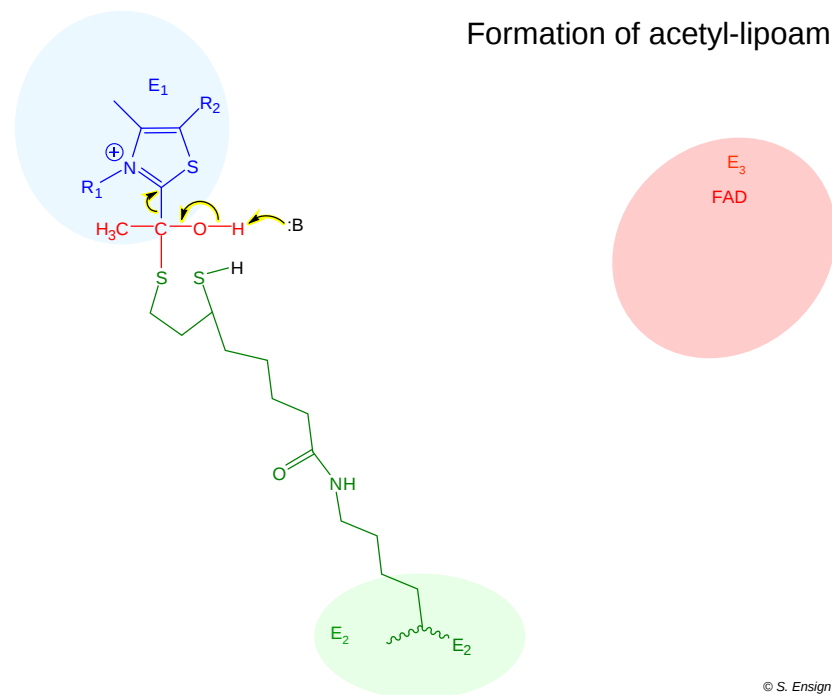
Mechanism starting with HETPP carbanion



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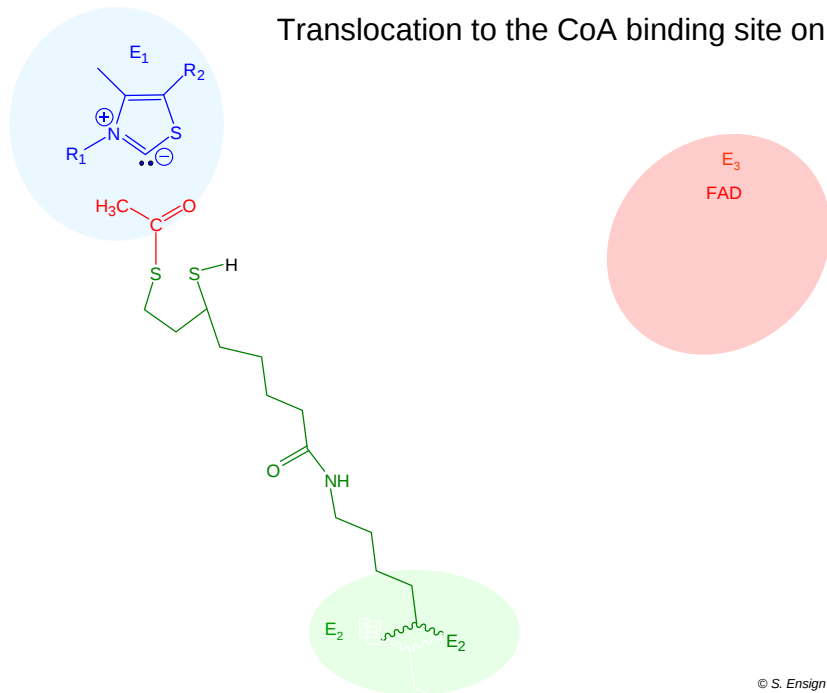
Formation of acetyl-lipoamide



36

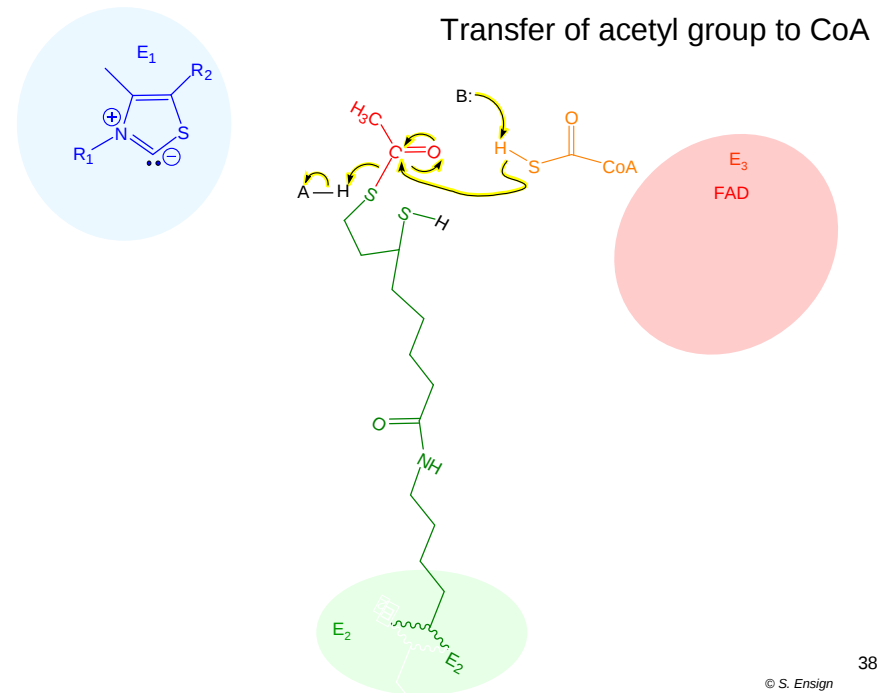
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Translocation to the CoA binding site on E₂



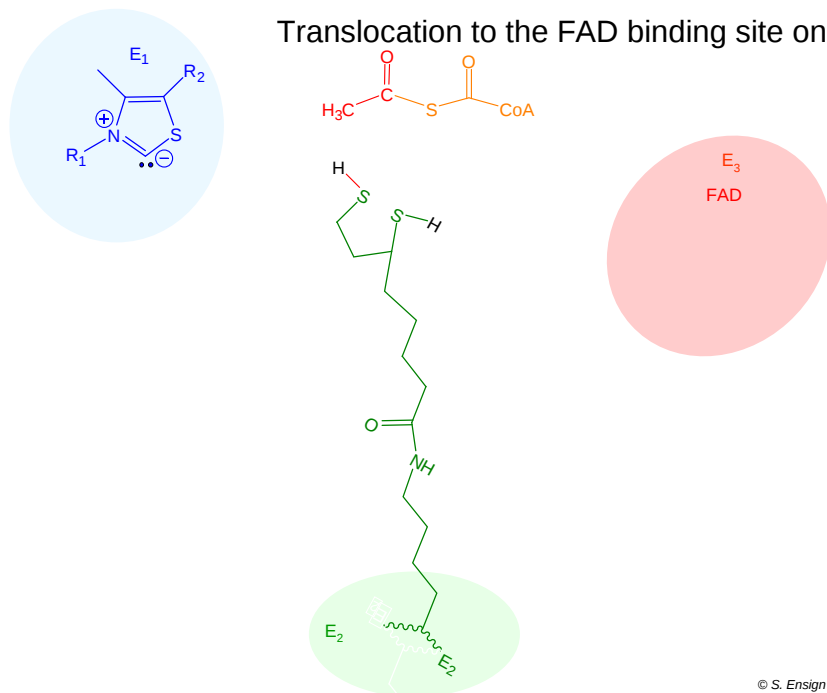
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Transfer of acetyl group to CoA



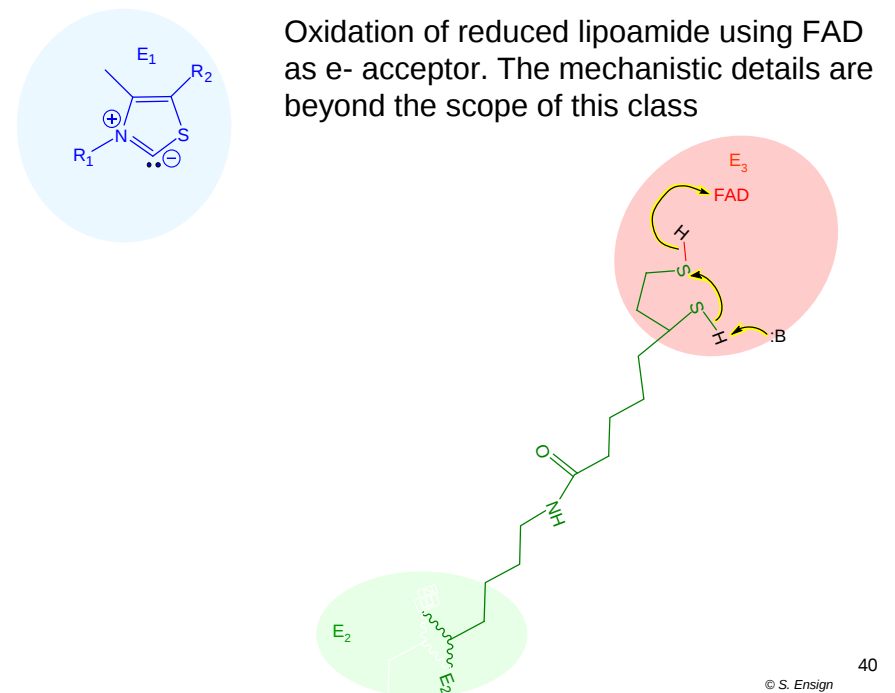
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Translocation to the FAD binding site on E₃

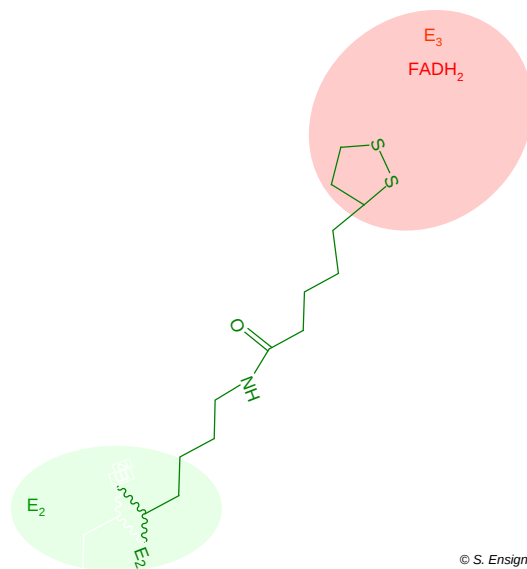


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Oxidation of reduced lipoamide using FAD as e⁻ acceptor. The mechanistic details are beyond the scope of this class

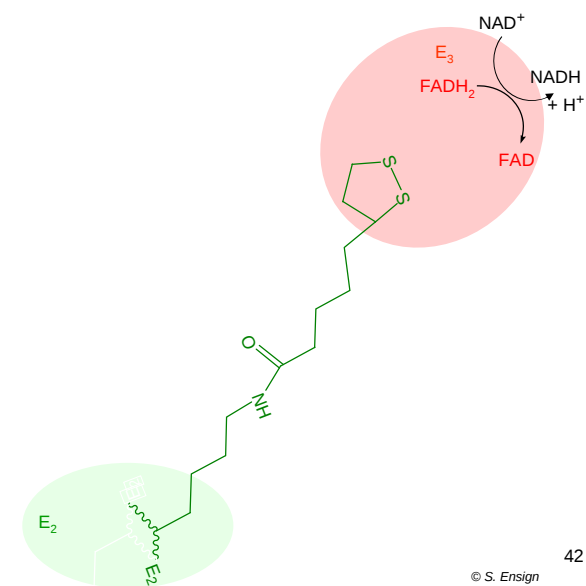
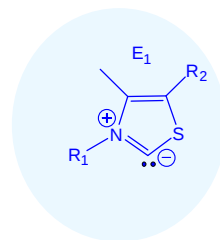


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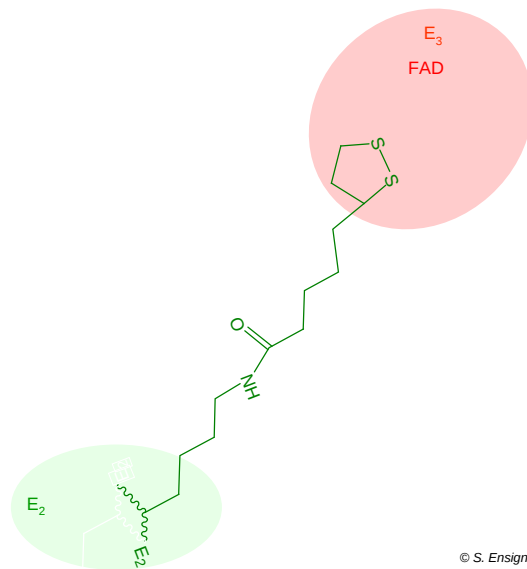
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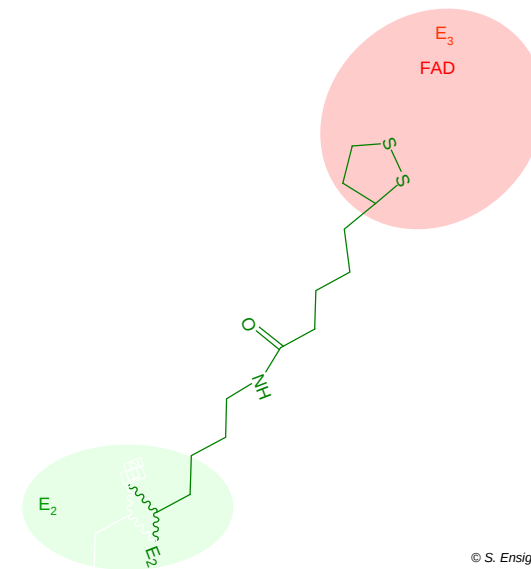
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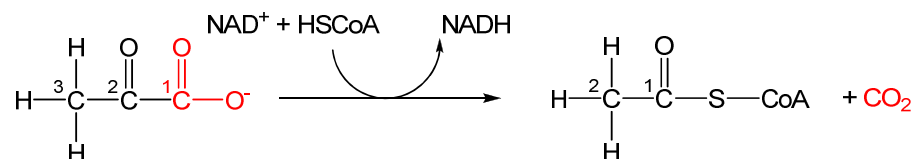
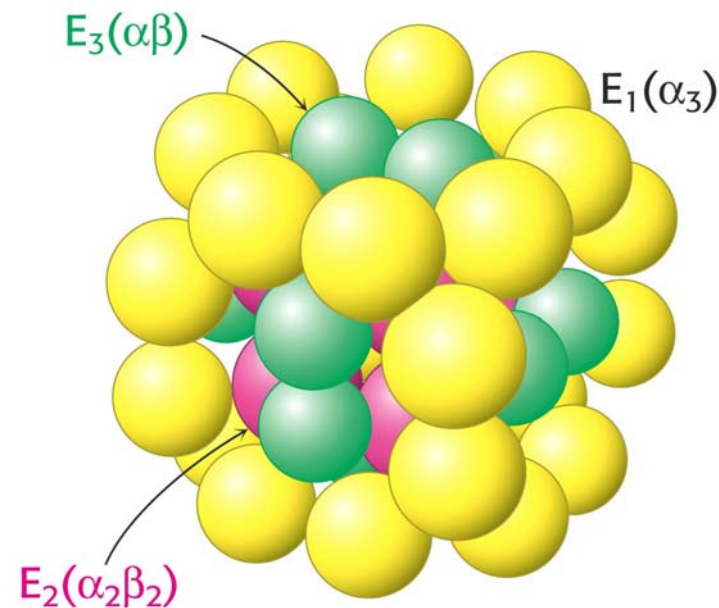
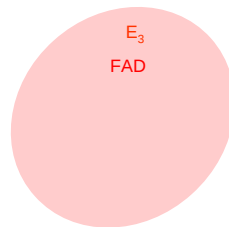
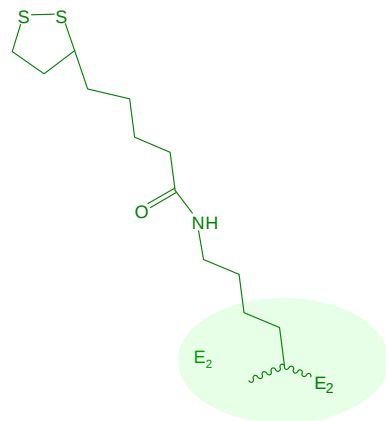
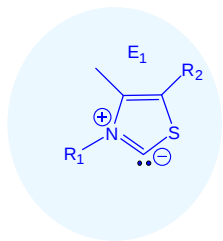
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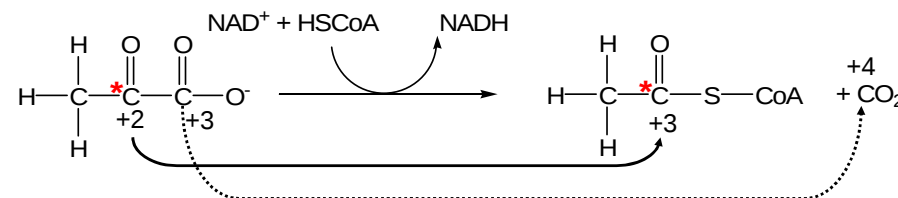
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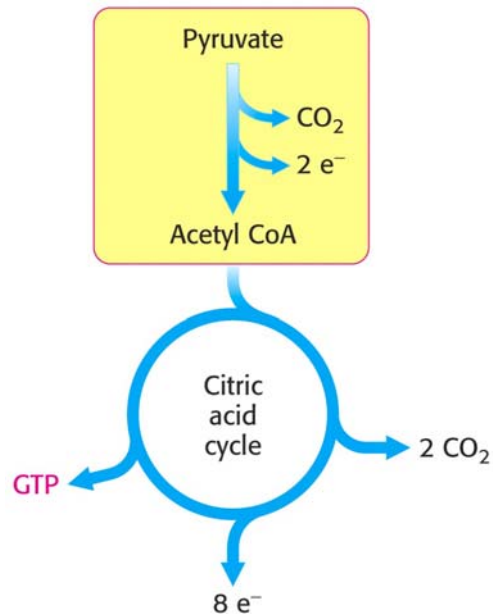
Reduction of NAD^+ by $FADH_2$



“Oxidative decarboxylation” because pyruvate lost two electrons (gained by NAD^+) in the course of the reaction

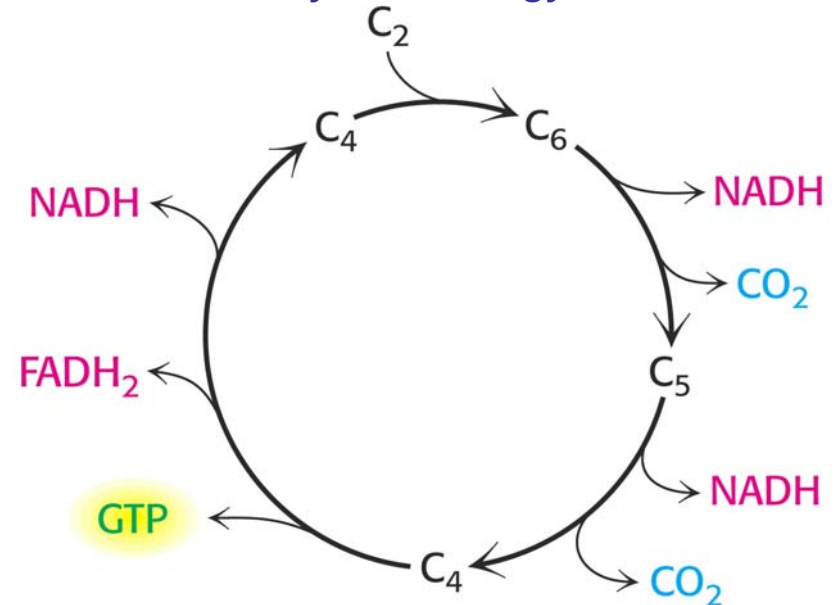
Where did the other electron for NADH come from??:



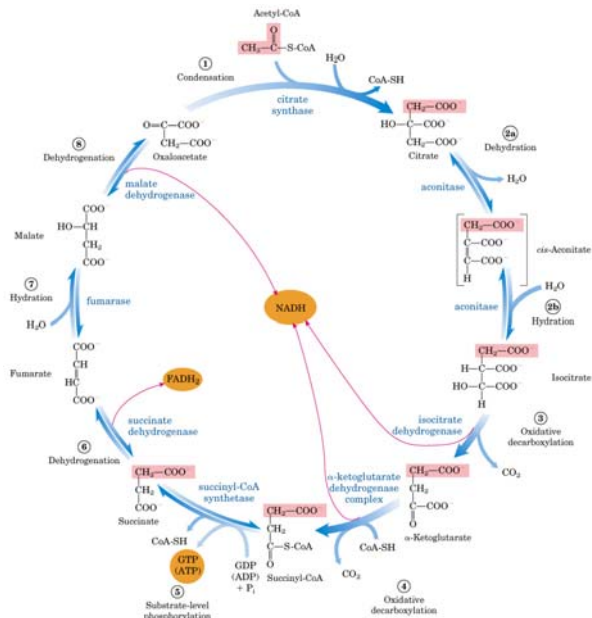


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Citric acid cycle strategy

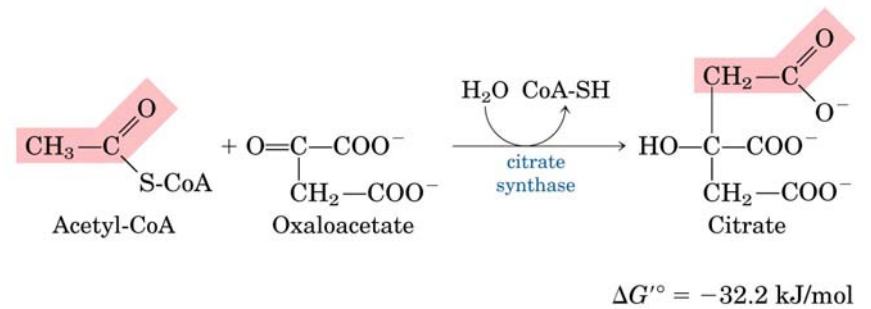


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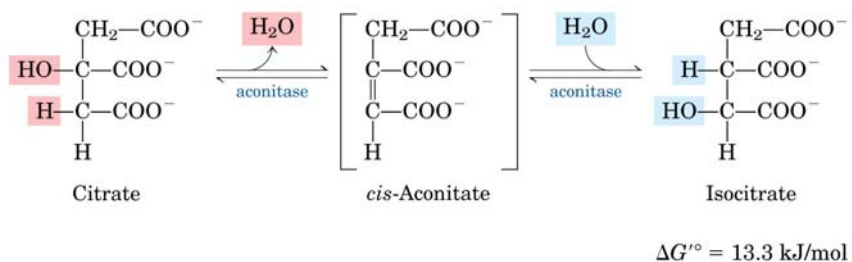
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(1) Claisen condensation of Acetyl-CoA and OAA



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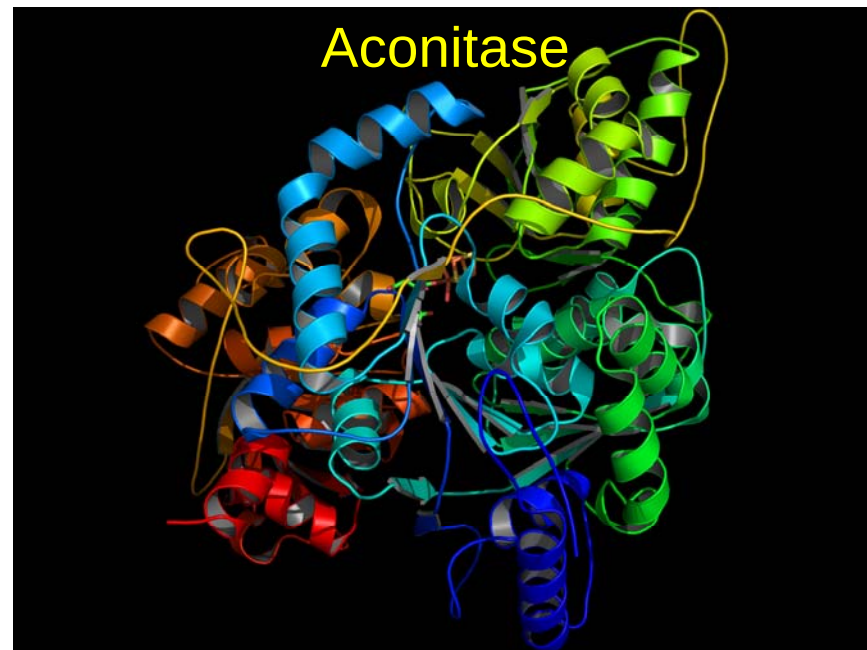
(2) Isomerization of citrate



53

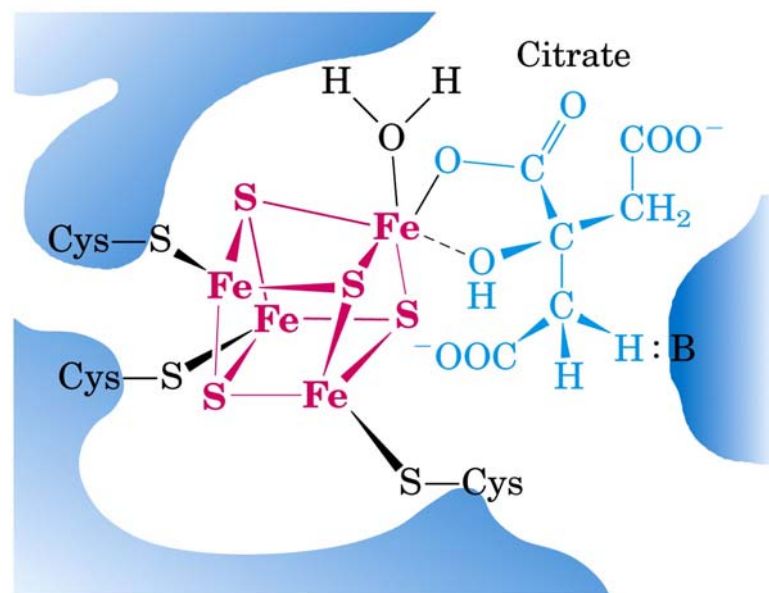
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Aconitase



54

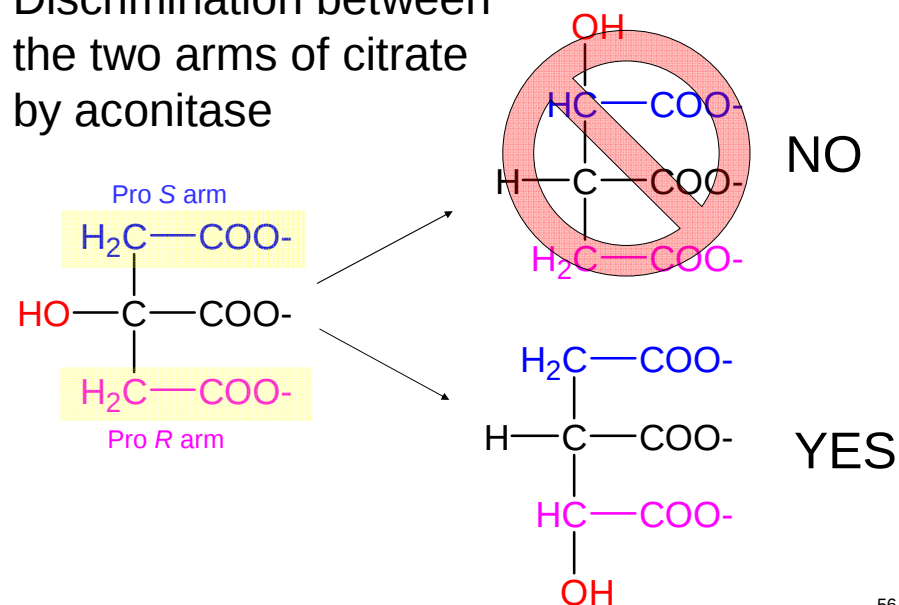
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55

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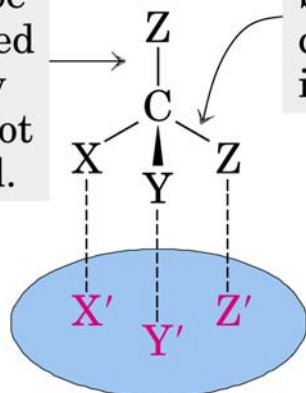
Discrimination between the two arms of citrate by aconitase



56

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This bond cannot be positioned correctly and is not attacked.



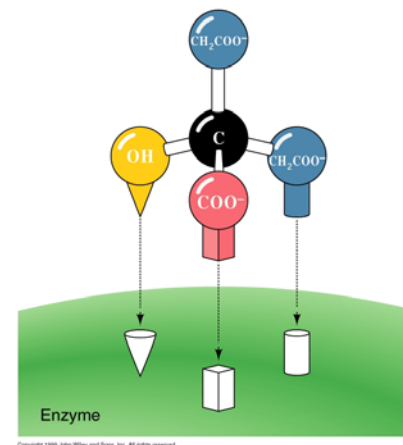
This bond can be positioned correctly and is attacked.

Active site has complementary binding points.

(c)

57

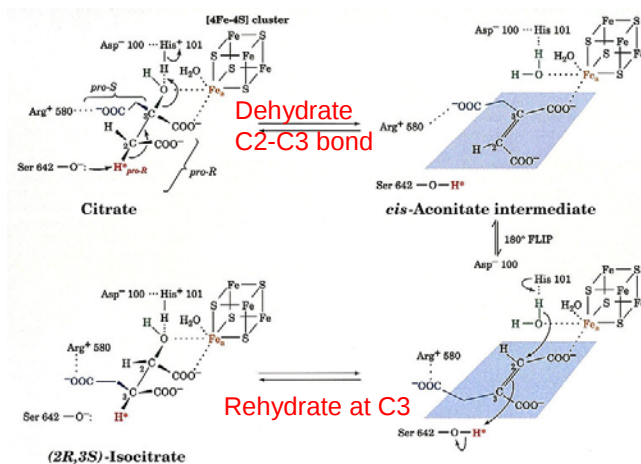
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Aconitase mechanism

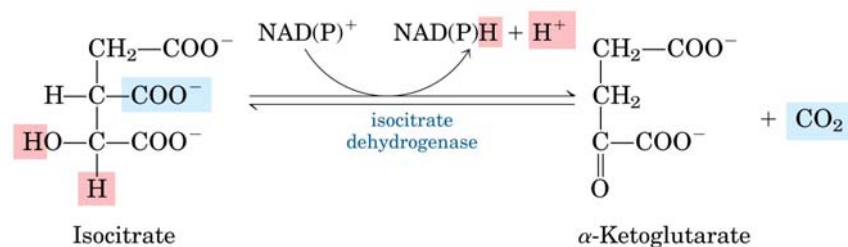


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(3) Oxidative decarboxylation of isocitrate

TPP is not used for this decarboxylation



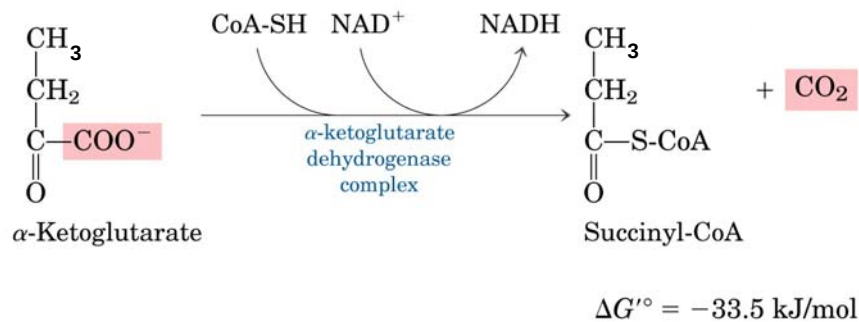
$$\Delta G'^{\circ} = -20.9 \text{ kJ/mol}$$

60

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(4) Oxidative decarboxylation of α -KG

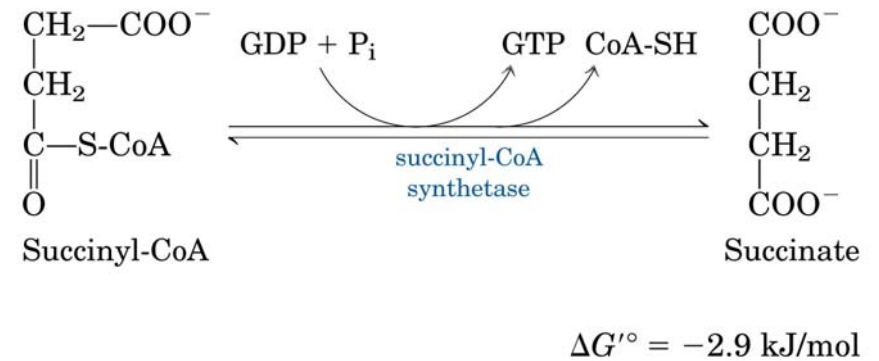
The α -KGDH complex is analogous to the PDH complex



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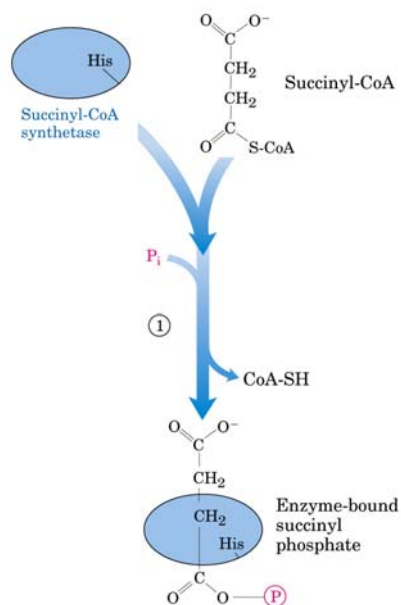
(5) Substrate level phosphorylation



A multistep reaction proceeding as follows...

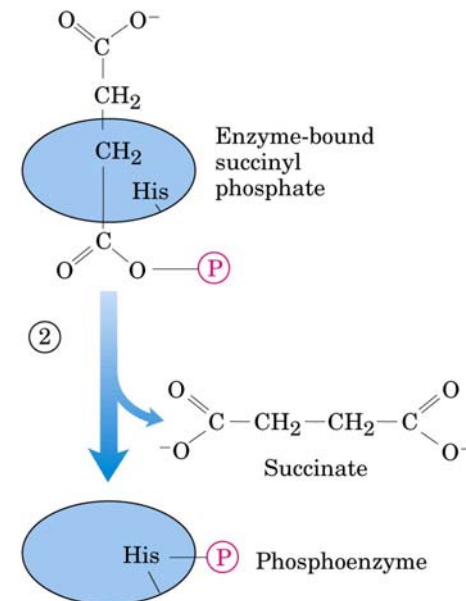
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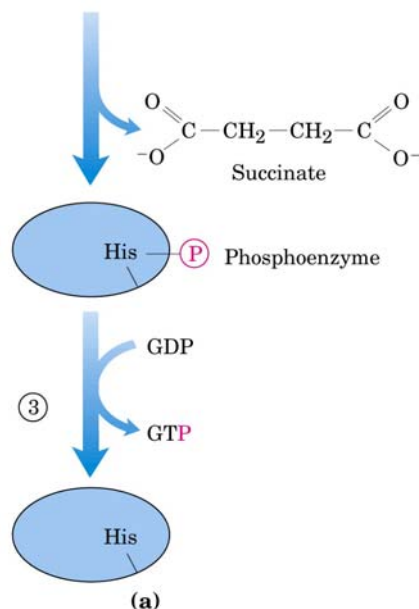
63

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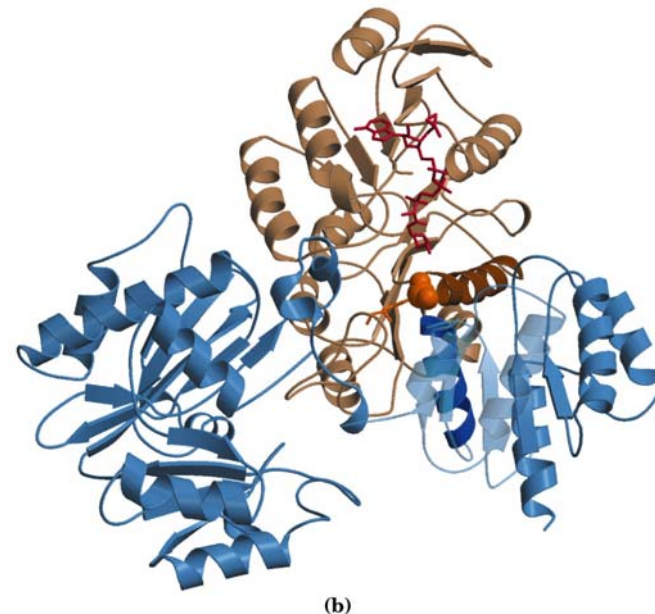
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65

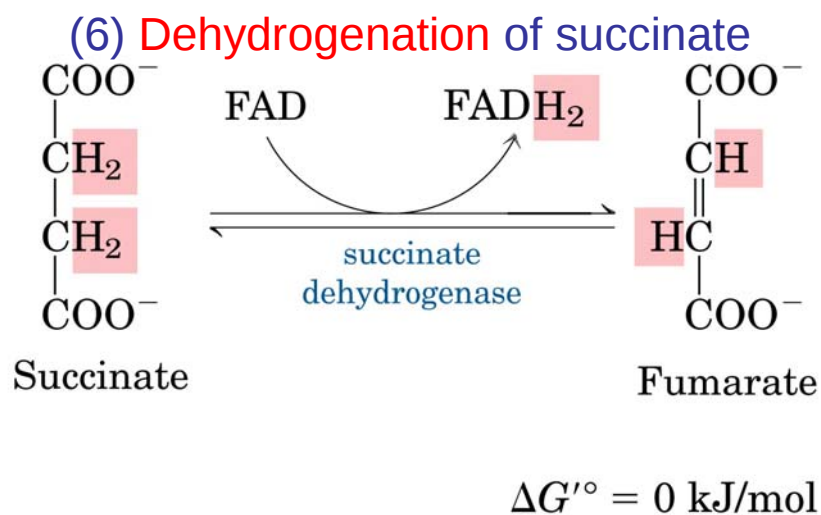
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(b)

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- For a redox reaction to be spontaneous, E'° for the overall reaction must have a positive value
- When writing a half reaction in the oxidative direction, the sign of E'° is switched
- The **oxidized form** of a given redox pair will react spontaneously with the **reduced form** of any redox pair that is below it in a table of standard reduction potentials (under standard conditions)

table 14-7

Standard Reduction Potentials of Some Biologically Important Half-Reactions, at 25 °C and pH 7

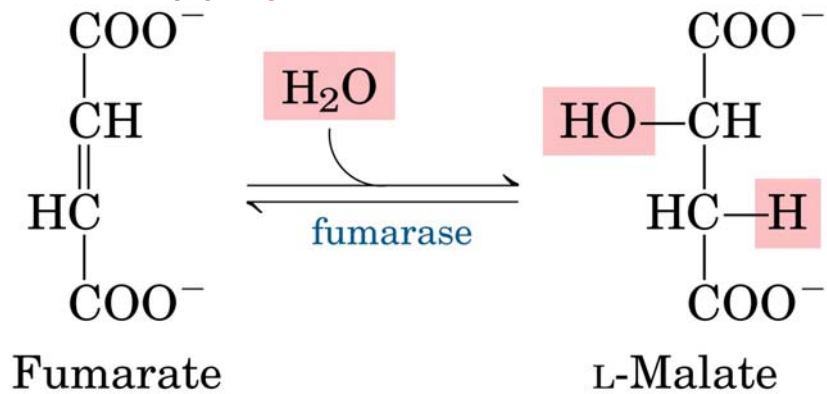
Half-reaction	E'° (V)
$\frac{1}{2}\text{O}_2 + 2\text{H}^+ + 2\text{e}^- \rightarrow \text{H}_2\text{O}$	0.816
$\text{Fe}^{3+} + \text{e}^- \rightarrow \text{Fe}^{2+}$	0.771
$\text{NO}_2^- + 2\text{H}^+ + 2\text{e}^- \rightarrow \text{NO}_2 + \text{H}_2\text{O}$	0.421
Cytochrome f (Fe^{3+}) + $\text{e}^- \rightarrow$ cytochrome f (Fe^{2+})	0.365
$\text{Fe}(\text{CN})_6^{3-}$ (ferricyanide) + $\text{e}^- \rightarrow \text{Fe}(\text{CN})_6^{4-}$	0.36
Cytochrome a_3 (Fe^{3+}) + $\text{e}^- \rightarrow$ cytochrome a_3 (Fe^{2+})	0.35
$\text{O}_2 + 2\text{H}^+ + 2\text{e}^- \rightarrow \text{H}_2\text{O}_2$	0.295
Cytochrome a (Fe^{3+}) + $\text{e}^- \rightarrow$ cytochrome a (Fe^{2+})	0.29
Cytochrome c (Fe^{3+}) + $\text{e}^- \rightarrow$ cytochrome c (Fe^{2+})	0.254
Cytochrome c_1 (Fe^{3+}) + $\text{e}^- \rightarrow$ cytochrome c_1 (Fe^{2+})	0.22
Cytochrome b (Fe^{3+}) + $\text{e}^- \rightarrow$ cytochrome b (Fe^{2+})	0.077
Ubiquinone + $2\text{H}^+ + 2\text{e}^- \rightarrow$ ubiquinol + H_2	0.045
Fumarate²⁻ + $2\text{H}^+ + 2\text{e}^- \rightarrow$ succinate²⁻	0.031
$2\text{H}^+ + 2\text{e}^- \rightarrow \text{H}_2$ (at standard conditions, pH 0)	0.000
Crotonyl-CoA + $2\text{H}^+ + 2\text{e}^- \rightarrow$ butyryl-CoA	-0.015
Oxaloacetate ²⁻ + $2\text{H}^+ + 2\text{e}^- \rightarrow$ malate ²⁻	-0.166
Pyruvate + $2\text{H}^+ + 2\text{e}^- \rightarrow$ lactate	-0.185
Acetaldehyde + $2\text{H}^+ + 2\text{e}^- \rightarrow$ ethanol	-0.197
FAD + $2\text{H}^+ + 2\text{e}^- \rightarrow$ FADH₂	-0.219*
Glutathione + $2\text{H}^+ + 2\text{e}^- \rightarrow$ 2 reduced glutathione	-0.23
$\text{S} + 2\text{H}^+ + 2\text{e}^- \rightarrow \text{H}_2\text{S}$	-0.243
Lipoic acid + $2\text{H}^+ + 2\text{e}^- \rightarrow$ dihydrolipoic acid	-0.29
NAD⁺ + $\text{H}^+ + 2\text{e}^- \rightarrow$ NADH	-0.320
NADP ⁺ + $\text{H}^+ + 2\text{e}^- \rightarrow$ NADPH	-0.324
Acetoacetate + $2\text{H}^+ + 2\text{e}^- \rightarrow$ β -hydroxybutyrate	-0.346
α -Ketoglutarate + $\text{CO}_2 + 2\text{H}^+ + 2\text{e}^- \rightarrow$ isocitrate	-0.38
$2\text{H}^+ + 2\text{e}^- \rightarrow \text{H}_2$ (at pH 7)	-0.414
Ferredoxin (Fe^{3+}) + $\text{e}^- \rightarrow$ ferredoxin (Fe^{2+})	-0.432

Data mostly from Loach, P.A. (1976) In *Handbook of Biochemistry and Molecular Biology*, 3rd edn (Fasman, G.D., ed.), Physical and Chemical Data, Vol. 1, pp. 122-130, CRC Press, Boca Raton, FL.

*This is the value for free FAD; FAD bound to a specific flavoprotein (for example succinate dehydrogenase) has a different E'° .

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(7) Hydration of fumarate

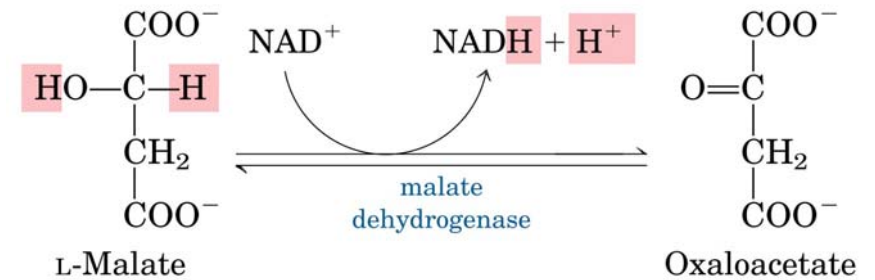


$$\Delta G'^{\circ} = -3.8 \text{ kJ/mol}$$

69

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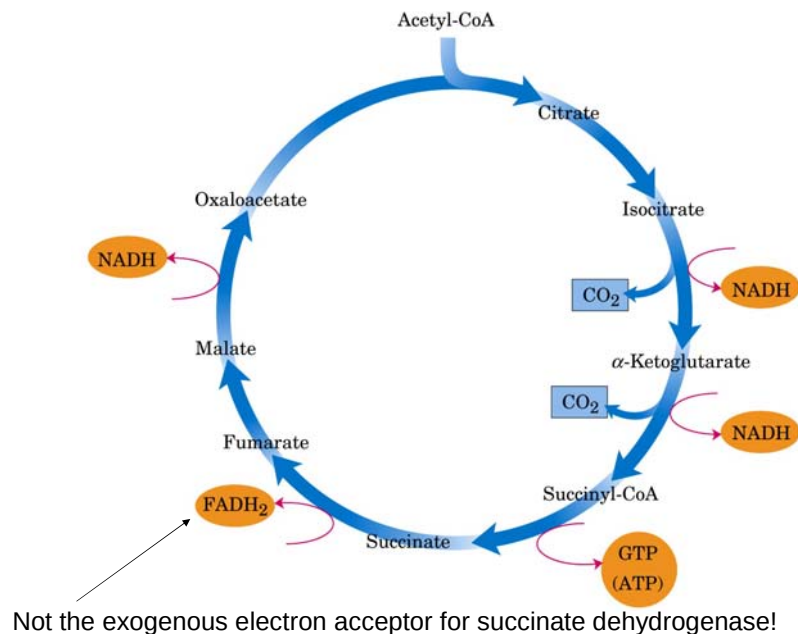
(8) Dehydrogenation of L-malate



$$\Delta G'^{\circ} = 29.7 \text{ kJ/mol}$$

70

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71

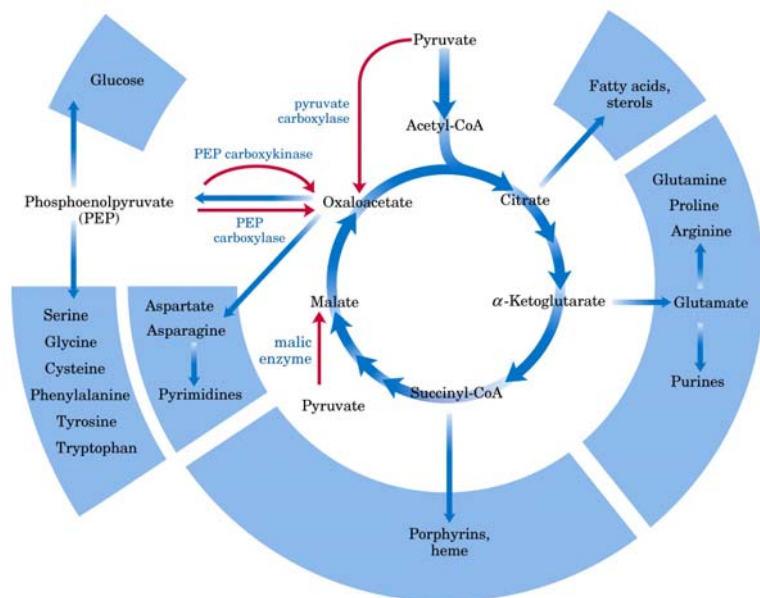
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Why a cyclic pathway?

- Feed in other than C2
- Take stuff out
- Avoid toxic C1 compounds

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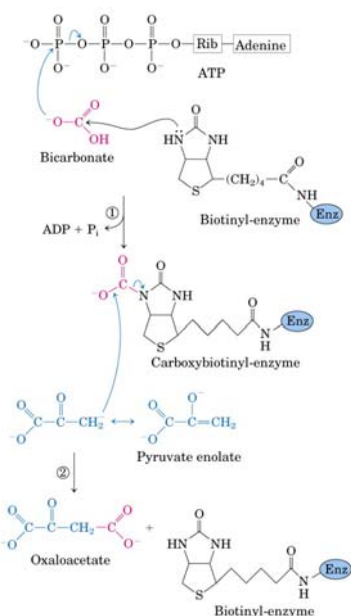
Replenishment of citric acid cycle intermediates: anaplerotic reactions



table 16-2

Anaplerotic Reactions

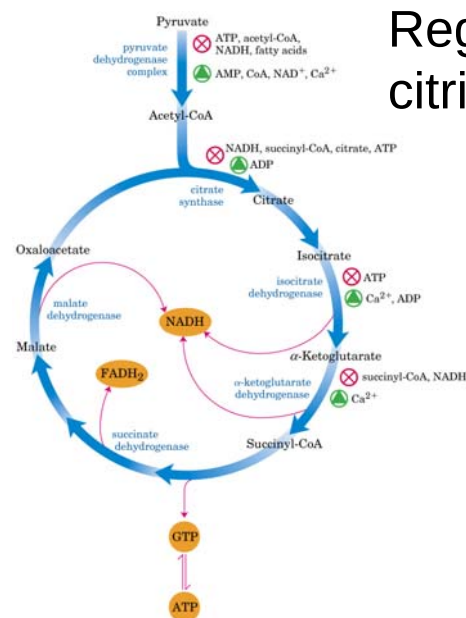
Reaction	Tissue(s)/organism(s)
Pyruvate + HCO_3^- + ATP $\xrightleftharpoons{\text{pyruvate carboxylase}}$ oxaloacetate + ADP + P_i	Liver, kidney
Phosphoenolpyruvate + CO_2 + GDP $\xrightleftharpoons{\text{PEP carboxykinase}}$ oxaloacetate + GTP	Heart, skeletal muscle
Phosphoenolpyruvate + HCO_3^- $\xrightleftharpoons{\text{PEP carboxylase}}$ oxaloacetate + P_i	Higher plants, yeast, bacteria
Pyruvate + HCO_3^- + NAD(P)H $\xrightleftharpoons{\text{malic enzyme}}$ malate + NAD(P) $^+$	Widely distributed in eukaryotes and prokaryotes



Biotin: the cofactor for many organic substrate carboxylation reaction

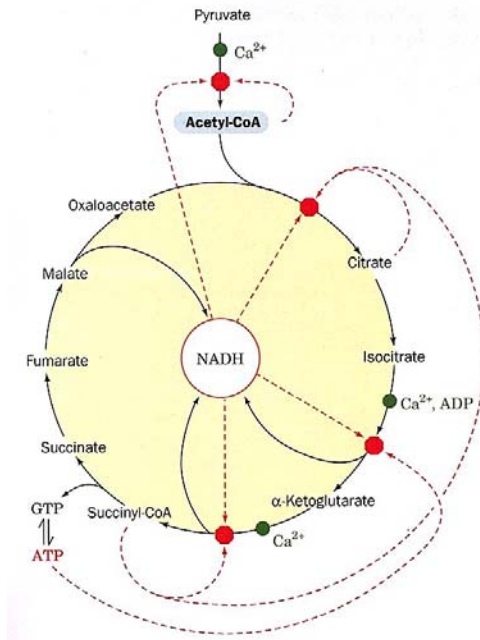
Biotin activates abundant bicarbonate to electrophilic CO_2

Regulation of the citric acid cycle

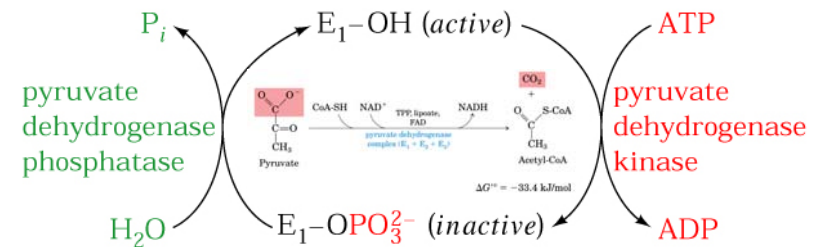


Four key sites:

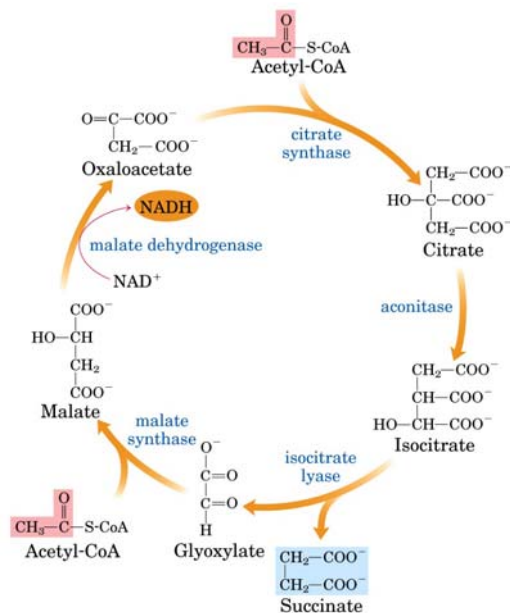
- (1) Pyr → Ac-CoA
- (2) Ac-CoA → citrate
- (3) Isocitrate → α-KG
- (4) α-KG → Succ-CoA



Regulation of PDH



- *Allosteric product inhibition*: Ac-CoA and NADH ⊗
- *Allosteric inhibition* by ATP ⊗
- *Allosteric activation* by AMP, CoA, NAD⁺ ⊕
- *On/off regulation* by PDH phosphatase and PDH kinase
 - How do you suppose these enzymes are regulated?



The Glyoxylate cycle: let's save this for next semester. Basically, it's the means by which plants and some bacteria carry out anaplerotic reactions from Ac-CoA. The main objective is to synthesize C3 and C4 units from Ac-CoA. Mammals cannot do this.