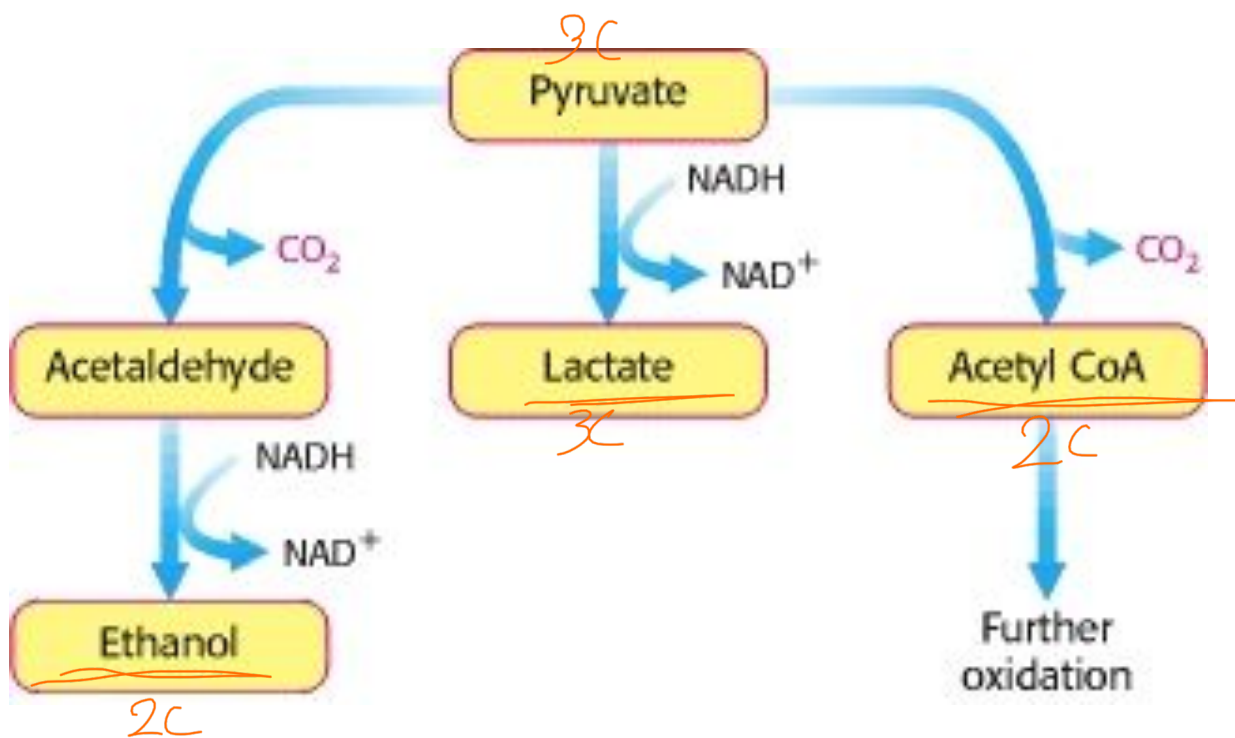
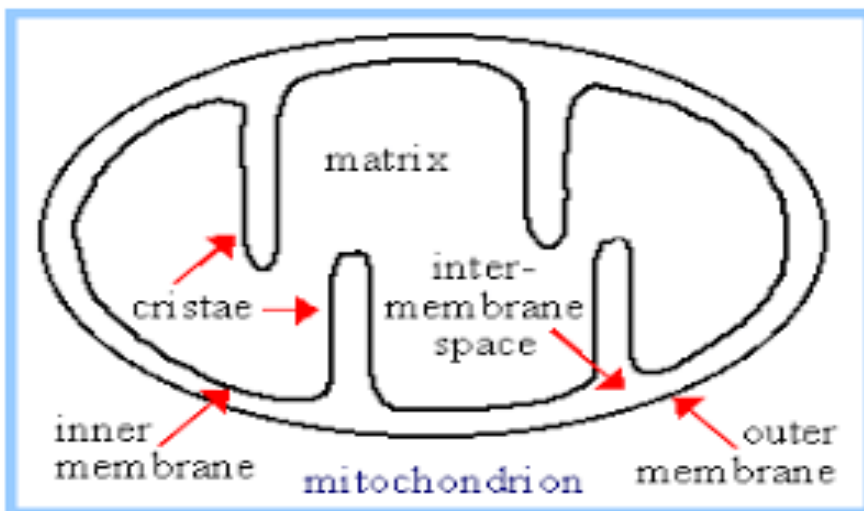


# Pyruvate metabolism

## Chapter 3



- Before entering the citric acid cycle, the carbon skeletons of sugars and fatty acids are degraded to the acetyl group of acetyl-CoA → ??
- pyruvate dehydrogenase (PDH) complex
- located in the mitochondria of eukaryotic cells



Inner membrane foldings called Cristae contain ETC

The matrix contains Pyruvate dehydrogenase enzymes, and enzymes of the Krebs cycle

- Five cofactors, four derived from vitamins, participate in the reaction mechanism
- combination of covalent modification and allosteric regulation ↳ under go conformational change
- PDH complex consists of multiple copies of three enzymes:
  - Pyruvate dehydrogenase (E1)
  - Dihydrolipoamide transacetylase (E2)
  - Dihydrolipoamide dehydrogenase (E3)
- Also part of the complex are two regulatory enzymes
  - A protein kinase
  - A phosphoprotein phosphatase

## The pyruvate dehydrogenase reaction involves multiple coenzymes

Coenzyme	Subunit	Role in catalysis
thiamine pyrophosphate	E <sub>1</sub>	provides a carbanion for nucleophilic attack on the substrate
lipoamide	E <sub>2</sub>	transfers substrate to coenzyme A, retains hydrogen
flavin adenine dinucleotide (FAD)	E <sub>3</sub>	transfers H <sub>2</sub> from lipoamide to NAD <sup>+</sup>

Step 1: E1



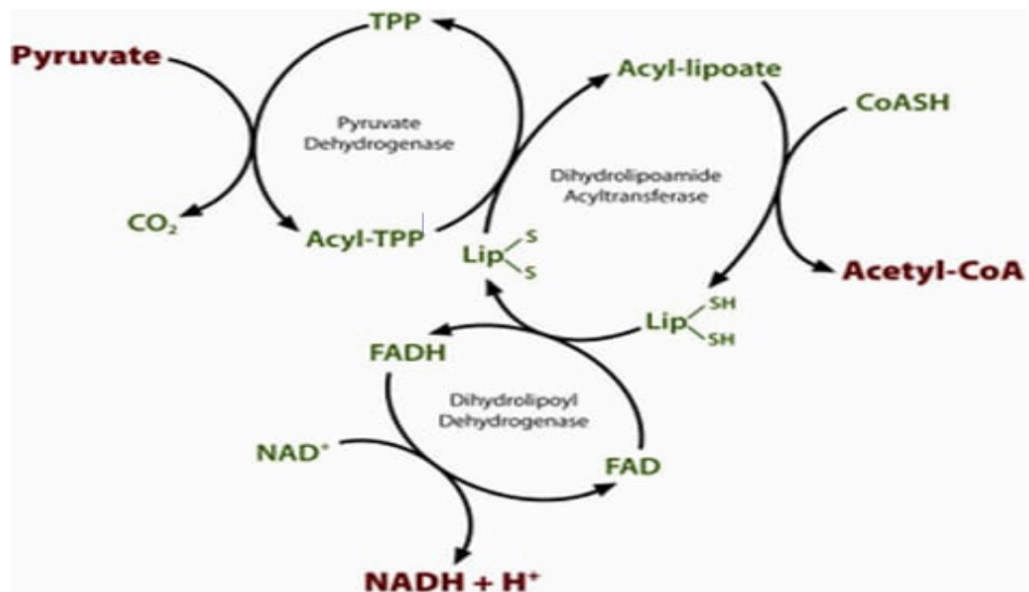
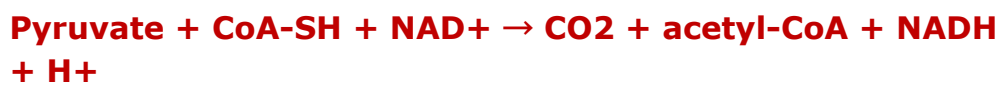
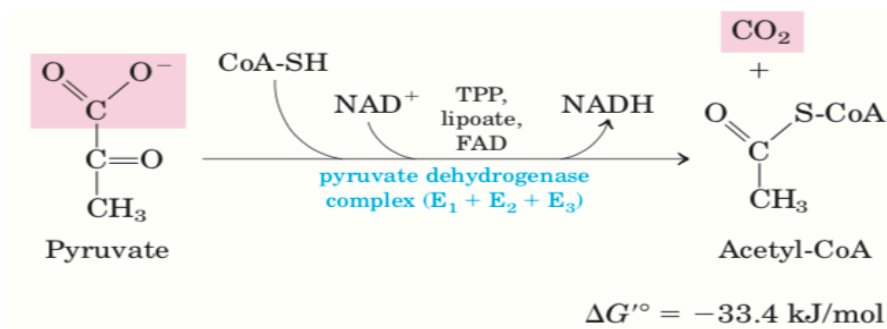
Step 2:



Step 3: E2



The overall reaction catalyzed by the pyruvate dehydrogenase complex is an **oxidative decarboxylation**, an **irreversible** oxidation process in which the **carboxyl group is removed from pyruvate as a molecule of CO<sub>2</sub>**



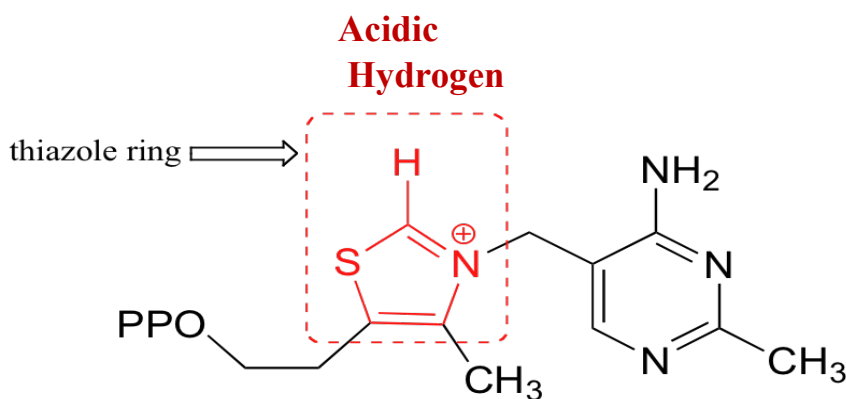


# PDH Complex

## five different coenzymes or prosthetic groups

- Thiamine pyrophosphate (TPP) □ Thiamine (B1)
- Flavin adenine dinucleotide (FAD) □ Riboflavin (B2)
- Coenzyme A (CoA, sometimes de- noted CoA-SH, to emphasize the role of the OSH group □ pantothenate (B5)
- Nicotinamide adenine dinucleotide (NAD) □ Niacin (B3)
- Lipoamide

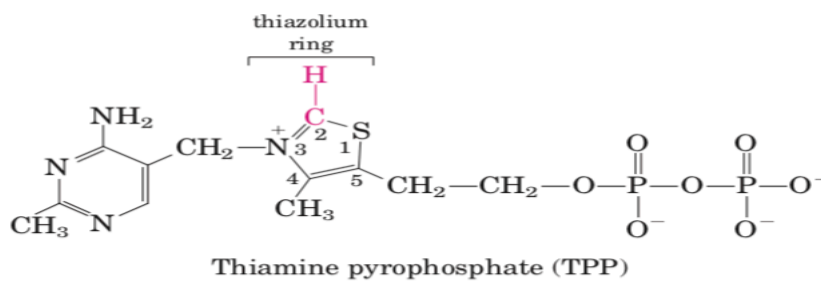
## Thiamine pyrophosphate



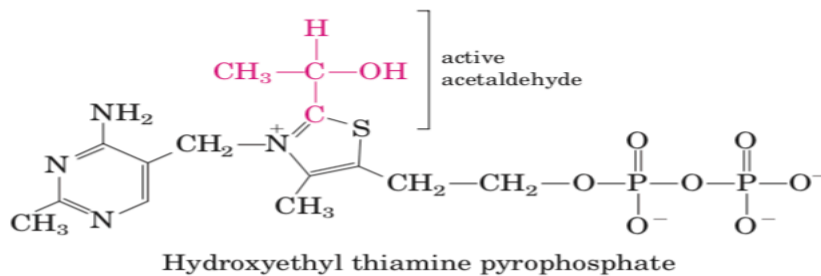
thiamine diphosphate (TPP)

The carbanion then acts as a strong nucleophile **carbanion** Initiates a nucleophilic attack on the carbonyl **carbon** of pyruvate

1. The keto C of pyruvate reacts with the carbanion of TPP on E1 to yield an addition compound. The electron-pulling (+) charged N of the thiazole ring promotes CO<sub>2</sub> loss. Hydroxyethyl-TPP remains.
2. The hydroxyethyl carbanion on TPP of E1 reacts with the disulfide of lipoamide on E2. What was the keto C of pyruvate is oxidized to a carboxylic acid, as the lipoamide disulfide is reduced to a dithiol
3. Acetate is transferred from the thiol of lipoamide to the thiol of coenzyme A, yielding acetyl CoA.
4. Dihydrolipoamide is reoxidized to the disulfide as 2 e<sup>-</sup> + 2 H<sup>+</sup> are transferred to FAD.
5. The resulting FADH<sub>2</sub> is reoxidized by electron transfer to NAD<sup>+</sup>, to yield NADH + H<sup>+</sup>.



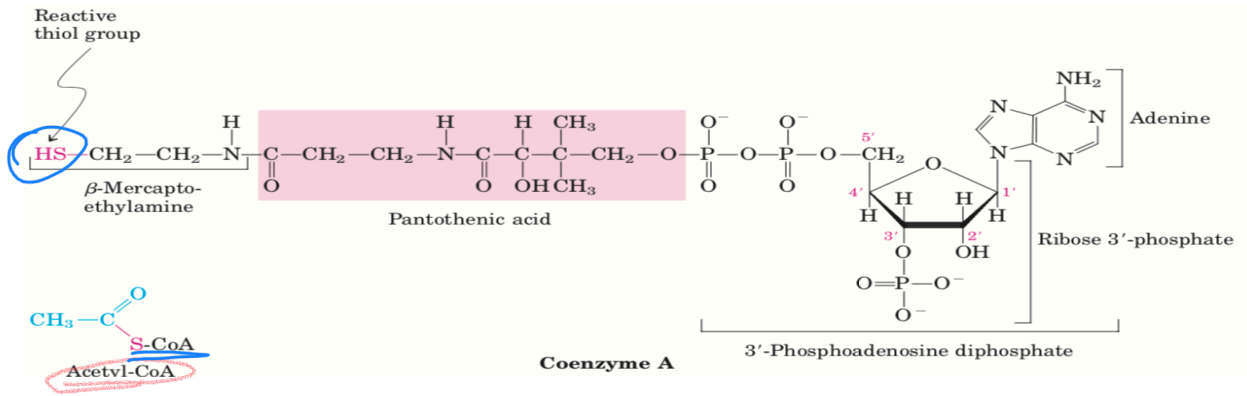
(a)



# Coenzyme A

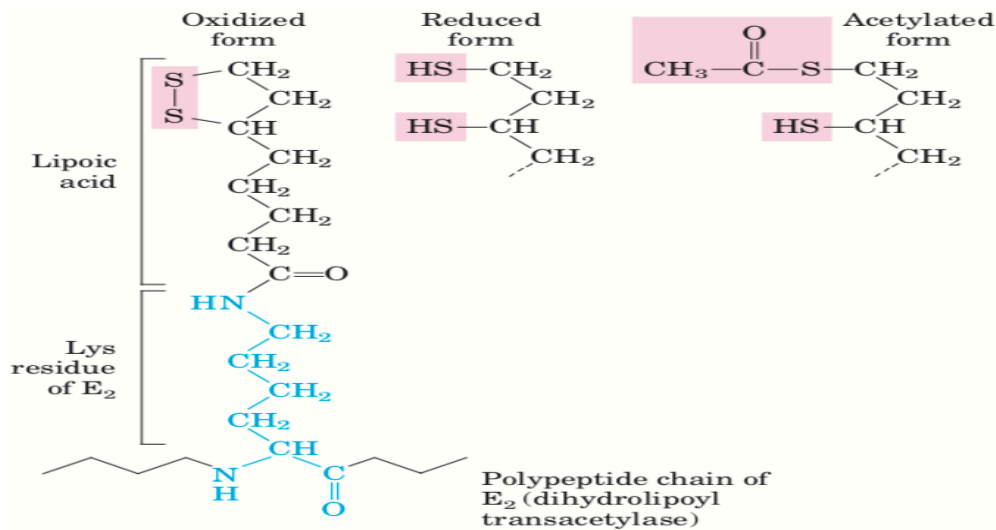
- Coenzyme A -has a reactive thiol (OSH)
- Role of CoA as an acyl carrier- Acyl groups are covalently linked to the thiol group, forming thioesters

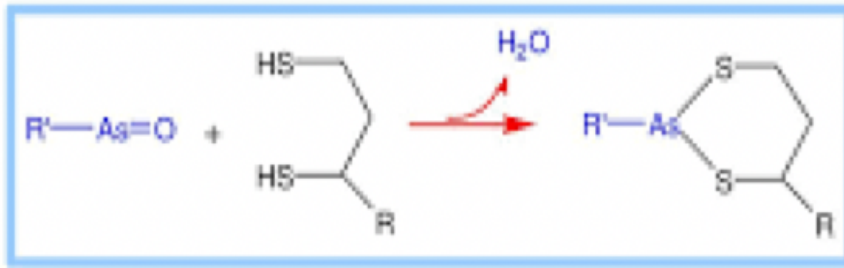
16.1 Production of Acetyl-CoA (Activated Acetate)



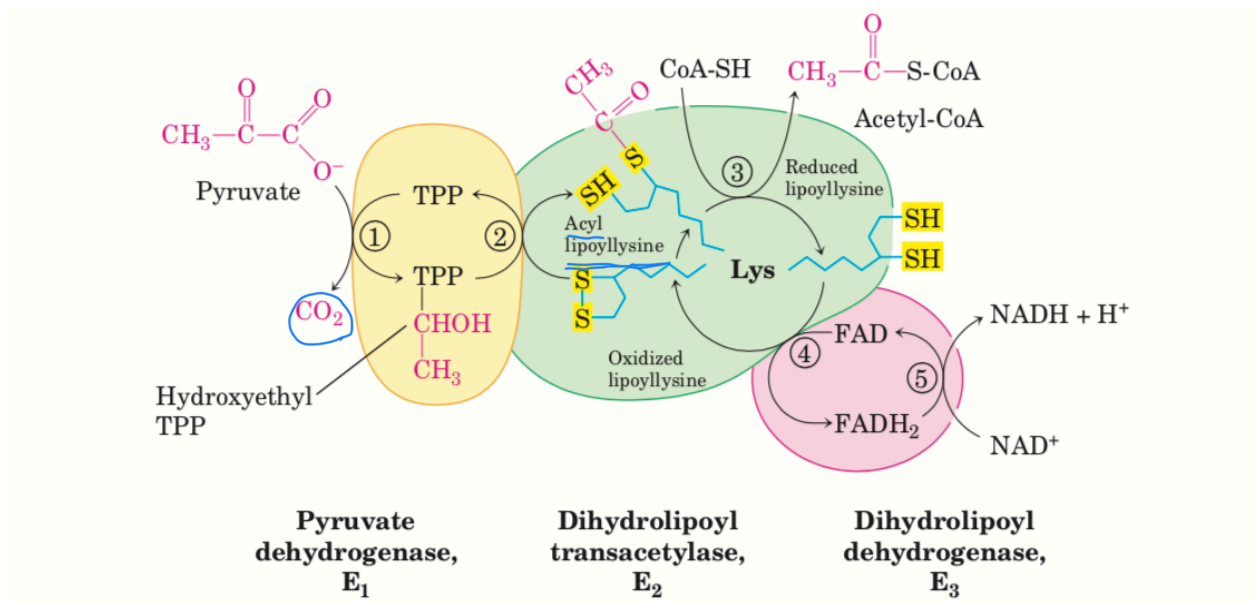
# lipoate

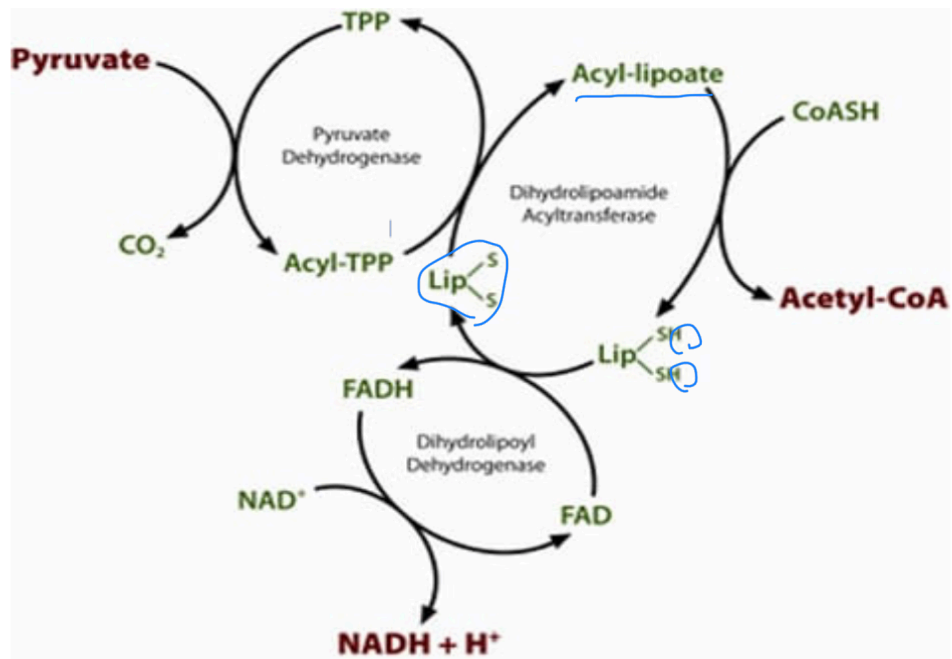
lipoate has two thiol groups that can undergo reversible oxidation to a disulfide bond ( $\text{--S--S--}$ )





Organic **arsenicals** are potent inhibitors of lipoamide-containing enzymes such as Pyruvate Dehydrogenase. These highly toxic compounds react with **dithiols** such as the functional group of lipoate.





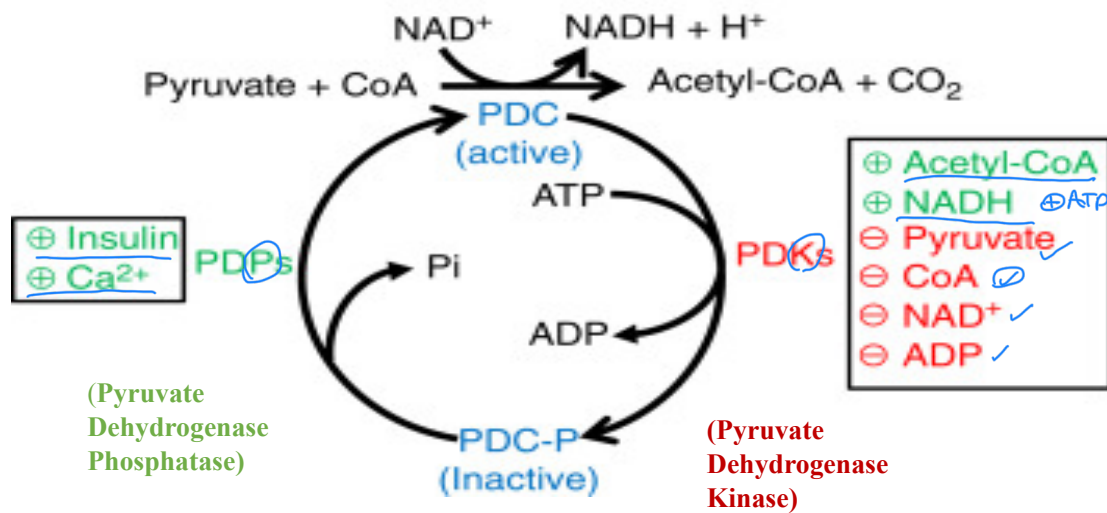
- FAD is a prosthetic group, permanently part of E3. Reaction:
- $\text{FAD} + 2 \text{e}^- + 2 \text{H}^+ \rightarrow \text{FADH}_2$

# Regulation of Pyruvate Dehydrogenase Complex:

## **Product inhibition by NADH & acetyl CoA:**

NADH competes with  $\text{NAD}^+$  for binding to E3. Acetyl CoA competes with CoA for binding to E2.

- Specific regulatory Kinases & Phosphatases associated with Pyruvate Dehydrogenase in the mitochondrial matrix:
  - **Pyruvate Dehydrogenase Kinases** catalyze phosphorylation of serine residues of E1, inhibiting the complex.
  - **Pyruvate Dehydrogenase Phosphatases** reverse this inhibition.



<https://www.sciencedirect.com/topics/agricultural-and-biological-sciences/pyruvate-dehydrogenase-phosphatase>

## Regulation of PDH activity

- PDH activity is inhibited by reversible phosphorylation of the E1
- The phosphorylation  $\square$  PDH kinase
- Dephosphorylation to restore PDH activity phosphatases

PDH Kinase ( a special regulatory enzyme which is part of the PDH multienzyme complex)

The PDH kinase enzyme is activated by NADH and acetyl-CoA and inhibited by ADP, NAD<sup>+</sup> and by free coenzyme A

# During starvation

- Pyruvate Dehydrogenase Kinase increases in amount in most tissues, including skeletal muscle, via increased gene transcription.
  - Under the same conditions, the amount of Pyruvate Dehydrogenase Phosphatase decreases
  - The resulting inhibition of Pyruvate Dehydrogenase prevents muscle and other tissues from catabolizing glucose & gluconeogenesis precursors.
  - Metabolism shifts toward fat utilization.
- \* Available glucose is spared for use by the brain.

in starvation  
CACP →  
↓ Acetyl  
CoA  
↓ NADH