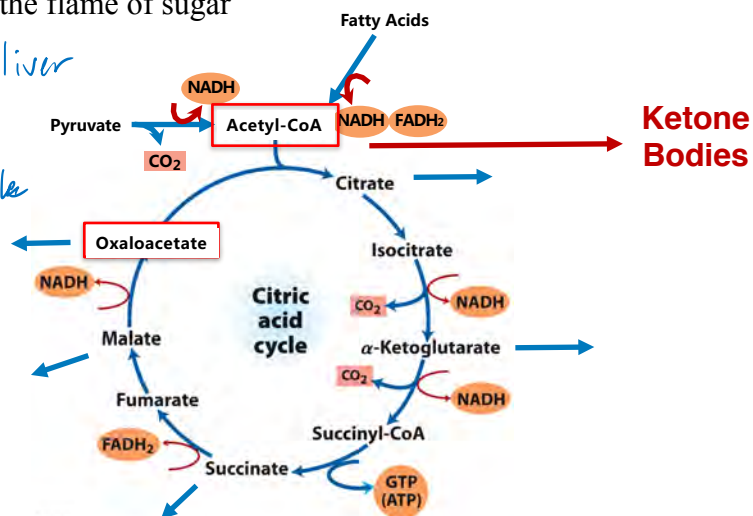


Ketone Bodies

Fatty Acid Degradation Ketone Bodies

“Fat burns in the flame of sugar”

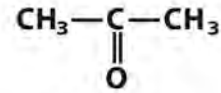
Made by liver
Used by heart, muscle
adapted brain



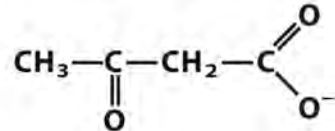
Fatty Acid Degradation

- Entry of acetyl-CoA into citric acid cycle requires **oxaloacetate**.
- When oxaloacetate is depleted, acetyl-CoA is converted into **ketone bodies**.
 - frees coenzyme A for continued β oxidation
- Three forms of ketone bodies can leave the liver: acetone, acetoacetate, and β -hydroxybutyrate.
- Therefore, the **anabolism** of Ketone Bodies is connect to **catabolism** of fatty acids and sugars
- For now, we will discuss Ketone Body **catabolism**

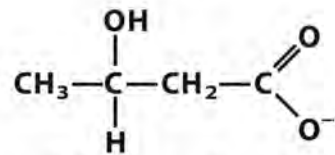
Ketone Bodies



Acetone



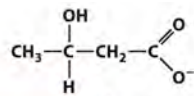
Acetoacetate



D- β -Hydroxybutyrate

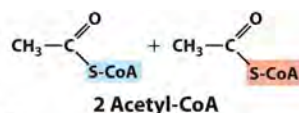
Fatty Acid Degradation

Ketone Bodies as Fuel



D- β -Hydroxybutyrate

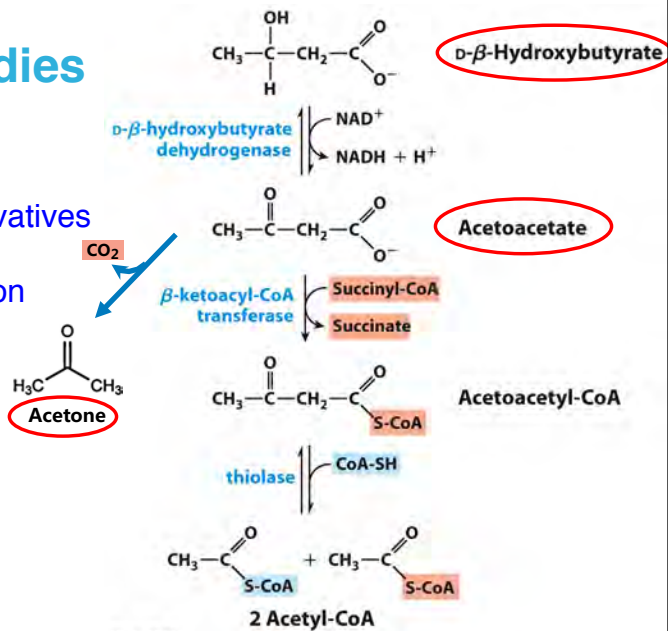
1. Make CoA derivatives
2. Oxidize alcohol
3. Thiolase reaction



Fatty Acid Degradation

Ketone Bodies as Fuel

1. Make CoA derivatives
2. Oxidize alcohol
3. Thiolase reaction



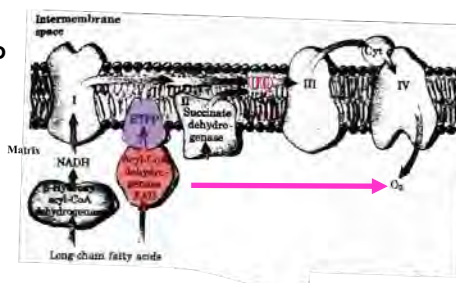
Fatty Acid Oxidation in Other Organelles

Fatty Acid Degradation

β Oxidation in Plants Occurs Mainly in Peroxisomes

- **Mitochondrial** acyl-CoA dehydrogenase passes electrons into **respiratory chain** via electron-transferring flavoprotein (ETFP).

– energy captured as ATP



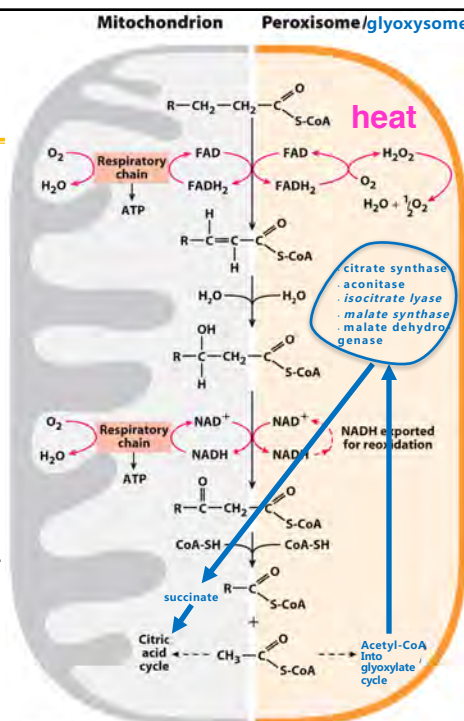
- **Peroxisomal/glyoxysomal** acyl-CoA dehydrogenase passes electrons directly to **molecular oxygen** and H_2O_2 .

– energy released as **heat**

– hydrogen peroxide eliminated by **catalase**

Fatty Acid Degradation

- Acetyl-CoA released from peroxisomal β oxidation is exported to cytosol and then imported to mitochondria.
- A peroxisome is also a **glyoxysome** when enzymes for **glyoxylate cycle** are present (e.g., germinating seeds).
- Acetyl-CoA released from peroxisomal β oxidation can be used in **glyoxylate cycle**.
- The **glyoxylate cycle** is found in most organisms, but not vertebrate animals.
- The **glyoxylate cycle** has an **anabolic function** to synthesize larger molecules from acetyl-CoA.

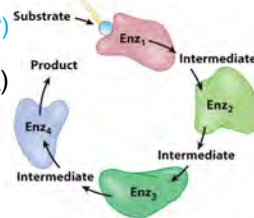


Fatty Acid Degradation

Fatty Acid Oxidation is Performed by a Single TriFunctional Protein (TFP)

- Hetero-octamer ($\alpha_4\beta_4$)
 - four α subunits
 - enoyl-CoA hydratase activity
 - β -hydroxyacyl-CoA dehydrogenase activity
 - responsible for binding to membrane
 - four β subunits
 - long-chain thiolase activity
- May allow substrate channeling between enzymes
- Associated with inner-mitochondrial membrane
- Processes fatty acid chains with 12 or more carbons
- Shorter chains processed by soluble enzymes in the matrix

(a) Gram-positive bacteria and mitochondrial short-chain-specific system

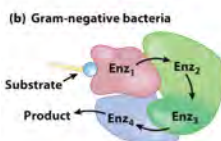
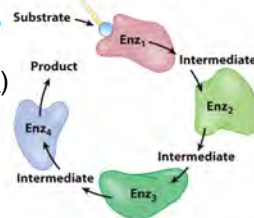


Fatty Acid Degradation

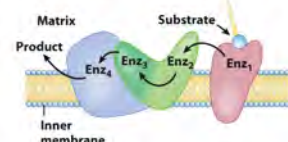
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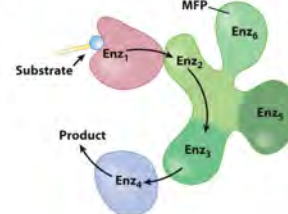
(a) Gram-positive bacteria and mitochondrial short-chain-specific system



(c) Mitochondrial very-long-chain-specific system



(d) Peroxisomal and glyoxysomal systems



Fatty Acid Degradation: Summary

We learned that:

- fats are an important **energy source** in animals
- two-carbon units in fatty acids are oxidized in a four-step **β oxidation** process into acetyl-CoA
- in the process, a lot of **NADH** and **FADH₂** forms; these can yield a lot of ATP in the electron-transport chain
- Mono- and poly-unsaturated fatty acids require additional enzymes and lose an FADH₂ for every double bond and cost an NADPH for every pair.
- acetyl-CoA formed in the liver can be either **oxidized via the citric acid cycle** or **converted to ketone bodies** that serve as fuels for other tissues
- Other organelles can perform fatty-acid oxidation; during peroxisomal oxidation, fats can be oxidized to generate heat