

BI/CH 422/622

OUTLINE:

Lipid Degradation (Catabolism)

FOUR stages in the catabolism of lipids:

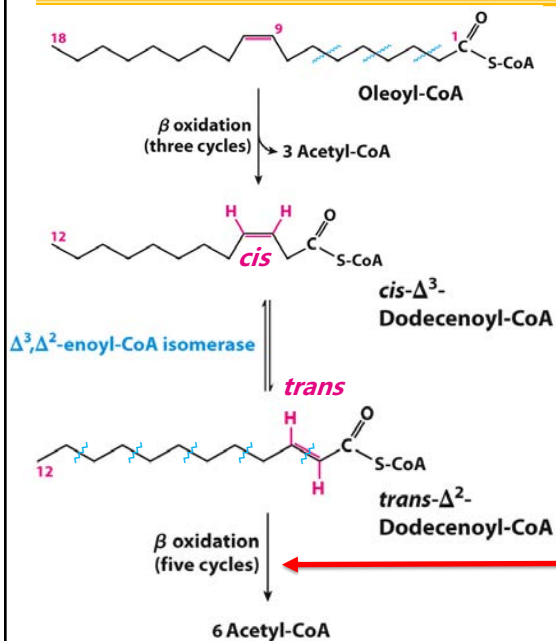
- Mobilization from tissues (mostly adipose)
 - hormone regulated
 - specific lipases
 - glycerol
- Activation of fatty acids
 - Fatty-acyl CoA Synthetase
- Transport into mitochondria
 - carnitine
- Oxidation
 - rationale
 - Saturated FA
 - 4 steps
 - dehydrogenation
 - hydration
 - oxidation
 - thiolase
 - energetics
 - Unsaturated FA
 - Odd-chain FA
 - Ketone Bodies
- Other organelles

OUTLINE:

Protein Degradation (Catabolism)

- Digestion
- Inside of cells
 - Protein turnover
 - Ubiquitin
 - Proteasome
- Amino-Acid Degradation

Fatty Acid Degradation



Monounsaturated Fatty Acids

During first of five remaining cycles, acyl-CoA dehydrogenase step is skipped, resulting in 1 fewer FADH_2 .

Fatty Acid Degradation

Energy from Fatty Acid Catabolism

TABLE 17-1a Yield of ATP during Oxidation of One Molecule of Oleoyl-CoA to CO₂ and H₂O

| Enzyme catalyzing the oxidation step | Number of NADH or FADH ₂ formed | Number of ATP ultimately formed ^a |
|--------------------------------------|--|--|
| β Oxidation | | |
| Acyl-CoA dehydrogenase | 7 FADH ₂ | 10.5 |
| β-Hydroxyacyl-CoA dehydrogenase | 8 NADH | 20 |
| Citric acid cycle | | 35 → 87.5 ATP |
| Isocitrate dehydrogenase | 9 NADH | 22.5 |
| α-Ketoglutarate dehydrogenase | 9 NADH | 22.5 |
| Succinyl-CoA synthetase | 16 → 24 ATP | 9 ^b |
| Succinate dehydrogenase | 9 FADH ₂ | 13.5 |
| Malate dehydrogenase | 9 NADH | 22.5 |
| Total | | 120.5 - 2 = 118.5* |

^aThese calculations assume that mitochondrial oxidative phosphorylation produces 1.5 ATP per FADH₂ oxidized and 2.5 ATP per NADH oxidized.

^bGTP produced directly in this step yields ATP in the reaction catalyzed by nucleoside diphosphate kinase (p. 516).

*These 2 "ATP" equivalents were expended in the activation by Fatty acyl-CoA synthetase.

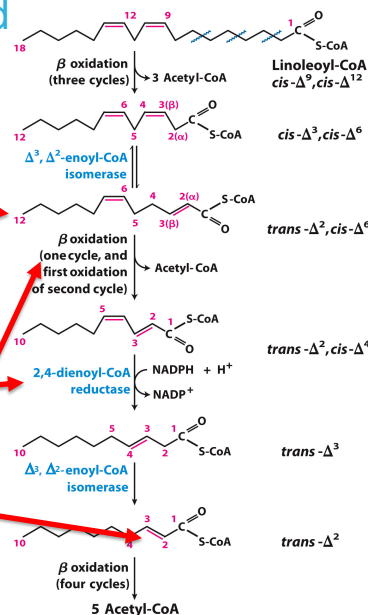
Fatty Acid Degradation

Oxidation of Polyunsaturated Fatty Acids

Results in 1 fewer FADH₂ after isomerization, as the acyl-CoA dehydrogenase step is skipped and goes right to the hydratase.

NADPH reduces the 2 conjugated unsaturated bonds to one trans-double bond between them; but it's still at an odd carbon

This bond already made with FADH₂ already generated before reductase and isomerization.



Fatty Acid Degradation

Energy from Fatty Acid Catabolism

TABLE 17-1a Yield of ATP during Oxidation of One Molecule of Linoleoyl-CoA to CO₂ and H₂O

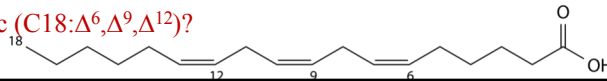
| Enzyme catalyzing the oxidation step | Number of NADH or FADH ₂ formed | Number of ATP ultimately formed ^a |
|---|--|--|
| β Oxidation | | |
| Acyl-CoA dehydrogenase | 7 FADH ₂ | 10.5 |
| β-Hydroxyacyl-CoA dehydrogenase (8-1=7; 1 used for reductase) | 7 NADH | 17.5 |
| Citric acid cycle | | |
| Isocitrate dehydrogenase | 9 NADH | 22.5 |
| α-Ketoglutarate dehydrogenase | 9 NADH | 22.5 |
| Succinyl-CoA synthetase | | 9 ^b |
| Succinate dehydrogenase | 9 FADH ₂ | 13.5 |
| Malate dehydrogenase | 9 NADH | 22.5 |
| Total | | 118 - 2 = 116* |

^aThese calculations assume that mitochondrial oxidative phosphorylation produces 1.5 ATP per FADH₂ oxidized and 2.5 ATP per NADH oxidized.

^bGTP produced directly in this step yields ATP in the reaction catalyzed by nucleoside diphosphate kinase (p. 516).

*These 2 "ATP" equivalents were expended in the activation by Fatty acyl-CoA synthetase.

What would it be for γ-linolenic (C18:Δ⁶,Δ⁹,Δ¹²)?



Degradation of Odd-Chain Fatty Acids

Fatty Acid Degradation

Oxidation of Odd-Numbered Fatty Acids

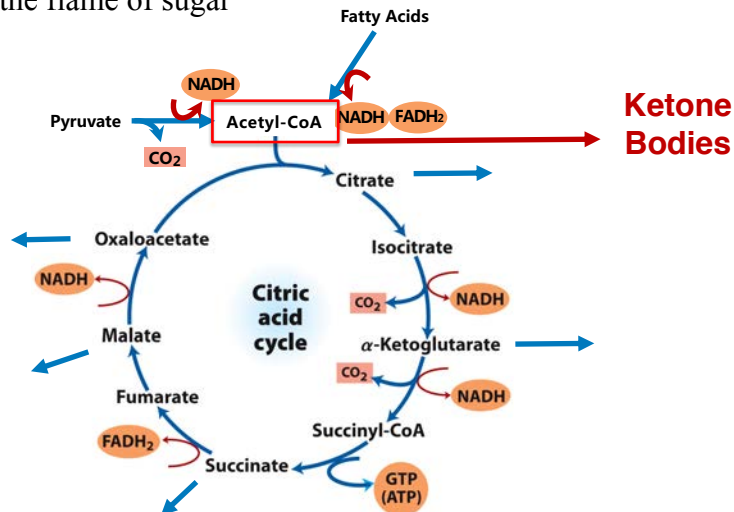
- Most dietary fatty acids are even-numbered.
- Many plants and some marine organisms also synthesize **odd-numbered fatty acids**.
- The metabolism of oxidation of odd-chain fatty acids CONVERGES with that of some amino acids
- Details will be discussed with amino acids

Ketone Bodies

Ketone Body Degradation

What are Ketone Bodies?

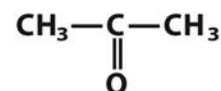
“Fat burns in the flame of sugar”



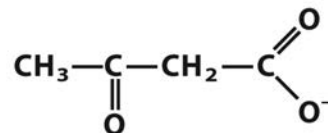
Ketone Body Degradation

- 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**

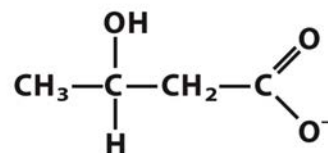
Ketone Bodies



Acetone



Acetoacetate

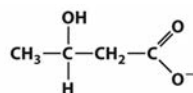


D- β -Hydroxybutyrate

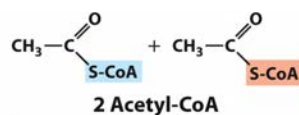
Ketone Body Degradation

Ketone Bodies as Fuel

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



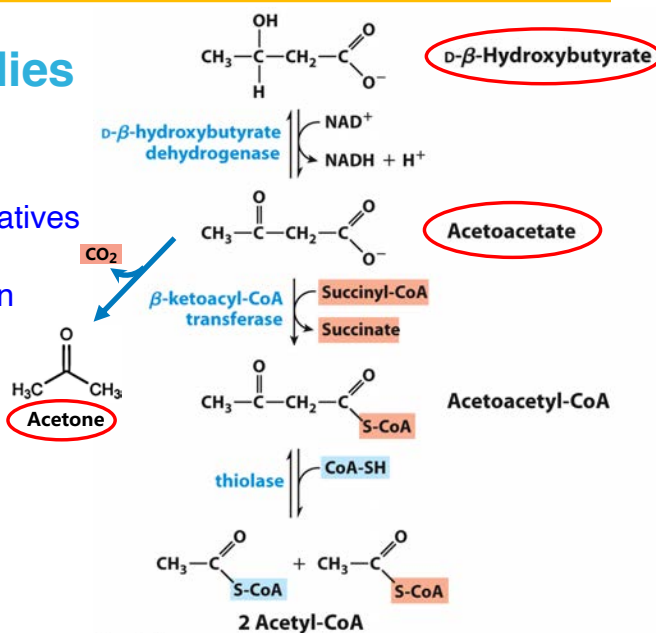
D-β-Hydroxybutyrate



Ketone Body Degradation

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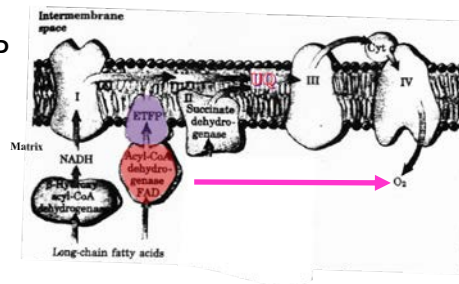


Fatty Acid Oxidation in Other Organelles

Fatty Acid Degradation

β Oxidation in Plants Occurs Mainly in Peroxisomes

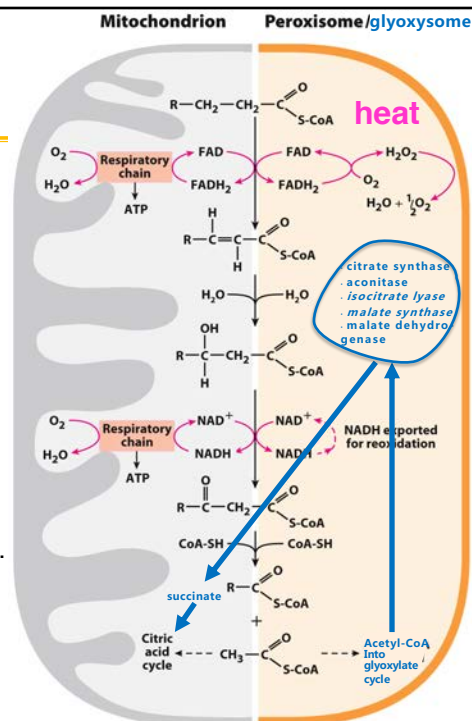
- **Mitochondrial** acyl-CoA dehydrogenase passes electrons **into respiratory chain** via electron-transferring flavoprotein (ETFP).
 - energy captured as ATP



- **Peroxisomal/glyoxysomal** acyl-CoA oxidase 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.

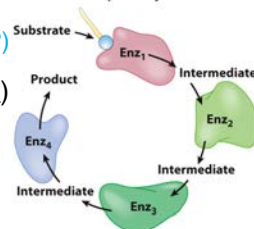


Fatty Acid Degradation

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

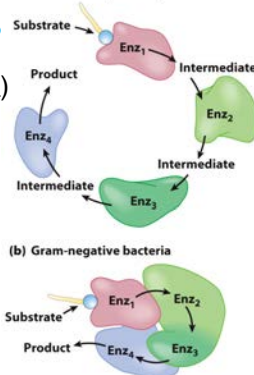


Fatty Acid Degradation

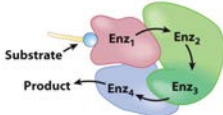
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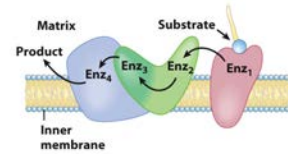
(a) Gram-positive bacteria and mitochondrial short-chain-specific system



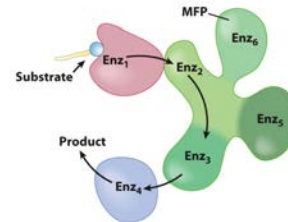
(b) Gram-negative bacteria



(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 $FADH_2$ 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 $FADH_2$ for every double bond at an odd carbon, 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