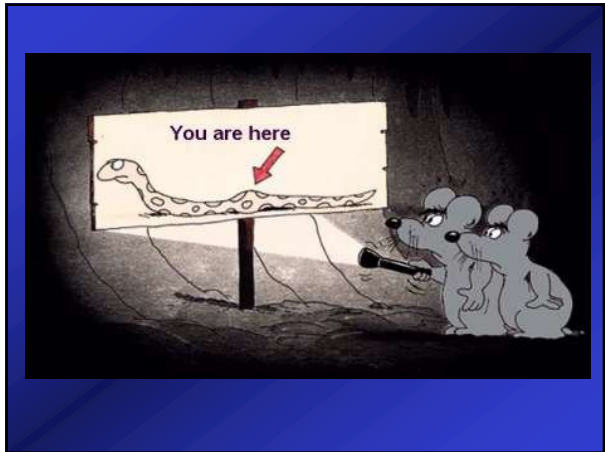


Fatty Acid Metabolism



Oxidation - even or odd

Fatty acid fed	Breakdown product	Excretion product
$\text{HOOC}-\text{CH}_2-(\text{CH}_2)_n-\text{CH}_2-\text{COOH}$ <p>Odd-chain fatty acid</p>	$\text{HOOC}-\text{CH}_2-\text{COOH}$ <p>Benzoic acid</p>	$\text{HOOC}-\text{CH}_2-\text{NH}-\text{CH}_2-\text{COOH}$ <p>Hippuric acid Glycine residue</p>
$\text{HOOC}-\text{CH}_2-\text{CH}_2-(\text{CH}_2)_n-\text{CH}_2-\text{COOH}$ <p>Even-chain fatty acid</p>	$\text{HOOC}-\text{CH}_2-\text{COOH}$ <p>Phenylacetic acid</p>	$\text{HOOC}-\text{CH}_2-\text{NH}-\text{CH}_2-\text{COOH}$ <p>Phenylacetic acid Glycine residue</p>

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Fatty Acid Oxidation (α , ω and β -oxidation)

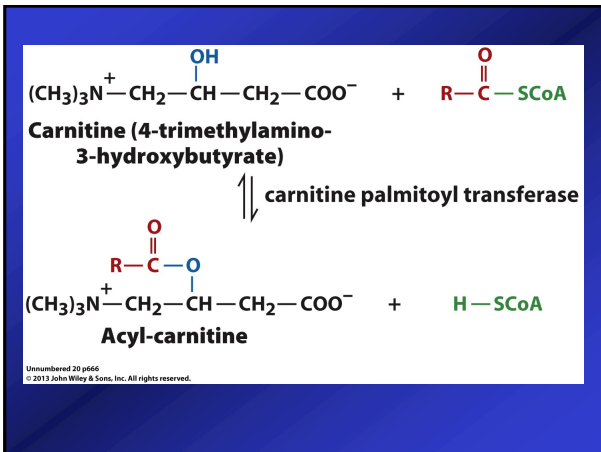
- Three phases to β oxidation
 - activation with Coenzyme A
 - transport into the mitochondria
 - oxidation of 2 carbon units

Activation acyl CoA synthetase aka thiokinase

- uses ATP (only energy requiring step)
- 2 part step forming acyl CoA and PPi (drives reaction)
- mixed anhydride forms attack site for CoA

$$\text{CH}_3-(\text{CH}_2)_n-\text{COO}^- + \text{ATP} + \text{HS-CoA} \rightarrow \text{acyl-CoA} + \text{AMP} + \text{PP}_i$$

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Transport

- FA greater than 12 C long need to be transported into mito
- three step reaction 2 transferases and a translocase
- carnitine replaces CoA
- has nearly the same E of hydrolysis and acylCoA
- little change of energy during translocation
- cell keeps separate pools of CoA

β-oxidation of fatty acids

occurs in the mitochondria (inner membrane)
 subtracts 2 carbons from COO- end of β carbon of FA

- four steps :
 1. oxidation -> acyl-CoA dehydrogenase
 - FAD linked via electron transferring protein
 - feeds to site 2 in ETS
 - double bond formed between C 2 & 3
 - acyl-CoA dehydrogenase isozymes

β-oxidation of fatty acids

occurs in the mitochondria (inner membrane)
 subtracts 2 carbons from COO- end of β carbon of FA

- four steps :
 2. hydration -> enoyl CoA hydratase
 - puts water across Δ2 carbon

β-oxidation of fatty acids

occurs in the mitochondria (inner membrane)
 subtracts 2 carbons from COO- end of β carbon of FA

- four steps :
 3. oxidation -> β hydroxyacyl CoA dehydrogenase
 - NADH₂ linked oxidation of OH from last step
 - NADH₂ free to be oxidized at site 1 of ETS
 - specific for the L isomer

β-oxidation of fatty acids

occurs in the mitochondria (inner membrane)
 subtracts 2 carbons from COO- end of β carbon of FA

- four steps :
 4. thiolase -> β keto thiolase
 - similar reaction to hydrolysis uses SH group of CoA
 - leads to new acylCoA (minus 2 carbons)

steps 1 thru 3 are similar to part of the TCA
 (succinate->fumarate->malate->oxalacetate)
 Repeat cycle until two or 3 Carbons (1 acyl CoA) are left

Energetics of β oxidation

palmitate (C₁₆) - CoA + 7 CoASH + 7 FAD + 7 NAD⁺ + 7 H₂O
 -> 8 acetyl CoA + 7 FADH₂ + 7 NADH

The net yield in ATP currency is:

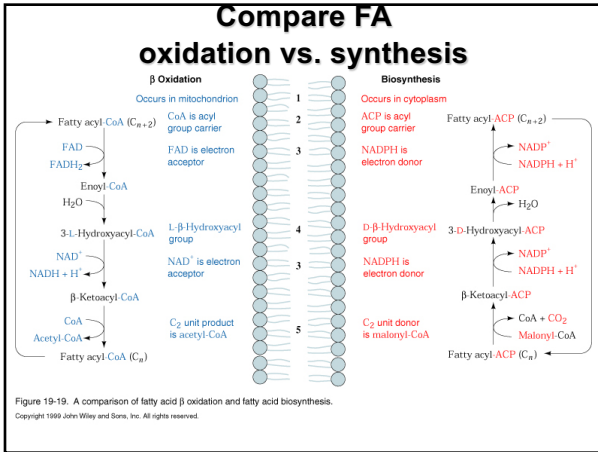
7 FADH ₂ - 14 ATP	8 Acyl-CoA	8 NADH = 72 ATP
7 NADH - 21 ATP		8 FADH ₂ = 16 ATP
		8 GTP = 8 ATP

with 100% efficiency - 131 ATP
 activation costs of 1 fatty acid - 2 ATP (7)
 total ATPs from palmitate 107 to 129

Long Chain FA shortened in peroxisomes -> O₂ is the e- acceptor

The double bond problem - remember 2nd step of β ox has a double bond at the C2

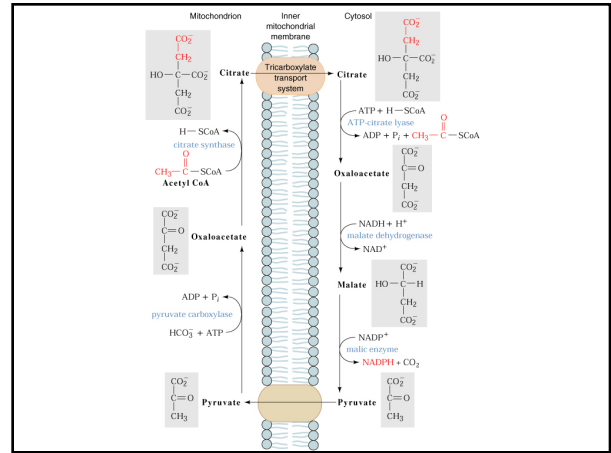
- Even # unsaturation - skip first step and result in one less FADH₂
- Odd # unsaturations undergo isomeraton when the double bond gets to delta 3.



Biosynthesis of fatty acids

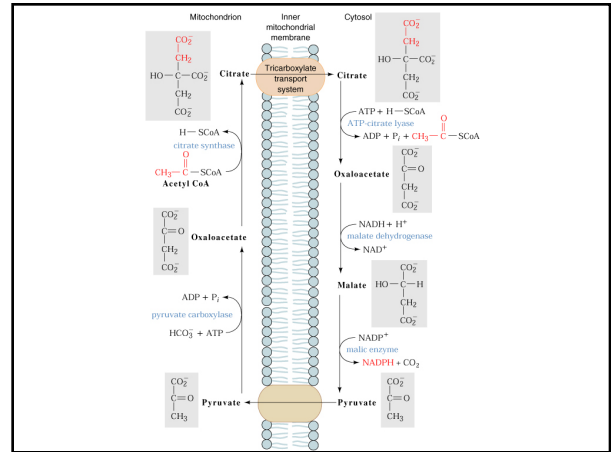
This pathway occurs in the cytosol. 2 carbon are added at a time to produce acetyl CoA. The precursors are from glucose and amino acids. This is distinct from β oxidation- it is a reductive process and uses NADPH. It takes place in the cytosol. A 3 carbon acid malonyl-CoA as the 2 carbon donor. The growing chain is attached to an acyl-carrier protein rather than Co-A. The enzymes are not shared between either

Acetyl CoA are produced by pyruvate carboxylase from both glycolytic and TCA intermediates. Acetyl CoA, which is generated by pyruvate dehydrogenase (PDH) in the mitochondrion cannot cross the inner mitochondrial membrane to reach the cytosol. Instead transport occurs by the tricarboxylate transport system.



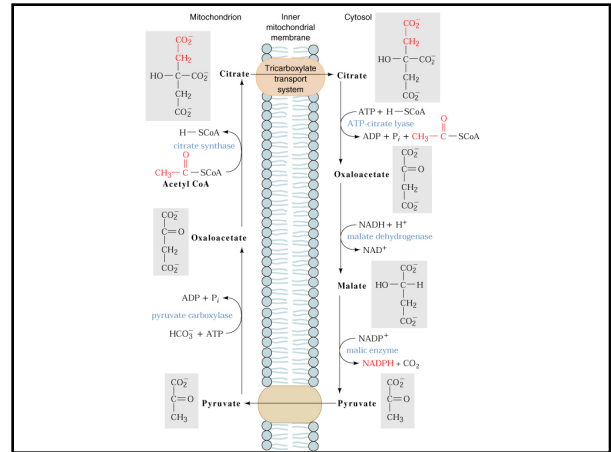
Acetyl CoA transport

Acetyl CoA produced in mito - transported out as citrate
glucose ->-> pyruvate -> OAA and Acetyl CoA
- pyruvate carboxylase and pyruvate dehydrogenase
OAA and Acetyl CoA -> Citrate (in mito)
citrate transfer into cytosol and cleaved to Acetyl CoA and OAA by ATP citrate lyase - cost of ATP to transport
- OAA shuffled back into mito via MDH, Malic enzyme (NADPH)
- pyruvate transport into mito converted to OAA
- produces both 1 NADPH /and Acetyl CoA



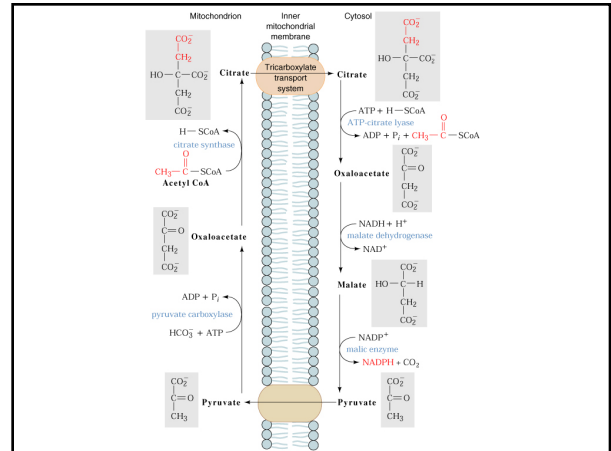
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Biosynthesis-

Two phases - malonyl CoA formation and FA complex

1) Production of malonyl CoA

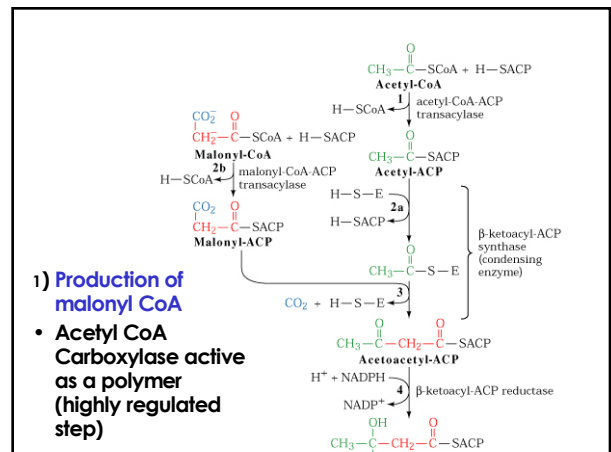
Acetyl CoA Carboxylase active as a polymer (highly regulated step)

1st committed step in fatty acid synthesis (FAS)

biotin cofactor - important point in activated one carbon carriers

Highly regulated step

allosteric effectors and covalent modification citrate activates by increasing the Vmax, where long chain FAs inhibit. AMP kinase inactivates (think of why) and insulin activates a phosphatase which relieves this inhibition by promoting the dephosphorylation of ACoA carboxylase



1) Production of malonyl CoA

- Acetyl CoA Carboxylase active as a polymer (highly regulated step)

2) Fatty acid synthesis transacylase

- the first reaction starts with a malonyl CoA and an ACoA
- transfer of acyl CoAs to ACP
- ACP - acyl carrier protein (derivative of CoA)
- long flexible arms w/ sulfhydryls
- allows flexibility to different active sites

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2) Fatty acid synthesis condensation

- acetyl CoA + malonyl CoA
- loss of CO₂ drives reaction (from ATP hydrolysis)
- ACoA added onto malonyl CoA
- forms new 2 carbon acetoacetyl group

2) Fatty acid synthesis reductions

- NADPH required
- similar reactions to reversal of beta oxidation
- end up with a new +2 carbon acyl-ACP group

recycle Reactions 2-6

continuation reactions

- the above is one full round of fatty acid synthase
- reactions continues 7 times until 16 carbon chain is formed (palmitate)
- termination - by thiolase

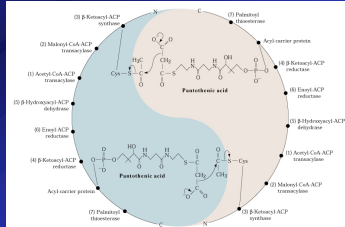
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Single enzymes or complexes

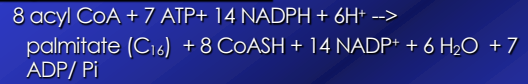
Two types of FAS systems bacterial and higher organisms. In *E. coli*, this takes place by several successive different enzymatic reactions. In eukaryotic cells, it takes place on one enzyme with several catalytic activities.

Reactions. The benefits of multi-enzyme systems:
 - channeling of substrates - faster synthesis
 - no loss of intermediates to side reactions

Source of reducing equivalents
 pentose phosphate pathway
 malic enzyme - 1 per acetyl CoA transport



Energetics of FA synthesis

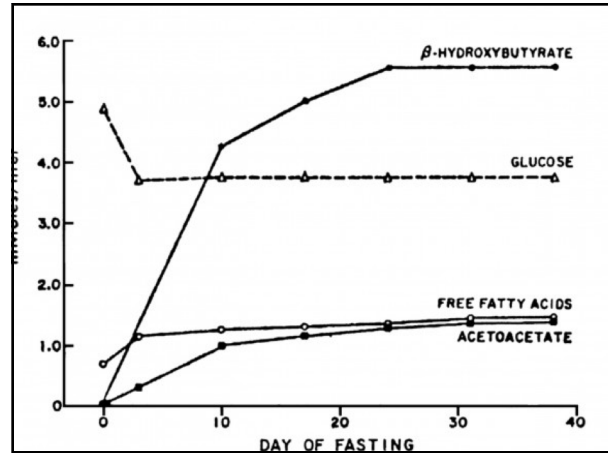


The net yield in ATP currency is from mito is
 - 8 ATP - citrate lyase
 - 7 ATP - Acetyl CoA Carboxylase
 - 7 rounds of FAS x NADPH \rightarrow 42 ATP
 total of 57 ATP to make palmitate / 129 to break down palmitate

Fruity Smelling Breath

Ketogenesis - Acetyl-CoA forms acetoacetate under high fatty acid oxidation (**loss of insulin**) and or prolonged starvation

- Acetoacetate can spontaneously decarboxylate into acetone OR reduce to β hydroxybutyrate
- These are collectively called ketone bodies
- Acetoacetate and β hydroxybutyrate serve as alternative fuel sources by the brain and muscle



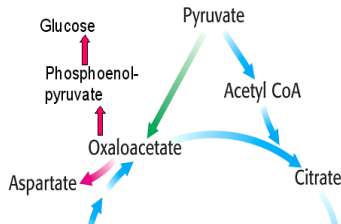
KETONE BODIES

The entry of acetyl CoA into the citric acid cycle depends on the availability of oxaloacetate.

The concentration of oxaloacetate is lowered if carbohydrate is unavailable (starvation) or improperly utilized (diabetes).

Oxaloacetate is normally formed from pyruvate by *pyruvate carboxylase* (anaplerotic reaction).

Fats burn in the flame of carbohydrates.



Fruity Smelling Breath

Synthesis

- Acetoacetyl-CoA forms by the condensation of two acetyl-CoA - thiolase in reverse
- A third Acetyl CoA is added by another condensation reaction to form HMG-CoA (β -hydroxy- β -methylglutaryl-CoA) - part of cholesterol synthesis
- HMG CoA is lysed to form acetyl CoA and acetoacetate

Fruity Smelling Breath

Acetoacetate and β HG formed in the liver are transported in the bloodstream to peripheral tissues and there are converted into two acetyl CoA units. This is still inefficient because succinyl CoA is used and therefore is not available for STK formation of GTP in the Krebs cycle.

B. Ketone bodies are a major fuel in some tissues

Ketone bodies diffuse from the liver mitochondria into the blood and are transported to peripheral tissues.

Ketone bodies are important molecules in energy metabolism.

Heart muscle and the renal cortex use acetoacetate in preference to glucose in physiological conditions.

The brain adapts to the utilization of acetoacetate during starvation and diabetes.

KETOSIS

The absence of insulin in diabetes mellitus

- liver cannot absorb glucose
 - inhibition of glycolysis
 - activation of gluconeogenesis
 - caused by a deficit of oxaloacetate
 - activation of fatty acid mobilization by adipose tissue
 - leading to large amounts of acetyl CoA which can not be utilized in Krebs cycle
 - large amounts of **ketone bodies** (moderately strong acids)
 - Giving rise to severe acidosis (**ketosis**)
- Impairment of the tissue function, most importantly in the central nervous system
-