

Fatty Acid Catabolism



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Outline

1. Mobilization of Fats from Dietary Intake and Adipose Tissue
2. Beta-Oxidation of Fatty Acids
3. Odd-Carbon Fatty Acids
4. Unsaturated Fatty Acids
5. Other Aspects of Fatty Acid oxidation
6. Ketone Bodies

Trends in Lipid Consumption

Lipid consumption has increased

- Luxury
- Make food taste better
- Satiation (provide fullness)

<u>Recommended</u>	<u>Intake</u>	<u>Fat</u>
30% of calories	40% of calories - mid 1960s 36% of calories - 1978 34% of calories - 1990	13% saturated fat 12% saturated fat

Composition of some known fats

Myristic acid	Coconut and palm oils, most animal and plant fats
Palmitic acid	Animal and plant fats
Stearic acid	Animal fats, some plant fats
Arachidic acid	Peanut oil
Lignoceric acid	Most natural fats, peanut oil in small amounts
Palmitoleic acid	Marine animal oils, small amounts in animal and plant fats
Oleic acid	Animal and plant fats
Linoleic acid	Corn, safflower, soybean, cottonseed, sunflower seed, and peanut oils
Linolenic acid	Linseed, soybean, and other seed oils
Arachidonic acid	Animal fats in small amounts

Food	Total lipid	Cholesterol	Saturated	Oleic	Linoleic	Linolenic
Milk	3.5	12	59	25	3	1.0
Egg	11.0	548	29	37	11	0.2
Beef -lean ground	22.0	70	50	41	3	0.7
Pork -lean	14.0	85	37	42	9-14	1.0
Chicken leg -flesh	3.5	74	27	47	22	2.0
Salmon	14.0	35	18	16	2	20.0
Whole wheat	2.0	0	21	14	55	4.0
Corn -whole	3.8	0	15	44	43	2.0
Soybeans -whole	18.0	0	13	22	54	5.0
Peanuts -butter	48.0	0	14	48	28	0.5
Coconut -fresh	38.0	0	83	5	2	0
Avocado -fresh	24.0	0	14	66	9	trace

Lipid Absorption from Lumen into Mucosal Cells

To get a lipid into the mucosal cell, it must be deesterified into free fatty acids (FFA) and monoacyl glycerols (MAG) and cholesterol. To get them out of the mucosa and into circulation, via the lymphatics, the lipids are packaged as chylomicrons. CMs are composed largely of TAGs, plus cholesterol esters and several proteins (called apoproteins).

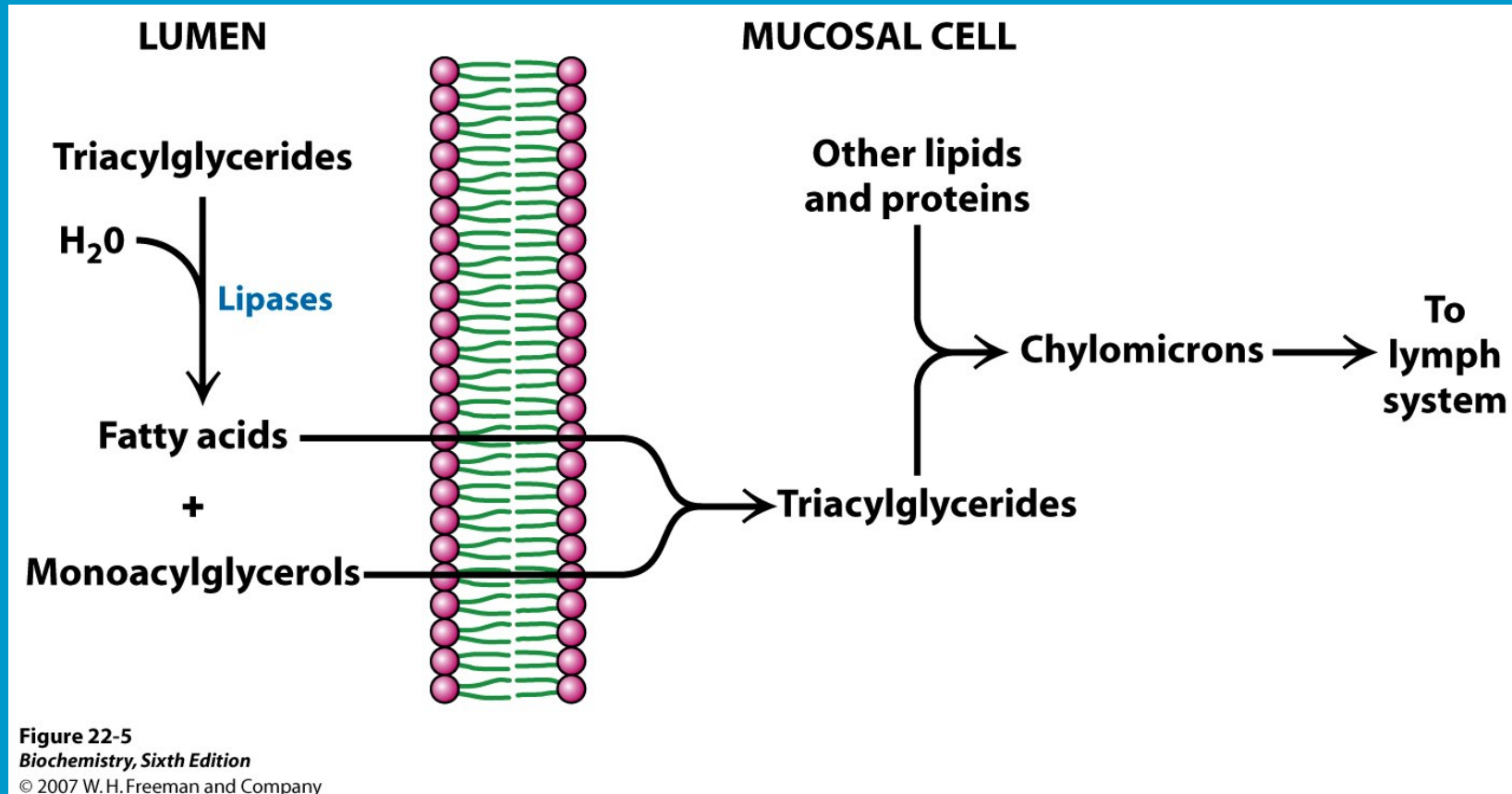


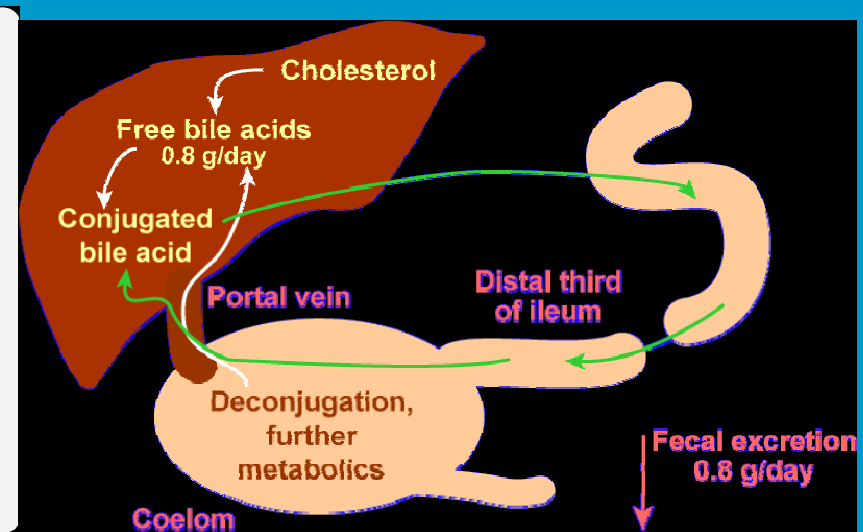
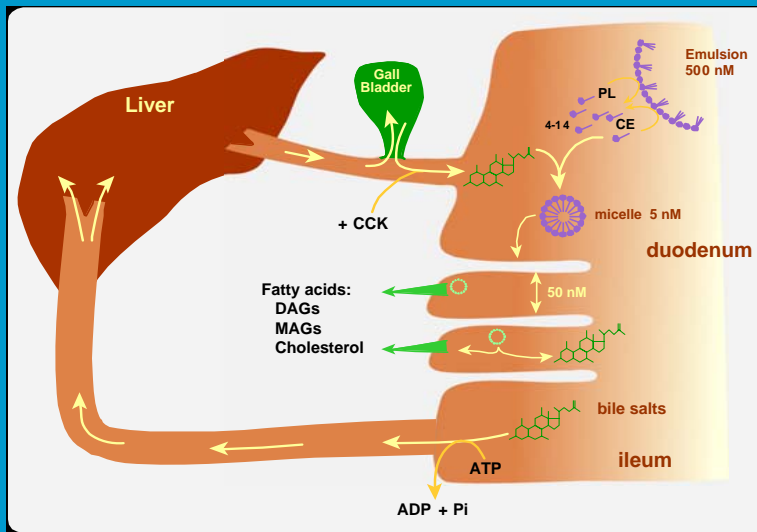
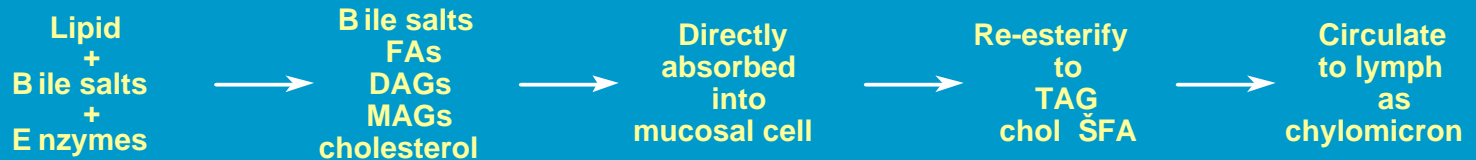
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Absorption

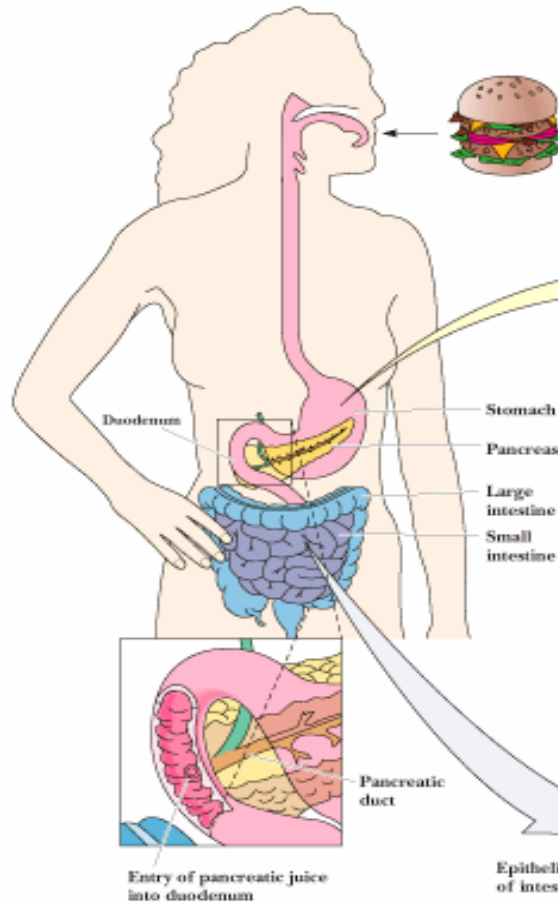
Liver: Synthesizes primary bile acids from cholesterol, sends them to storage in gall bladder. When food hits small intestine, gall bladder is stimulated to release bile acids into the sm. int. Most bile acids are derivatized with the amino acids glycine or taurine; this lowers their pKa so that they are soluble in the lower pH of the duodenum (pH 5.8 - 6.5).

Small intestine: At sufficiently high conc (critical micellar concentration = CMC), bile salts spontaneously aggregate to form micelles. When mixed with food, they trap lipid particles during their aggregation; this makes micelles from emulsion. The bile salt structure contains hydrophobic (rings) and hydrophilic (hydroxy group, amino acid) groups; this allows the lipid to be in contact with the aqueous environment required by degradative enzymes.

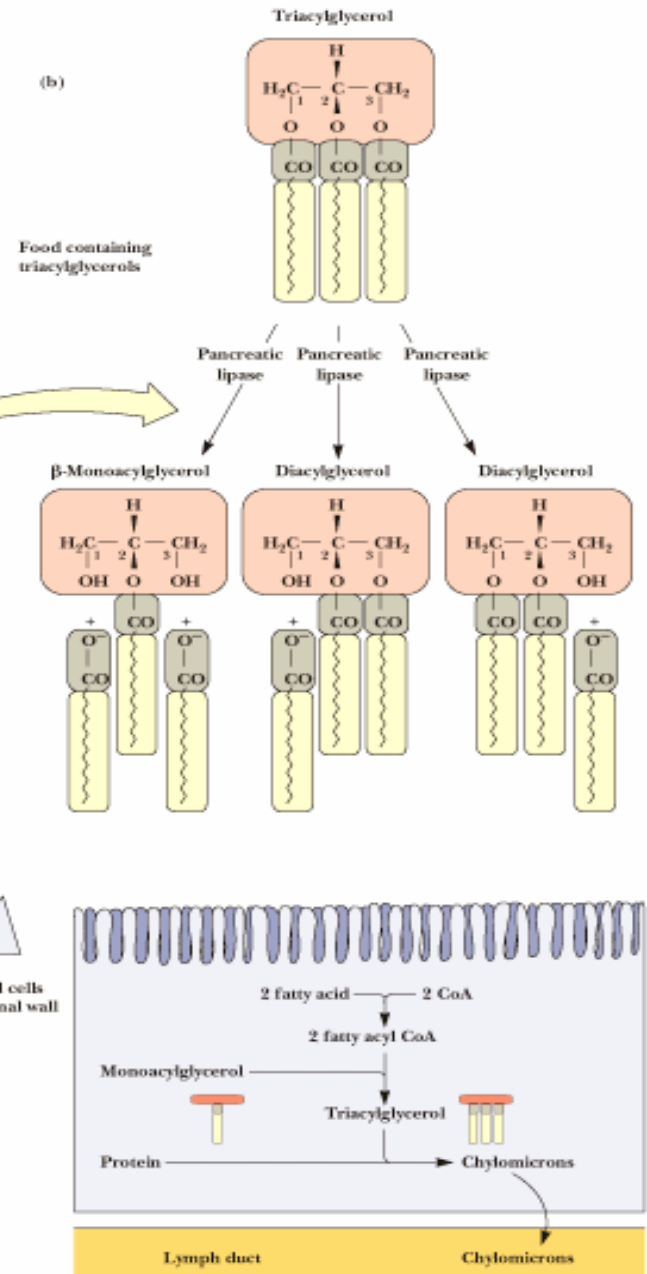
Lower small intestine, colon: Intestinal bacteria further derivatize bile acids by removing hydroxyl groups at the C-7 position. These secondary bile acids are associated with higher risks for colonic cancer (mechanism unclear).



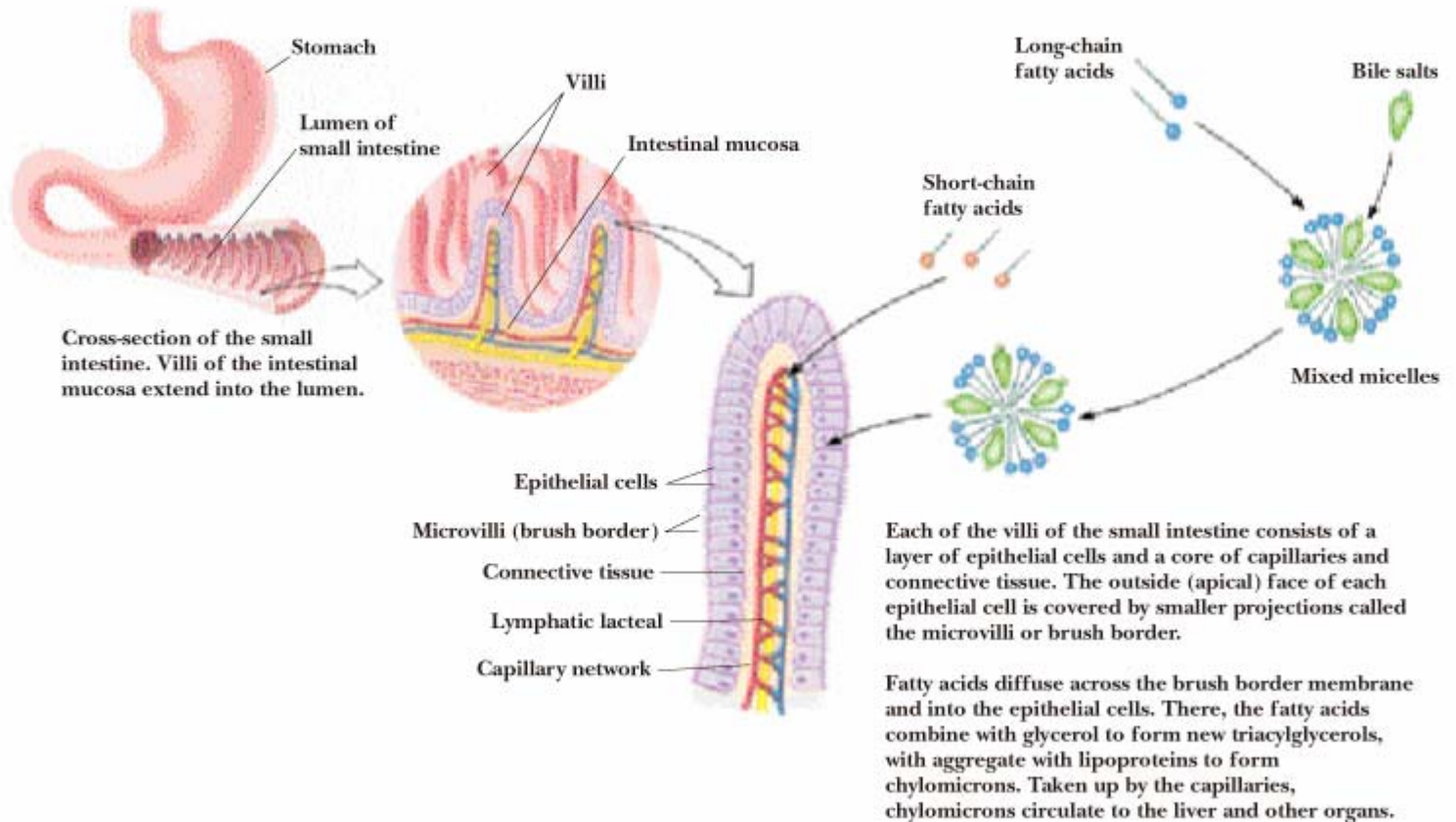
(a)



(b)



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 Figure 24.4

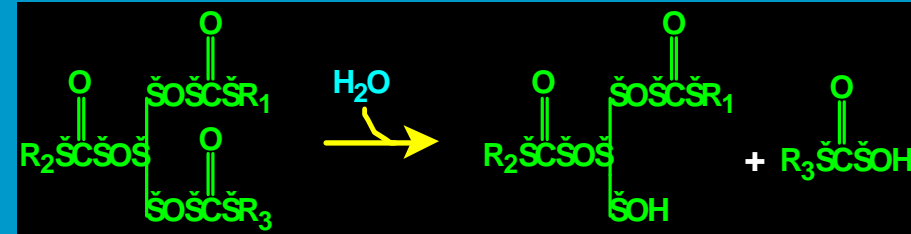


Lipid Digestion

Mouth:

- Important largely for neonates
- Acts on milk fats - preemulsified fats

Lingual Lipase:



Stomach:

- Little to none
- Churning creates a coarse lipid emulsion
- Fat slows the release of food from the stomach

Small intestine

- Performs 90% of all lipid digestion
- Primarily performs hydrolysis and de-esterification
- Mix bile salts to convert coarse emulsion into micelles

- **Lipoprotein Lipase:** found on endothelial (vessel) walls lining tissues such as adipose and muscle. Releases FFA from TAGs in CM/VLDL for cellular uptake and usage as either energy (muscle) or storage (adipocyte). Thus insulin & glucagon differentially regulate this enzyme on muscle vs. adipose cells.



- **Hormone-sensitive lipase:** Only found INSIDE adipocyte. Releases FFA from adipocyte TAG stores, sends to serum. Incr by glucagon, epinephrine.



- **Phosphatidate Phosphatase:** in all cells, synthesizes TAG from glycerol-PO4.



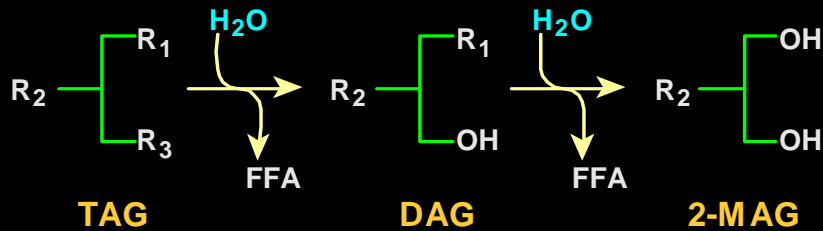
- **Regulation of LPL Activity:**

factor	adipose	muscle
lipoprotein CII	up	up
cholesterol	down	down
starvation	up	up
excess glucose	up	down
insulin	up	down

Lipid Digestion

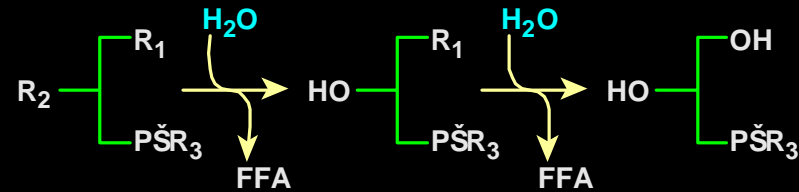
Pancreatic Lipase

long > short, unsaturated > saturated
inhibited by bile salts
enhanced by Ca ++ Š Colipase



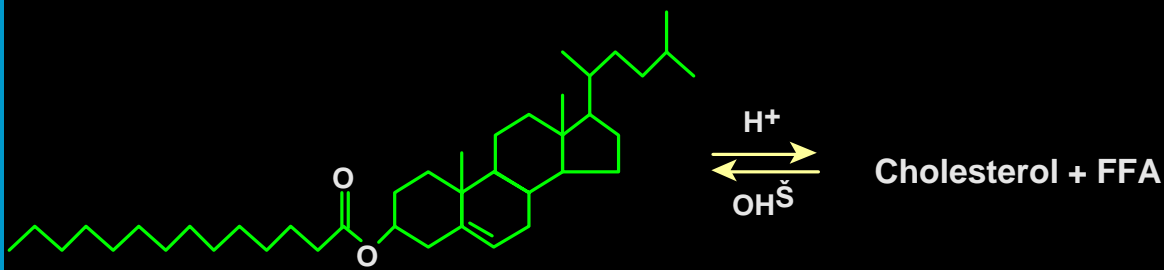
Phospholipase A2 = Lecithinase

acts at C1, 2 position of PL



Cholesterol Esterase

removes FA at low pH - lumen
adds FA at higher pH - mucosa





Why Fatty Acids?

(For energy storage?)

Two reasons:

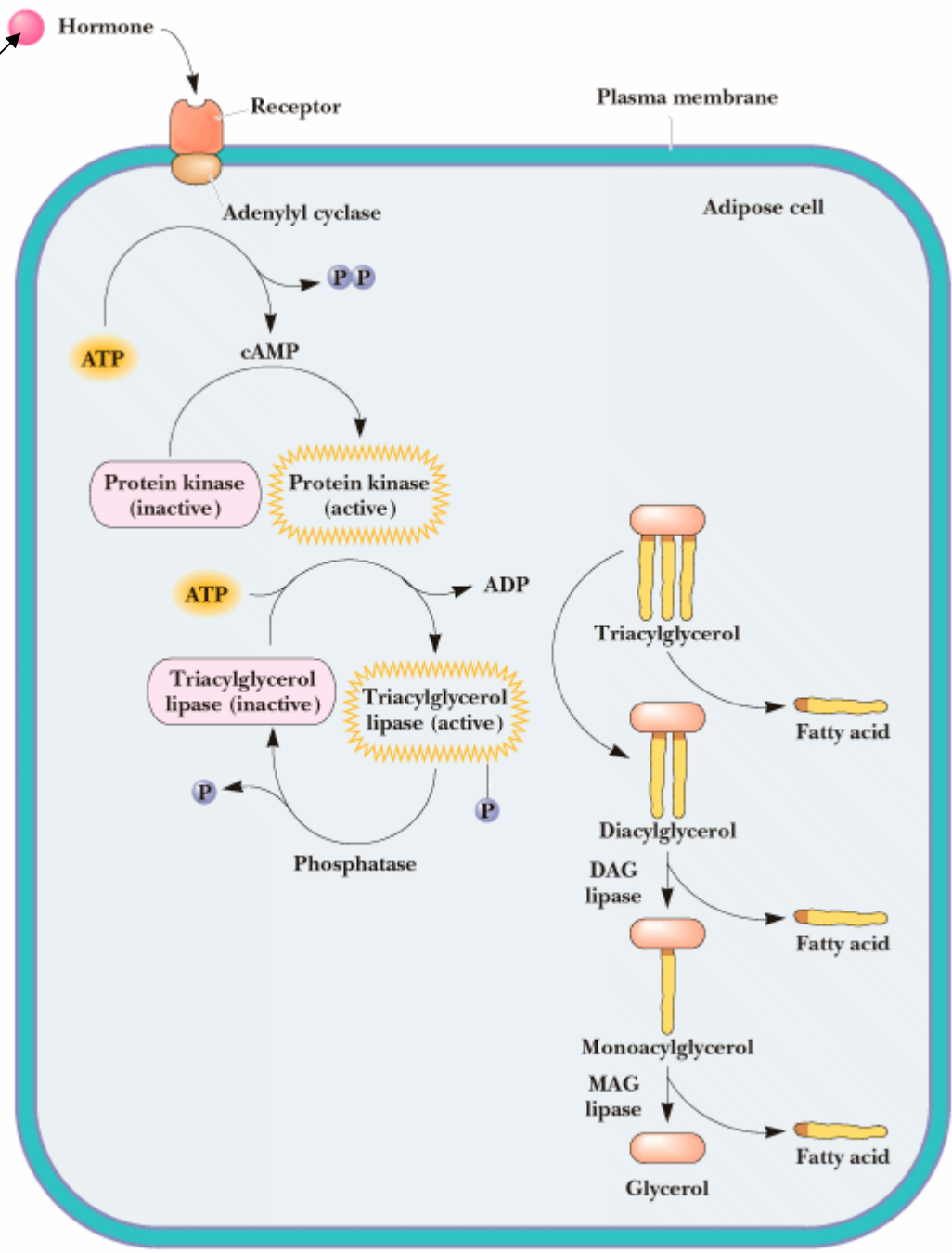
- The carbon in fatty acids (mostly CH_2) is almost **completely reduced** (so its oxidation yields the most energy possible).
- Fatty acids are **not hydrated** (as mono- and polysaccharides are), so they can pack more closely in storage tissues

Fat from Diet & Adipose Cells

Triacylglycerols either way

- Triglycerides represent the major energy input in the modern American diet
- Triglycerides are also the major form of stored energy in the body
- Hormones (**glucagon, epinephrine, ACTH**) trigger the release of fatty acids from adipose tissue

**glucagon,
epinephrine,
ACTH**



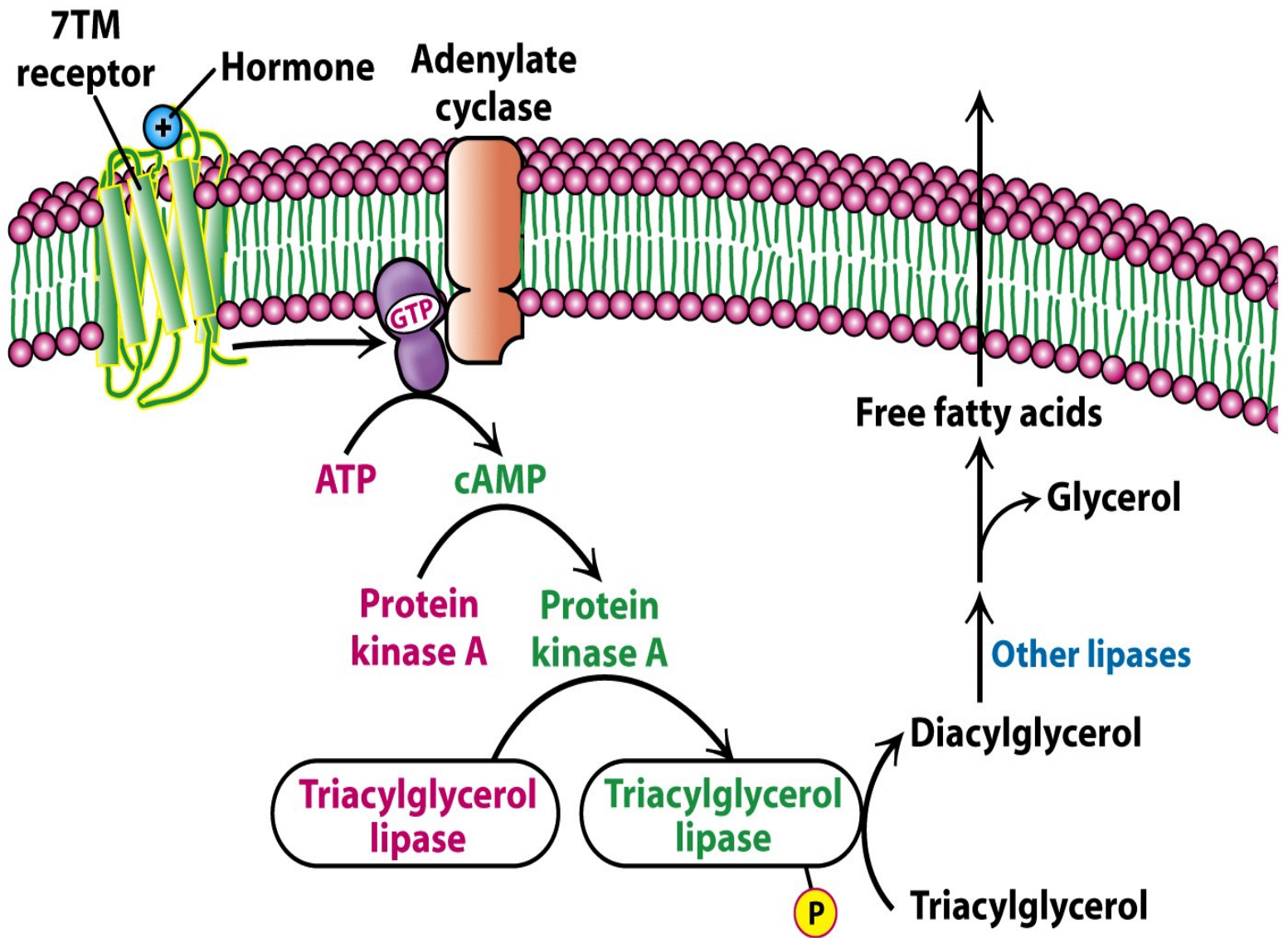


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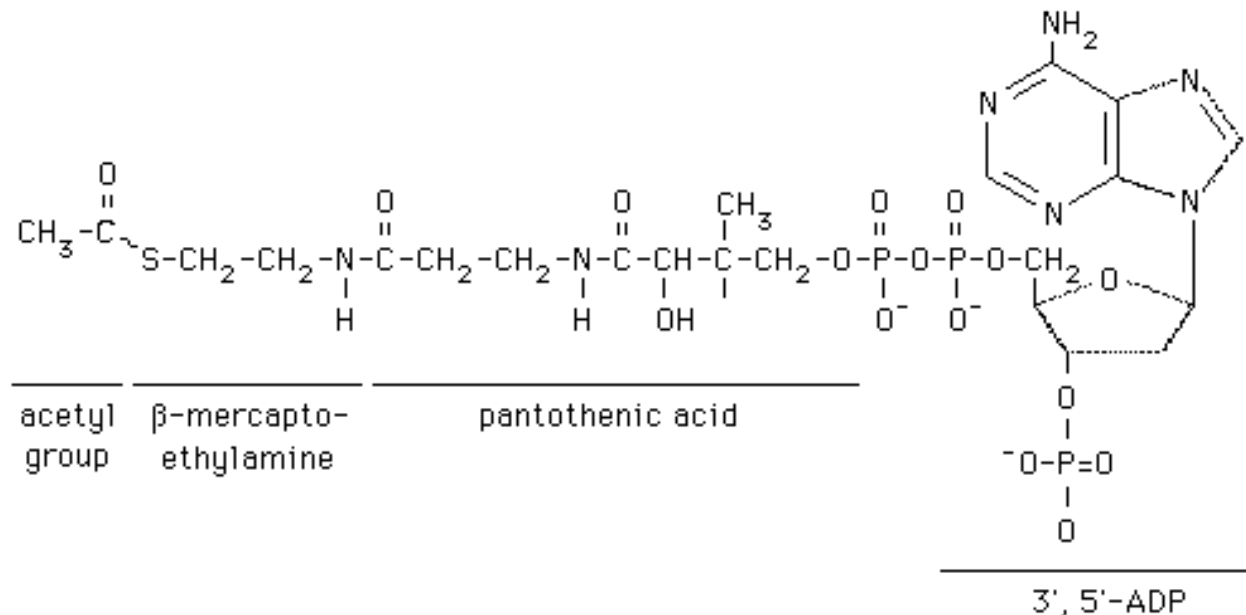
Beta Oxidation of Fatty Acids

- **Knoop** showed that fatty acids must be degraded by removal of 2-C units
- **Albert Lehninger** showed that this occurred in the mitochondria
- **F. Lynen and E. Reichart** showed that the 2-C unit released is **acetyl-CoA**, not free acetate
- The process begins with oxidation of the carbon that is "beta" to the carboxyl carbon, so the process is called "beta-oxidation"

Structure of Acetyl CoA

The structure of Acetyl CoA consists of two parts.

1. Acetyl group
2. Coenzyme A
 - Beta-mercaptoethylamine
 - Pantothenic acid (not synthesized in man -- an essential nutrient)
 - Phosphate
 - 3', 5'-adenosine diphosphate



Acetyl coenzyme A, showing its constituents

FATTY ACID DEGRADATION

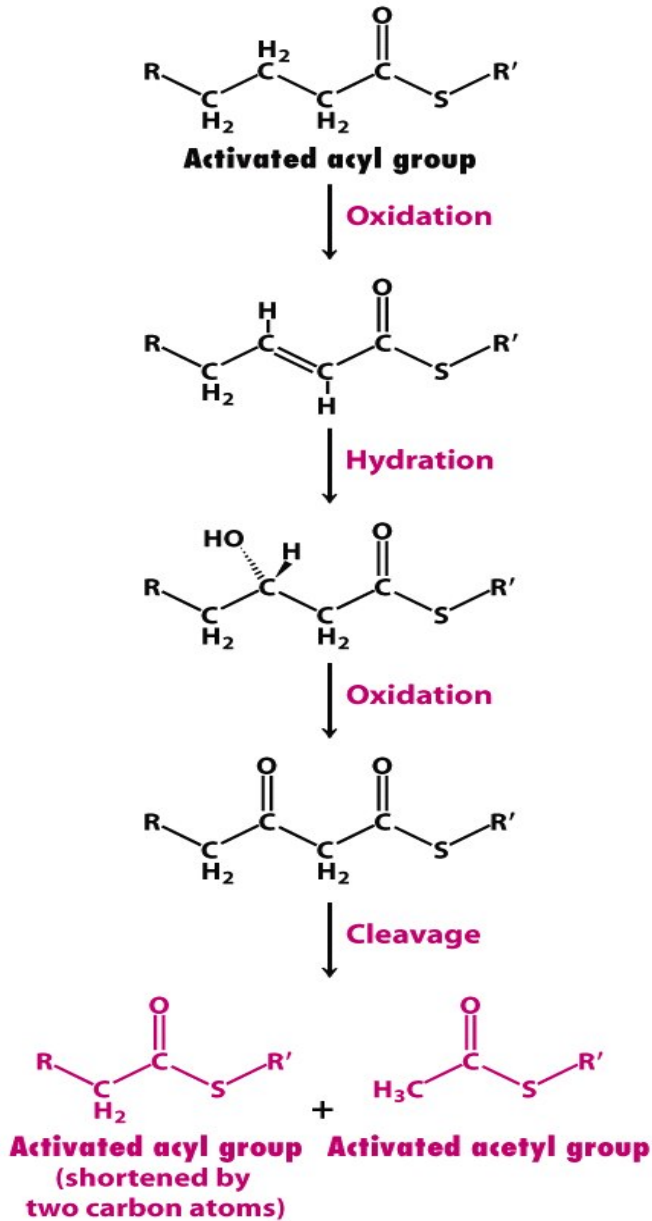
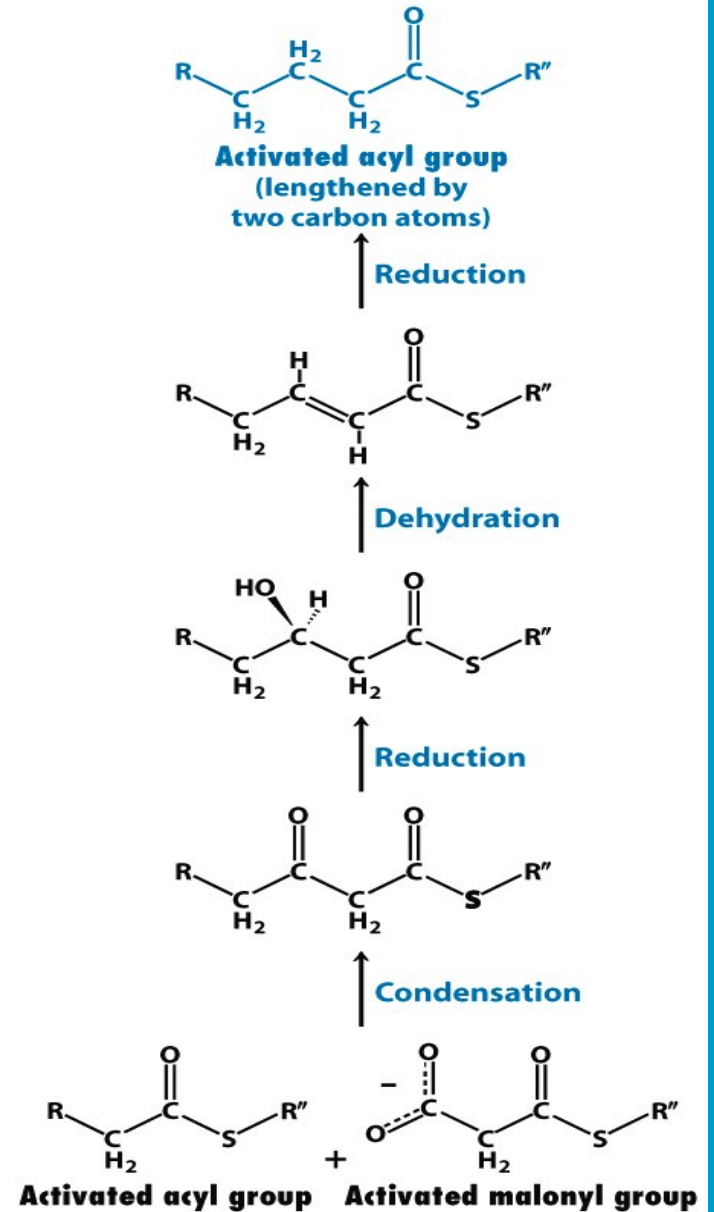


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FATTY ACID SYNTHESIS



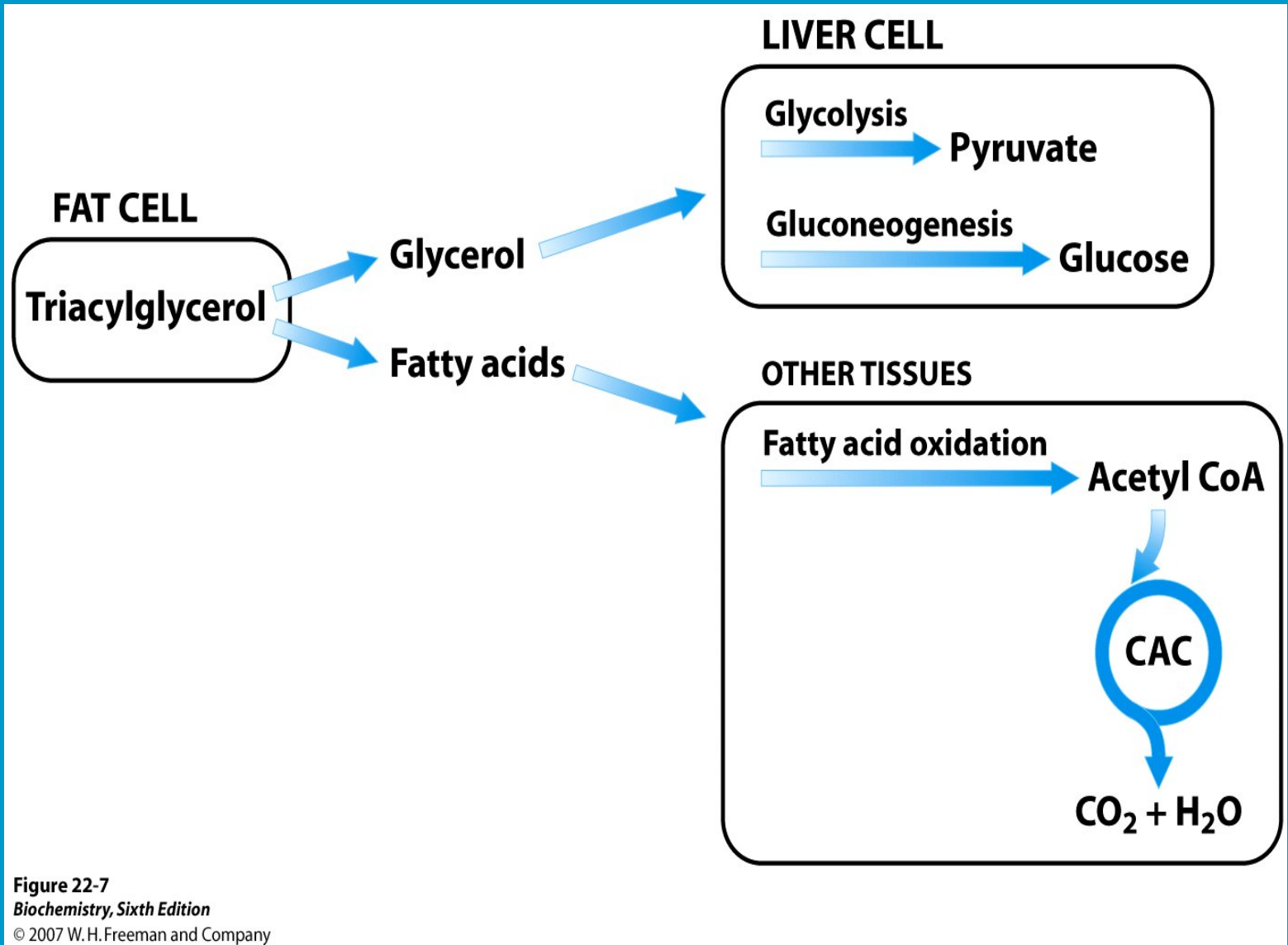


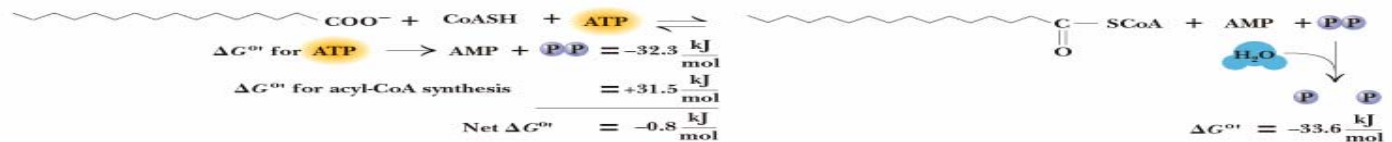
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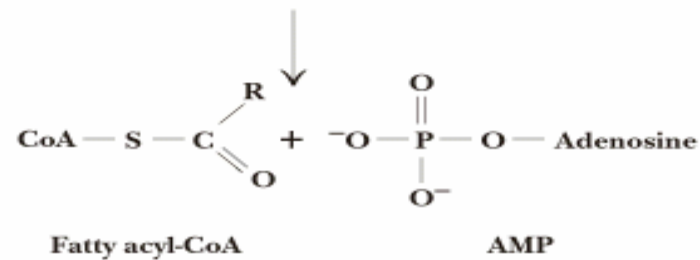
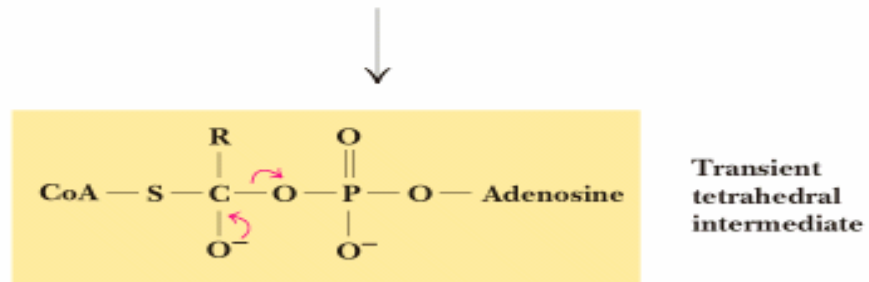
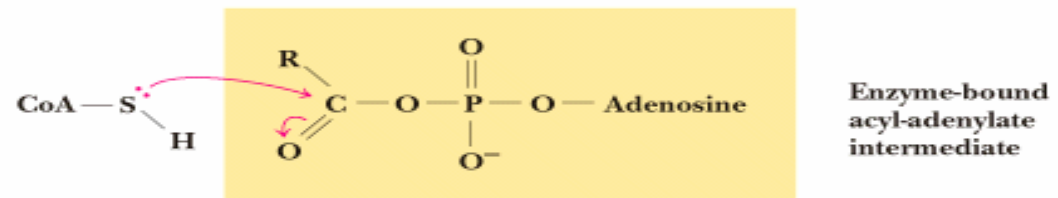
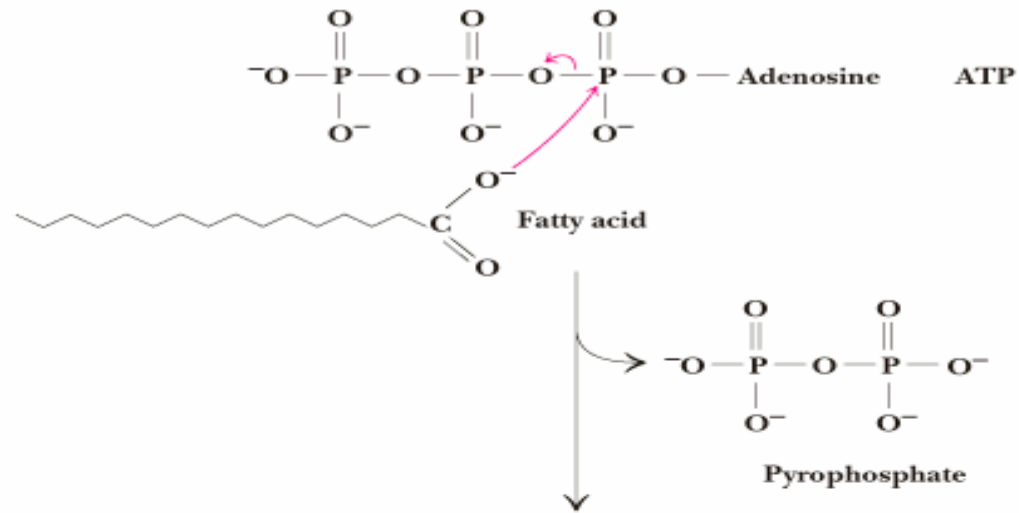
CoA activates FAs for oxidation

Acyl-CoA synthetase condenses fatty acids with CoA, with simultaneous hydrolysis of ATP to AMP and PP_i

- Formation of a CoA ester is expensive energetically
- Reaction just barely breaks even with ATP hydrolysis
- But subsequent hydrolysis of PP_i drives the reaction strongly forward
- Note the **acyl-adenylate** intermediate in the mechanism!

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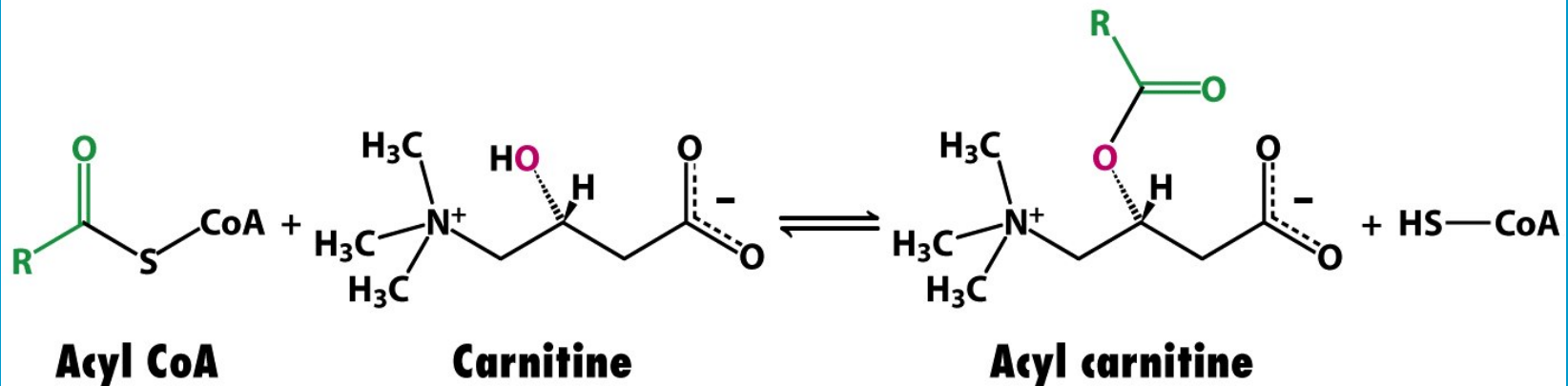




Carnitine as a Carrier

Carnitine carries fatty acyl groups across the inner mitochondrial membrane

- Short chain fatty acids are carried directly into the mitochondrial matrix
- Long-chain fatty acids cannot be directly transported into the matrix
- Long-chain FAs are converted to acyl carnitines and are then transported in the cell
- Acyl-CoA esters are formed inside the inner membrane in this way



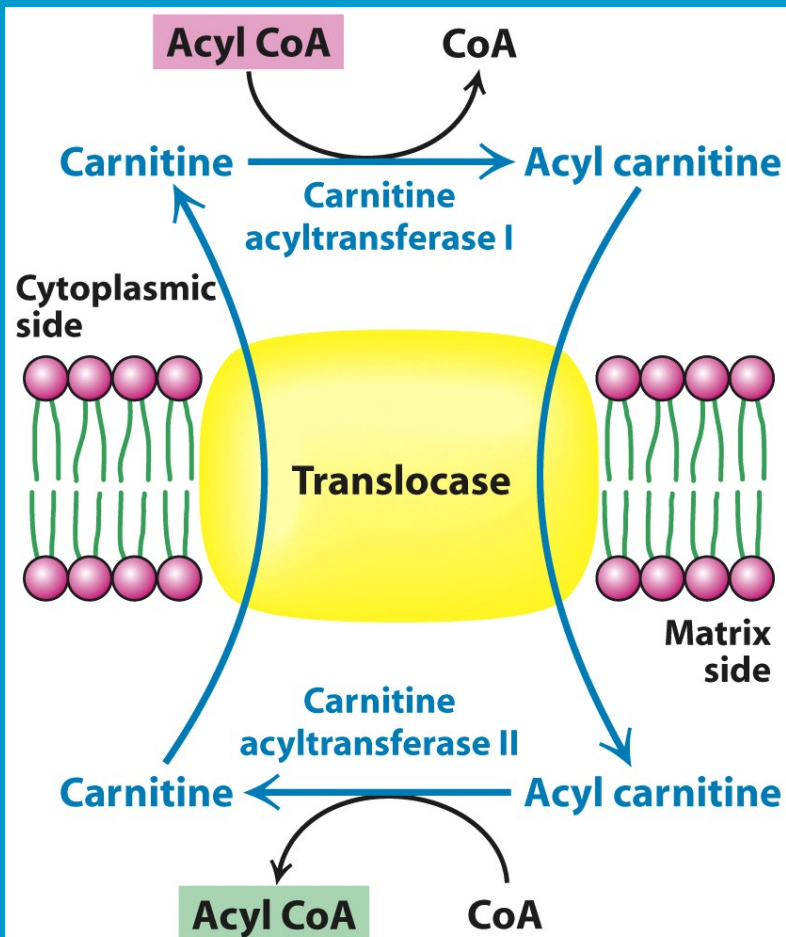
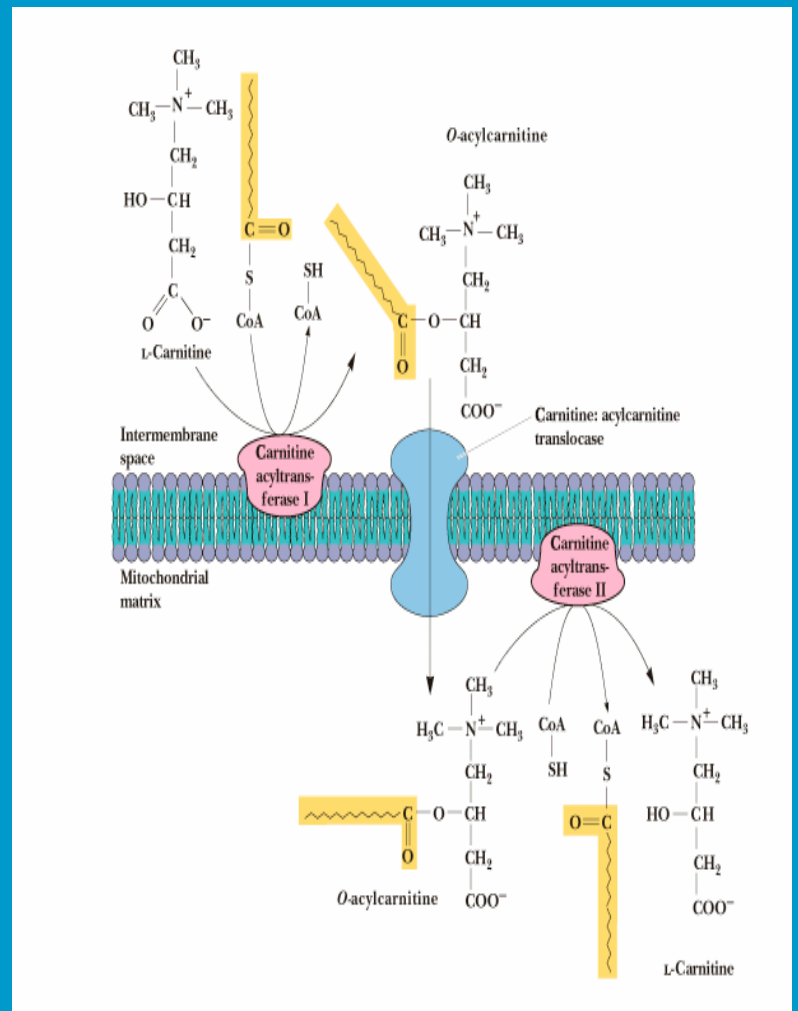


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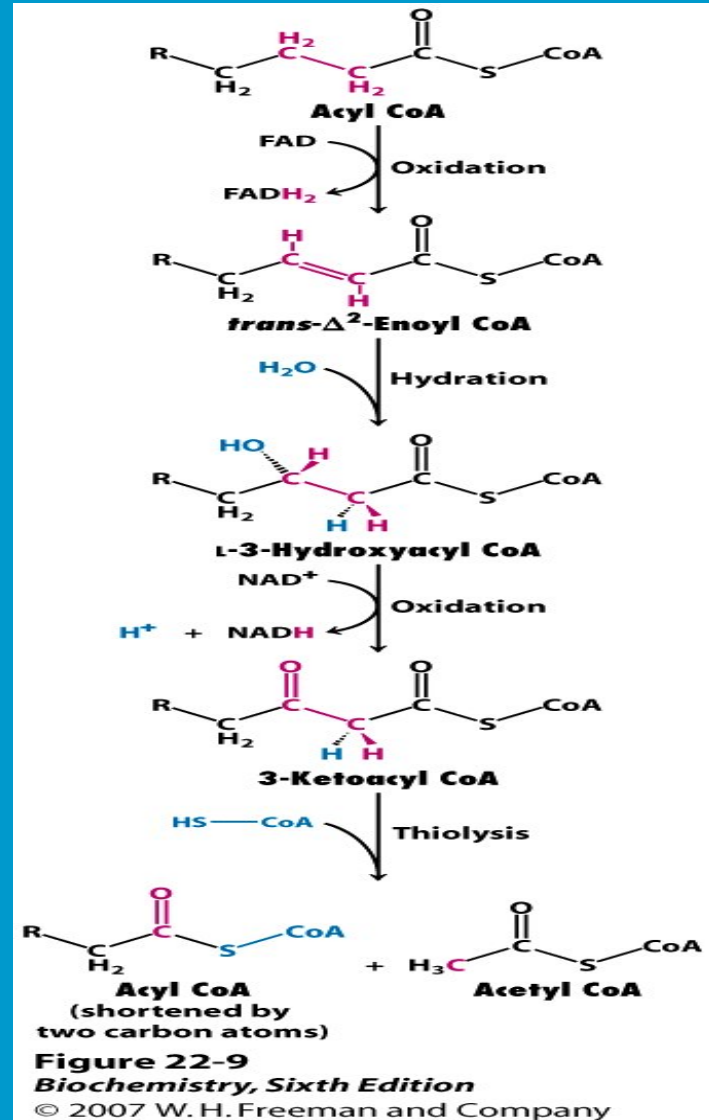
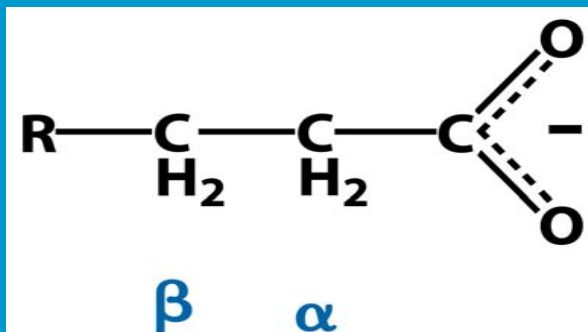
Carnitine acyltransferase 1 forms a O-acyl intermediate (which has a group transfer potential).

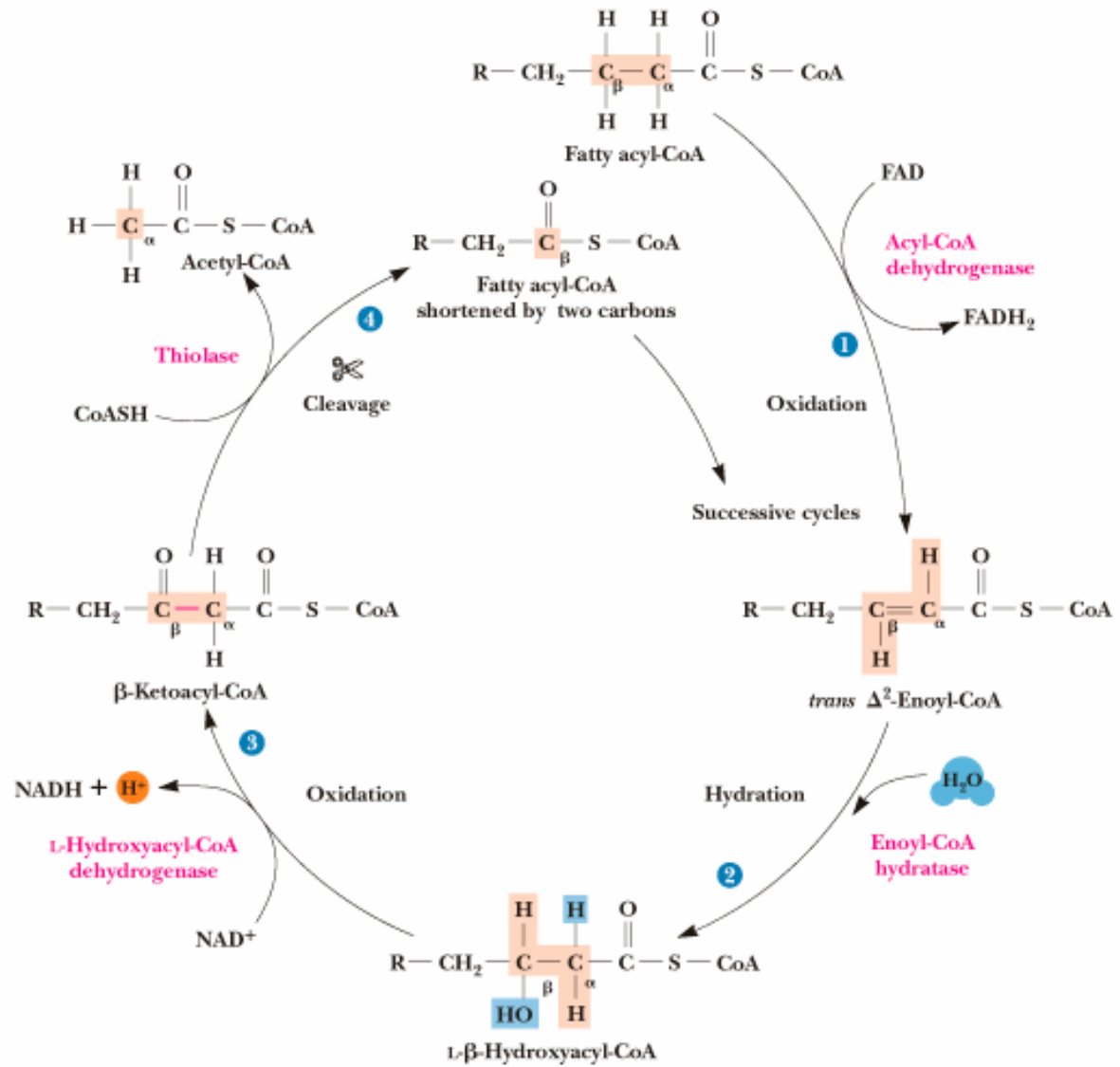
The eukaryotic cells maintain separate pools of CoA in the mitochondria (used in the oxidation of fatty acids, pyruvate and some amino acids) and in the cytosol (used principally in fatty acid biosynthesis)

β -Oxidation of Fatty Acids

A Repeated Sequence of 4 Reactions

- Strategy: create a carbonyl group on the β -C
- First 3 reactions do that; fourth cleaves the " β -keto ester" in a reverse Claisen condensation [CC – involves attack by a nucleophilic agent on a carbonyl carbon to yield a beta-keto acid]
- Products: an acetyl-CoA and a fatty acid two carbons shorter
- The first three reactions are crucial and classic - we will see them again and again in other pathways





Acyl-CoA Dehydrogenase

Oxidation of the C_{α} - C_{β} bond

- A family of three soluble matrix enzymes [that differ in specificity for either long, medium or short-chain acyl-CoAs]
- Mechanism involves proton abstraction, followed by double bond formation and hydride removal by FAD
- Electrons are passed to an **electron transfer flavoprotein**, and then to the electron transport chain

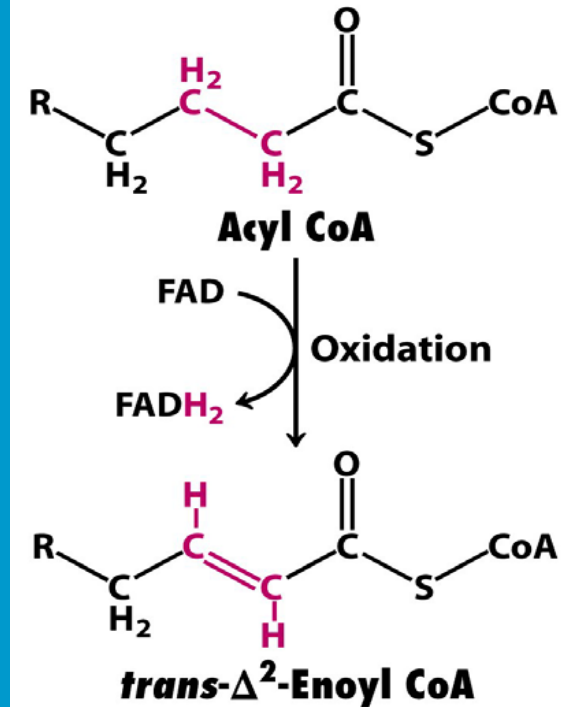
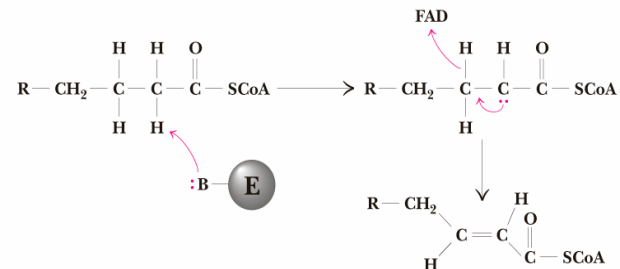
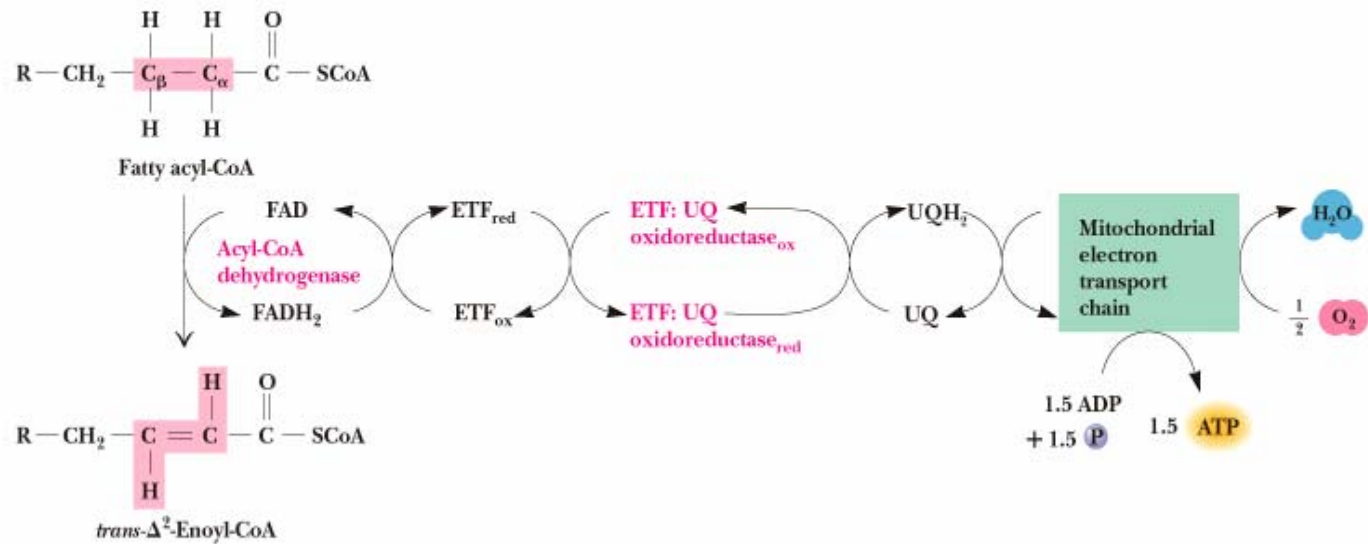


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Figure 24.11



Enoyl-CoA Hydratase

Adds water across the double bond

- at least three forms of the enzyme are known
- aka crotonases
- Normal reaction converts **trans**-enoyl-CoA to **L**- β -hydroxyacyl-CoA

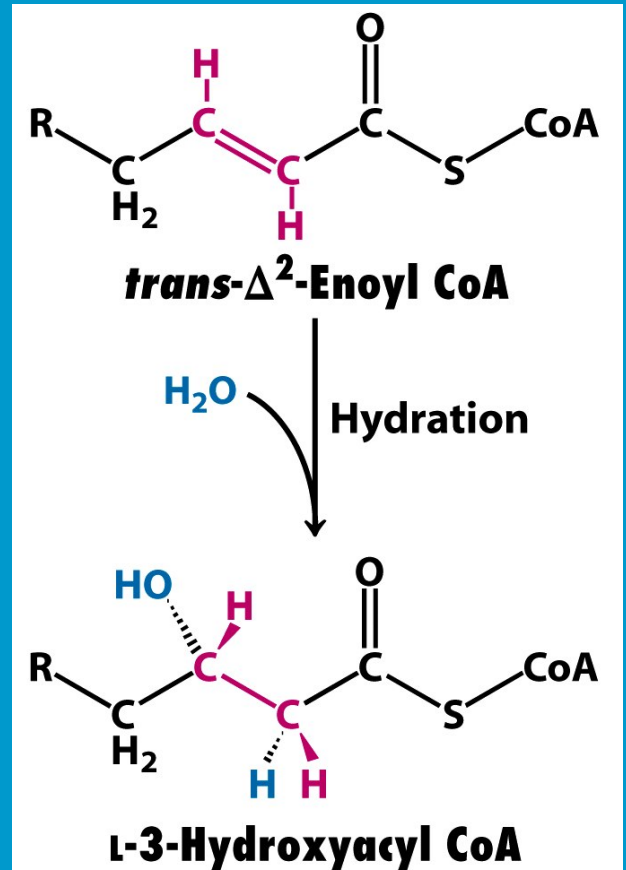


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Hydroxyacyl-CoA Dehydrogenase

Oxidizes the β -Hydroxyl Group

- This enzyme is completely specific for L-hydroxyacyl-CoA
- D-hydroxyacyl-isomers are handled differently

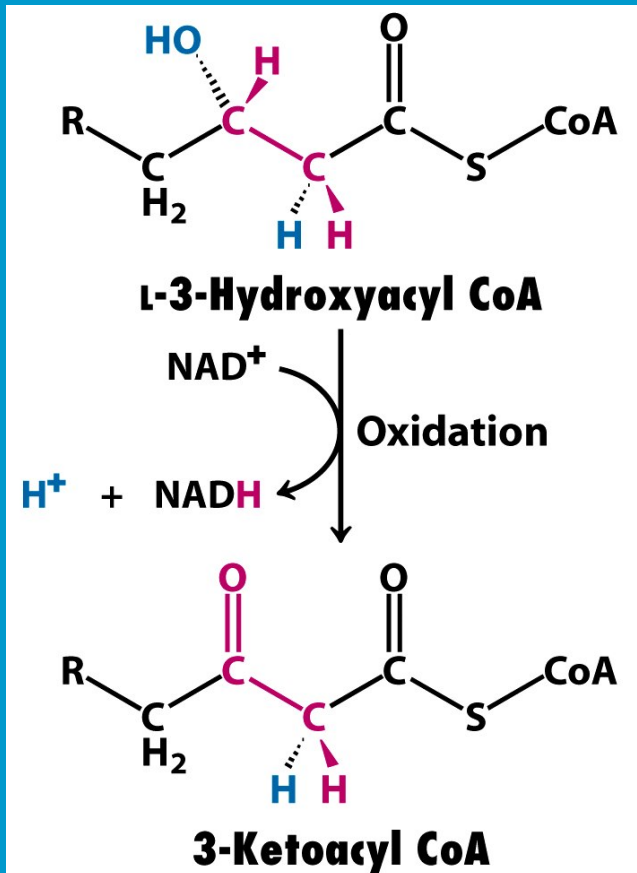
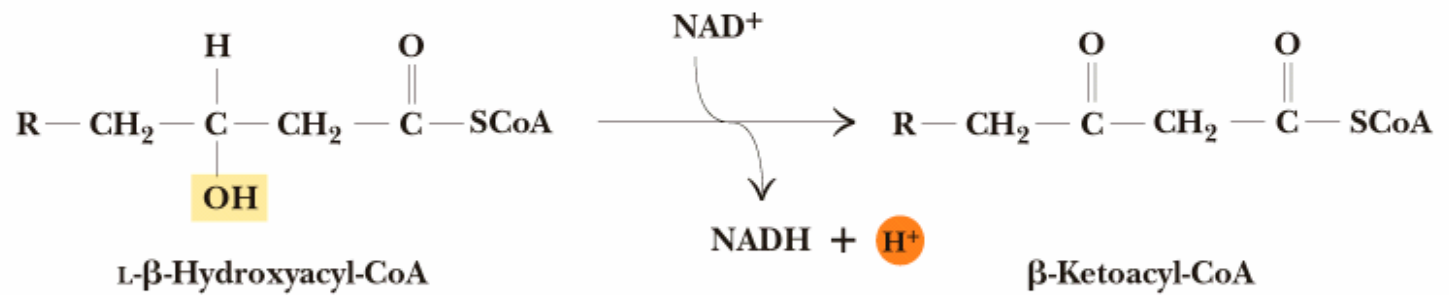


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Fourth reaction: thiolase

aka β -ketothiolase

- Cysteine thiolate on enzyme attacks the β -carbonyl group
- Thiol group of a new CoA attacks the shortened chain, forming a new, shorter acyl-CoA
- This is the reverse of a Claisen condensation: **attack of the enolate of acetyl-CoA on a thioester**
- Even though it forms a new thioester, the reaction is favorable and drives other three

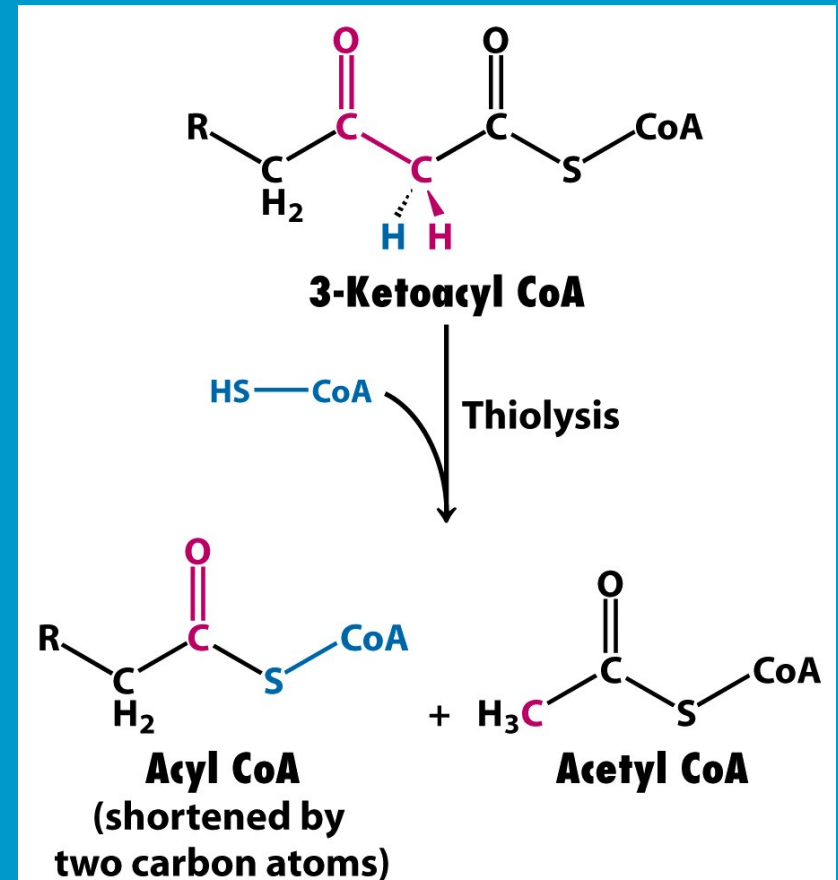
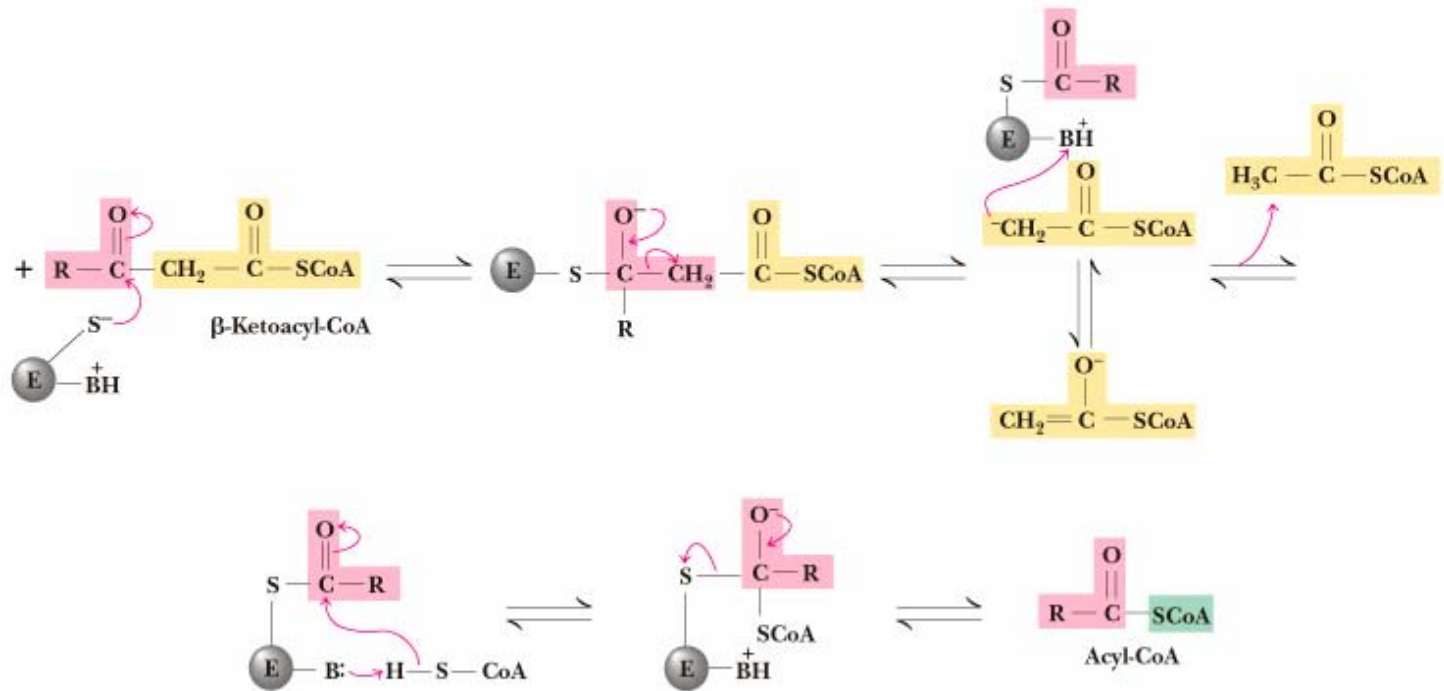


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Figure 24.17



Summary of β -Oxidation

Repetition of the cycle yields a succession of acetate units

- Thus, **palmitic acid yields eight acetyl-CoAs**
- Complete β -oxidation of one palmitic acid yields **106 molecules of ATP**
- Large energy yield is consequence of the **highly reduced state of the carbon** in fatty acids
- This makes fatty acids the fuel of choice for migratory birds and many other animals

TABLE 22.1 Principal reactions in fatty acid oxidation

Step	Reaction	Enzyme
1	Fatty acid + CoA + ATP \rightleftharpoons acyl CoA + AMP + PP _i	Acyl CoA synthetase [also called fatty acid thiokinase and fatty acid:CoA ligase]*
2	Carnitine + acyl CoA \rightleftharpoons acyl carnitine + CoA	Carnitine acyltransferase (also called carnitine palmitoyl transferase)
3	Acyl CoA + E-FAD \longrightarrow <i>trans</i> - Δ^2 -enoyl CoA + E-FADH ₂	Acyl CoA dehydrogenases (several isozymes having different chain-length specificity)
4	<i>trans</i> - Δ^2 -Enoyl CoA + H ₂ O \rightleftharpoons L-3-hydroxyacyl CoA	Enoyl CoA hydratase (also called crotonase or 3-hydroxyacyl CoA hydrolyase)
5	L-3-Hydroxyacyl CoA + NAD ⁺ \rightleftharpoons 3-ketoacyl CoA + NADH + H ⁺	L-3-Hydroxyacyl CoA dehydrogenase
6	3-Ketoacyl CoA + CoA \rightleftharpoons acetyl CoA + acyl CoA (shortened by C ₂)	β -Ketothiolase (also called thiolase)

* An AMP-forming ligase.

Table 22-1

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Energy Yield from β -Ox of Palmitate

Palmitoyl CoA + 7 CoASH + 7 FAD + 7 NAD + 7 H₂O



8 acetyl CoA + 7 FADH₂ + 7 NADH + 7 H⁺

FADH₂ = 1.5 ATP X 7 = 10.5 ATP

NADH = 2.5 ATP X 7 = 17.5 ATP

Acetyl CoA = 10 ATP X 8 = 80 ATP

108 ATP

- 2 ATP (activation of FA)

106 ATP

Odd-Carbon Fatty Acids

β -Oxidation yields propionyl-CoA

- Odd-carbon fatty acids are metabolized normally, until the last three-C fragment - propionyl-CoA - is reached
- Three reactions convert propionyl-CoA to succinyl-CoA
- An initial carboxylation at the α -carbon of propionyl-CoA to produce D-methylmalonyl-CoA, catalyzed by a biotin-dependent enzyme, **propionyl-CoA carboxylase**.
- D-methylmalonyl-CoA is converted to the L-isomer by **methylmalonyl-CoA epimerase**.
- The third reaction, catalyzed by **methylmalonyl-CoA mutase**, involves the migration of the carbonyl-CoA group from one carbon to its neighbor. This reaction is vitamin B12-dependent.

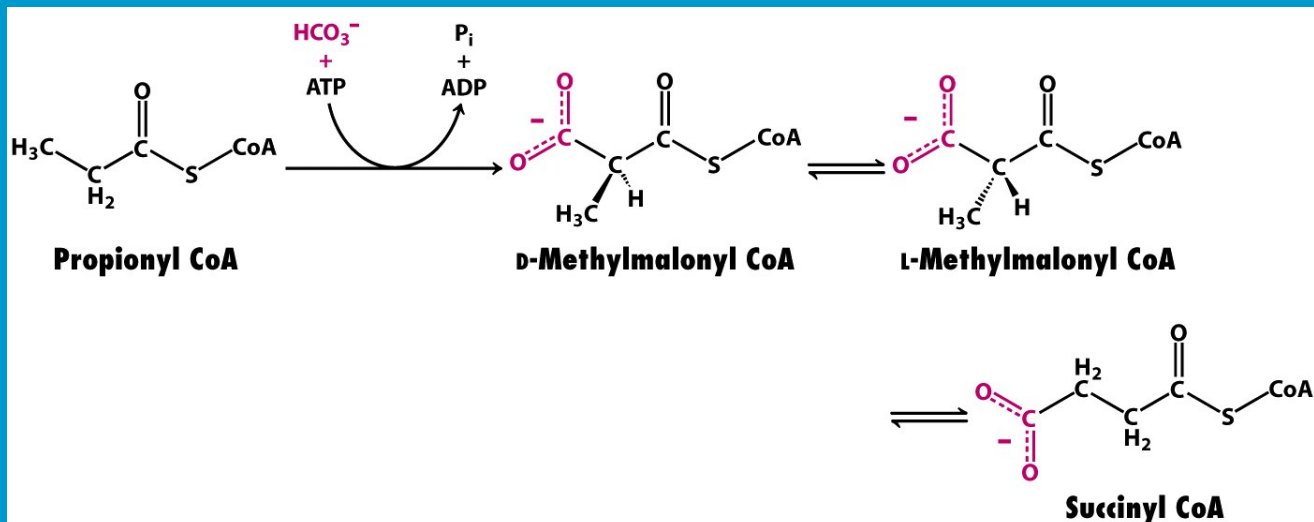
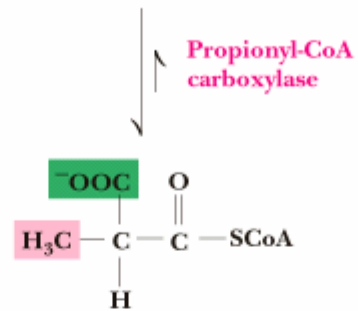
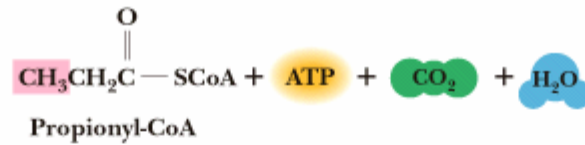
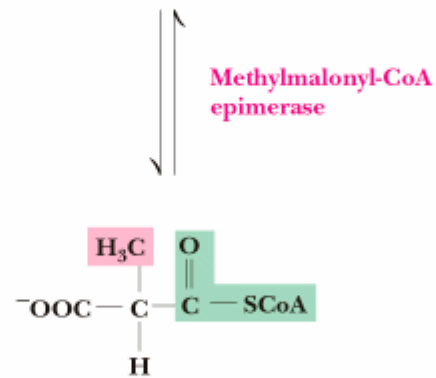


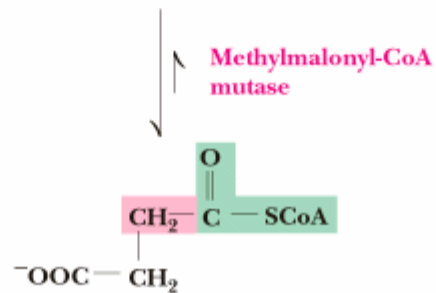
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D-Methylmalonyl-CoA



L-Methylmalonyl-CoA



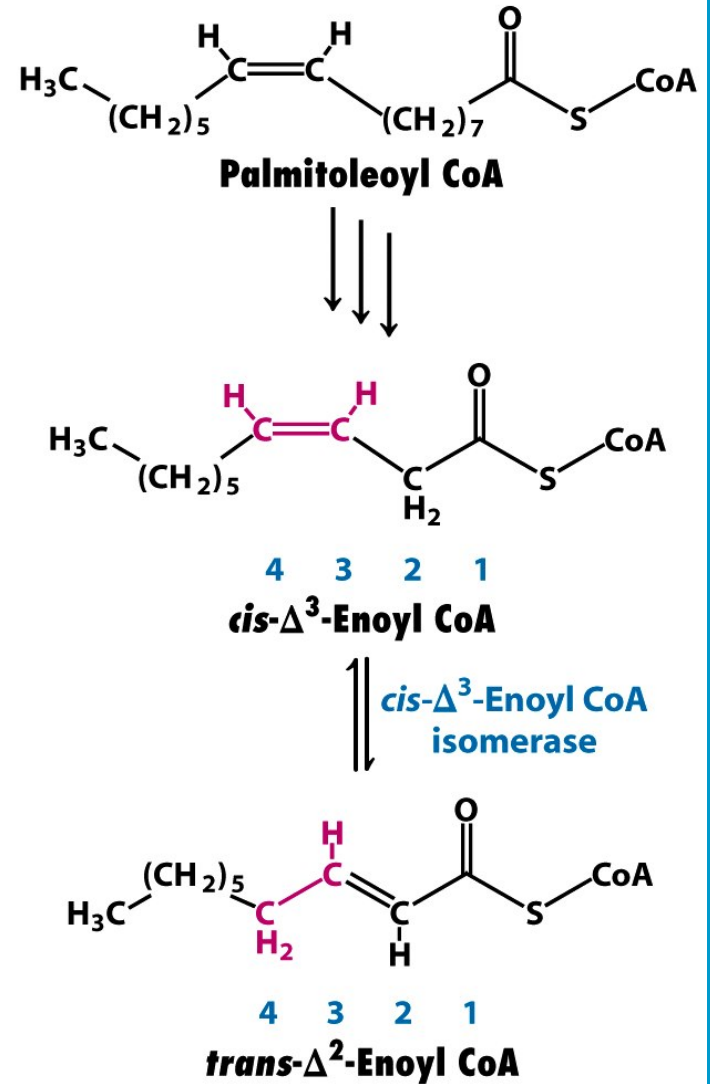
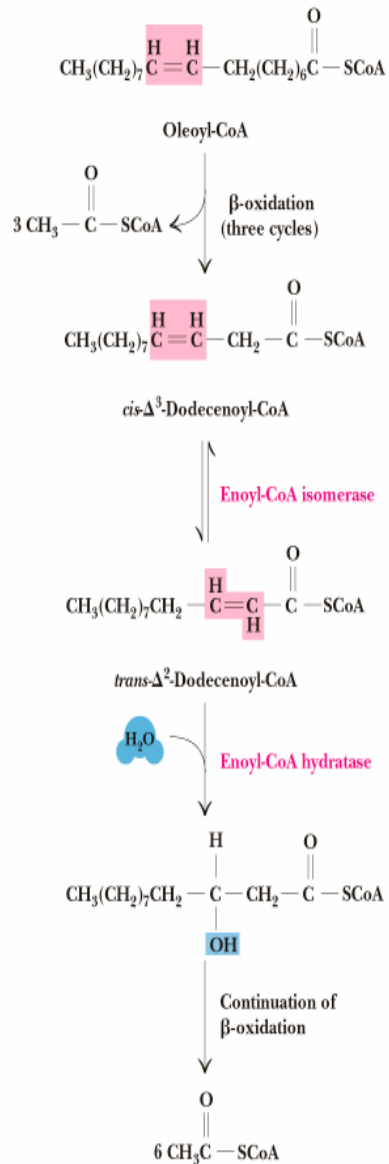
Succinyl-CoA

Succinyl CoA then enters the TCA cycle {succinyl CoA to malate to pyruvate to TCA cycle}

Unsaturated Fatty Acids

Consider monounsaturated fatty acids:

- Oleic acid, palmitoleic acid
- Normal β -oxidation for three cycles
- $\text{cis-}\Delta^3$ acyl-CoA cannot be utilized by acyl-CoA dehydrogenase
- **Enoyl-CoA isomerase** converts this to $\text{trans-}\Delta^2$ acyl CoA
- β -oxidation continues from this point

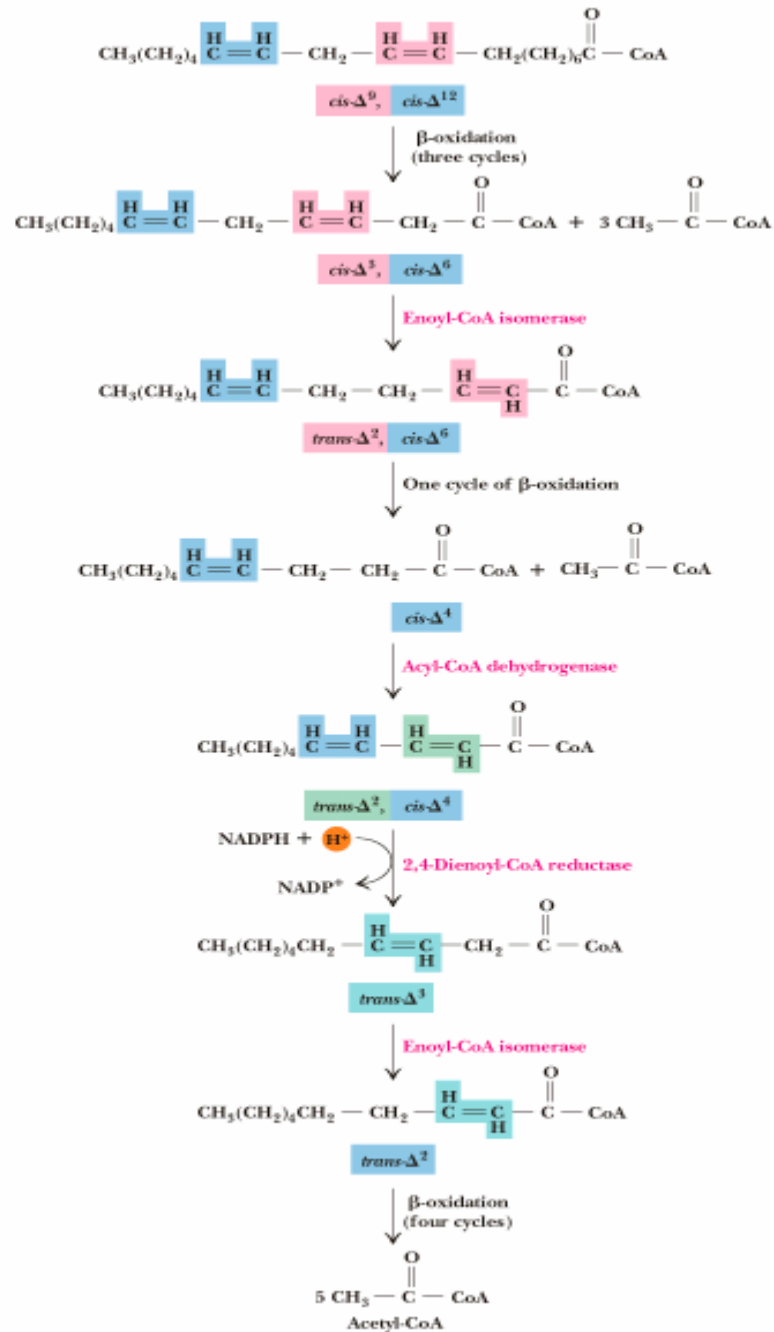


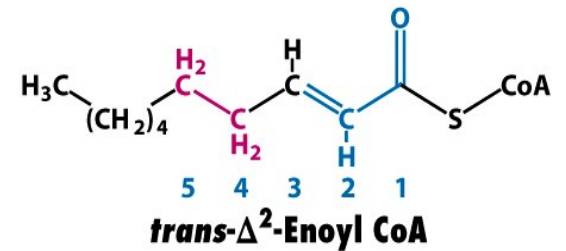
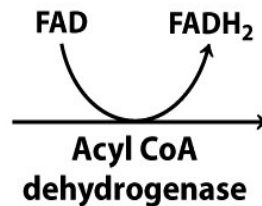
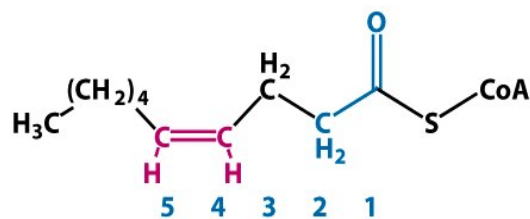
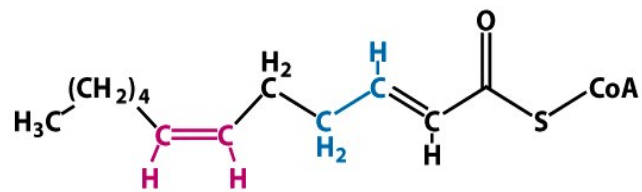
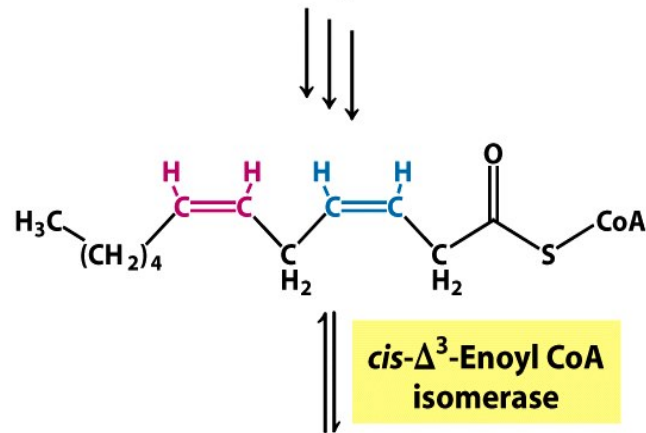
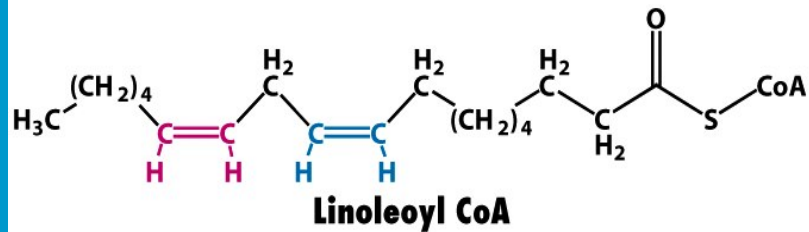
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Polyunsaturated Fatty Acids

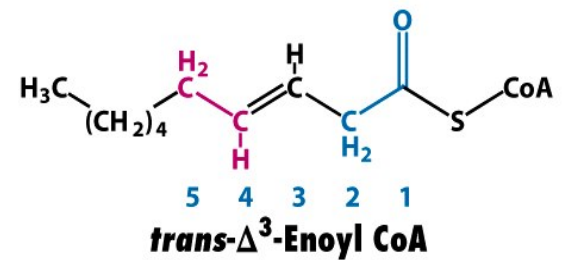
Slightly more complicated

- Same as for oleic acid, but only up to a point:
 - 3 cycles of β -oxidation
 - enoyl-CoA isomerase
 - 1 more round of β -oxidation
 - trans- Δ^2 , cis- Δ^4 structure is a problem!
- **2,4-Dienoyl-CoA reductase** to the rescue! — produces a trans- Δ^3 enoyl product. This enoyl product can be converted by an **enoyl-CoA isomerase** to the trans- Δ^2 enoyl CoA, which then proceeds normally through the beta-oxidation pathway.





cis- Δ^3 -Enoyl CoA isomerase



2,4-Dienoyl CoA reductase

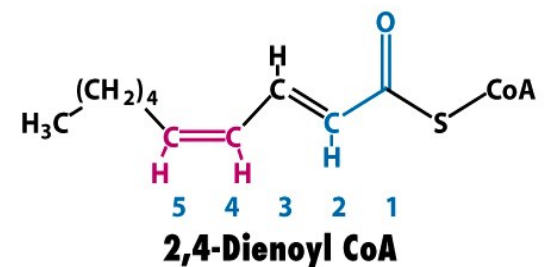
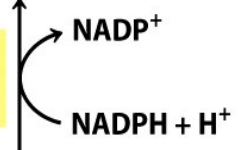


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Peroxisomal β -Oxidation

Peroxisomes - organelles that carry out flavin-dependent oxidations, regenerating oxidized flavins by reaction with O_2 to produce H_2O_2

- Similar to mitochondrial β -oxidation, but initial double bond formation is by acyl-CoA oxidase
- Electrons go to O_2 rather than e- transport
- Fewer ATPs result

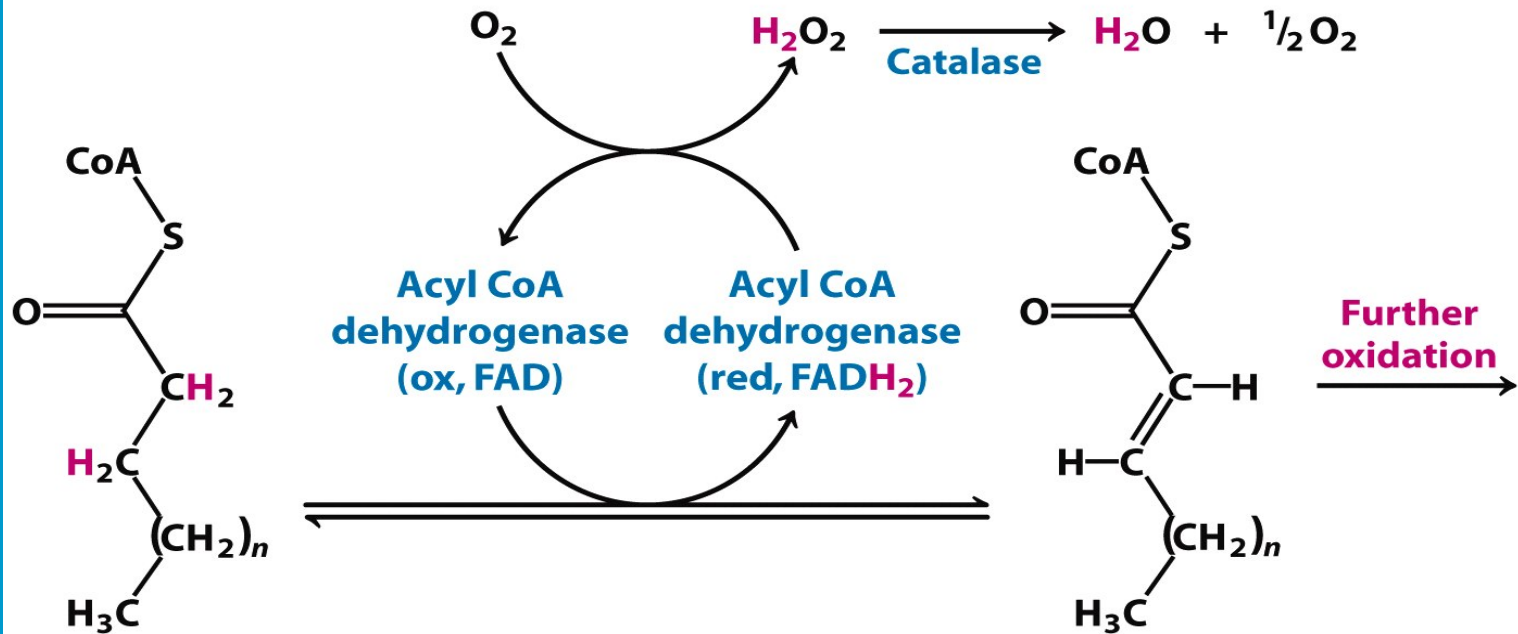
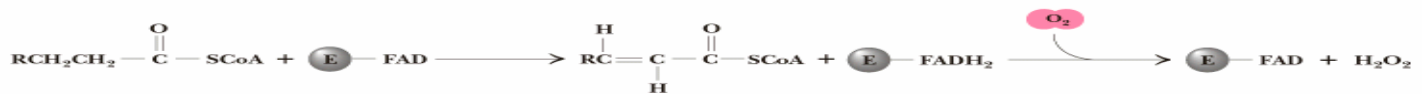


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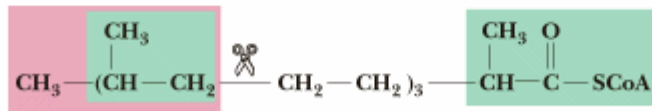
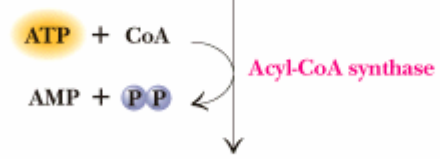
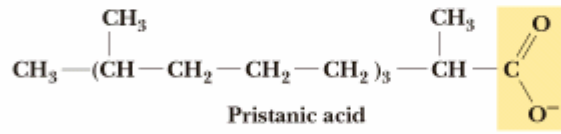
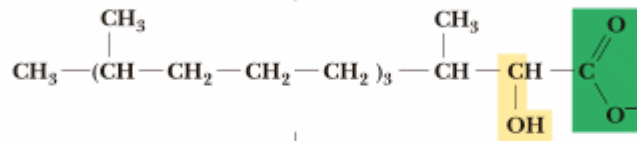
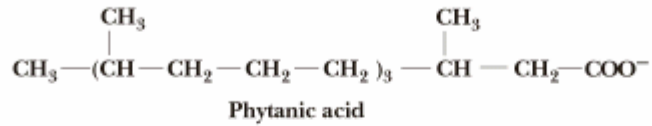
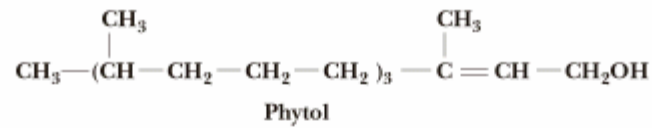
Garrett & Grisham: *Biochemistry*, 2/e
 Figure 24.25



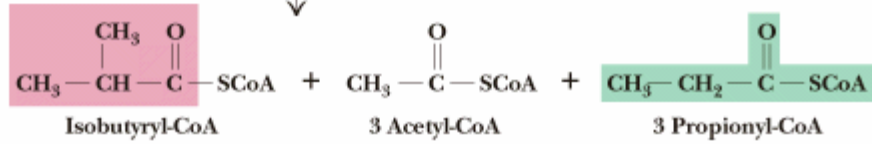
Branched-Chain Fatty Acids

An alternative to β -oxidation is required

- Branched chain FAs with branches at odd-number carbons are not good substrates for β -oxidation
- α -oxidation is an alternative
- **Phytanic acid α -oxidase** decarboxylates with oxidation at the alpha position
- β -oxidation occurs past the branch



Six cycles of β-oxidation

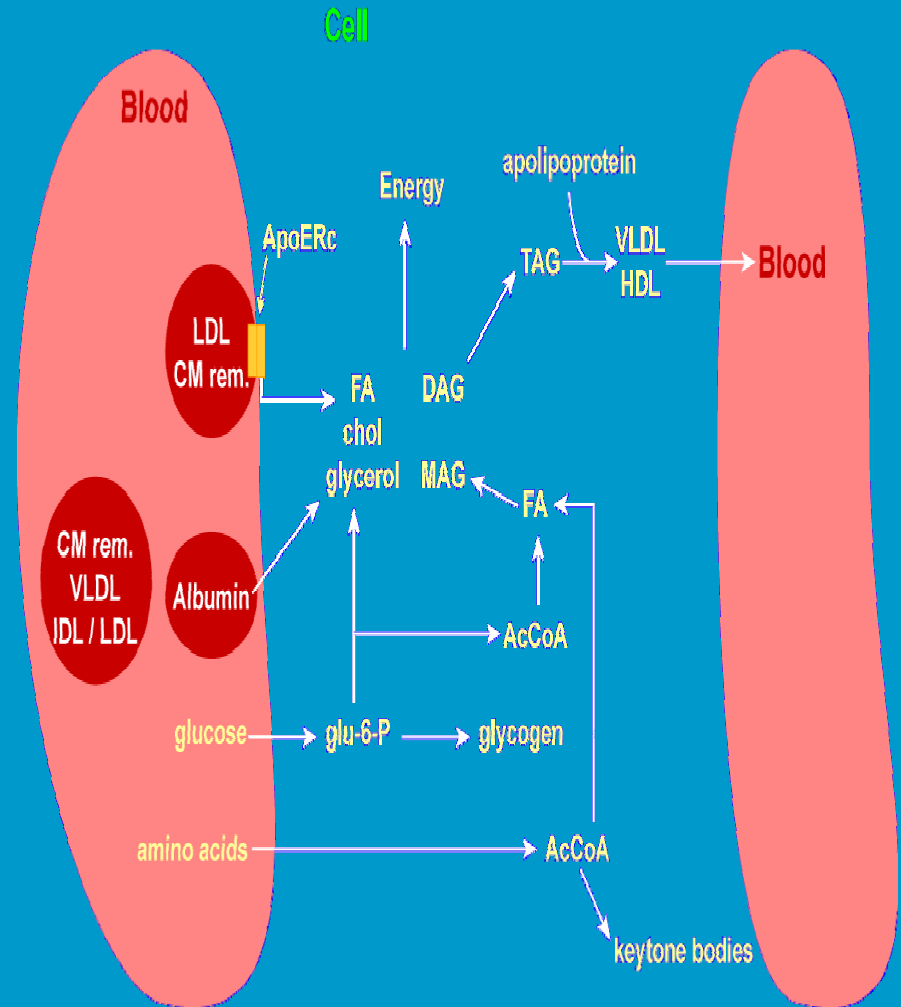


Liver Clearance of Lipids

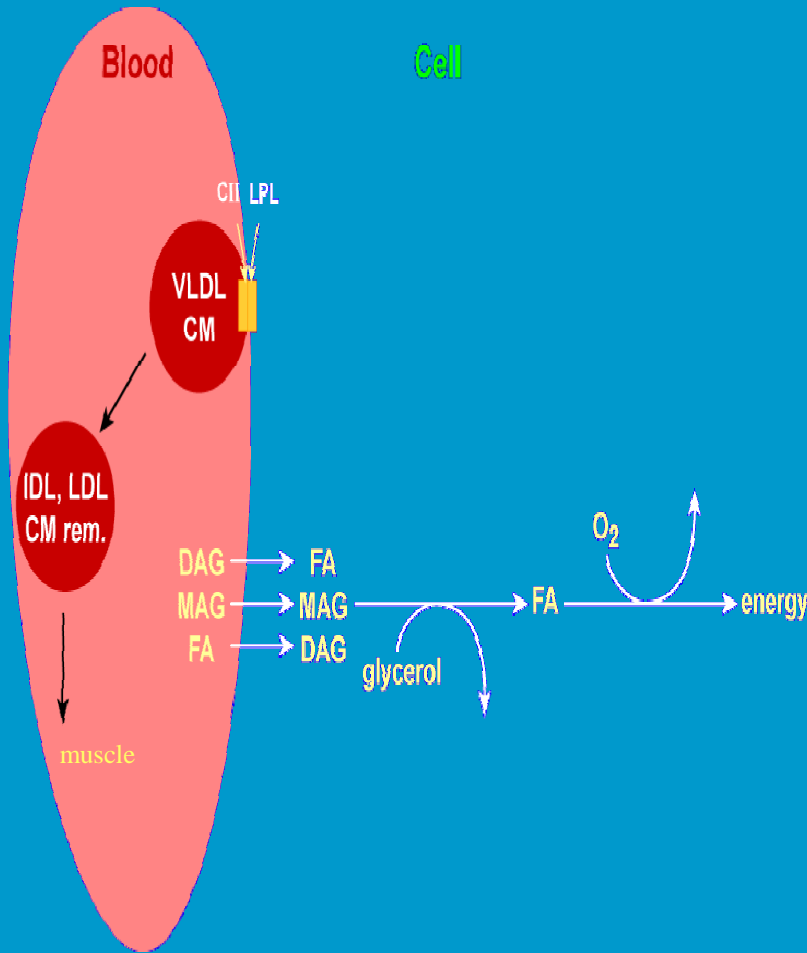
Liver is the major regulator of serum lipid levels. After a meal, it engulfs the chylomicron remnants and IDL particles, using its receptor for Apolipoprotein E. It also clears the short chain fatty acid bound with albumin. Once inside, the CM/IDL lipids are hydrolyzed to their constituent components: FFA, DAG, MAG, glycerol, and cholesterol.

Their fate in liver is to be repackaged into new lipoprotein particles (VLDL) and sent back into circulation as the TAG/FA/Cholesterol source between meals. The glycerol backbone of TAG comes from glycerol (via glycolysis); the fatty acids come from dietary fats, and from excess amino acids and glucose (via Acetyl-CoA, the fatty acid precursor).

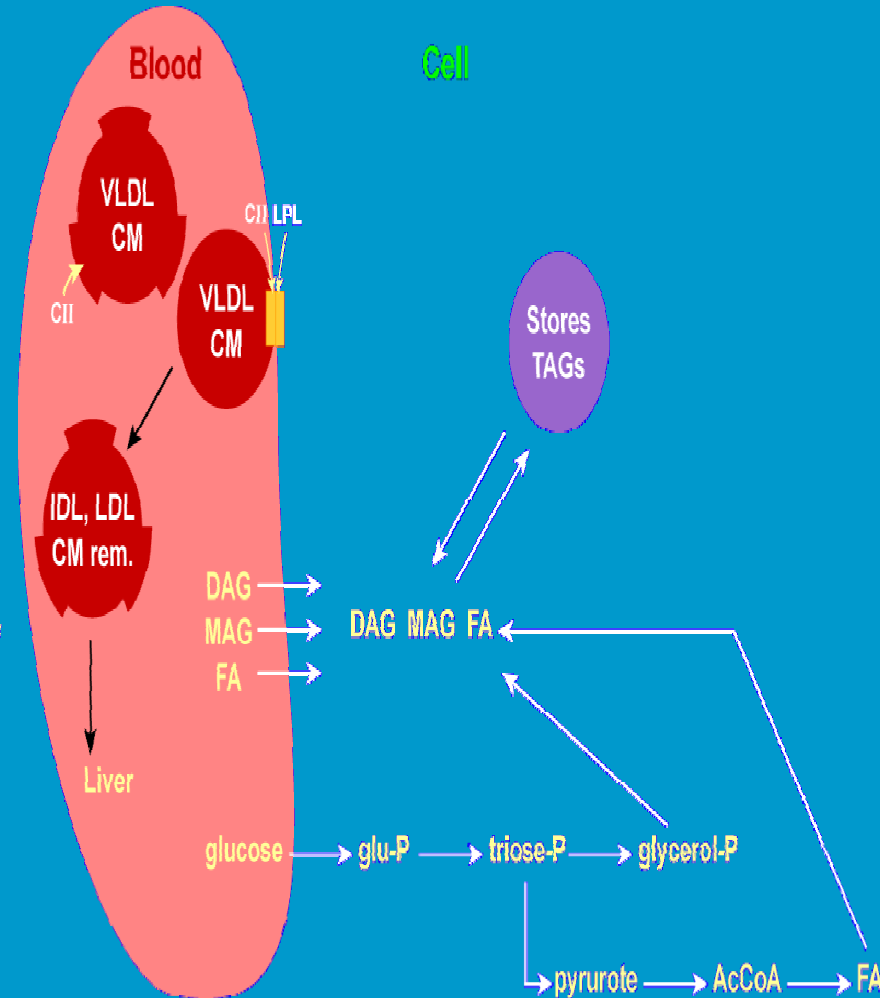
Liver is also the major regulator of the body's cholesterol. Liver takes up cholesterol from IDL or CM (not from LDL/HDL), and either repackages into VLDL lipoproteins or converts it into bile acids and excretes it out the gall bladder. Bile acids are the only way to remove cholesterol from the body (recall it's not oxidized for energy!)



Muscle Pathway



Adipose Pathway



Ketone Bodies

A special source of fuel and energy for certain tissues

- Some of the acetyl-CoA produced by fatty acid oxidation in liver mitochondria is converted to acetone, acetoacetate and β -hydroxybutyrate
- These are called "ketone bodies"
- Source of fuel for brain, heart and muscle
- Major energy source for brain during starvation
- They are transportable forms of fatty acids!

Ketone Bodies - II

Interesting Aspects of Their Synthesis

- Occurs only in the mitochondrial matrix
- First step - is reverse thiolase
- Second reaction makes **HMG-CoA**
- These reactions are mitochondrial analogues of the (cytosolic) first two steps of cholesterol synthesis
- Third step - HMG-CoA lyase - is similar to the reverse of citrate synthase

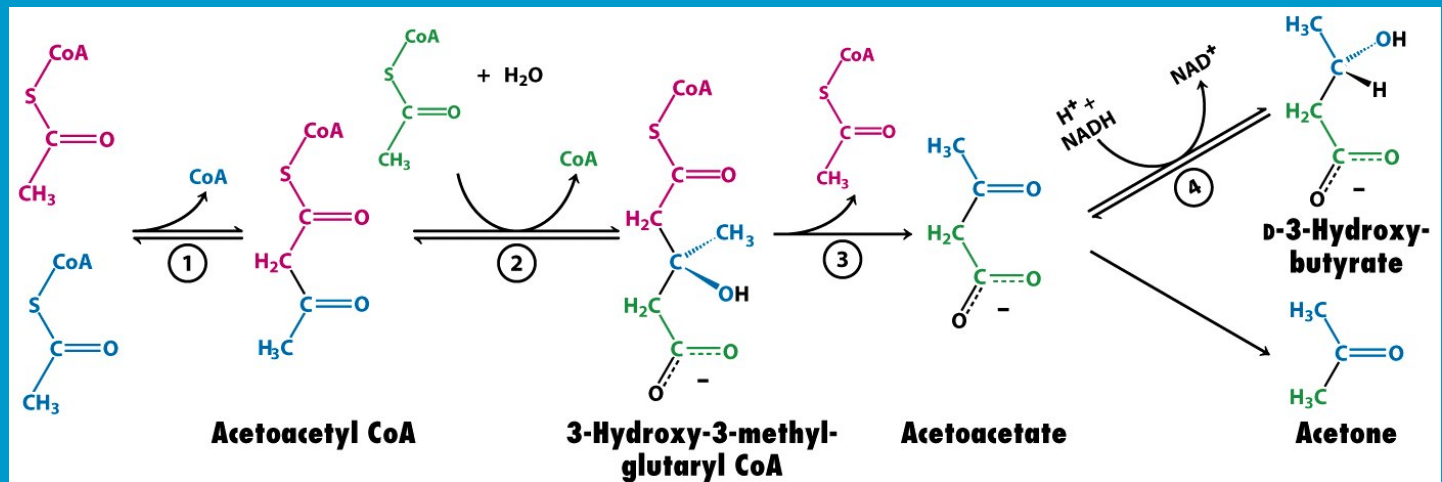
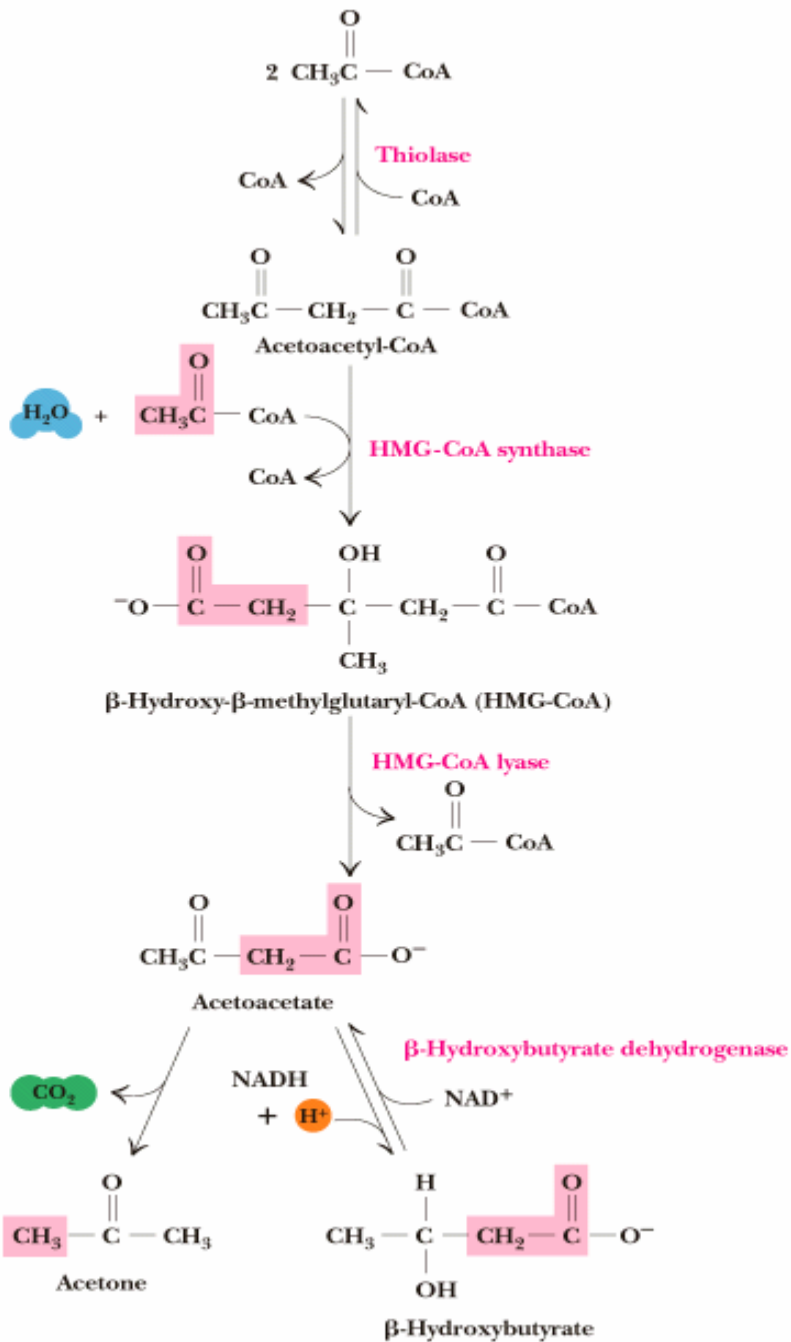


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Ketone Bodies and Diabetes

"Starvation of cells in the midst of plenty"

- Glucose is abundant in blood, but uptake by cells in muscle, liver, and adipose cells is low
- Cells, metabolically starved, turn to gluconeogenesis and fat/protein catabolism
- In type I diabetics, OAA is low, due to excess gluconeogenesis, so Ac-CoA from fat/protein catabolism does not go to TCA, but rather to ketone body production
- Acetone can be detected on breath of type I diabetics

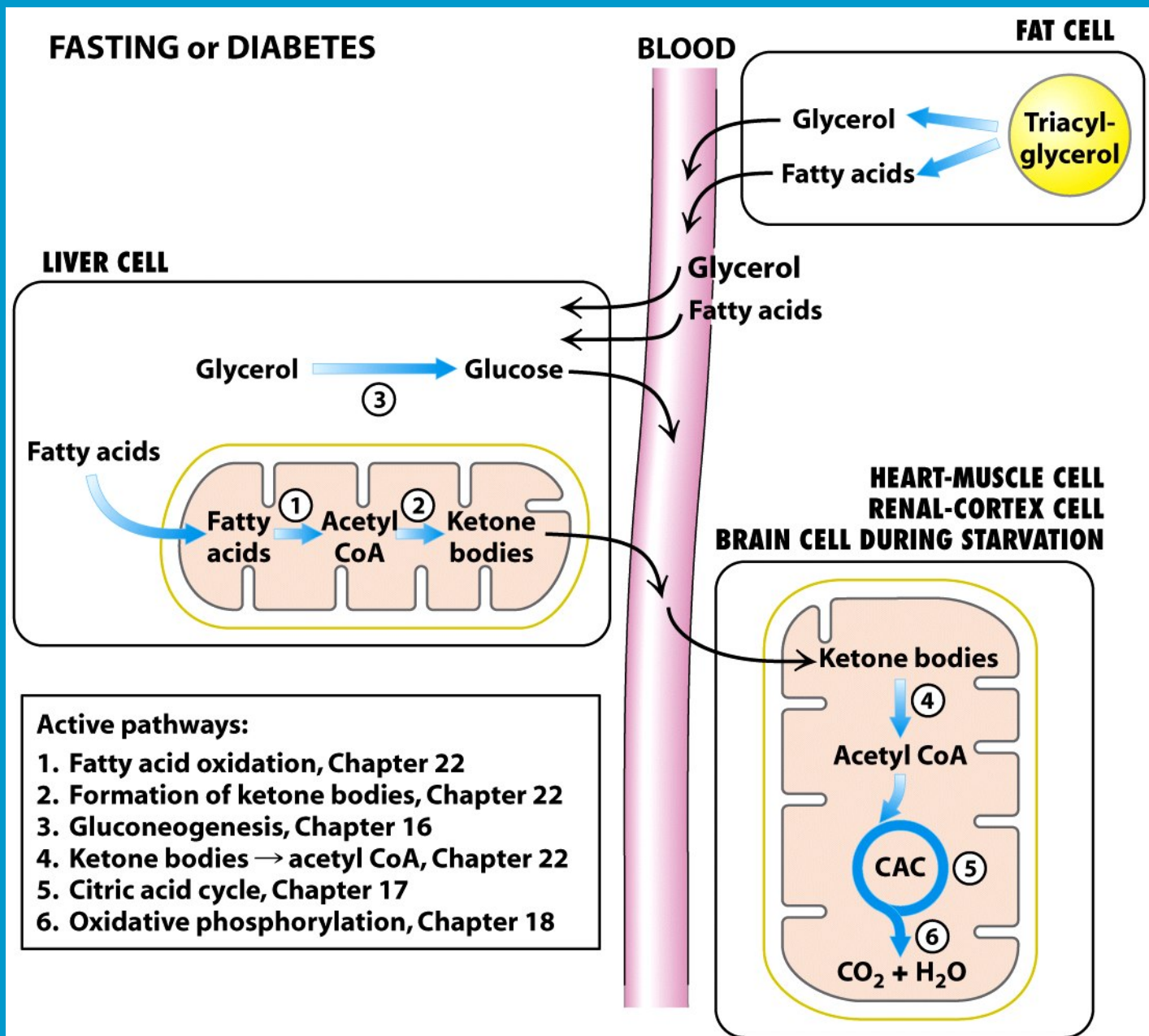


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Utilization of acetoacetate as fuel

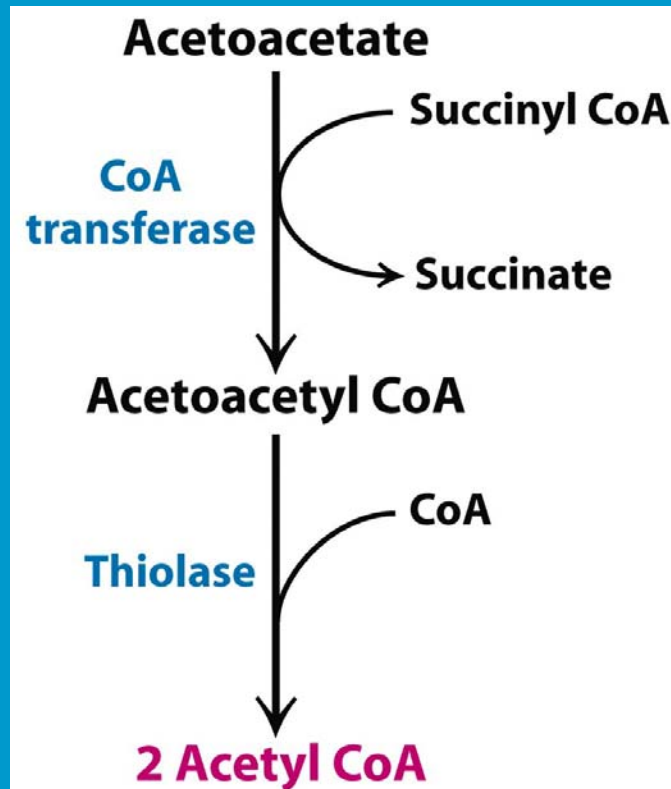


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Diabetic ketosis results in the absence of insulin

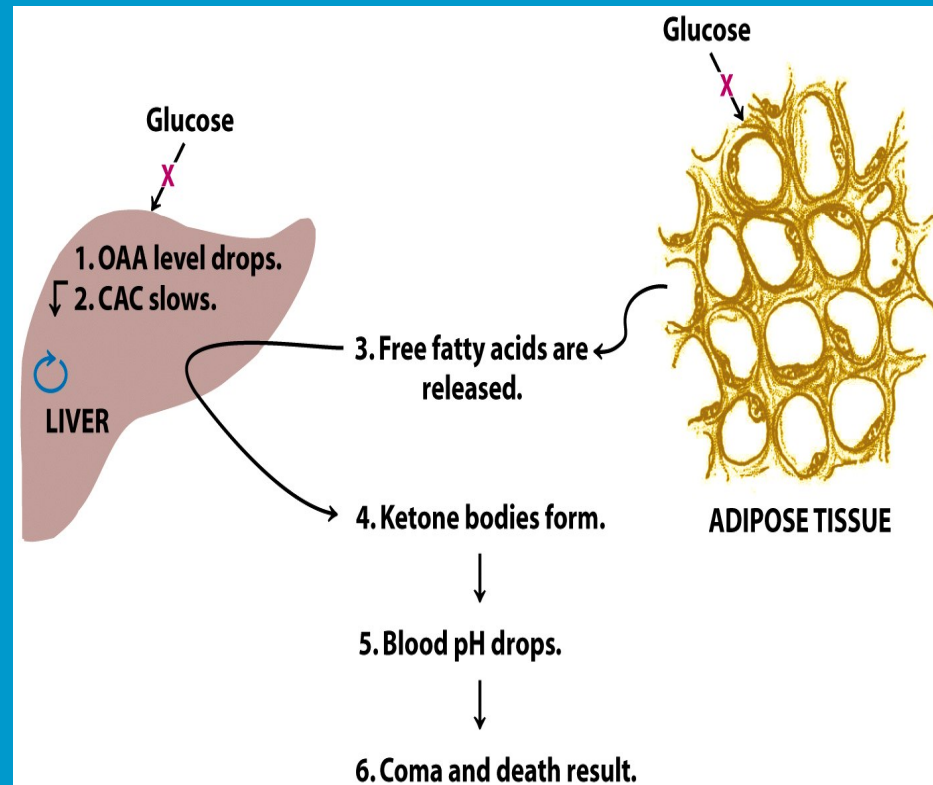


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