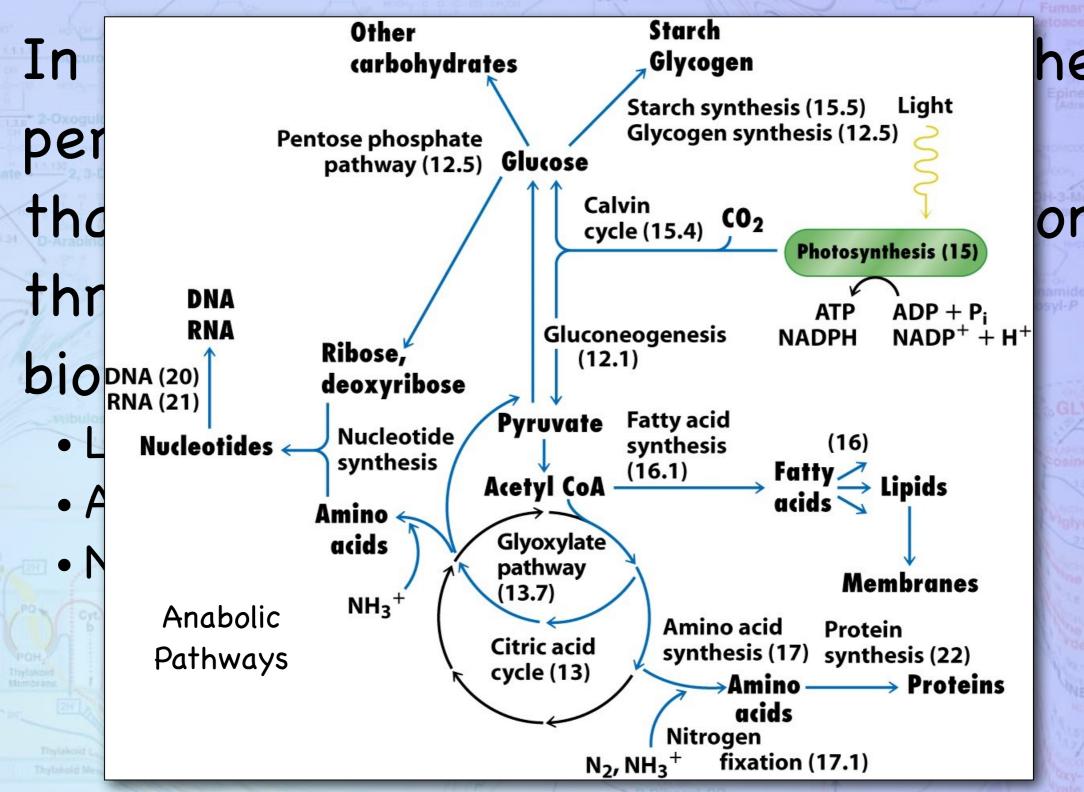


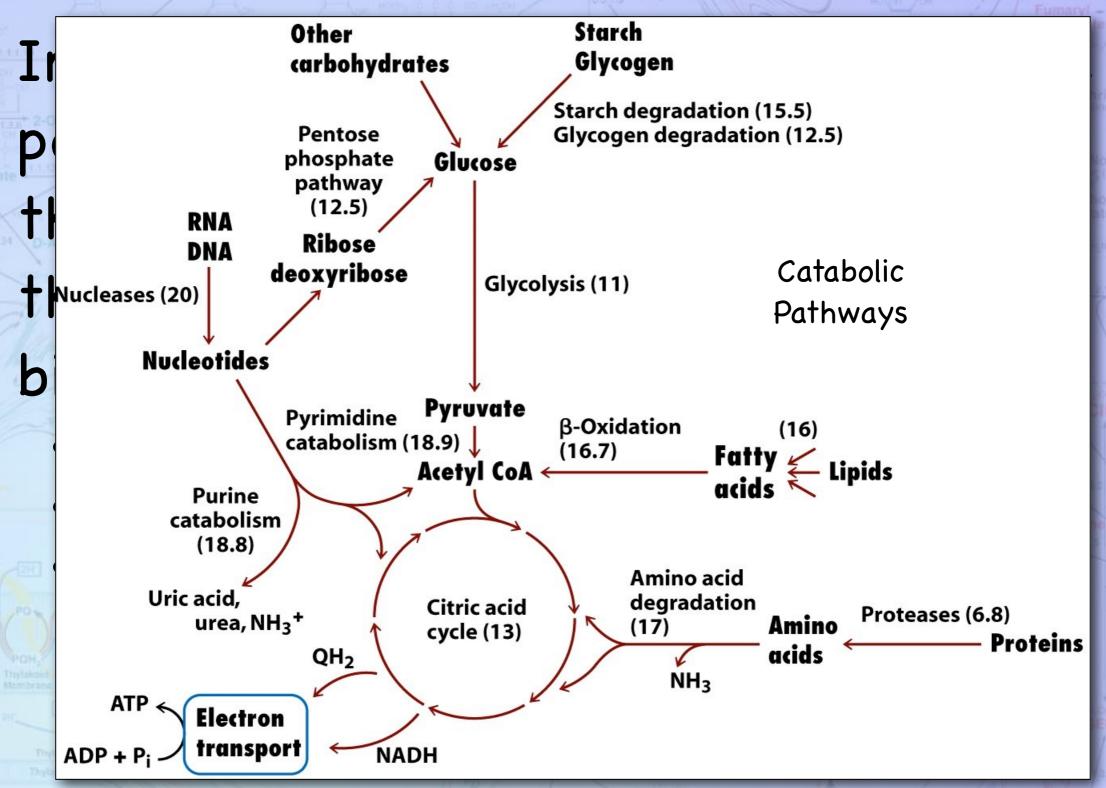
In Lecture 10 we will add some of the peripheral pathways in metabolism that lead to synthesis and degradation three important classes of biomolecules:

- Lipids
- Amino acids
- Nucleotides



In Lecture 10 we will add some of the peripheral pathways in metabolism that lead to synthesis and degradation three important classes of biomolecules:

- Lipids
- Amino acids
- Nucleotides



In Lecture 10 we will add some of the peripheral pathways in metabolism that lead to synthesis and degradation three important classes of biomolecules:

- Lipids
- Amino acids
- Nucleotides

Lipids play many important cellular roles

- Membrane components (phospholipids, et al. and cholesterol)
- · Fuels (Triacylglycerides)
 - · Meet long term energy needs in mammals
- · Regulators (steroids, eicosanoids)

We will focus on just a couple of key metabolic pathways.

We will focus on the following sections from Chapter 16

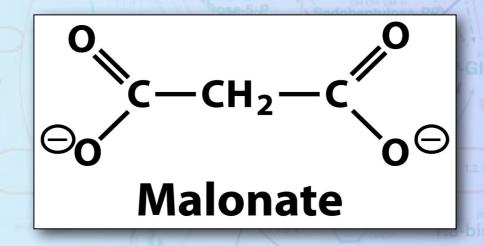
- 16.1: Fatty Acid Synthesis
- 16.6: Synthesis of Cholesterol
- 16.7: Fatty Acid Oxidation
- 16.9: Lipid Metabolism is Regulated by Hormones in Mammals
- 16.10: Absorption and Mobilization of Fuel Lipids in Mammals
- 16.11: Ketone Bodies Are Fuel Molecules

Fatty acids are synthesized by the repetitive addition of 2 carbon units to a growing chain.

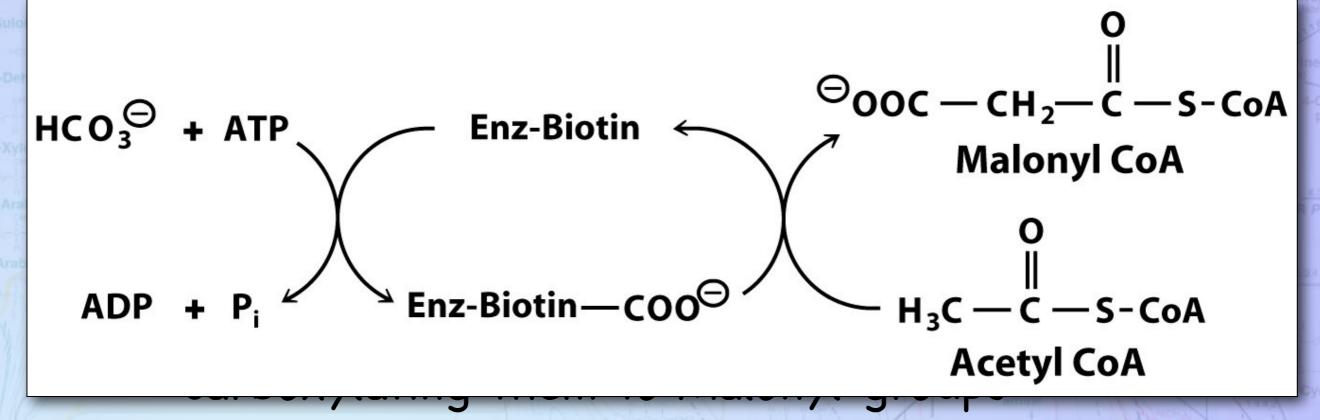
- Acetyl-CoA is the source of the 2 carbon units.
- The Acetyl groups are activated by carboxylating them to Malonyl groups

Fatty acids are synthesized by the repetitive addition of 2 carbon units to a growing chain.

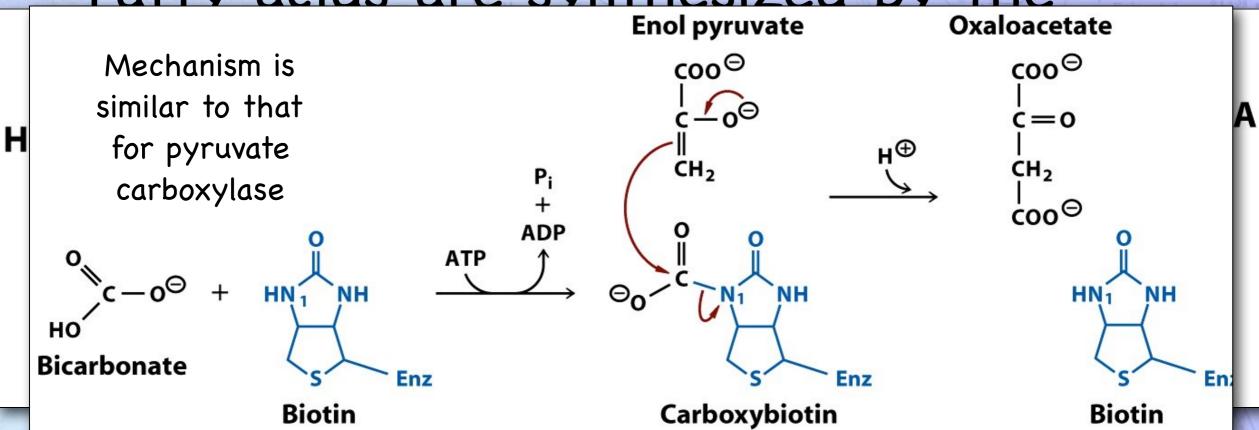
- Acetyl-CoA is the source of the 2 carbon units.
- The Acetyl groups are activated by carboxylating them to Malonyl groups



Fatty acids are synthesized by the



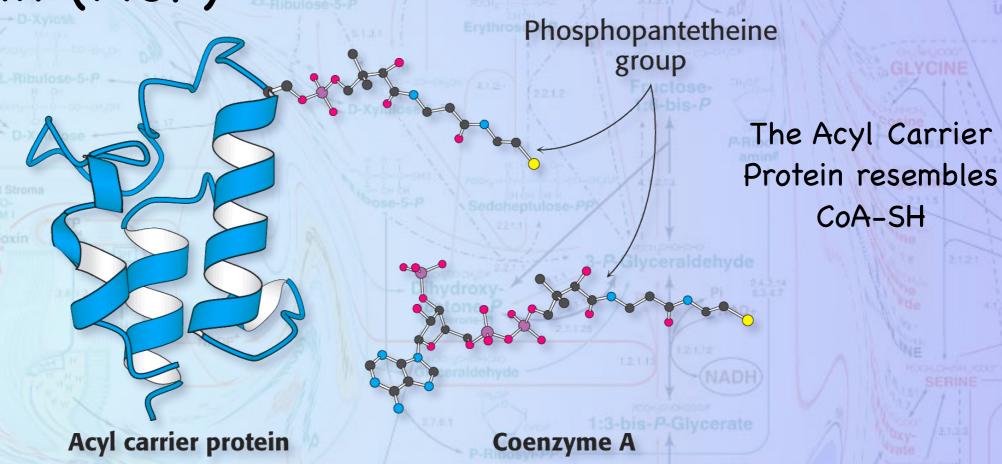
Fatty acids are synthesized by the



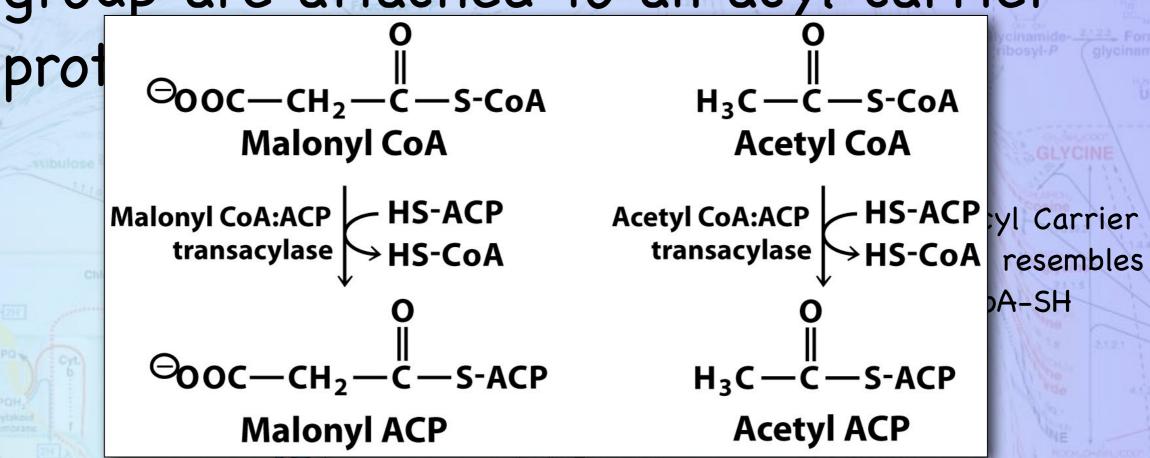
Fatty acids are synthesized by the repetitive addition of 2 carbon units to a growing chain.

- Acetyl-CoA is the source of the 2 carbon units.
- The Acetyl groups are activated by carboxylating them to Malonyl groups

In eukaryotes, both the growing chain, the acetyl group, and the malonyl group are attached to an acyl carrier protein (ACP)



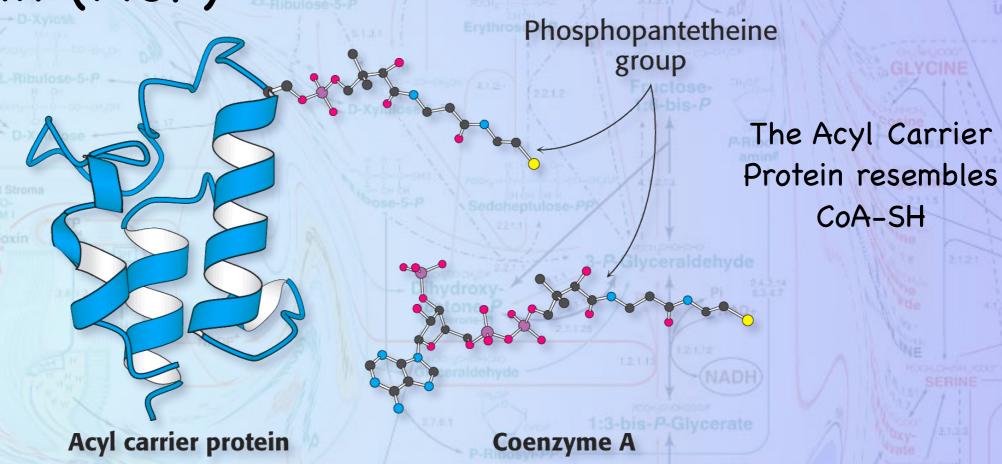
In eukaryotes, both the growing chain, the acetyl group, and the malonyl group are attached to an acyl carrier



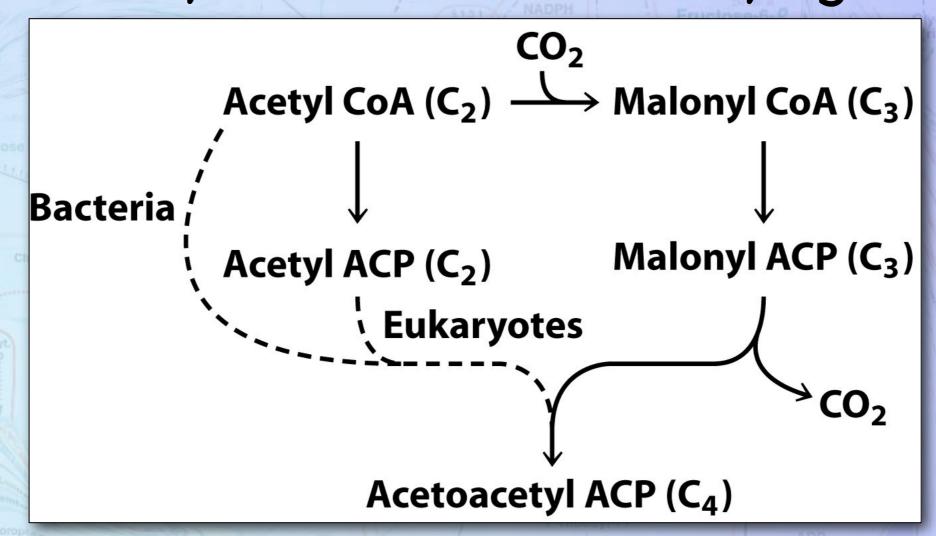
Acyl carrier protein

Coenzyme A

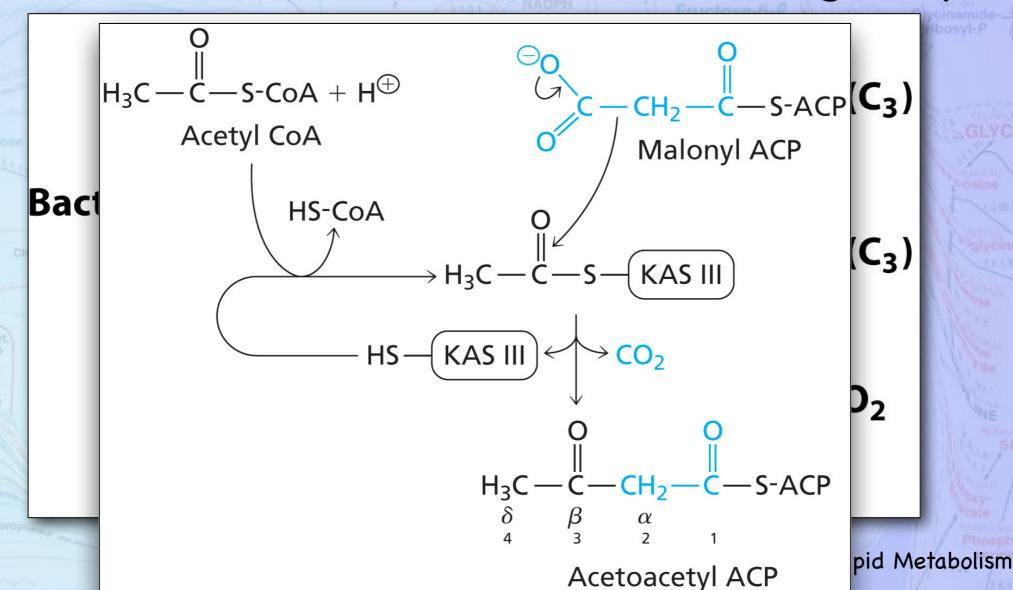
In eukaryotes, both the growing chain, the acetyl group, and the malonyl group are attached to an acyl carrier protein (ACP)



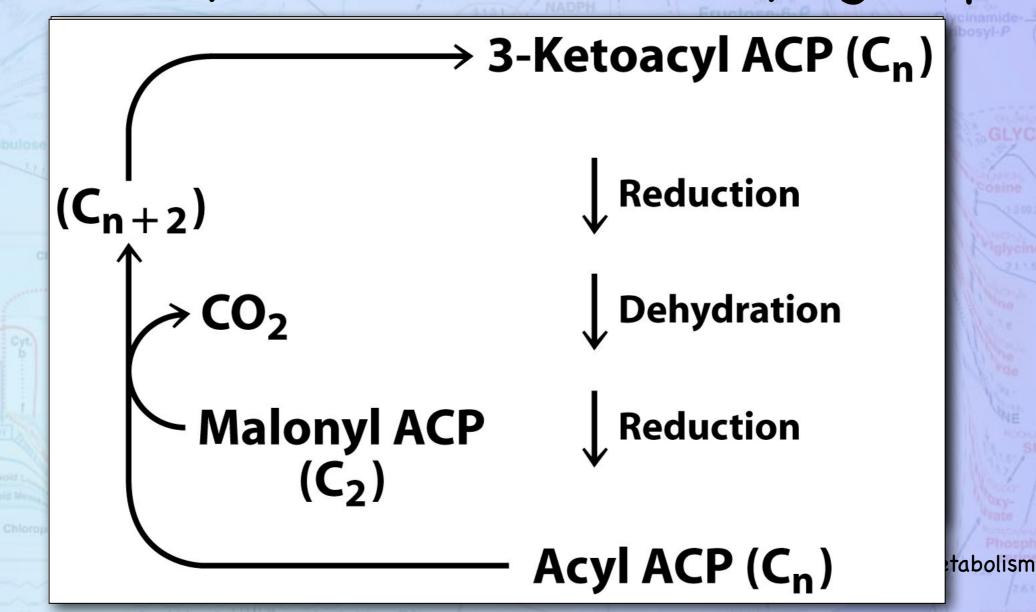
The ligation of an acetyl group to the growing chain is coupled to the decarboxylation of the malonyl group.



The ligation of an acetyl group to the growing chain is coupled to the decarboxylation of the malonyl group.



The ligation of an acetyl group to the growing chain is coupled to the decarboxylation of the malonyl group.



The reduction/dehydration/reduction steps similar to a series of reactions found in the citric acid cycle, but in reverse order.

$$\begin{bmatrix} O & OH & H \\ \| & | & | \\ R_1-C-CH_2-R_2 & \longrightarrow R_1-C-CH_2-R_2 & \longrightarrow R_1-C=C-R_2 & \longrightarrow R_1-CH_2-CH_2-R_2 \\ | & | & | & | \\ Reduction & Dehydration & Reduction \end{bmatrix}$$

The reduction/dehydration/reduction $R - C - CH_2 - C - S - ACP$ steps s 3 Ketoactyl ACP Elongation Stage found i reductase

D-3-Hydroxyacyl ACP

3-Hydroxyacyl-ACP dehydratase
$$\rightarrow$$
 H₂O
$$R-C=\begin{matrix} H & 0 \\ C & C - S-ACP \end{matrix}$$

 $R_1 - C - CH_2 - R_2$

Red

trans-∆2-Enoyl ACP

Enoyl-ACP NADPH + H
$$\oplus$$
 NADP \oplus NADP \oplus NADP \oplus Acyl-ACP

> CO2 3-Ketoactyl-ACP synthase

 Θ_{OOC-CH_2} - \ddot{C}_{-S-ACP} R_1 - CH_2 - CH_2 - R_2

tions

The reduction/dehydration/reduction steps similar to a series of reactions found in the citric acid cycle, but in reverse order.

$$\begin{bmatrix} O & OH & H \\ \| & | & | \\ R_1-C-CH_2-R_2 & \longrightarrow R_1-C-CH_2-R_2 & \longrightarrow R_1-C=C-R_2 & \longrightarrow R_1-CH_2-CH_2-R_2 \\ | & | & | & | \\ Reduction & Dehydration & Reduction \end{bmatrix}$$

The elongation continues until reaching 16 carbons (palmitic acid).

 The palmitoyl group is cleaved from the ACP by a thioesterase.

The elongation continues until reaching 16 carbons (palmitic acid).

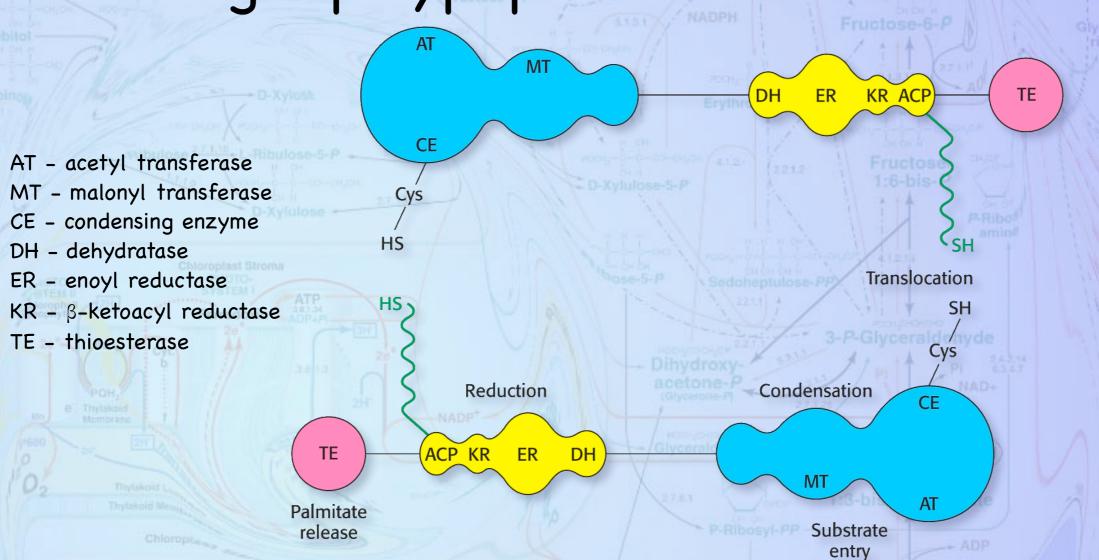
 The palmitoyl group is cleaved from the ACP by a thioesterase.

Acetyl CoA + 7 Malonyl CoA + 14 NADPH + 14 H
$$^{\oplus}$$
 \longrightarrow Palmitate+ 7 CO₂ + 14 NADP $^{\oplus}$ + 8 HS-CoA + 6 H₂O

The elongation continues until reaching 16 carbons (palmitic acid).

 The palmitoyl group is cleaved from the ACP by a thioesterase.

In eukaryotes, all of the active sites for fatty acid synthesis are located on a single polypeptide.



Chem 352, Lecture 10, Part I: Lipid Metabolism

In eukaryotes, all of the active sites for fatty acid synthesis are located on

a single

AT - acetyl transferase

MT - malonyl transferase

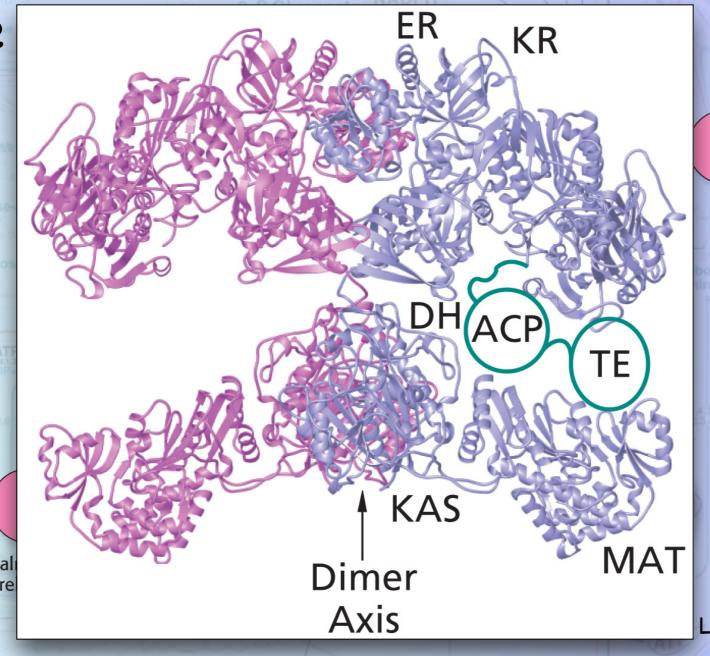
CE - condensing enzyme

DH - dehydratase

ER - enoyl reductase

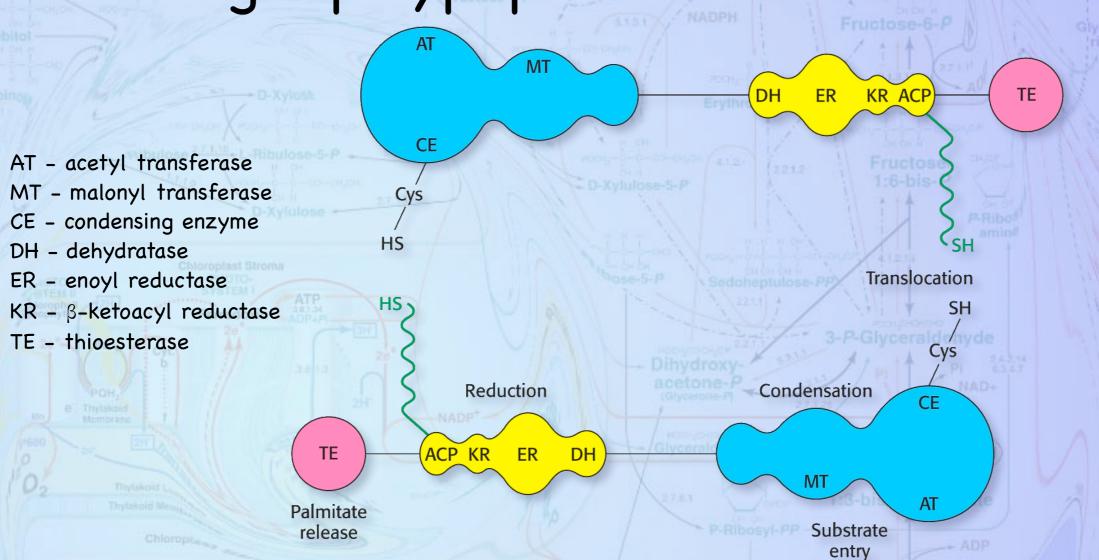
 $KR - \beta$ -ketoacyl reductase

TE - thioesterase



TE

In eukaryotes, all of the active sites for fatty acid synthesis are located on a single polypeptide.



Chem 352, Lecture 10, Part I: Lipid Metabolism

In eukaryotes, all of the active sites

for fatty Spiral Pathway a single

AT - acetyl transferase

MT - malonyl transferase

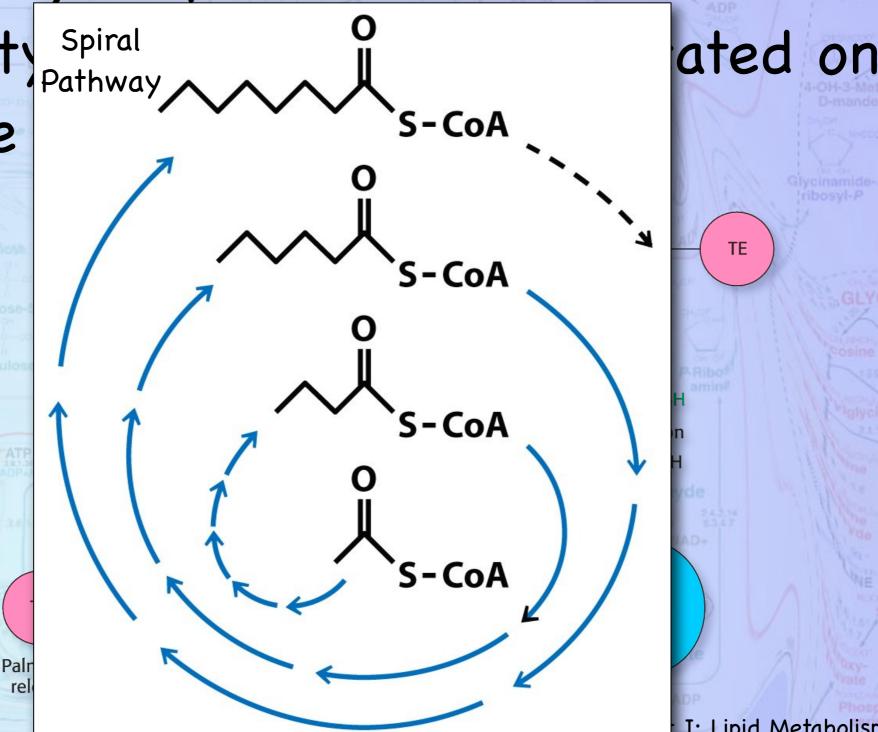
CE - condensing enzyme

DH - dehydratase

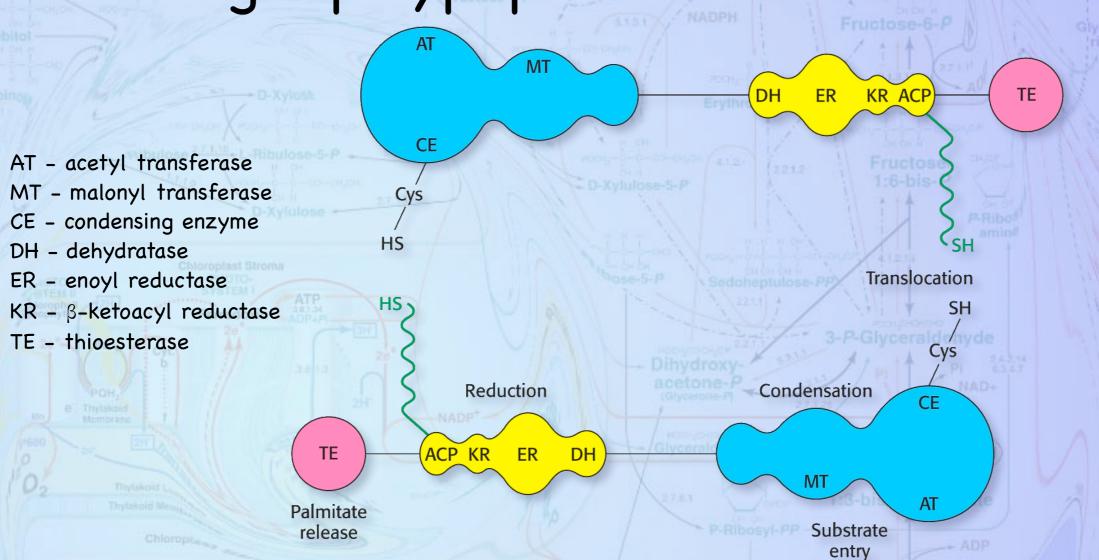
ER - enoyl reductase

KR - β-ketoacyl reductase

TE - thioesterase

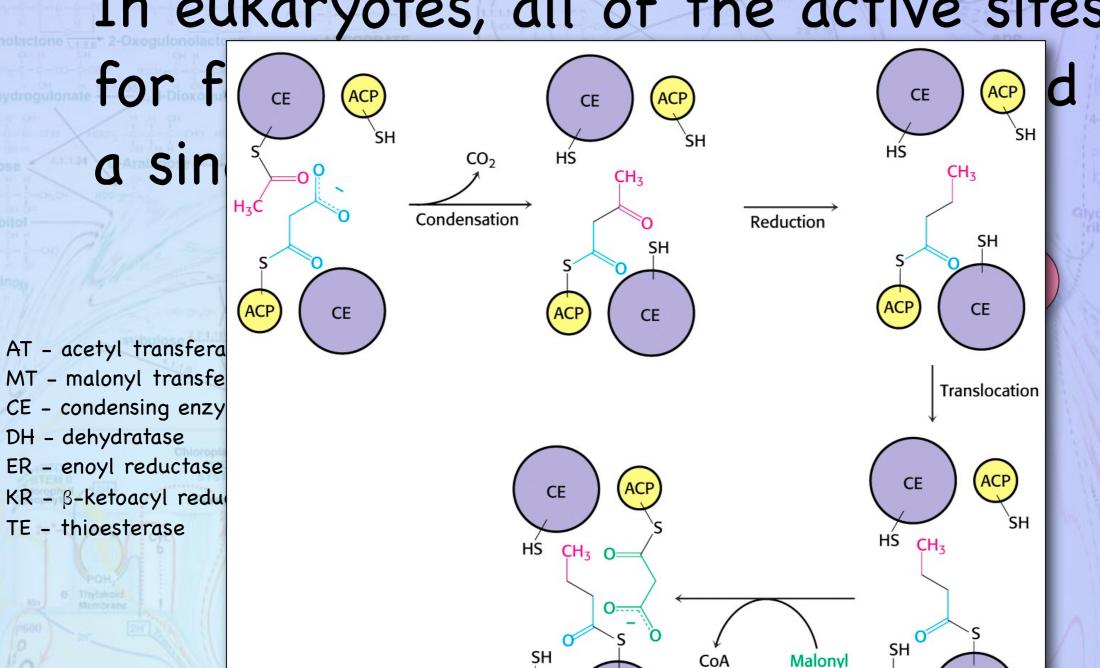


In eukaryotes, all of the active sites for fatty acid synthesis are located on a single polypeptide.



Chem 352, Lecture 10, Part I: Lipid Metabolism

In eukaryotes, all of the active sites

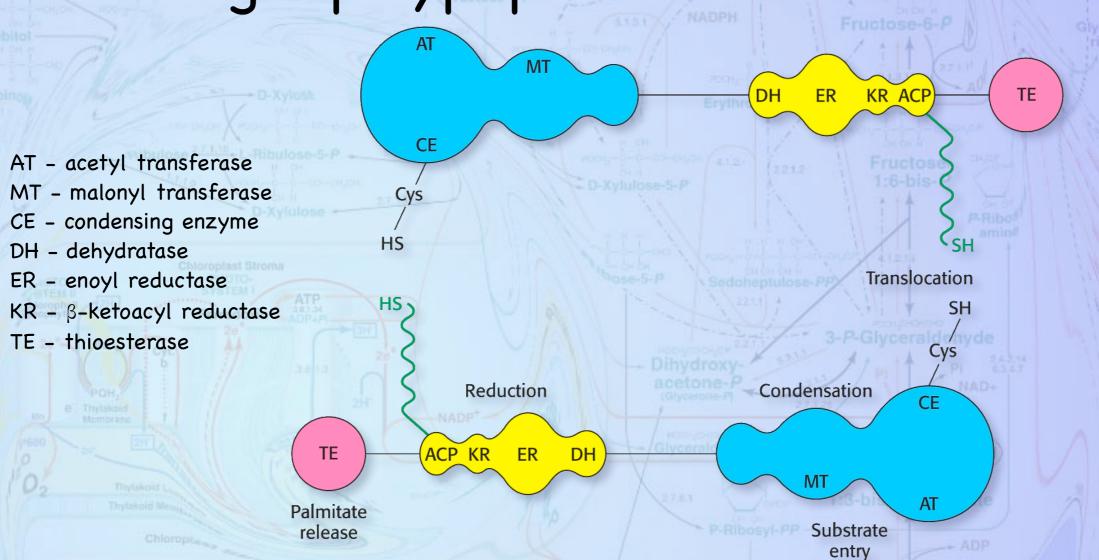


CE

CE

CoA

In eukaryotes, all of the active sites for fatty acid synthesis are located on a single polypeptide.



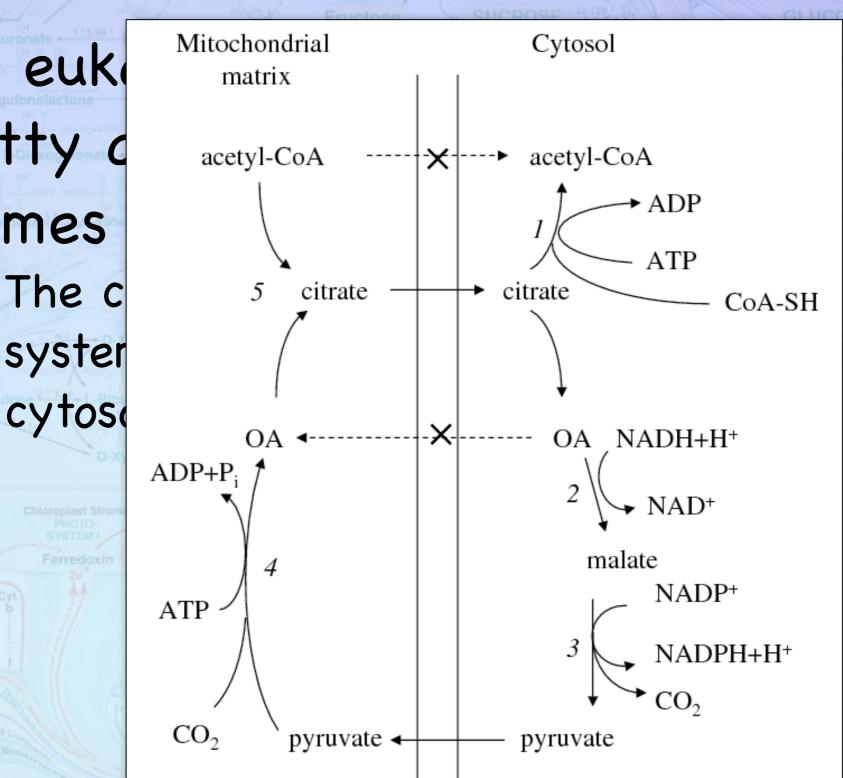
Chem 352, Lecture 10, Part I: Lipid Metabolism

After palmitic acid is synthesized it is elongated and desaturated to form other fatty acids

In eukaryotes, the acetyl-CoA for fatty acid synthesis in the cytosol comes from the mitochonrial matrix.

• The citrate/pyruvate shuttle is one of the systems used to move acetyl-CoA out into the cytosol.

In euk fatty o comes • The c syster



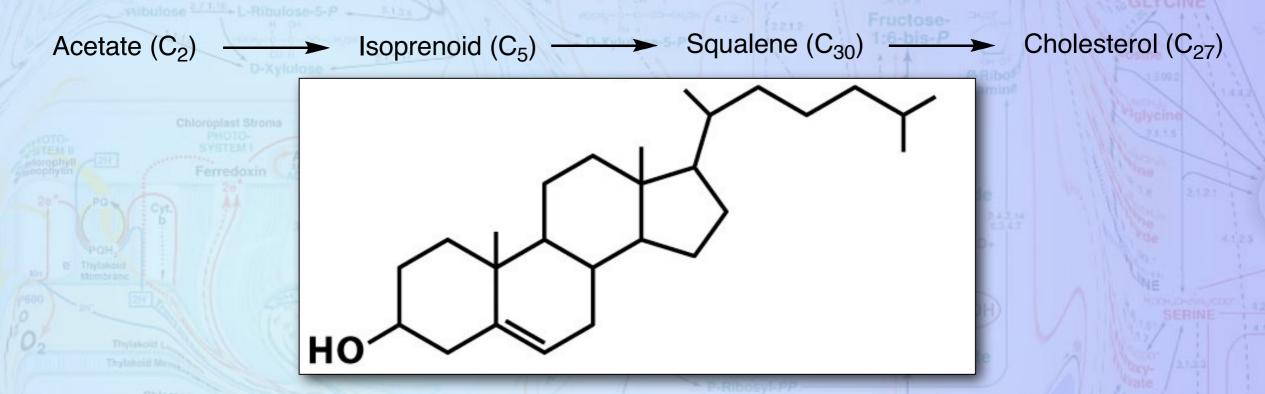
rix. the into the

In eukaryotes, the acetyl-CoA for fatty acid synthesis in the cytosol comes from the mitochonrial matrix.

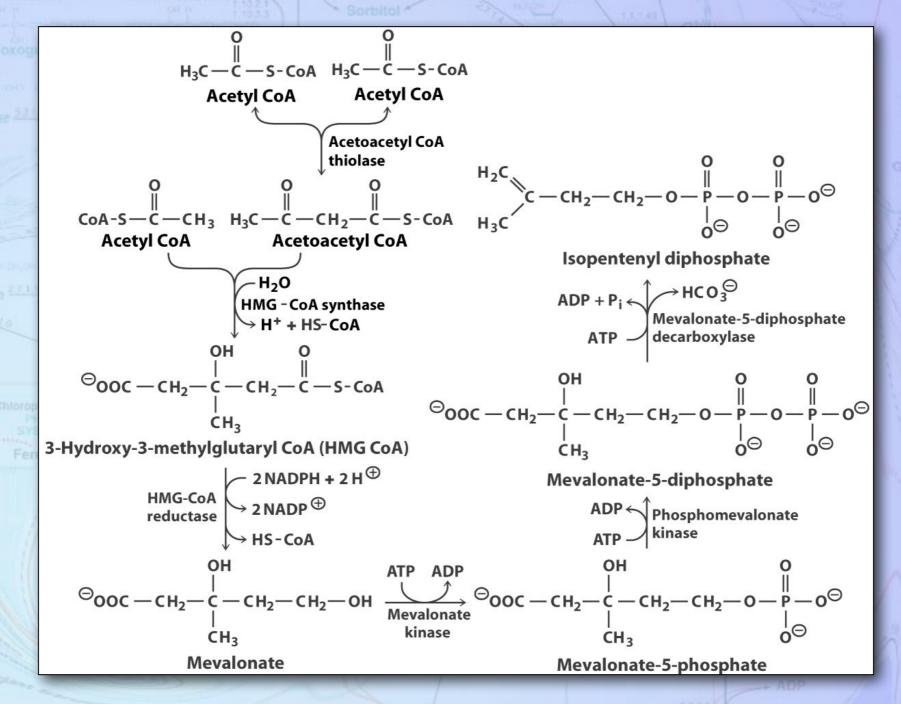
• The citrate/pyruvate shuttle is one of the systems used to move acetyl-CoA out into the cytosol.

Cholesterols is used to modulate the physical properties of membranes in animals

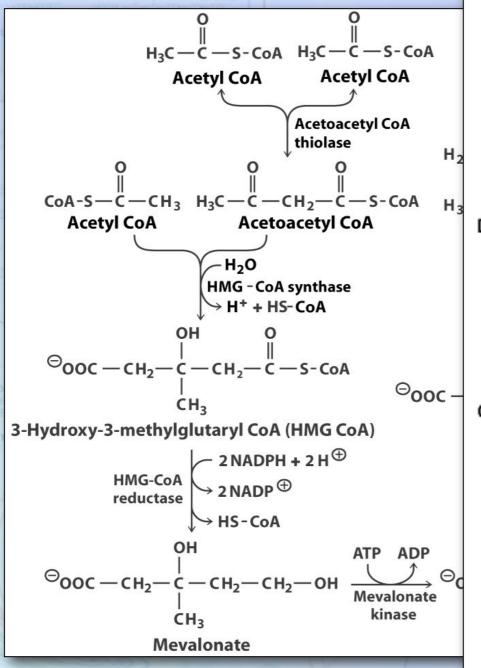
 It is also the starting point for the synthesis of all other steroid molecules

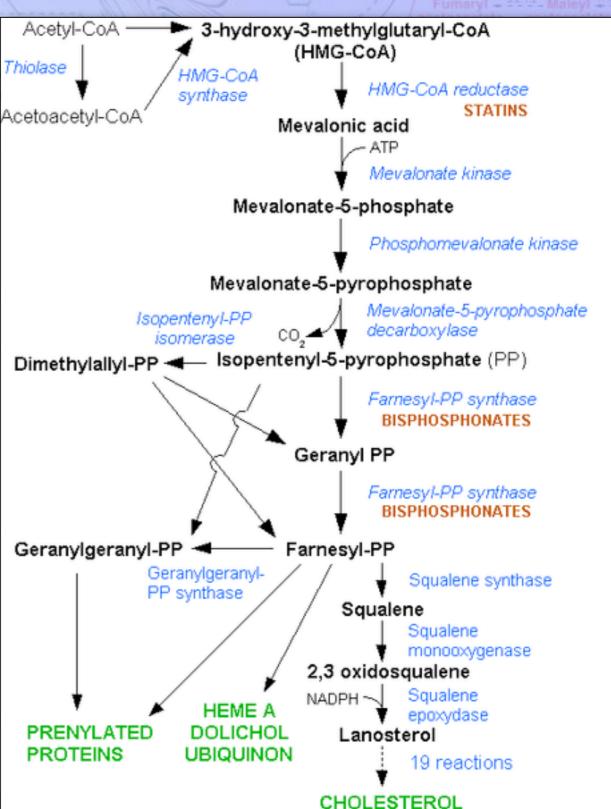


Acetate to Isoprenoid

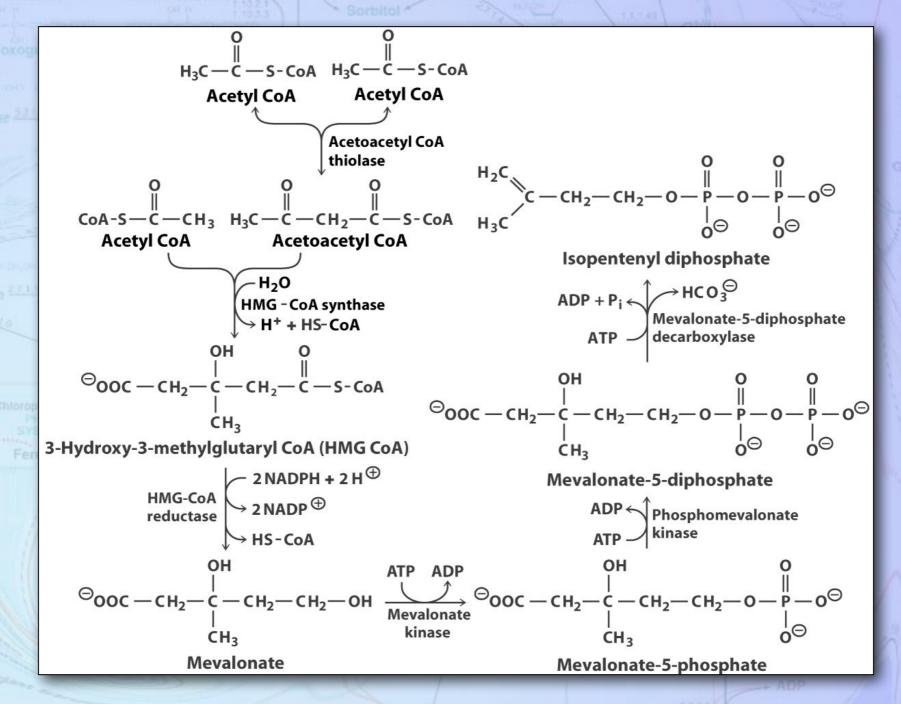


Acetate to Isoprenoic





Acetate to Isoprenoid



Statin (anticholesterol) drugs.

Isoprenoid to Squalene

Chem 352, Lecture 10, Part I: Lipid Metabolism

16

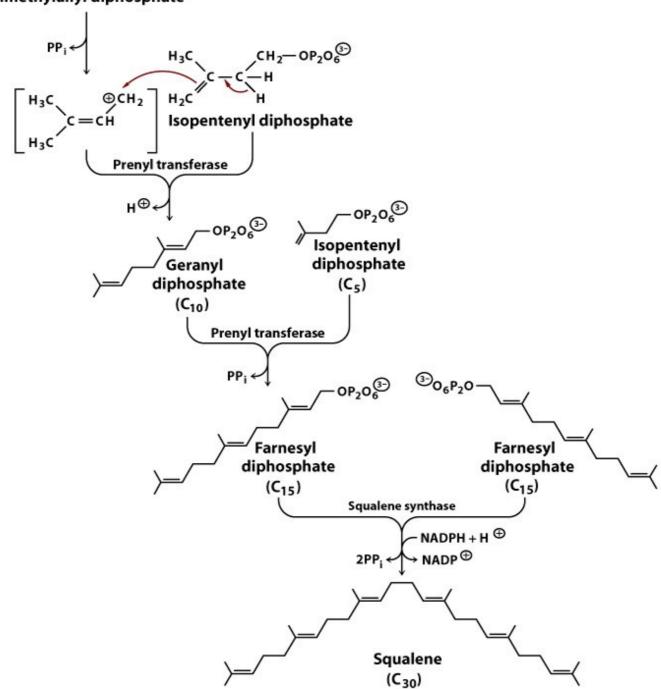
Chole H₃C C-CH₂-CH₂-O-

Isopentenyl diphosphate

Isopentenyl diphosphate

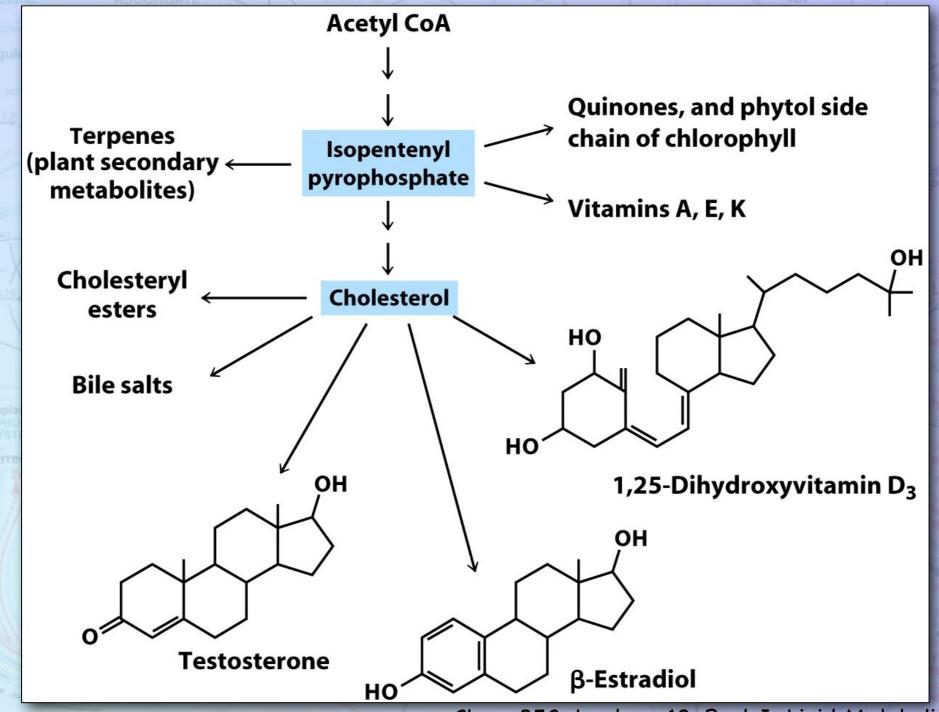
Isopre

Dimethylallyl diphosphate



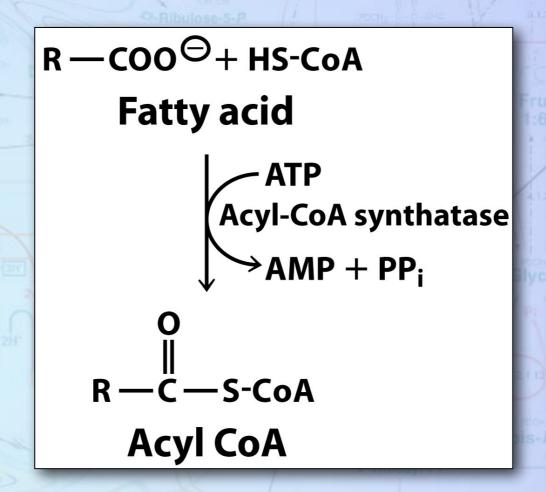
Squalene to Cholesterol

Cholesterol to other steroids

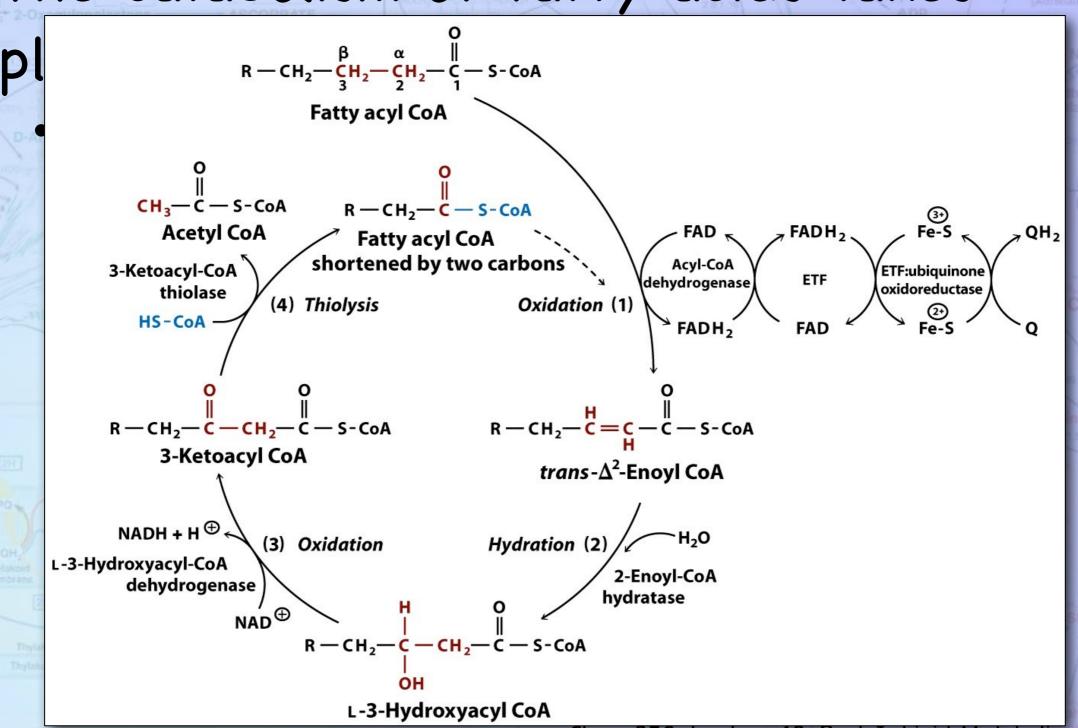


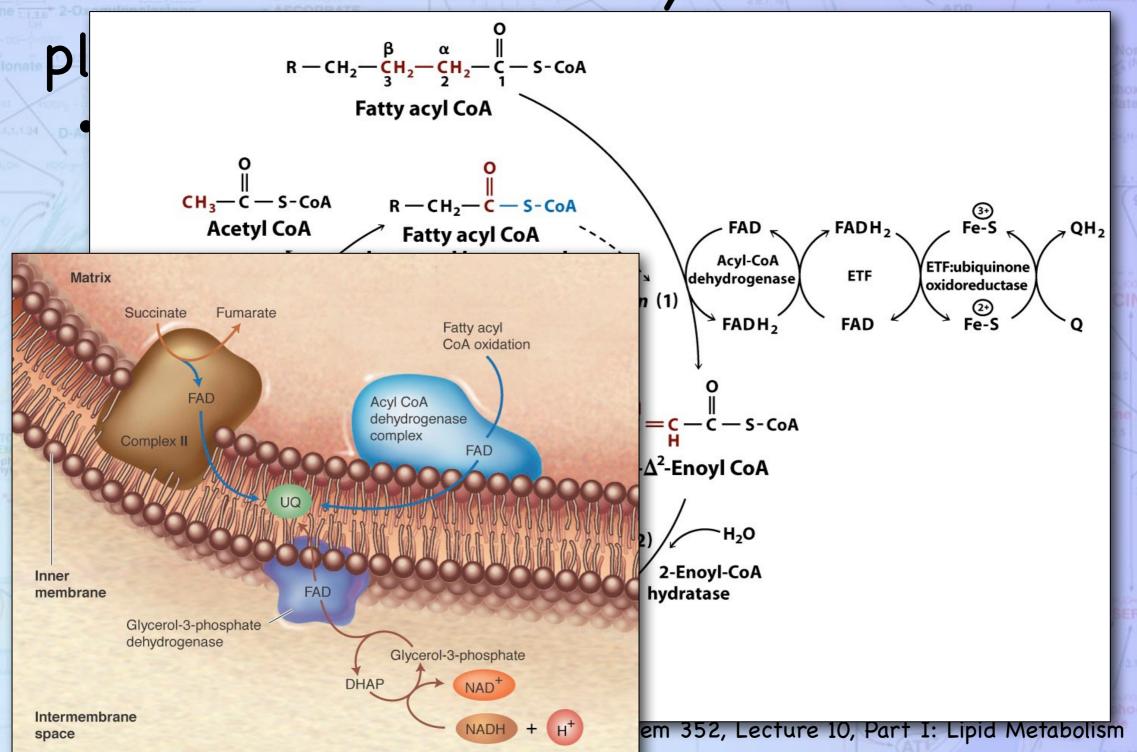
The catabolism of fatty acids takes place in the mitochondria

The fatty acid must first be activated to an acyl-CoA

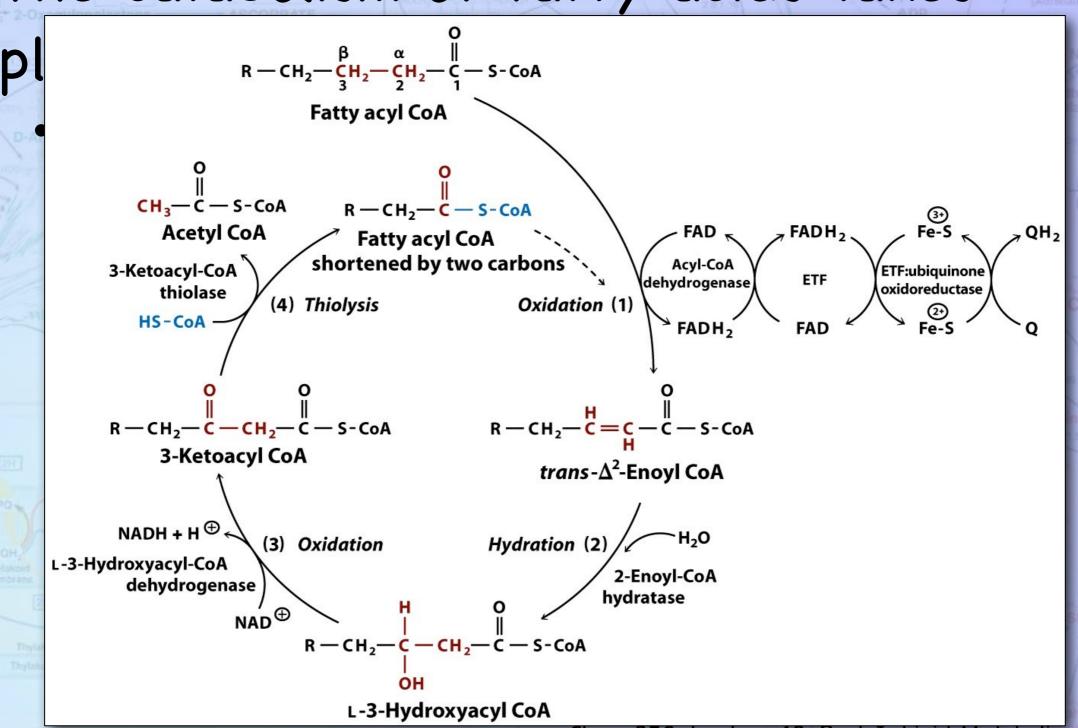


The catabolism of fatty acids takes

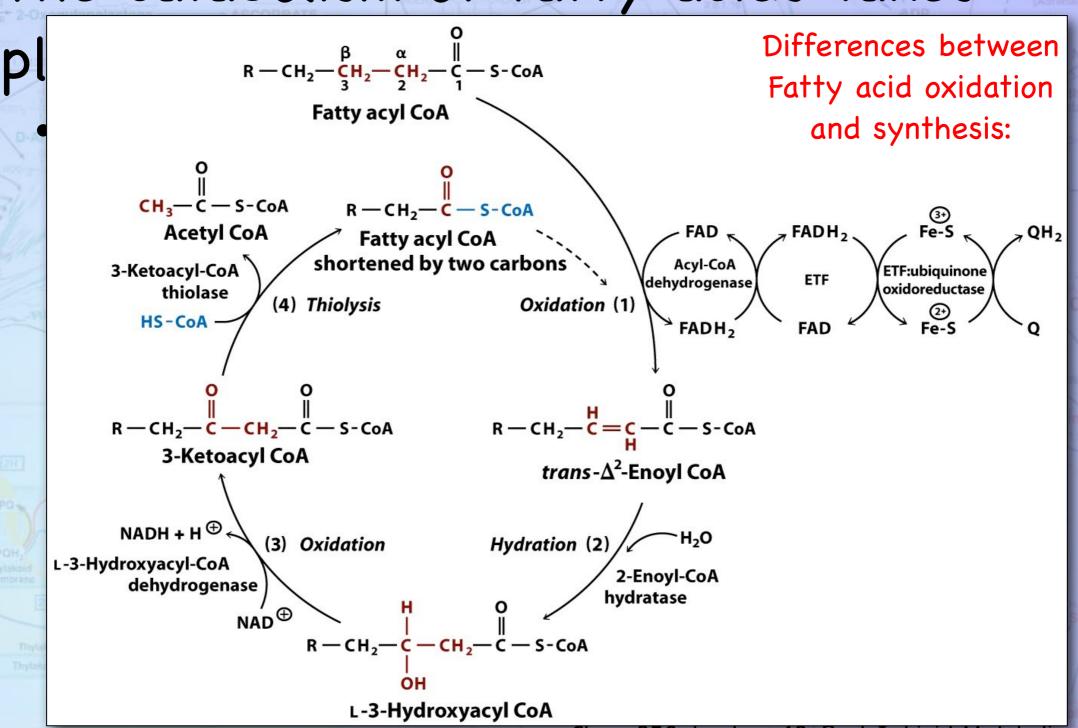




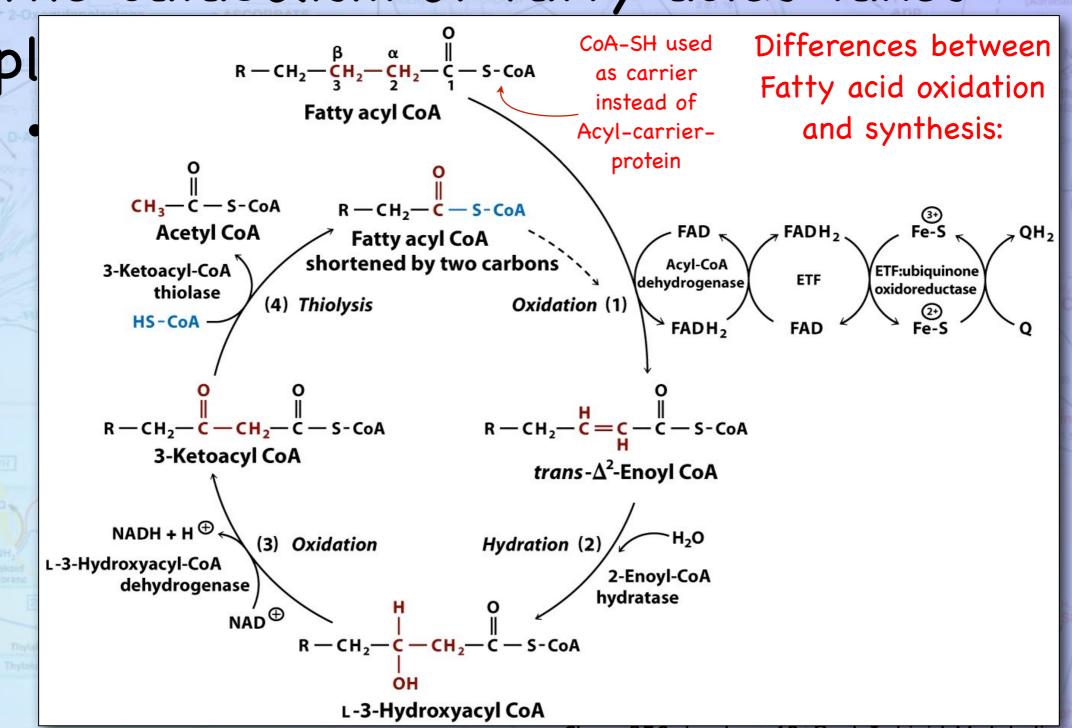
The catabolism of fatty acids takes

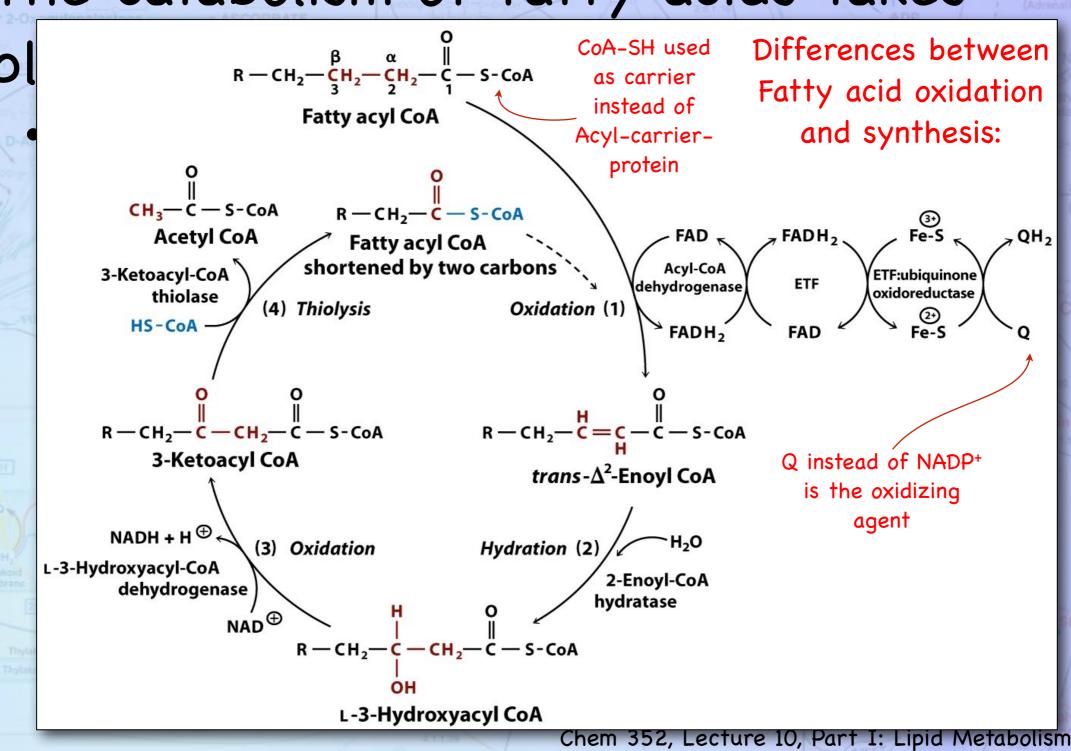


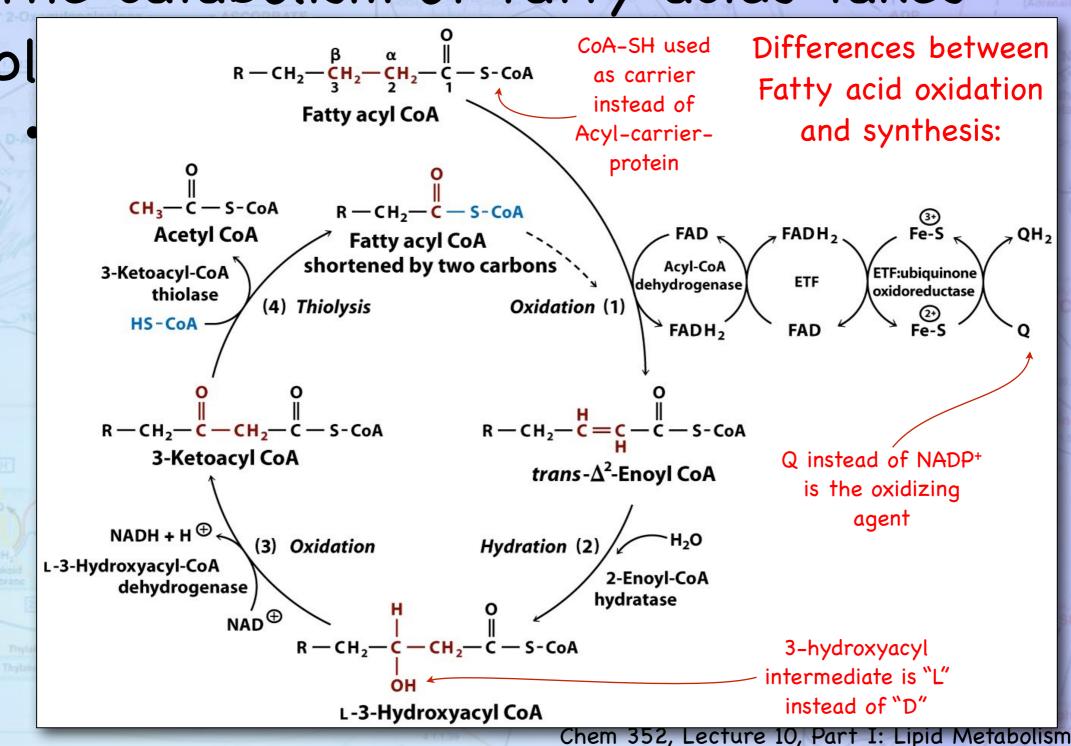
The catabolism of fatty acids takes

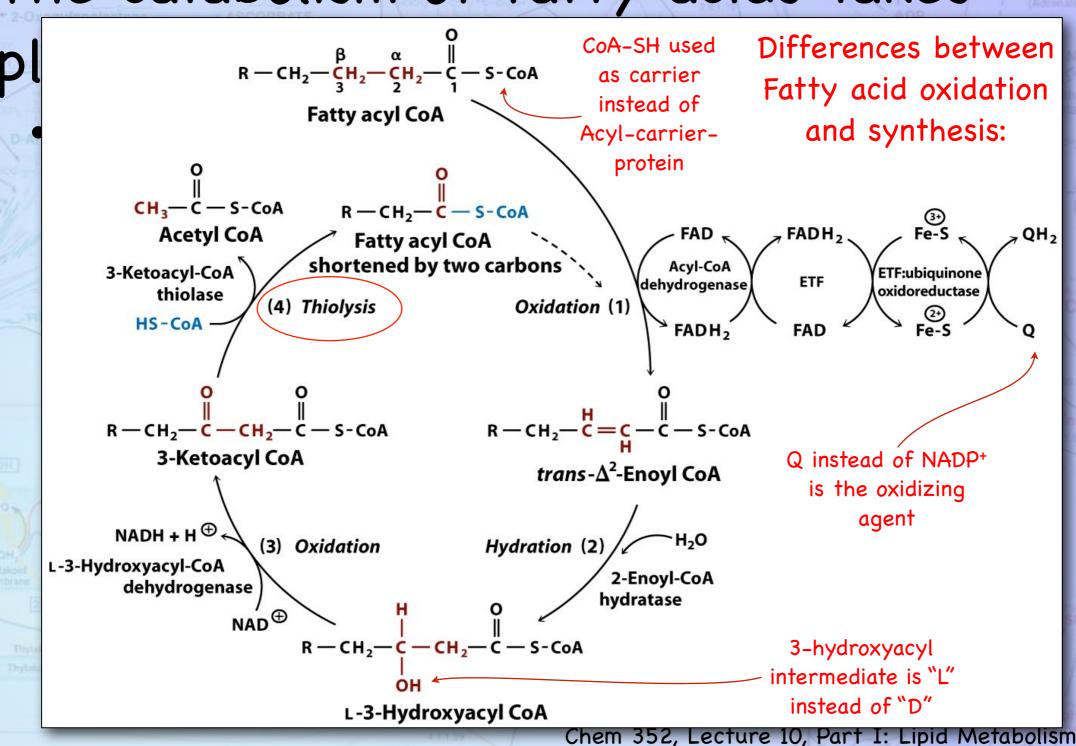


The catabolism of fatty acids takes



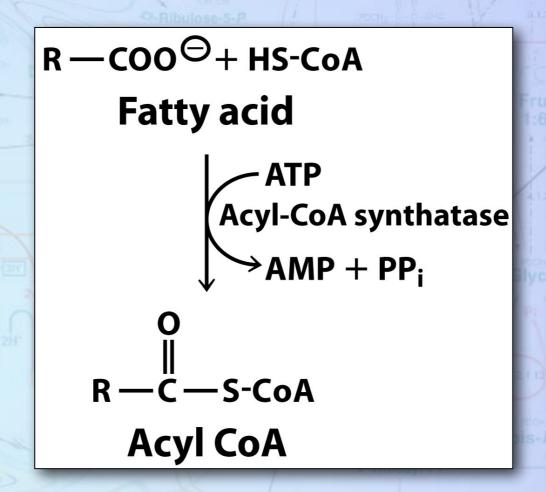




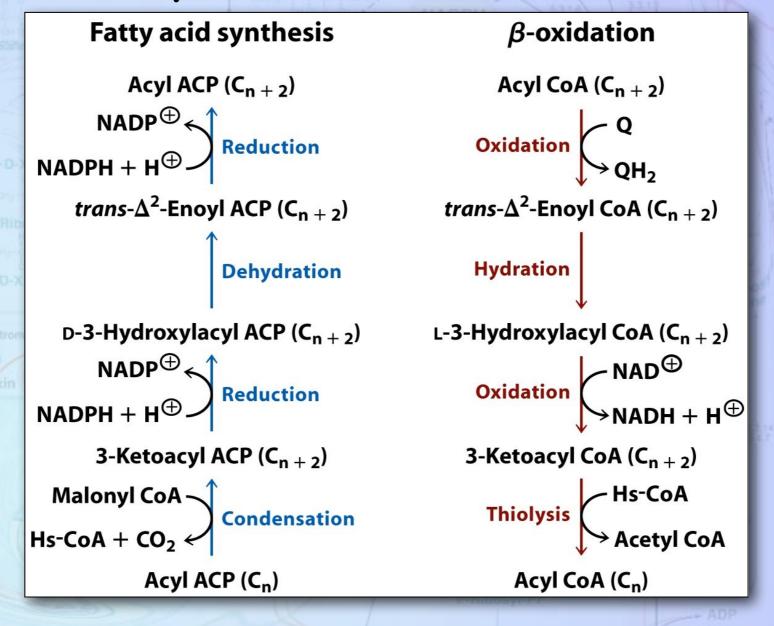


The catabolism of fatty acids takes place in the mitochondria

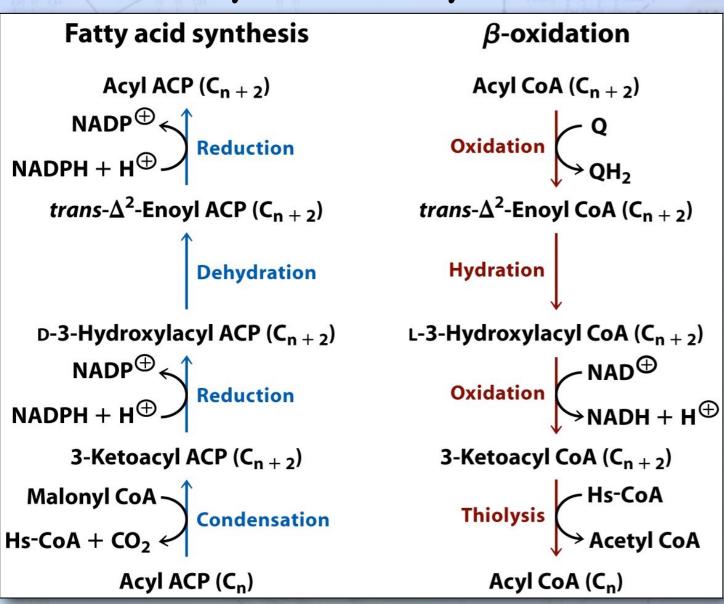
The fatty acid must first be activated to an acyl-CoA



In many respects it is the reverse of fatty acid synthesis.

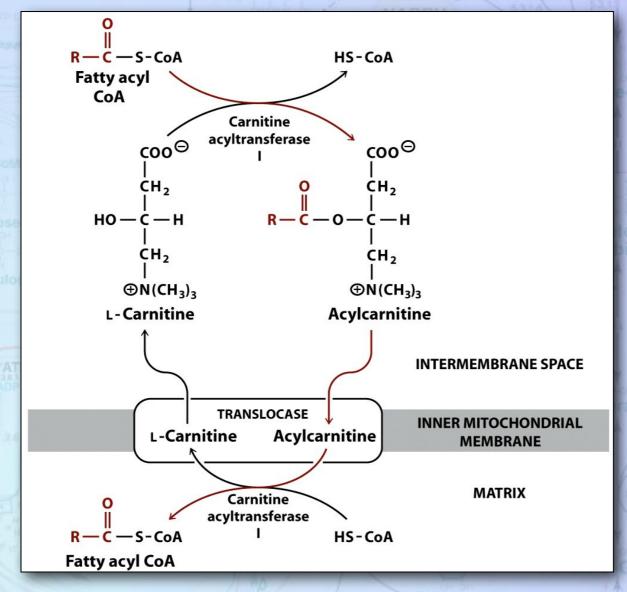


In many respects it is the reverse of fatty acid synthesis.



Synthesis	Oxidation
Cytosol	Mitochondria
NADPH	NAD+ and Q
Acyl-ACP	Acyl-CoA
D-3-Hydroxy-	L-3-Hydroxy-

Fatty acids enter the mitochondria by way of the carnitine shuttle.



ATP generation of the the complete oxidation of stearic acid (18:0)

Stearoyl-CoA + 8 CoA-SH + 8 Q + 8 NAD+ \rightarrow 9 Acetyl-CoA + 8 QH₂ + 8 NADH + 8 H+

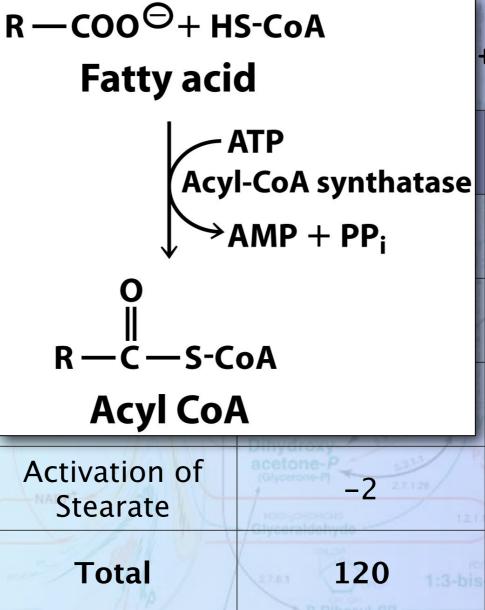
ATP generation of the the complete oxidation of stearic acid (18:0)

Stearoyl-CoA + 8 CoA-SH + 8 Q + 8 NAD+ -> 9 Acetyl-CoA + 8 QH2 + 8 NADH + 8 H+

Source	ATP's
8 QH ₂	12 F
8 NADH	20
9 Acetyl-CoA	90
Activation of Stearate	acetone-P -2 22128
Total	120 13-bis

ATP generation of the the complete oxidation of stearic acid (18:0)

Stearoyl-CoA + 8 CoA-SH



Chem 352, Lecture 10, Part I: Lipid Metabolism

8 QH₂ + 8 NADH + 8 H⁺

ATP generation of the the complete oxidation of stearic acid (18:0)

Stearoyl-CoA + 8 CoA-SH + 8 Q + 8 NAD+ -> 9 Acetyl-CoA + 8 QH2 + 8 NADH + 8 H+

Source	ATP's
8 QH ₂	12 F
8 NADH	20
9 Acetyl-CoA	90
Activation of Stearate	acetone-P -2 22128
Total	120 13-bis

ATP generation of the the complete oxidation of steric acid (18:0)

· Compared to Glucose (on a per C basis)

ATP generation of the the complete oxidation of steric acid (18:0)

· Compared to Glucose (on a per C basis)

	Source	ATP's
1	3 x Glucose	3 x 32 = 96
	Stearate	Dihydroxy acetone-P 120

ATP generation of the the complete oxidation of stearic acid (18:0)

• Compared to Fatty Acid Synthesis

ATP generation of the the complete oxidation of stearic acid (18:0)

Compared to Fatty Acid Synthesis

Source	ATP's
8 Acetyl-CoA → 8 Malonyl-CoA	8
8 Rounds 16 NADPH	40
9 Acetyl-CoA (Calvin cycle)	9 x 17 = 153
Total	201

ATP generation of the the complete oxidation of stearic acid (18:0)

Compared to Fatty Acid Synthesis

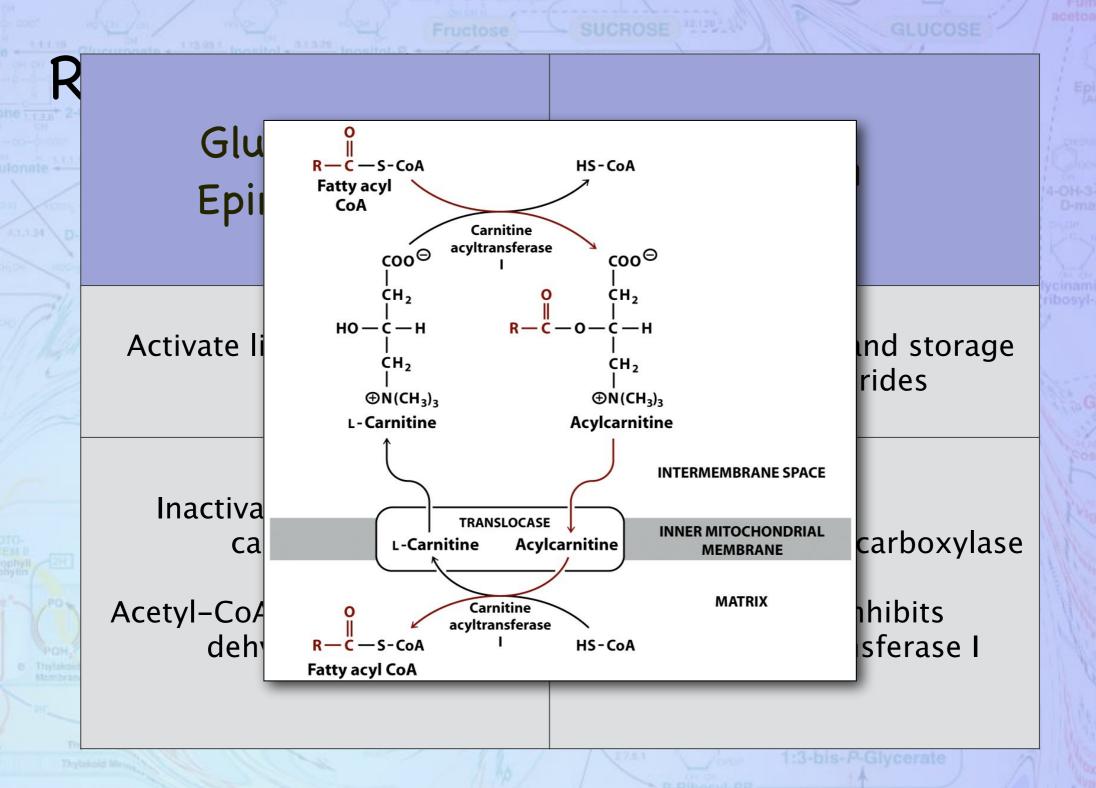
Source	ATP's
8 Acetyl-CoA → 8 Malonyl-CoA	8
8 Rounds 16 NADPH	40
9 Acetyl-CoA (Calvin cycle)	9 x 17 = 153
Total	201

Yield = 120/201 = 60%

Regulation of Lipid Metabolism

- Involves same hormones as carbohydrate metabolism
 - · Glucogon (fasting state)
 - · Epinephrin (excited state)
 - · Insulin (fed state)

Glucagon & Insulin Epinephrine Activate lipases in adipose Activate formation and storage of triacylglycerides tissues Inactivates acetyl-CoA carboxylase Activate acetyl-CoA carboxylase Acetyl-CoA inhibits pyruvate Malonyl-CoA inhibits Carnitine acyltransferase I dehydrogenase



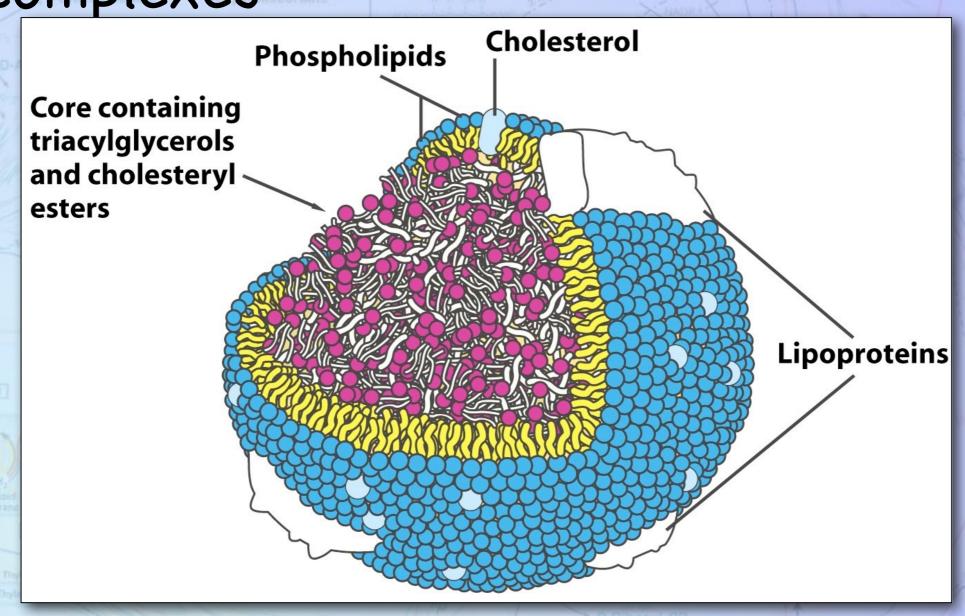
Glucagon & Insulin Epinephrine Activate lipases in adipose Activate formation and storage of triacylglycerides tissues Inactivates acetyl-CoA carboxylase Activate acetyl-CoA carboxylase Acetyl-CoA inhibits pyruvate Malonyl-CoA inhibits Carnitine acyltransferase I dehydrogenase

Transport of lipds in Blood

Lipids are transported by lipoprotein complexes

- Chylomicrons
- · VLDH (Very Low Density Lipoproteins)
- IDH (Intermediate Density Lipoproteins)
- · LDL (Low Density Lipoproteins)
- HDL (High Density Lipoproteins)

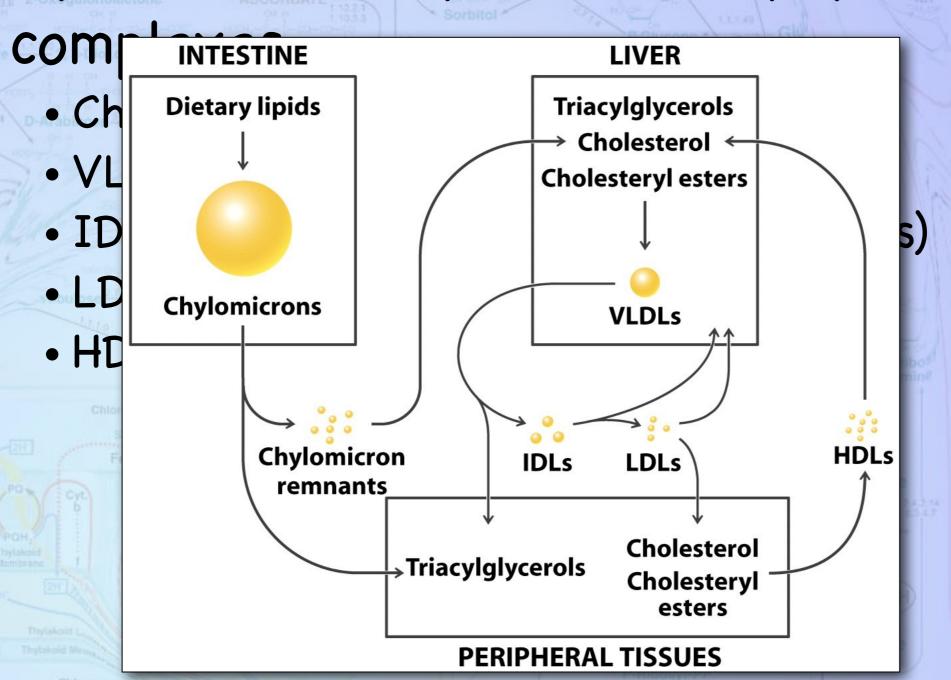
Lipids are transported by lipoprotein complexes



Lipids are transported by lipoprotein complexes

- Chylomicrons
- VLDH (Very Low Density Lipoproteins)
- IDH (Intermediate Density Lipoproteins)
- LDL (Low Density Lipoproteins)
- HDL (High Density Lipoproteins)

Lipids are transported by lipoprotein



Lipids are transported by lipoprotein complexes

- Chylomicrons
- VLDH (Very Low Density Lipoproteins)
- IDH (Intermediate Density Lipoproteins)
- LDL (Low Density Lipoproteins)
- HDL (High Density Lipoproteins)

Ketone bodies are formed from acetyl-CoA as a soluble circulating source of fat-derived energy.

· Produce under conditions of long-term fasting

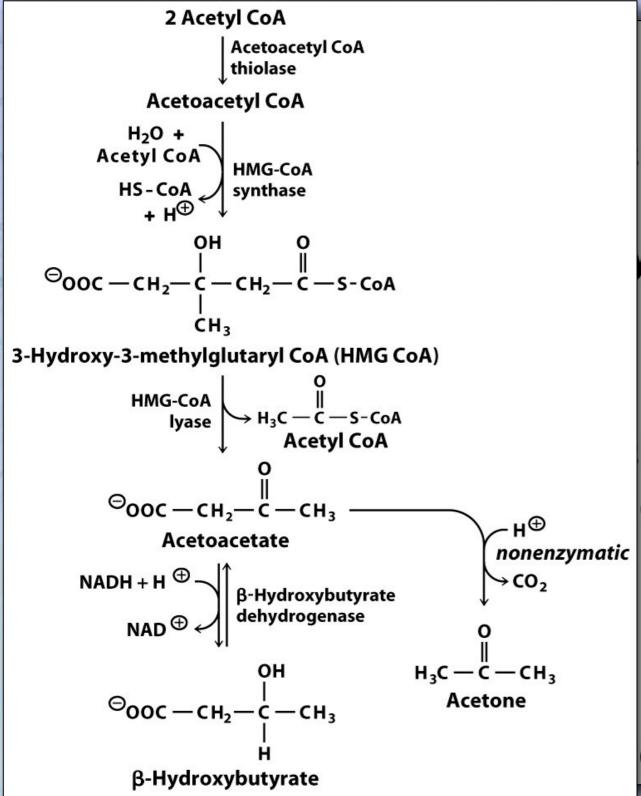
Ketone be source of Produce

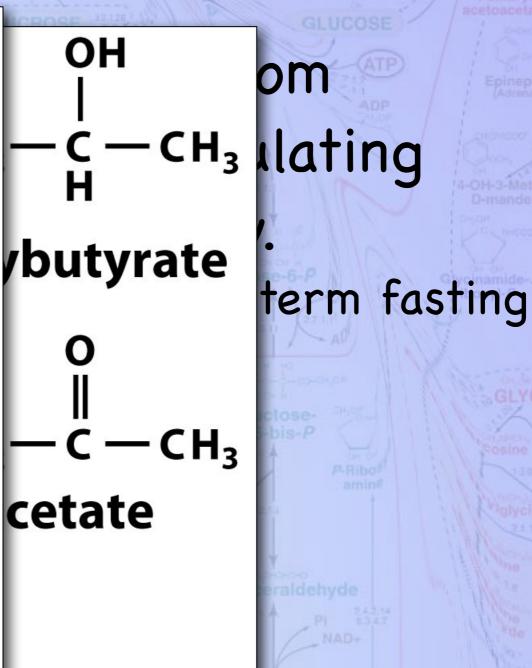
OH acetyl-Co Ooc-CH2-C-C-CH3 llating **β-Hydroxybutyrate**

> Θ 00C - CH₂- C - CH₃ Acetoacetate

$$0$$
 \parallel
 $H_3C-C-CH_3$
Acetone

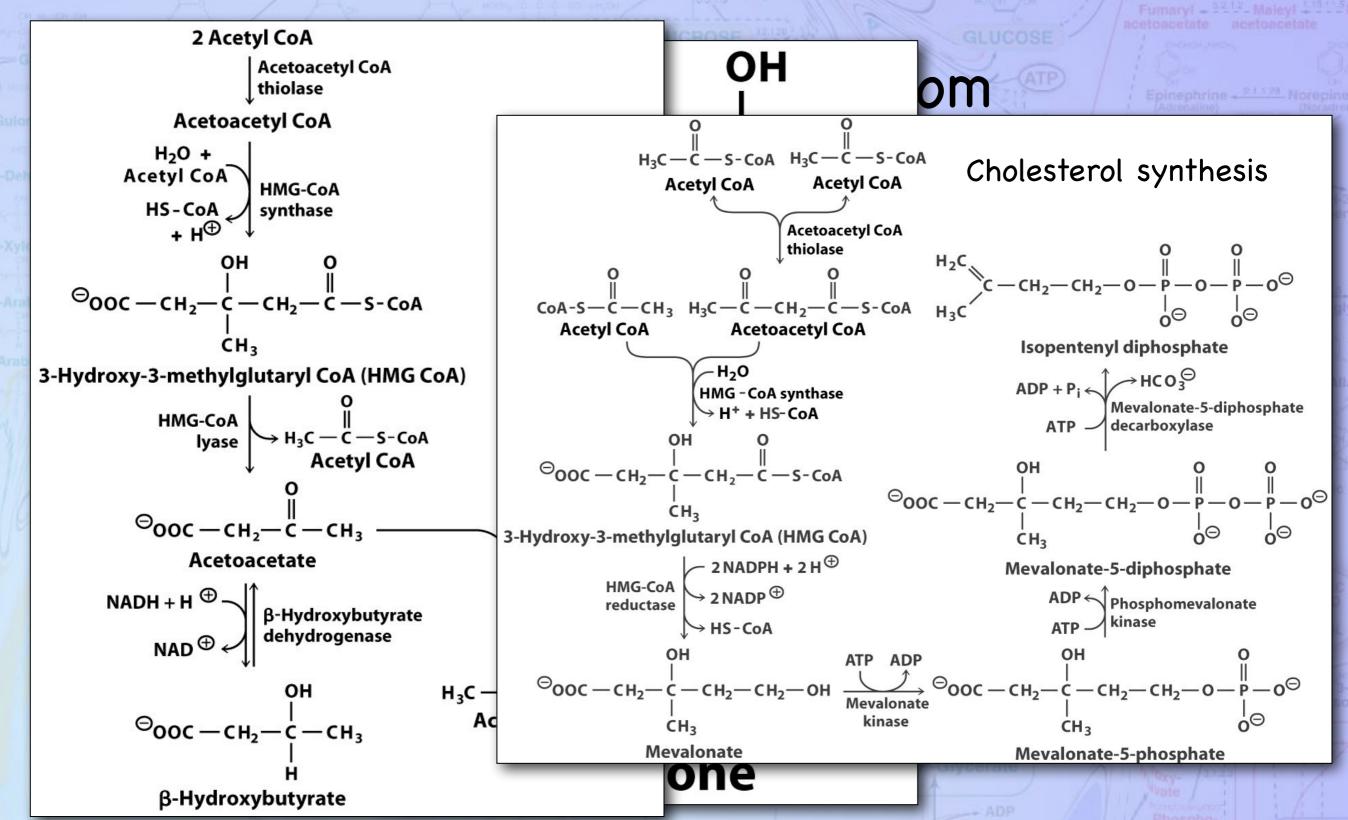
term fasting

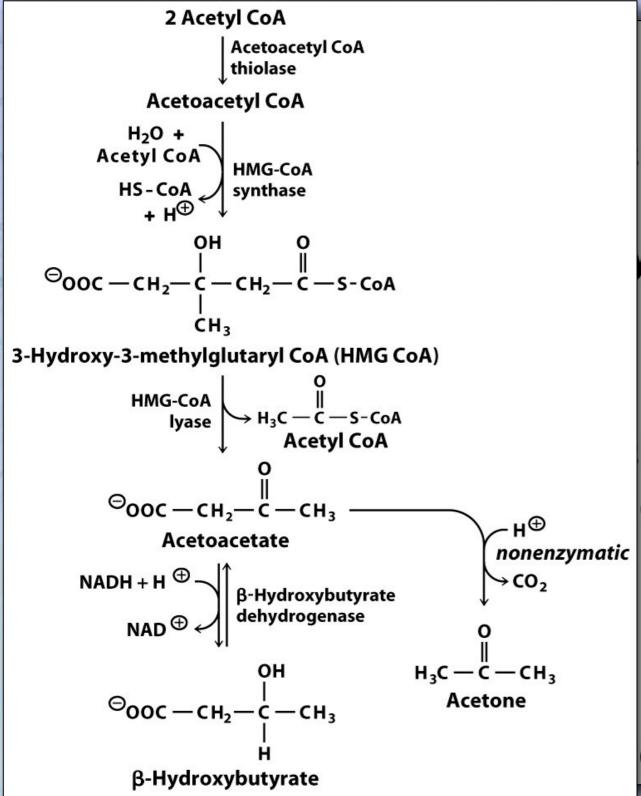


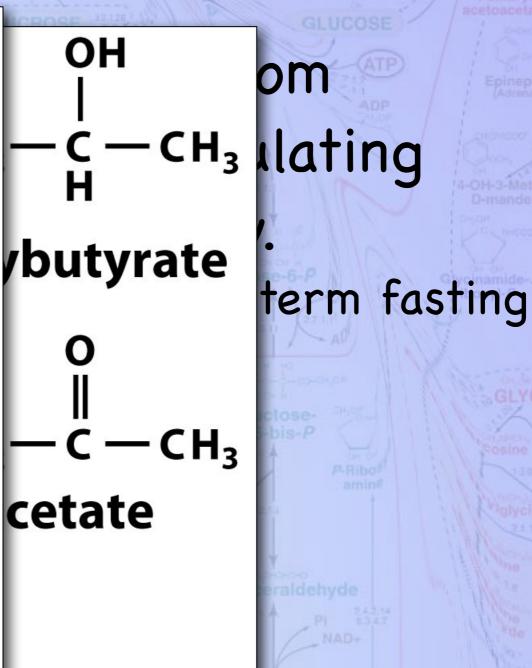


CH₃

one







CH₃

one

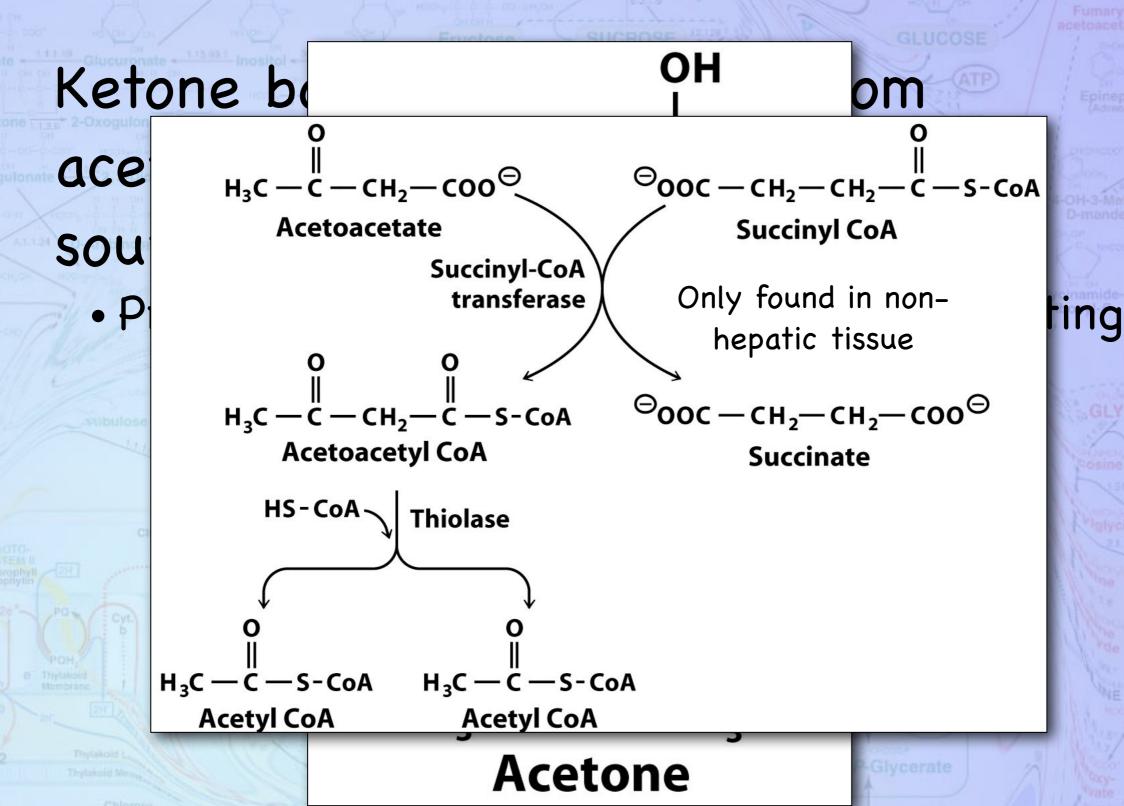
Ketone be source of Produce

OH acetyl-Co Ooc-CH2-C-C-CH3 llating **β-Hydroxybutyrate**

> Θ 00C - CH₂- C - CH₃ Acetoacetate

$$0$$
 \parallel
 $H_3C-C-CH_3$
Acetone

term fasting



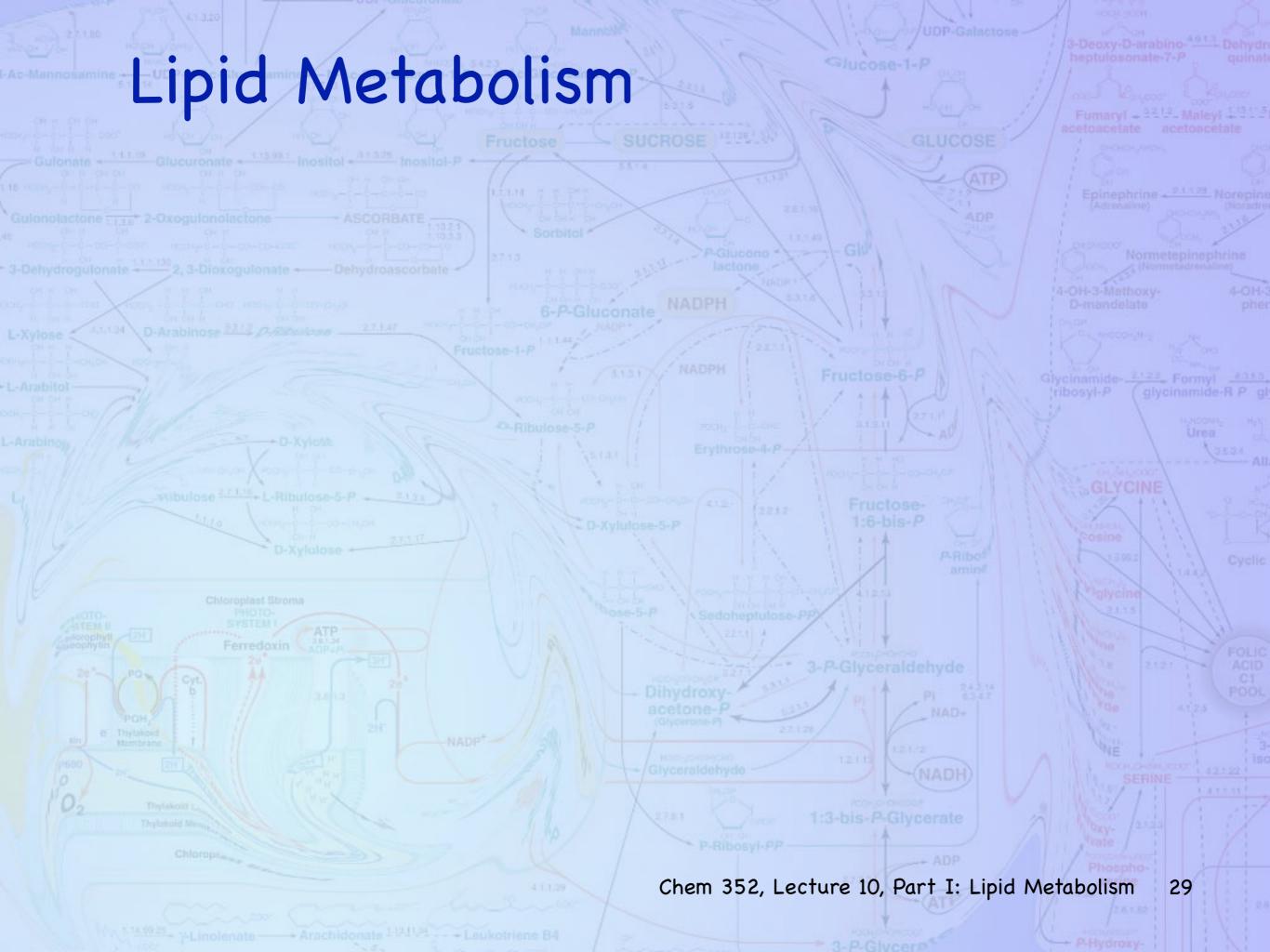
Ketone be source of Produce

OH acetyl-Co Ooc-CH2-C-C-CH3 llating **β-Hydroxybutyrate**

> Θ 00C - CH₂- C - CH₃ Acetoacetate

$$0$$
 \parallel
 $H_3C-C-CH_3$
Acetone

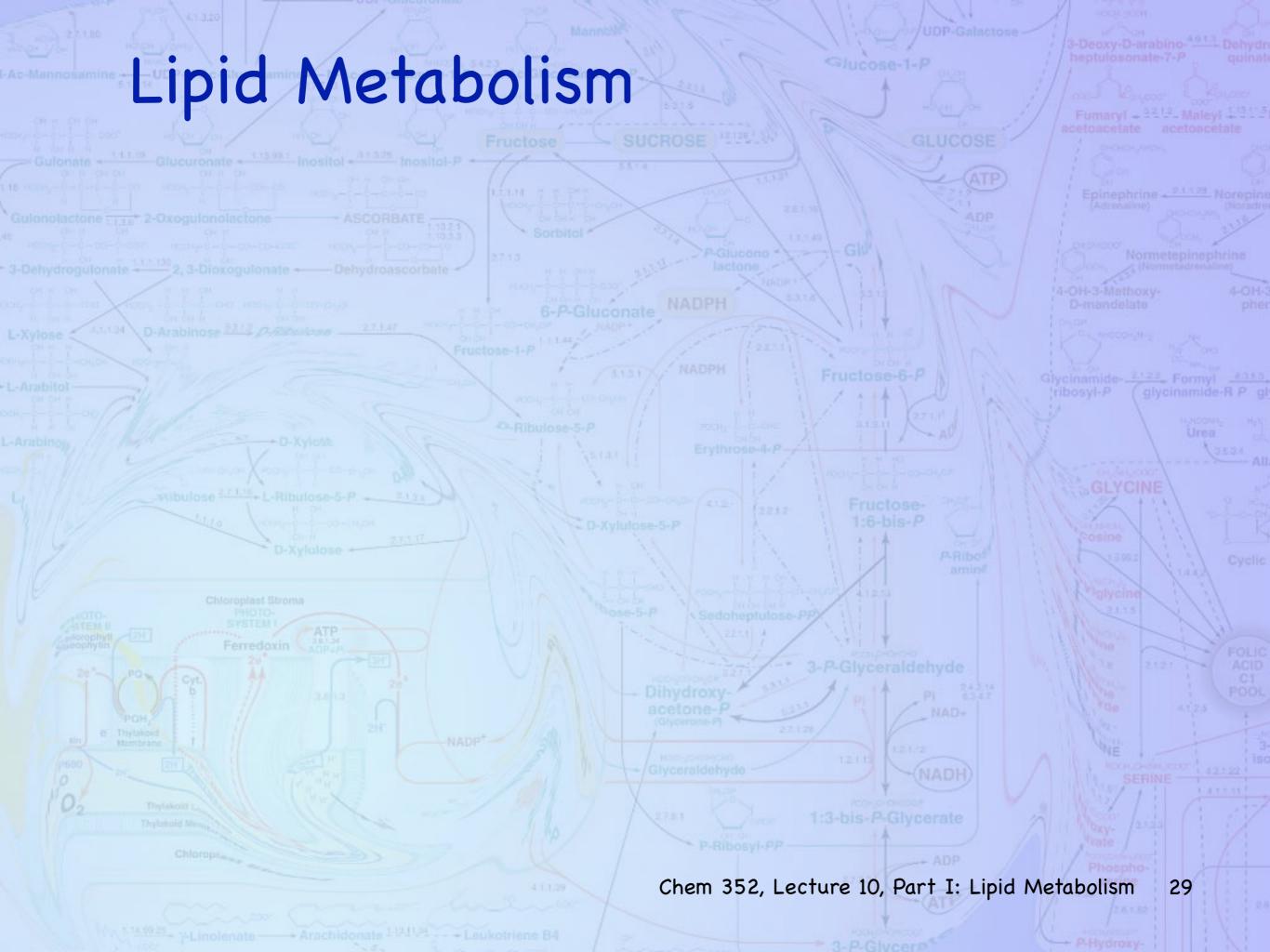
term fasting



Lipid Metabolism

Question:

Draw a general pathway for converting carbohydrates to fatty acids in a liver cell, and indicate which processes occur in the cytosol and which occur in mitochondria.



Next Up

Lecture 10 - Part II, Amino acid metabolism (Moran et al., Chapter 17)