

# Lecture 4

## **BT 203**

# **Biochemistry**

### **3-0-0-6**

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# Key Concepts

- How are fatty acids synthesized ?
- What is fatty acid synthesis?
- What is the role of Acetyl-CoA Carboxylase?
- Where does fatty acid synthesis takes place?

# Key Concepts

## FATTY ACID BIOSYNTHESIS

### + Key Features

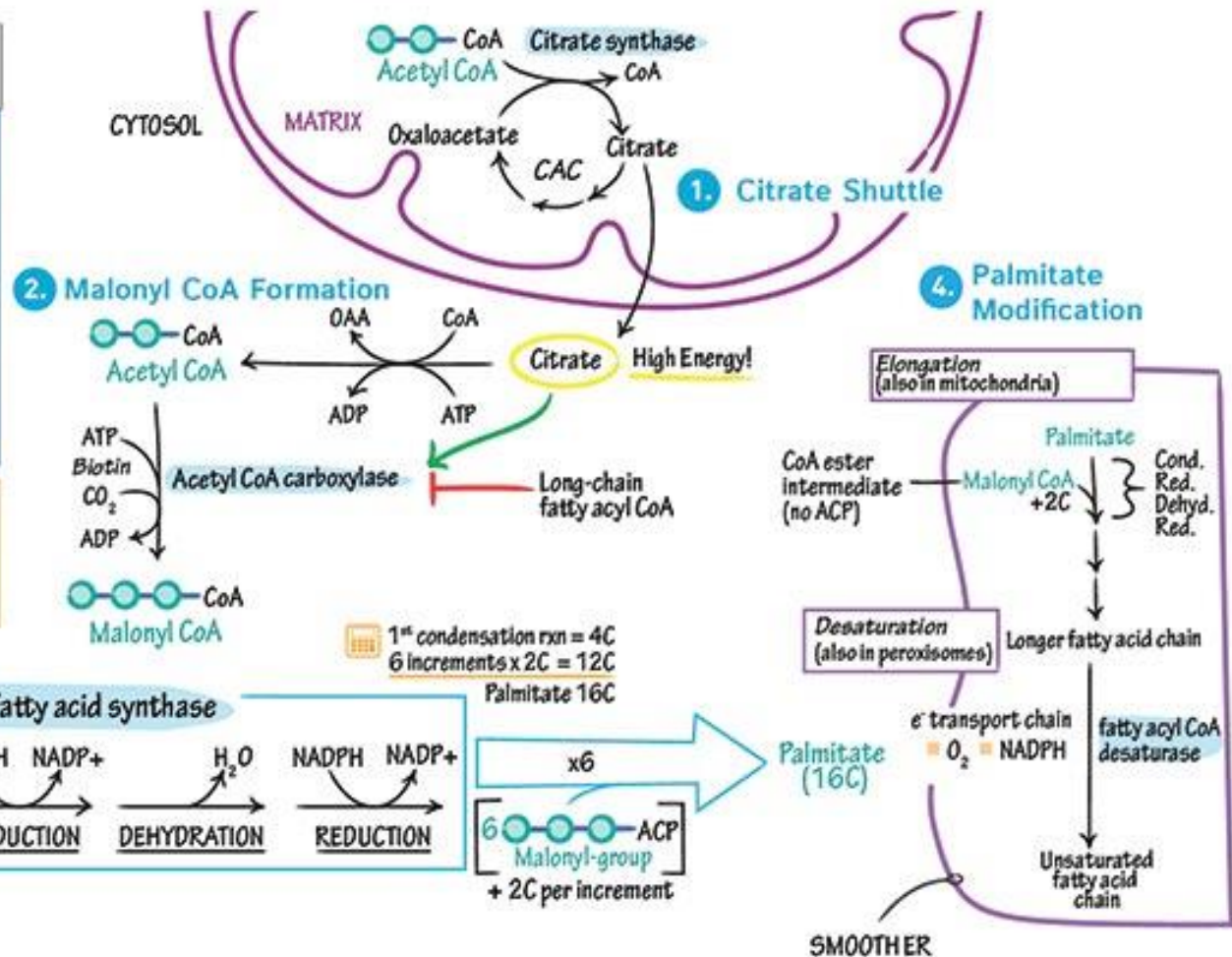
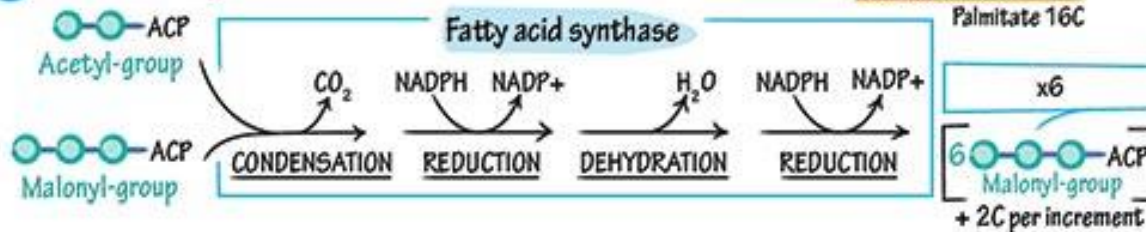
- ✓ Occurs in the liver & adipose (cytosol)
- ✓ After carb-rich meal (insulin: glucagon is high)
- ✓ Not reverse of beta-oxidation
- ✓ Energetically expensive

### § Clinical Correlation

- ✓ Linoleic & linolenic acid—  
Essential FA: can't be produced endogenously; must come from diet. Mammals can't induce d.b. beyond C9.

**ABC Carboxylase Rxns**  
 Malonyl CoA formation  
 Gluconeogenesis  
 Odd chain fatty acid oxidation

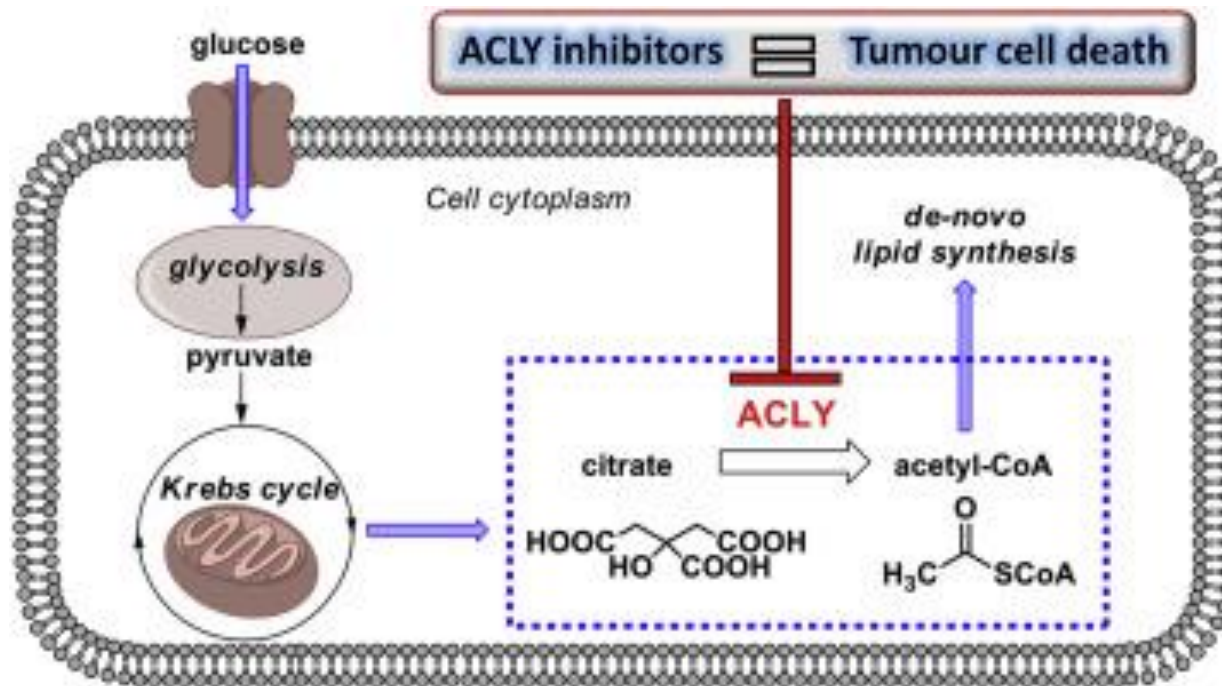
### 3. Palmitate Synthesis



# Citrate shuttle

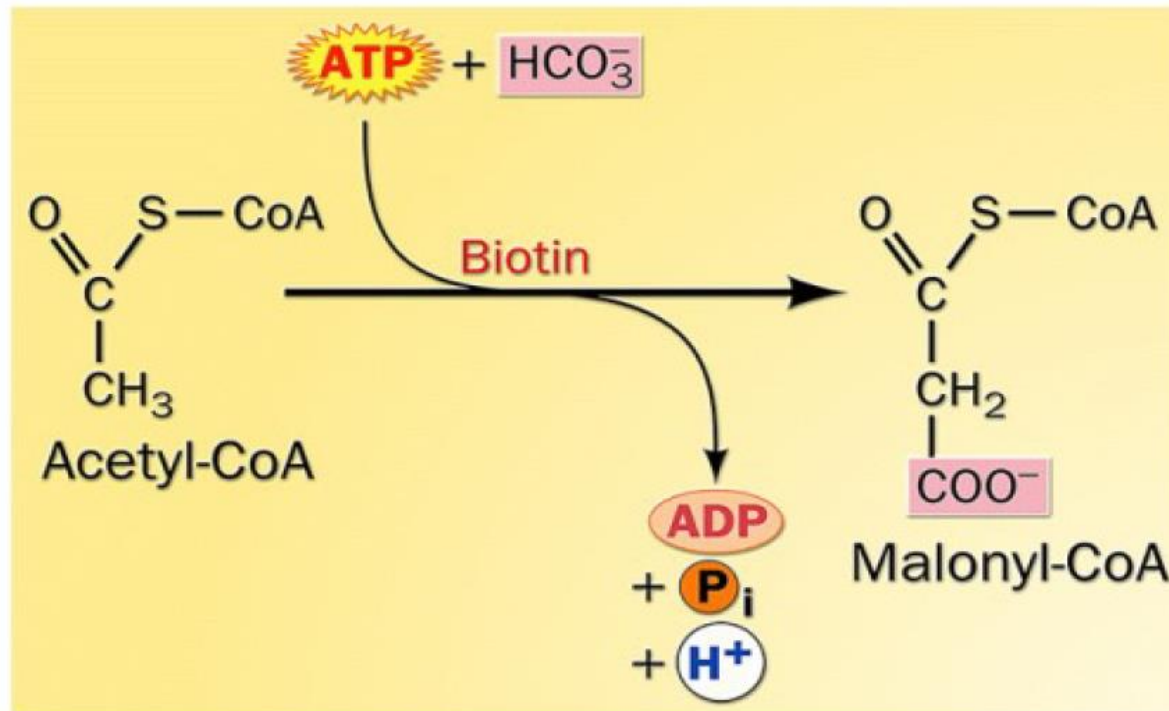
- FAs are synthesized in the cytoplasm from acetyl CoA
- Acetyl CoA generated from pyruvate by the action of PDH and by  $\beta$ -oxidation of fatty acids is in the mitochondria.
- For fatty acid biosynthesis, acetylCoA has to be transported from the mitochondria to the cytoplasm. This is done via a shuttle system called the Citrate Shuttle.
- Acetyl CoA reacts with oxaloacetate to give citrate. A tricarboxylate translocase transports citrate from mitochondria to cytosol.
- In the cytosol, citrate is cleaved back to oxaloacetate and acetylCoA. This reaction is catalyzed by ATP-citrate lyase and requires the hydrolysis of one molecule of ATP.

# Fatty acid synthesis



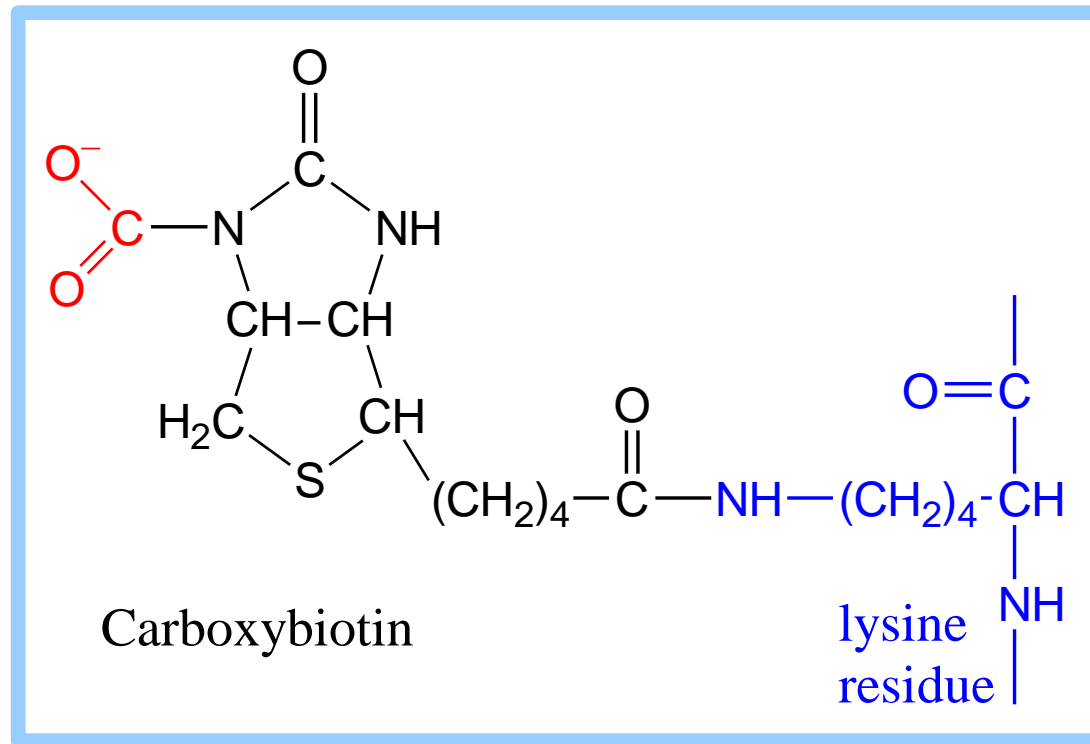
# Fatty acid synthesis

Reaction catalyzed by Acetyl CoA Carboxylase



Activation of acetate : Acetyl-CoA to malonyl CoA

# What is biotin?



**Biotin** is linked to the enzyme by an amide bond between the terminal carboxyl of the biotin side chain and the ε-amino group of a **lysine** residue.

The combined biotin and lysine side chains act as a **long flexible arm** and it allows the biotin ring to translocate between the 2 active sites.

# Acetyl-CoA Carboxylase

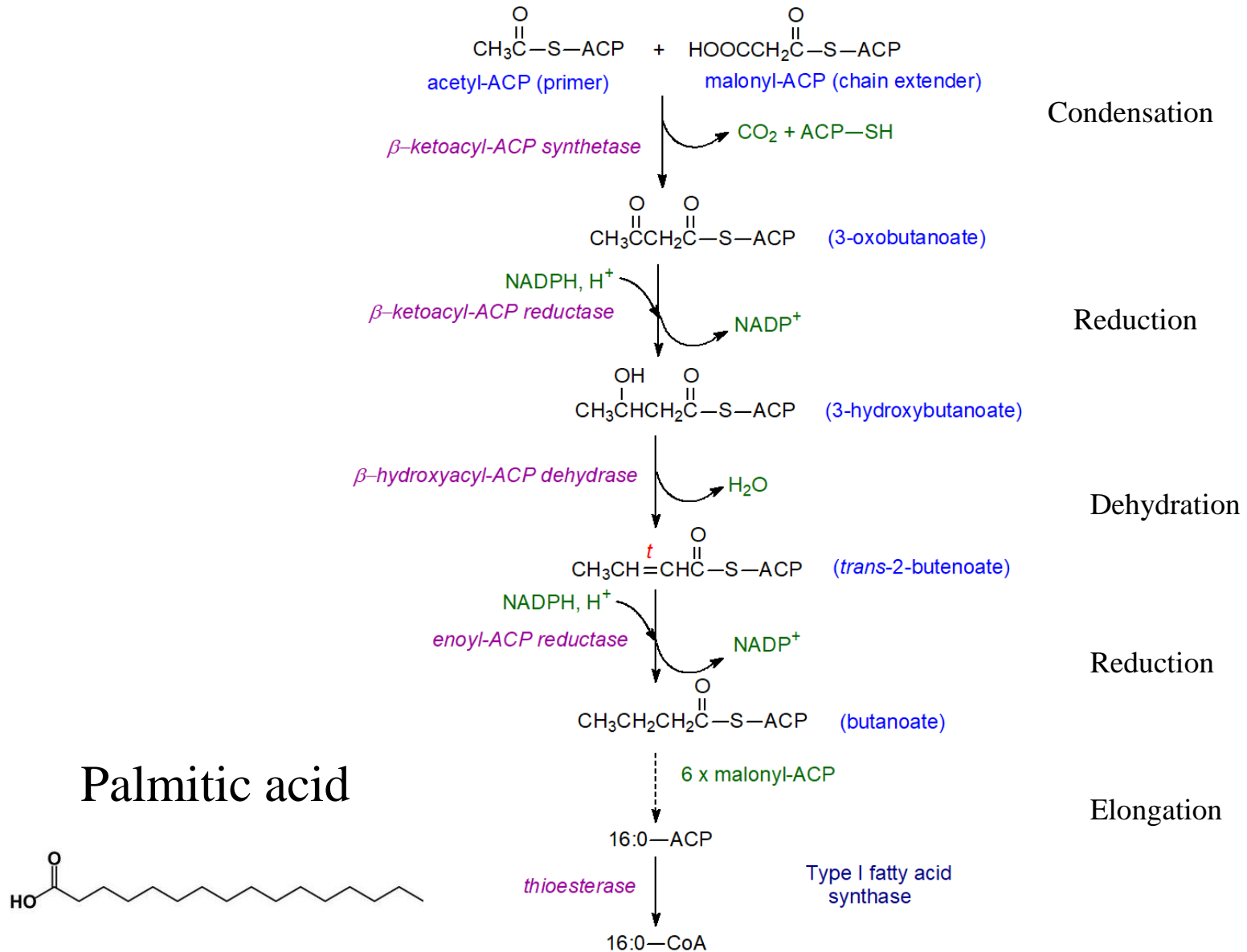
**Acetyl-CoA Carboxylase**, which converts acetyl-CoA to malonyl-CoA, is the **committed step** of the fatty acid synthesis pathway.

The mammalian enzyme is required to be **regulated**, and it is achieved by

- ♦ Phosphorylation  
and
- ♦ Allosteric control by local metabolites.

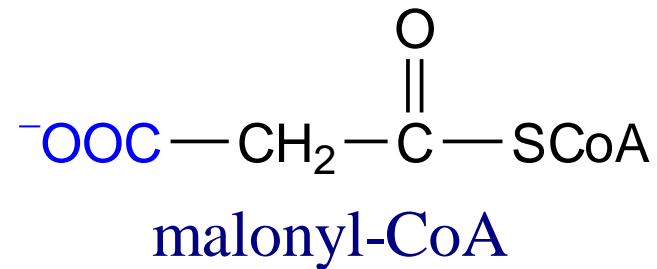
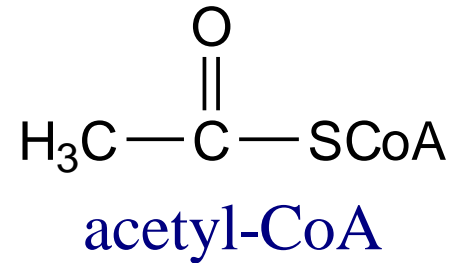


# Key Concepts



# Fatty acid precursor

The input to fatty acid synthesis is **acetyl-CoA**, a two carbon compound, which is carboxylated to three carbon compound **malonyl-CoA**.



ATP-dependent carboxylation provides energy input.

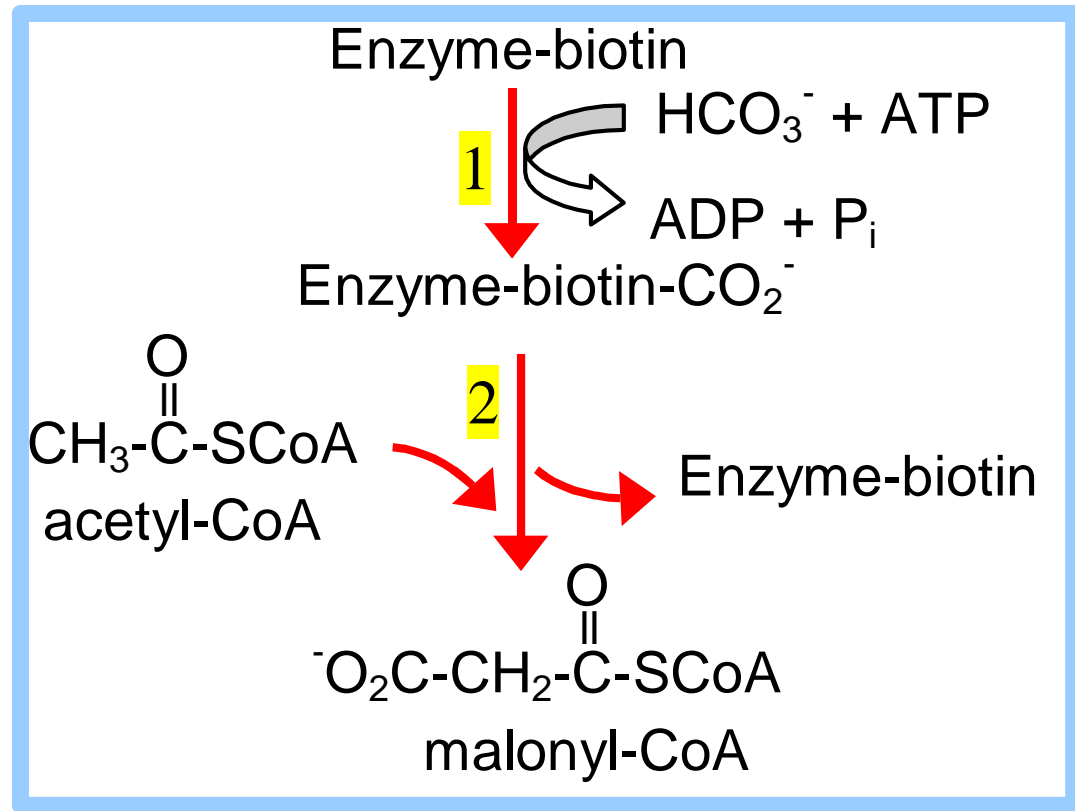
The  $\text{CO}_2$  is lost later during condensation with the growing fatty acid.

The spontaneous decarboxylation drives the condensation reaction.

# Fatty acid precursor

## Acetyl-CoA

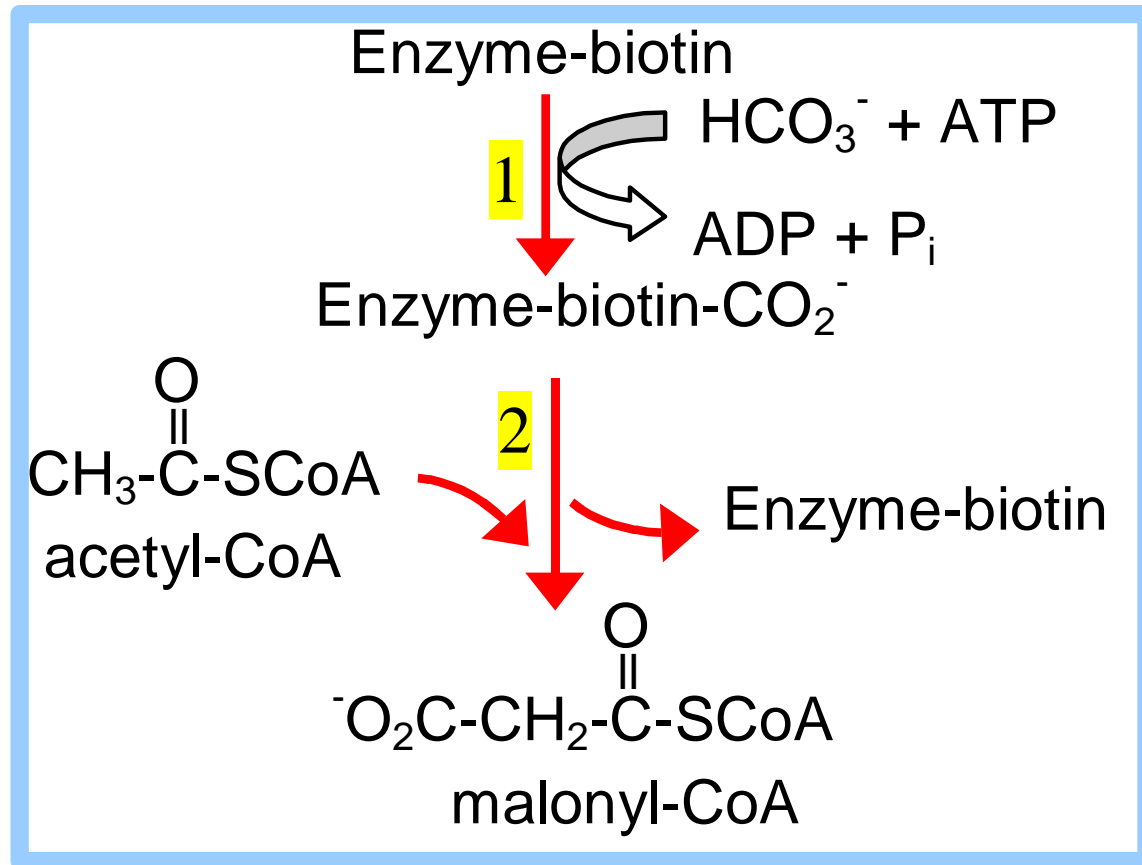
**Carboxylase** catalyzes the **2-step** reaction by which acetyl-CoA is carboxylated to form **malonyl-CoA**.



As with other carboxylation reactions, the enzyme prosthetic group is **biotin**.

ATP-dependent carboxylation of the biotin, carried out at one active site , is followed by transfer of the carboxyl group to acetyl-CoA at a second active site 2 .

# Fatty acid precursor



The overall reaction, which is **spontaneous**, is summarized as:  $\text{HCO}_3^- + \text{ATP} + \text{Acetyl-CoA} \rightarrow \text{ADP} + \text{P}_i + \text{Malonyl-CoA}$

# Fatty acid synthesis

**Fatty acid synthesis** from acetyl-CoA & malonyl-CoA occurs by a series of reactions that are:

- ♦ in **bacteria** catalyzed by 6 different enzymes plus a separate acyl carrier protein (ACP)
- ♦ in **mammals** catalyzed by individual domains of a very large polypeptide that includes an ACP domain.

Evolution of the mammalian Fatty Acid Synthase apparently has involved **gene fusion**.

**NADPH** serves as **electron donor** in the two reactions involving substrate reduction.

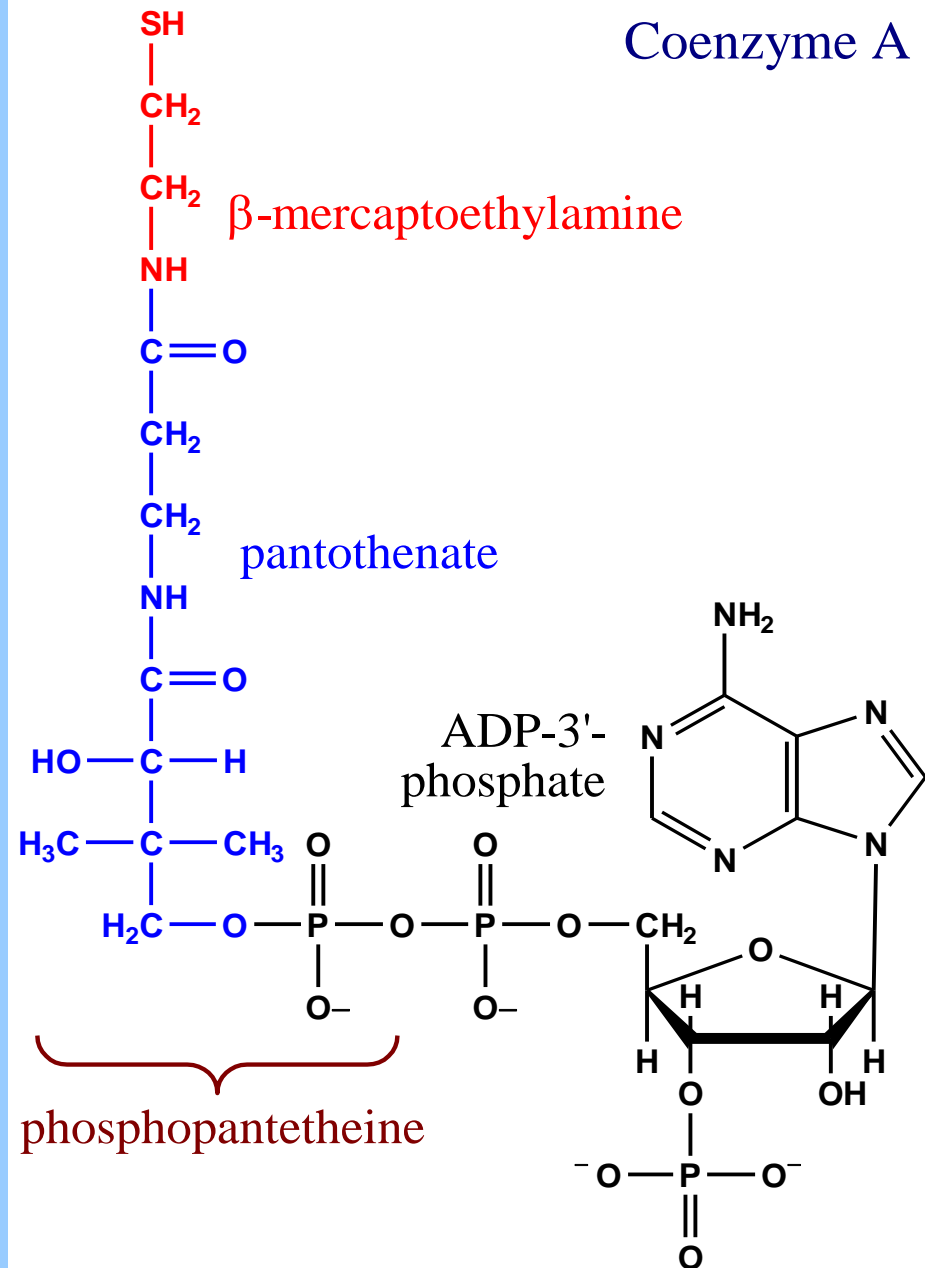
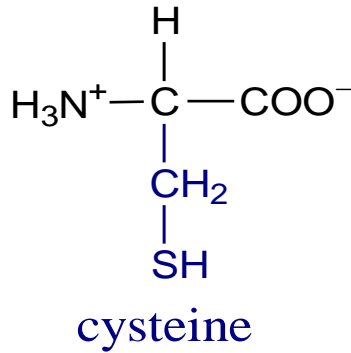
The NADPH is produced mainly by the Pentose Phosphate Pathway.

# Fatty acid synthesis

## Fatty Acid Synthase

prosthetic groups:

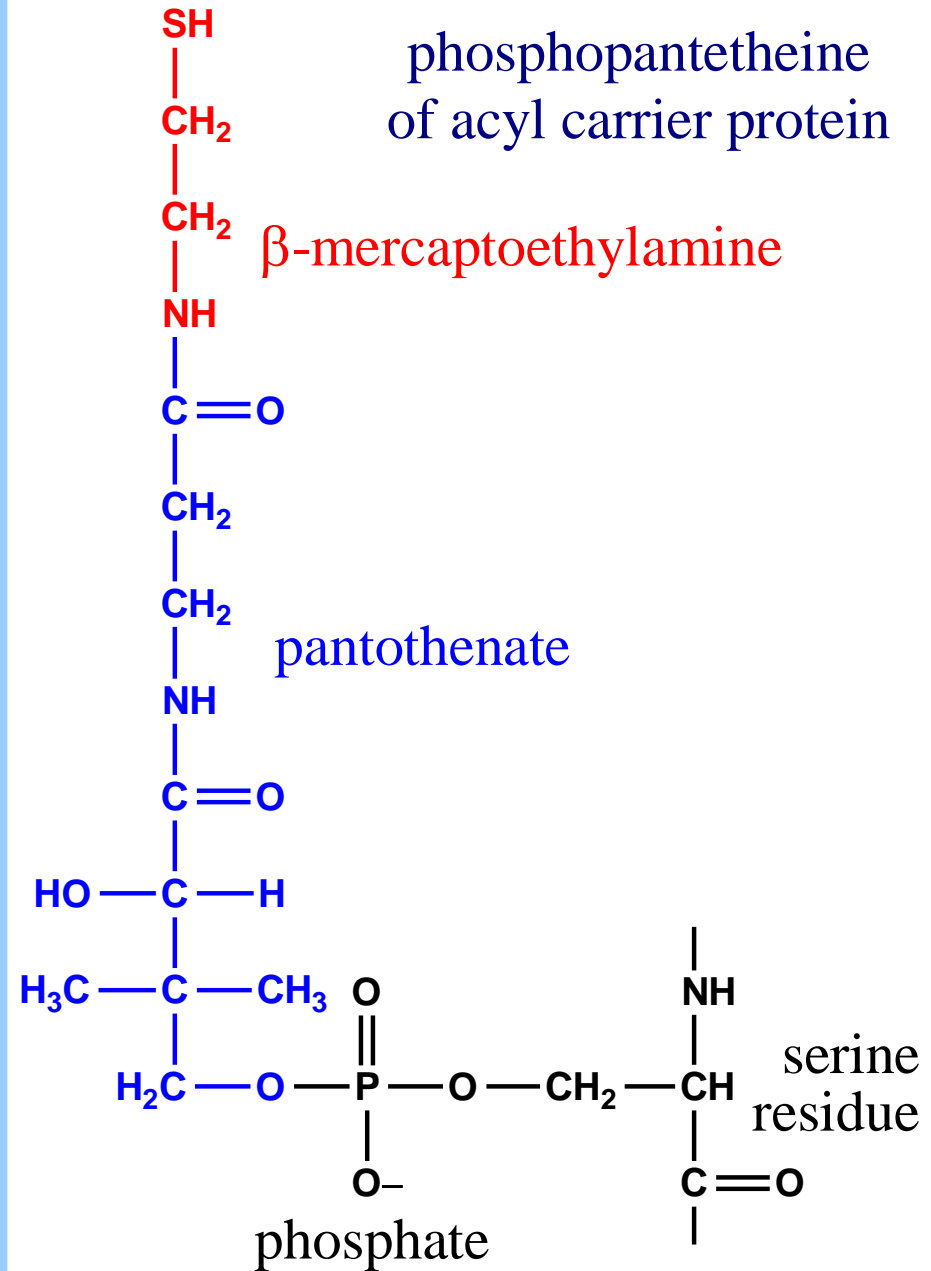
- ♦ the **thiol** of the side-chain of a **cysteine** residue of Condensing Enzyme domain.
- ♦ the **thiol** of **phosphopantetheine**, equivalent in structure to part of coenzyme A.



# Fatty acid synthesis

**Phosphopantetheine** (Pant) is covalently linked via a phosphate ester to a serine OH of the **acyl carrier protein** domain of Fatty Acid Synthase.

The **long flexible arm** of phosphopantetheine helps its thiol to move from one active site to another within the complex.



# Fatty acid synthesis

N-	Condensing Enzyme ( <b>Cys</b> )	Malonyl/acetyl-CoA Transacylase ( <b>Ser</b> )	Dehydratase	Enoyl Reductase	$\beta$ -Ketoacyl Reductase	ACP ( <b>Pant</b> )	Thioesterase	-C
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Order of domains in primary structure of mammalian Fatty Acid Synthase

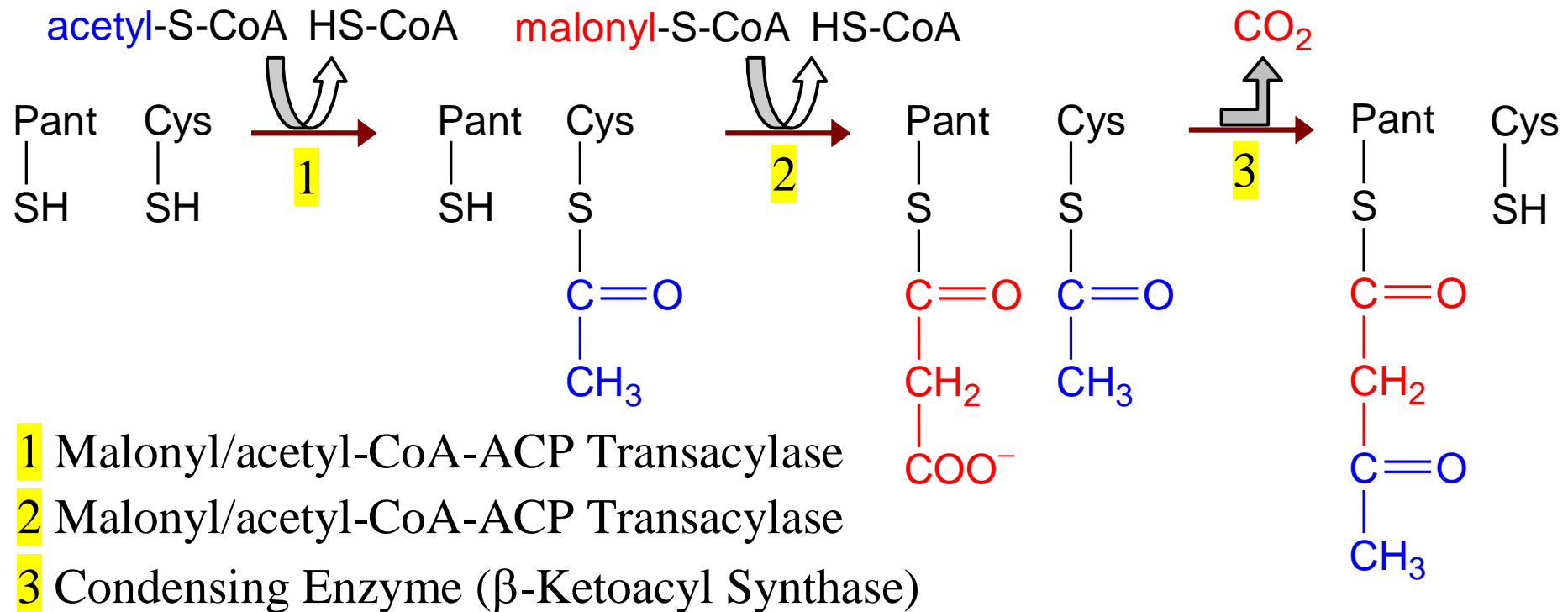
As each of the substrates acetyl-CoA & malonyl-CoA bind to the complex, the initial attacking group is the oxygen of a **serine hydroxyl** group of the Malonyl/acetyl-CoA Transacylase enzyme domain.

Each acetyl or malonyl moiety is transiently in ester linkage to this serine hydroxyl, before being transferred into thioester linkage with the **phosphopantetheine thiol** of the acyl carrier protein (ACP) domain.

Acetate is subsequently transferred to a **cysteine thiol** of the Condensing Enzyme domain.

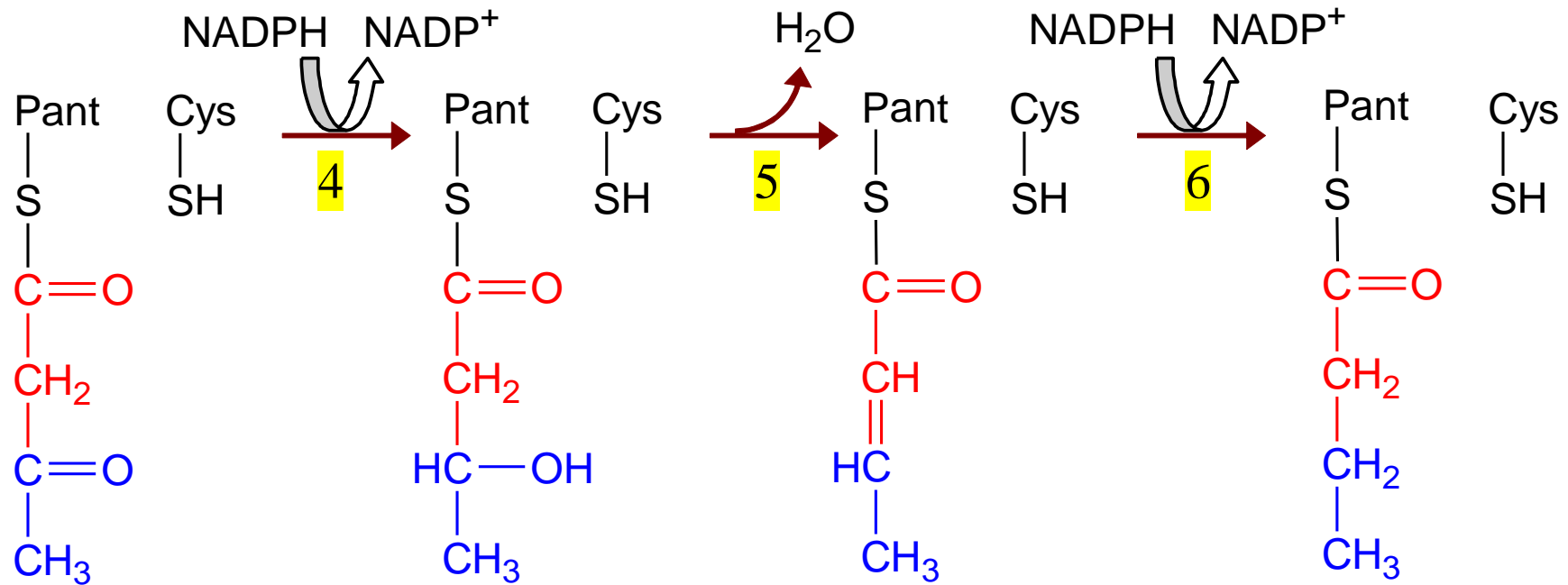


# Fatty acid synthesis



The **condensation** reaction (step 3) involves decarboxylation of the malonyl moiety, followed by attack of the resultant carbanion on the carbonyl carbon of the acetyl (or acyl) moiety.

# Fatty acid synthesis



4  $\beta$ -Ketoacyl-ACP Reductase

5  $\beta$ -Hydroxyacyl-ACP Dehydratase

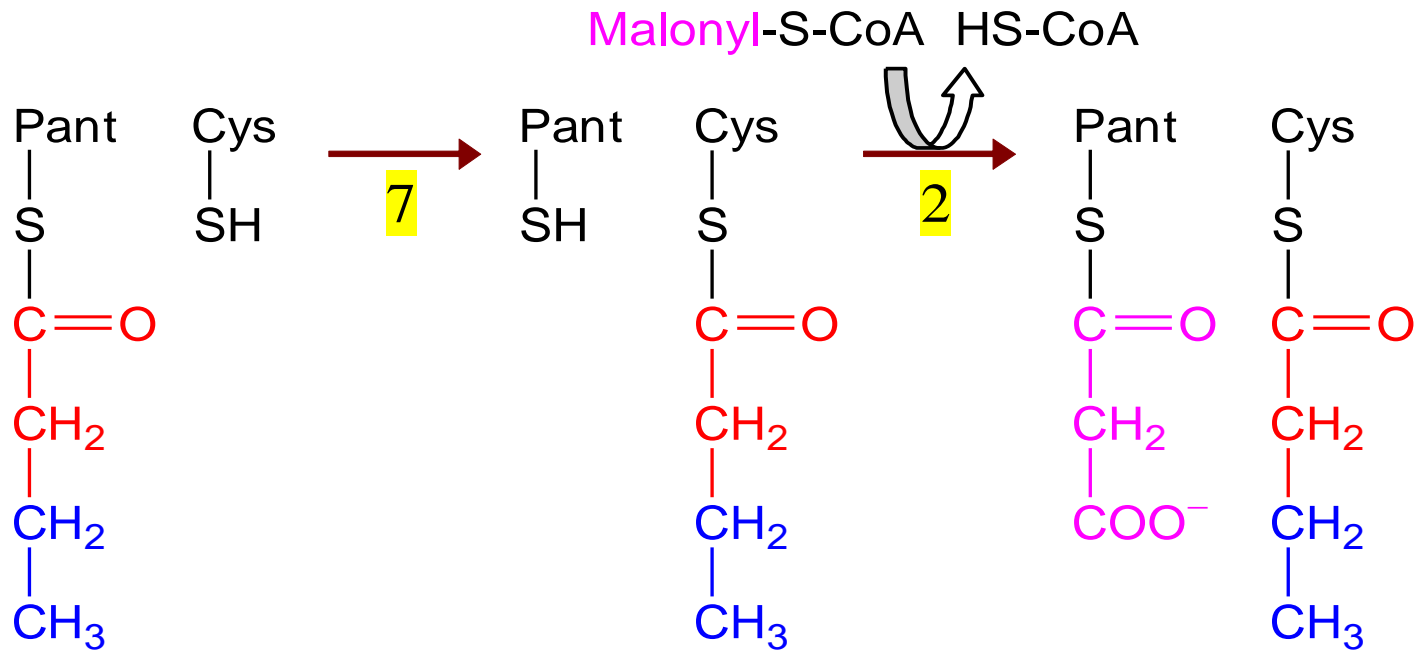
6 Enoyl-ACP Reductase

4. The  $\beta$ -ketone is **reduced** to an alcohol by e<sup>-</sup> transfer from NADPH.

5. **Dehydration** yields a trans double bond.

6. **Reduction** by NADPH yields a saturated chain.

# Fatty acid synthesis



Following **transfer** of the growing fatty acid from phosphopantetheine to the Condensing Enzyme's cysteine sulfhydryl, the **cycle begins again**, with another malonyl-CoA.

# Fatty acid synthesis

## Product release:

When the fatty acid is 16 carbon atoms long, a **Thioesterase** domain catalyzes hydrolysis of the thioester linking the fatty acid to phosphopantetheine.

The **16-C** saturated fatty acid **palmitate** is the final product of the Fatty Acid Synthase complex.

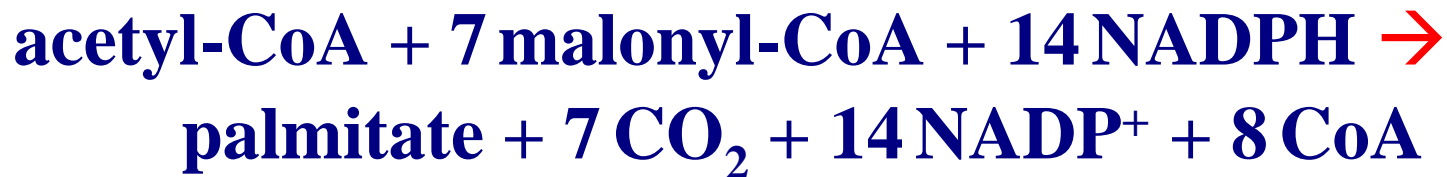
# Fatty acid synthesis

**Summary** (ignoring  $H^+$  & water):

Write a balanced equation for synthesis of palmitate from acetyl-CoA, listing net inputs and outputs:



Summary based on malonate as an input:



Fatty acid synthesis occurs in the **cytosol**. Acetyl-CoA generated in mitochondria is transported to the cytosol via a shuttle mechanism involving **citrate**.

# Fatty acid synthesis

Fatty Acid Synthase is **transcriptionally regulated**.

## **In liver:**

- ♦ **Insulin**, a hormone produced when blood glucose is high, **stimulates** Fatty Acid Synthase expression.

Thus excess glucose is stored as fat.

Transcription factors that mediate the stimulatory effect of insulin include **USFs** (upstream stimulatory factors) and **SREBP-1**.

**SREBPs** (sterol response element binding proteins) were first identified for their regulation of cholesterol synthesis.

- ♦ Polyunsaturated **fatty acids diminish** transcription of the Fatty Acid Synthase gene in liver cells, by suppressing production of SREBPs.

# Fatty acid synthesis

## In fat cells:

Expression of SREBP-1 and of Fatty Acid Synthase is **inhibited** by **leptin**, a hormone that has a role in regulating food intake and fat metabolism.

**Leptin** is produced by fat cells in response to excess fat storage.

Leptin regulates body weight by decreasing food intake, increasing energy expenditure, and inhibiting fatty acid synthesis.

# Fatty acid synthesis

**Elongation** beyond the 16-C length of the palmitate product of Fatty Acid Synthase is mainly catalyzed by enzymes associated with the **endoplasmic reticulum** (ER).

ER enzymes lengthen fatty acids produced by Fatty Acyl Synthase as well as dietary polyunsaturated fatty acids.

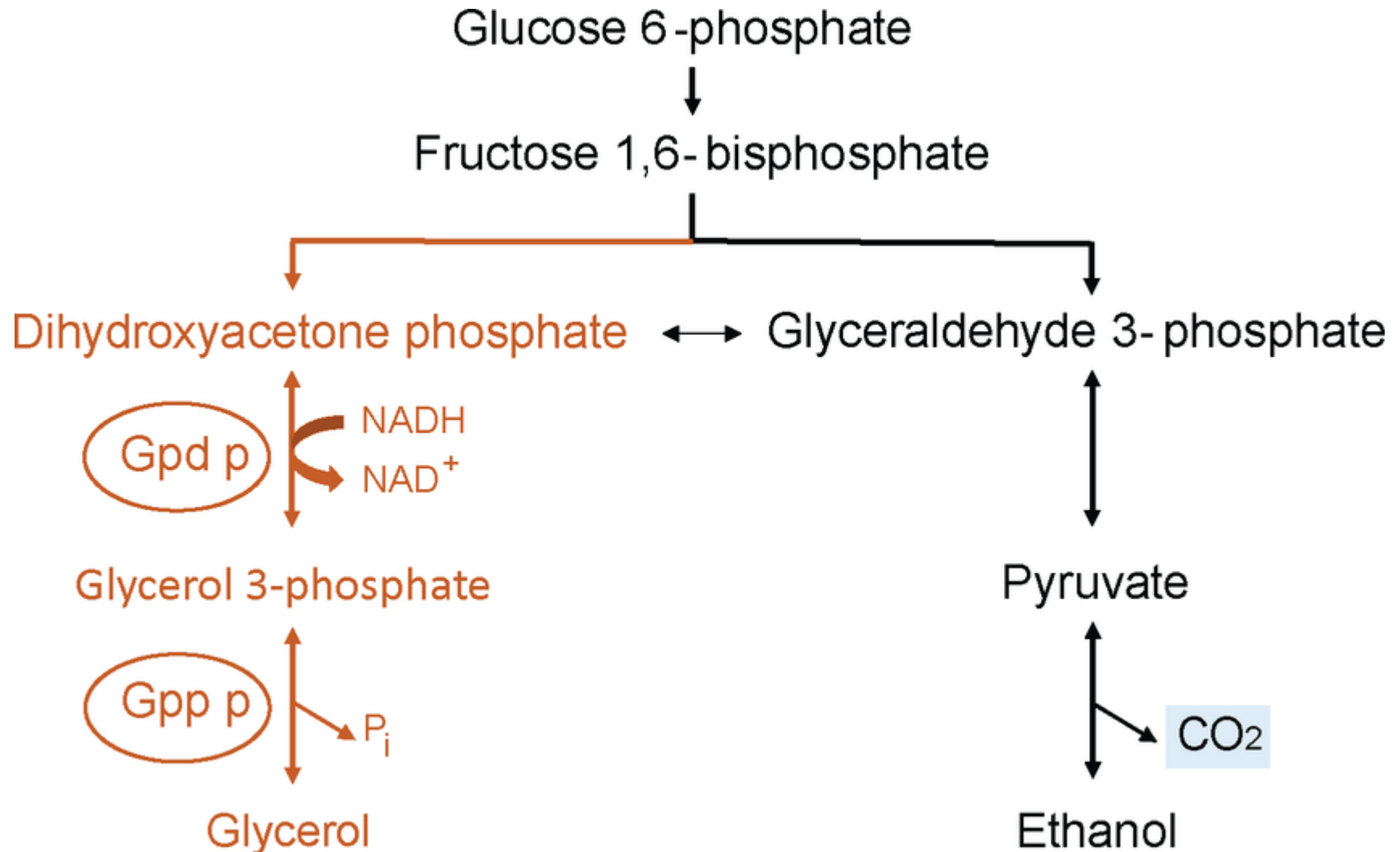
**Fatty acids** esterified to **coenzyme A** serve as substrates.

**Malonyl-CoA** is the donor of 2-carbon units in a reaction sequence **similar** to that of Fatty Acid Synthase except that individual steps are catalyzed by **separate proteins**.

A family of enzymes designated **Fatty Acid Elongases** or **ELOVL** (elongation of very long chain fatty acid) catalyze the initial condensation step.



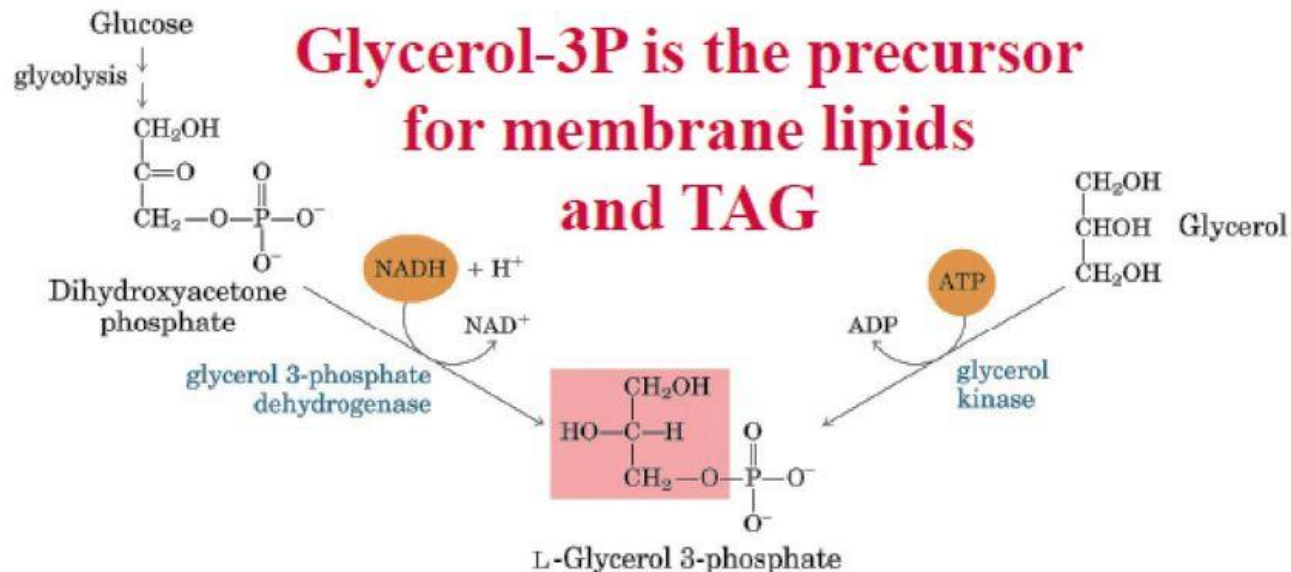
# Glycerol synthesis



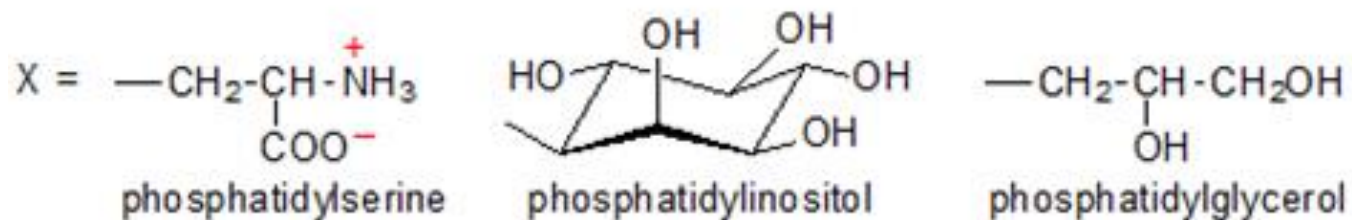
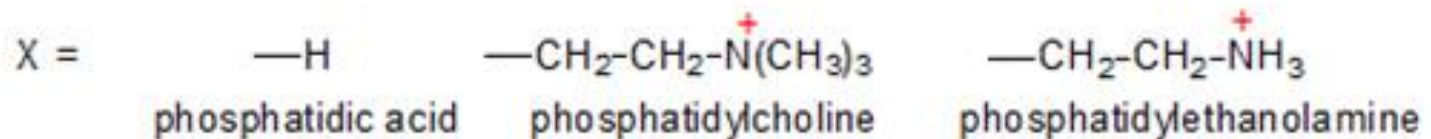
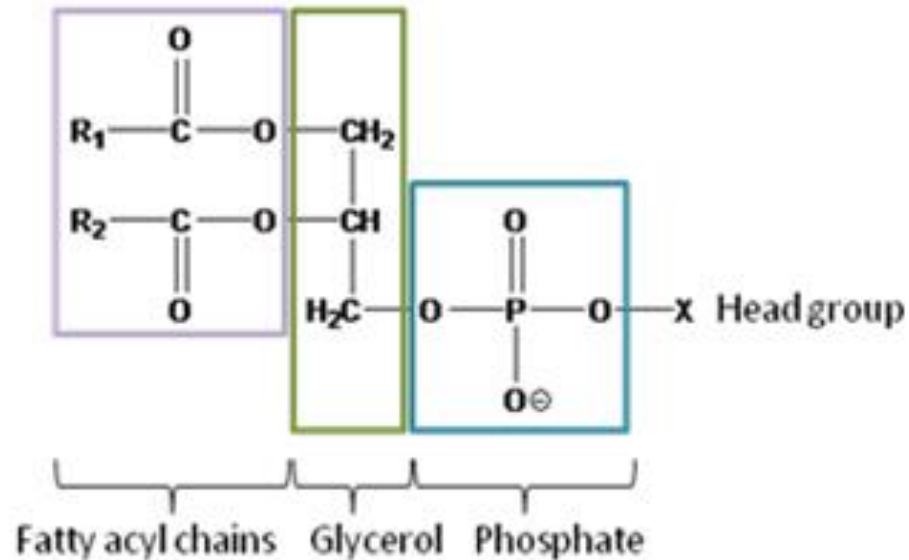
# Glycerol synthesis

## Biosynthesis of glycerol

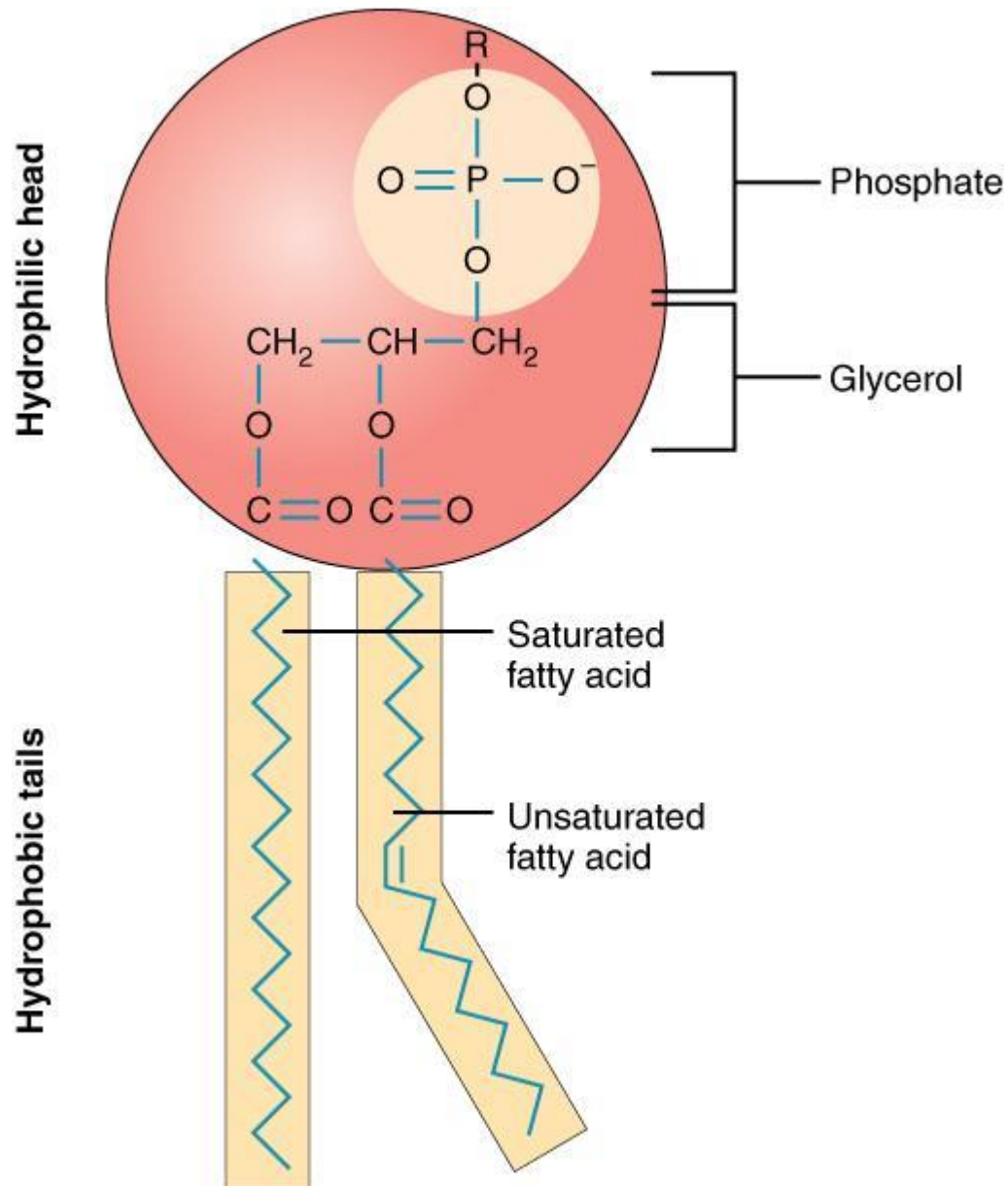
- Glucose is oxidized via glycolysis to **dihydroxy acetone phosphate**  
↓  
reduced to **glycerol-3 phosphate** by the enzyme glycerol-3 phosphate dehydrogenase.



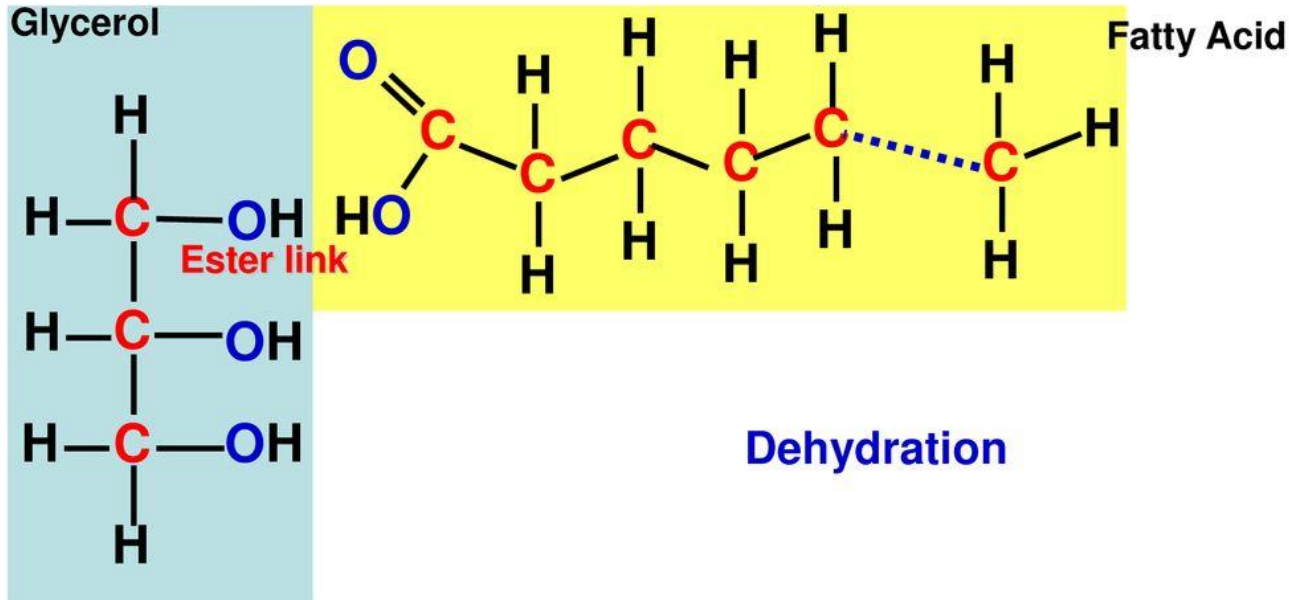
# Glycerol synthesis



# Phospholipid synthesis



# Phospholipid synthesis



A fat is composed of three fatty acids linked with glycerol.

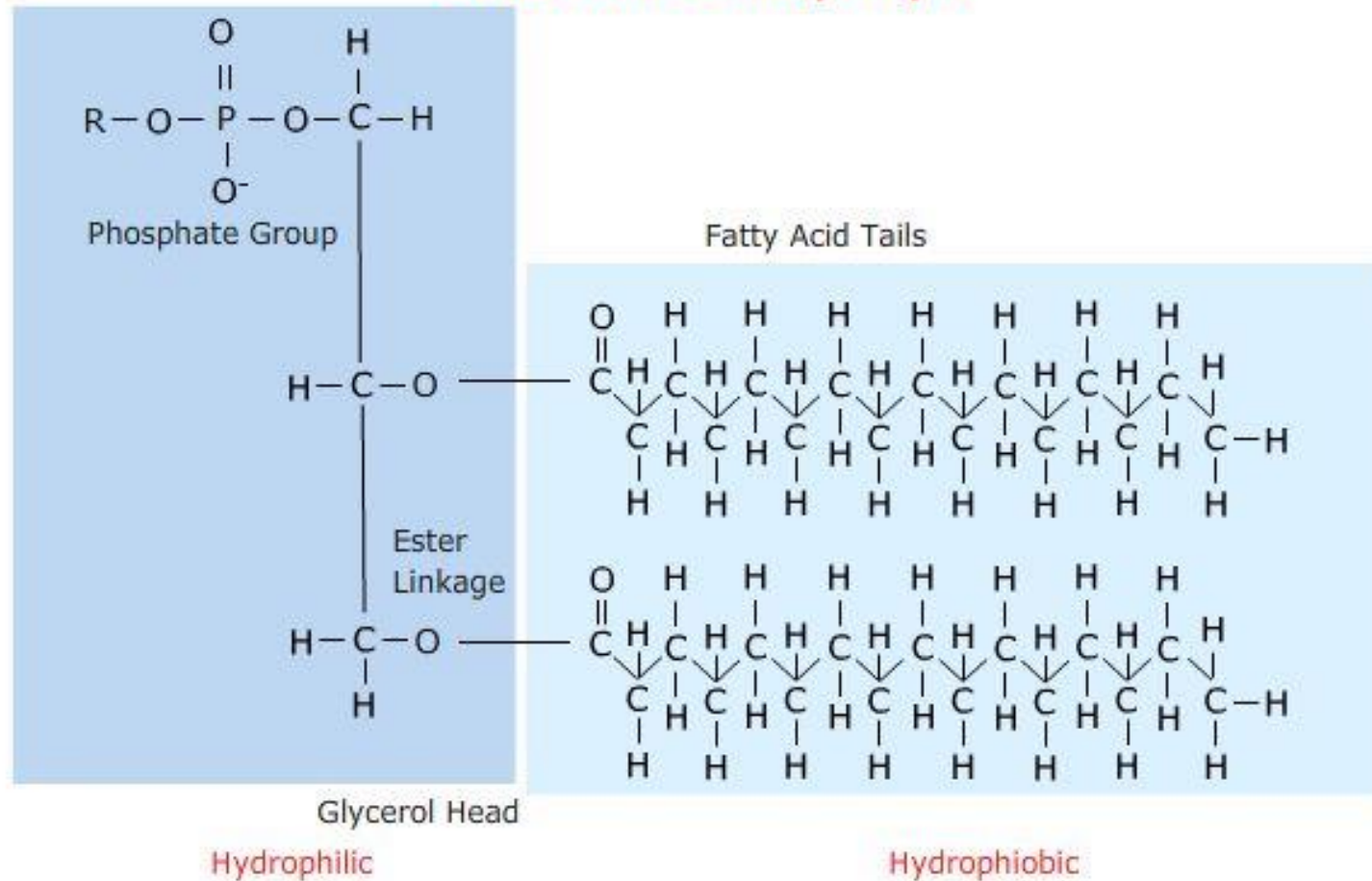
Fats are classified into **Saturated** and **Un-saturated** fats

Glycerol consists of a three **C** skeleton with an **OH** group attached to each.

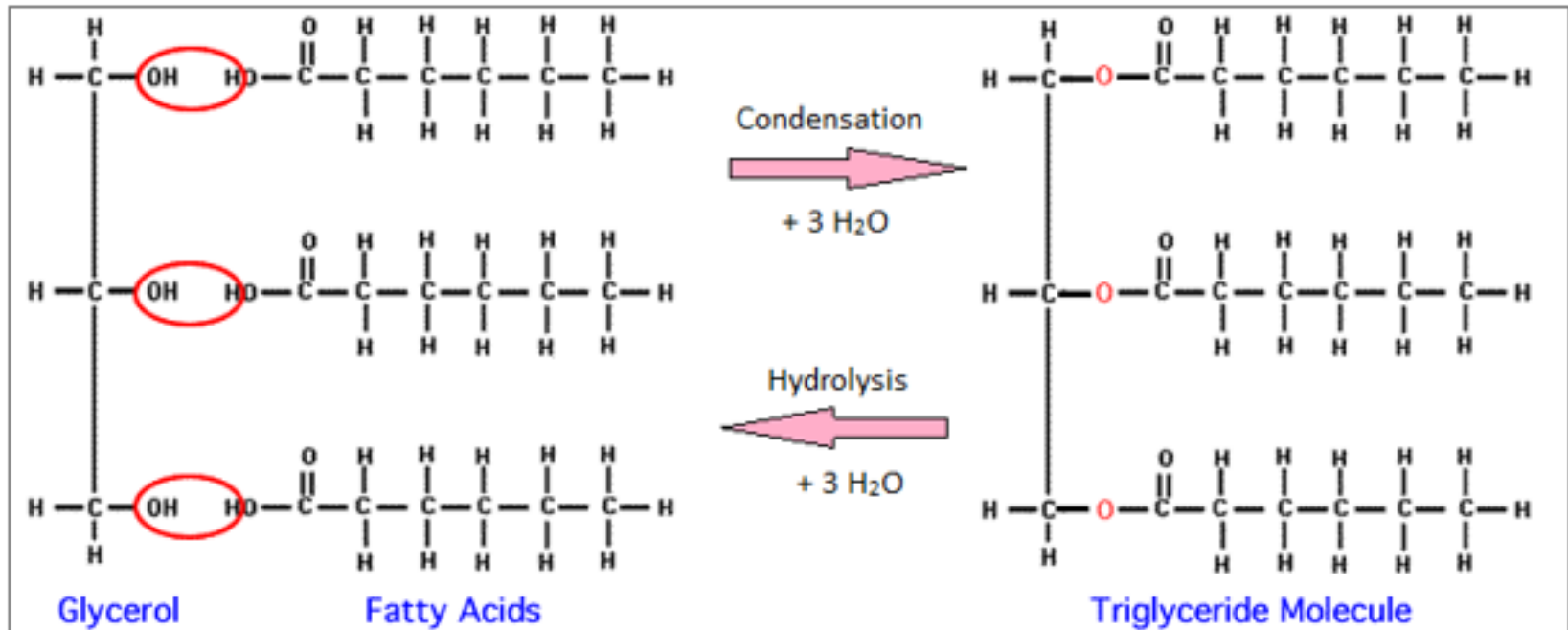
A fatty acid consists of a carboxyl group (**C=O**) attached to a long carbon skeleton, often 16 to 18 carbons long.

# Phospholipid synthesis

## Structure of a Phospholipid



# Triglyceride synthesis





# Biosynthesis of Triacyl Glycerols

