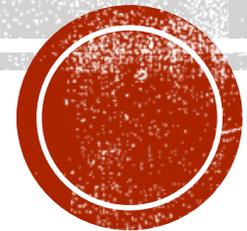


Adipose tissue metabolism

I. Lipogenesis & FA Synthesis



LIPOGENESIS

1. Definition

2. Site

3. Process

- A. **Synthesis of glycerol 3 phosphate (active glycerol).**
- B. **Synthesis & activation of long chain fatty acids (active FAs).**
- C. **Synthesis of the triacylglycerol**

Lipogenesis

Definition of lipogenesis:

It is the biosynthesis of **triacylglycerols (TGs)** principally from **excess glucose to be stored** after a carbohydrate rich meal.

Site

Cytoplasm of **most tissues** especially:

- Adipose tissue,
- Liver,
- lactating mammary gland
- Brain.

Lipogenesis

Processes for lipogenesis:

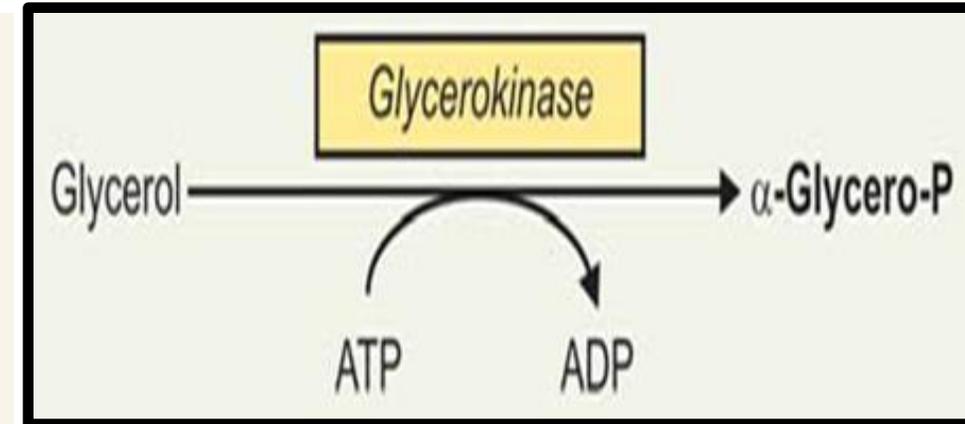
- **Lipogenesis can be divided into 3 processes:**
 - A. Synthesis of glycerol 3 phosphate (active glycerol).**
 - B. Synthesis & activation of long chain fatty acids (active FAs).**
 - C. Synthesis of the triacylglycerol (esterification of both active glycerol & active FA).**

A. Synthesis of Glycerol-3-P

Synthesis of Glycerol-3-P (active glycerol)

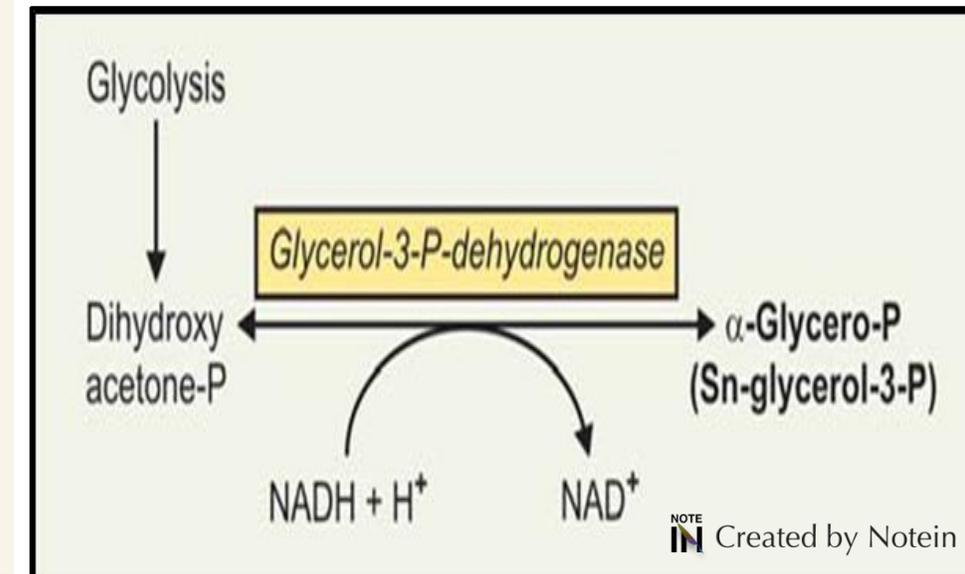
A) In tissues (except adipose tissue)

- Glycerol is activated by **glycerokinase** to glycerol-3-P (active glycerol)
- **Need ATP**



A) In Adipose tissue

- **No glycerokinase** is present in adipose tissue.
- Instead, glucose \rightarrow glycolysis \rightarrow **DHAP** (dihydroxy-acetone-phosphate) Which gives glycerol 3 P by **dehydrogenase** enzyme



B. Synthesis of fatty acids

Cytoplasmic (Extramitochondrial) FA synthesis Pathway

- Definition, site & building block
- Sources of acetyl CoA
- Translocation of acetyl CoA to cytoplasm
- Enzyme of extramitochondrial Pathway
- Steps of extramitochondrial Pathway
- Fate of palmitate

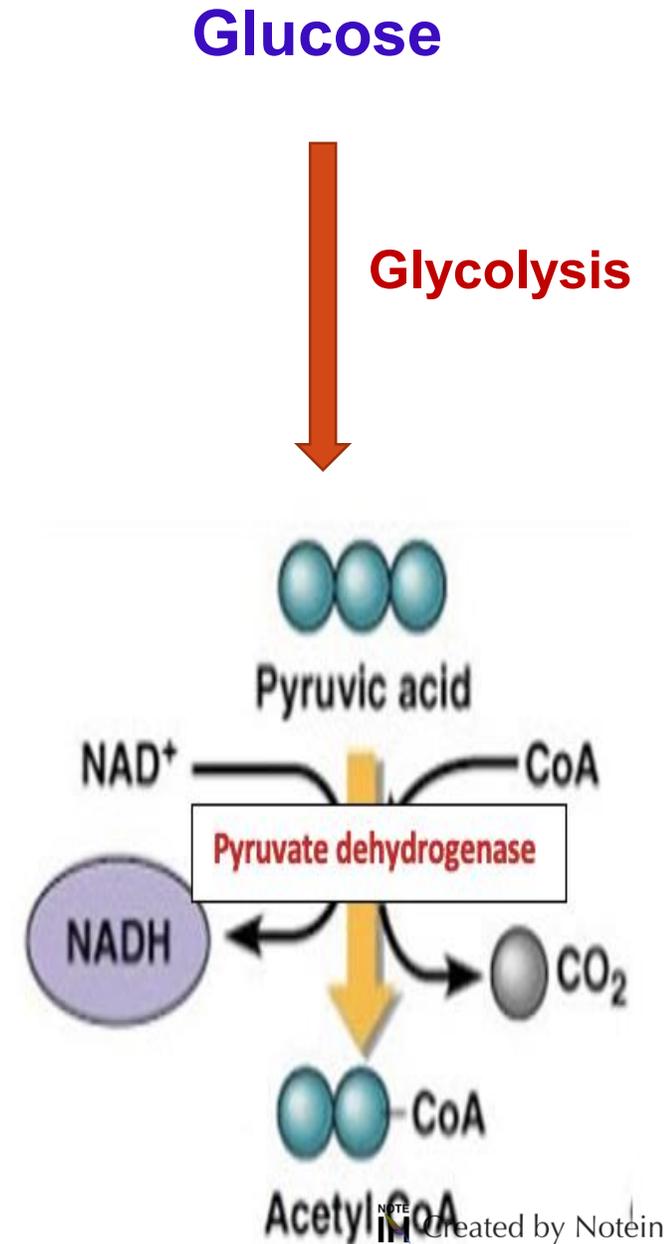
Cytoplasmic (Extramitochondrial) De novo Synthesis of long chain FA synthesis

- ❖ **Definition**: It is the only system responsible for **de novo** synthesis of long chain FA from **active acetate** (acetyl CoA).
- ❖ **Site**: **cytoplasm** of many tissues especially adipose tissue , liver, brain, lactating mammary gland.
- ❖ **Building block for FA synthesis**: Acetyl CoA
- ❖ **Product of FA synthesis**: long chain FA **Palmitic acid (16 carbons)**

❖ Steps

1. Source of acetyl-CoA for FA synthesis:

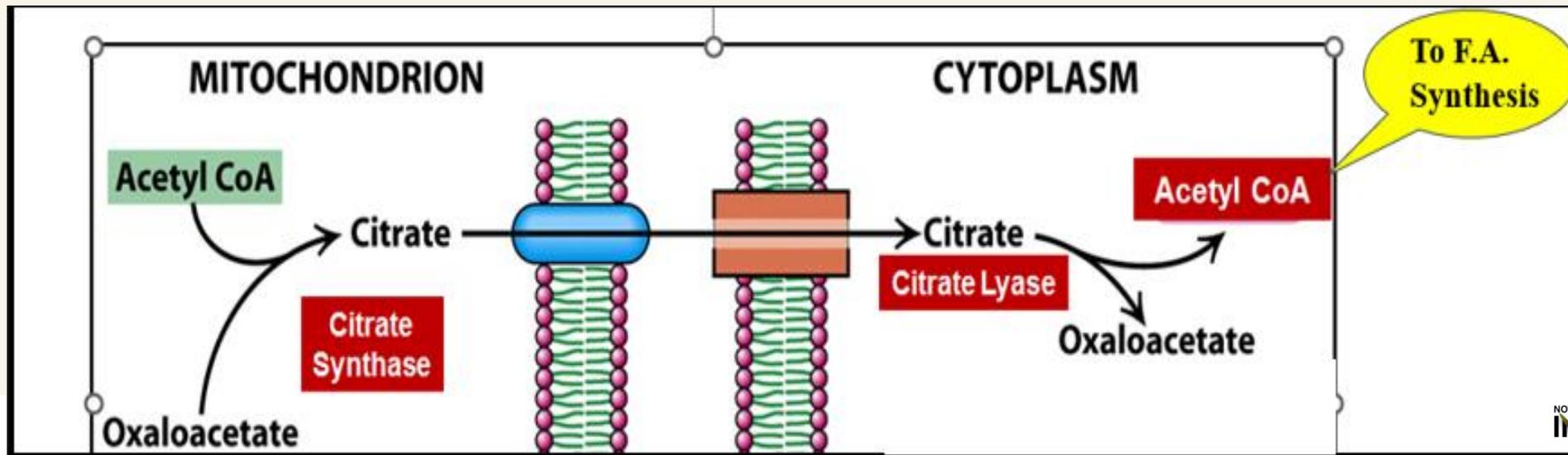
- **Mainly from Glucose (carbohydrates):**
(glucose → glycolysis → pyruvate → oxidative decarboxylation by **pyruvate dehydrogenase** enzyme in the **mitochondria** → acetyl CoA)
- **To less extent from ketogenic amino acids (proteins).**
- **Never from fats**



2. Translocation of Acetyl-CoA from mitochondria to cytoplasm

(Citrate shuttle)

- **Acetyl-CoA**, formed within mitochondria, **does not** penetrate the mitochondrial membrane to cytoplasm. 1st it must **condense with oxaloacetate** to form **citrate**, which can pass out mitochondrial membrane **(Citrate shuttle)**.
- In cytoplasm **citrate splits** again by **citrate lyase** enzyme into
 - a) **Acetyl-CoA** and
 - b) **oxaloacetate**.



3. Enzymes of cytoplasmic Fatty acid synthesis:

2 Enzymes are needed for this process:

1-Acetyl CoA carboxylase

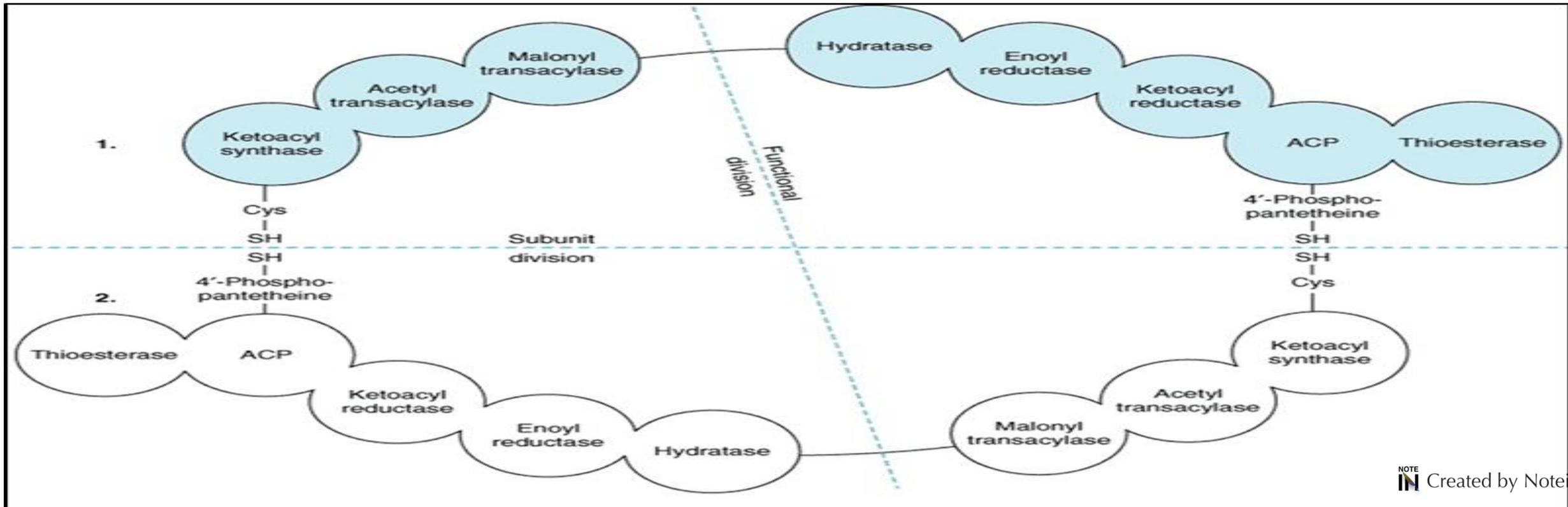
- (The key regulatory enzyme for Lipogenesis)
- This enzyme is responsible of synthesis of **malonyl Co A** from the acetyl CoA

2- The fatty acid synthase complex

- It is a multi-enzyme complex, formed of **2 subunits** (dimers)
- Each subunit contains all **7 enzymes** of FA synthase & **2 SH groups**
- This enzyme is responsible of adding **malonyl Co A** to the acetyl CoA in **7 cycles** to form the **palmitic acid** (16 Carbons FA).

2- The fatty acid synthase complex

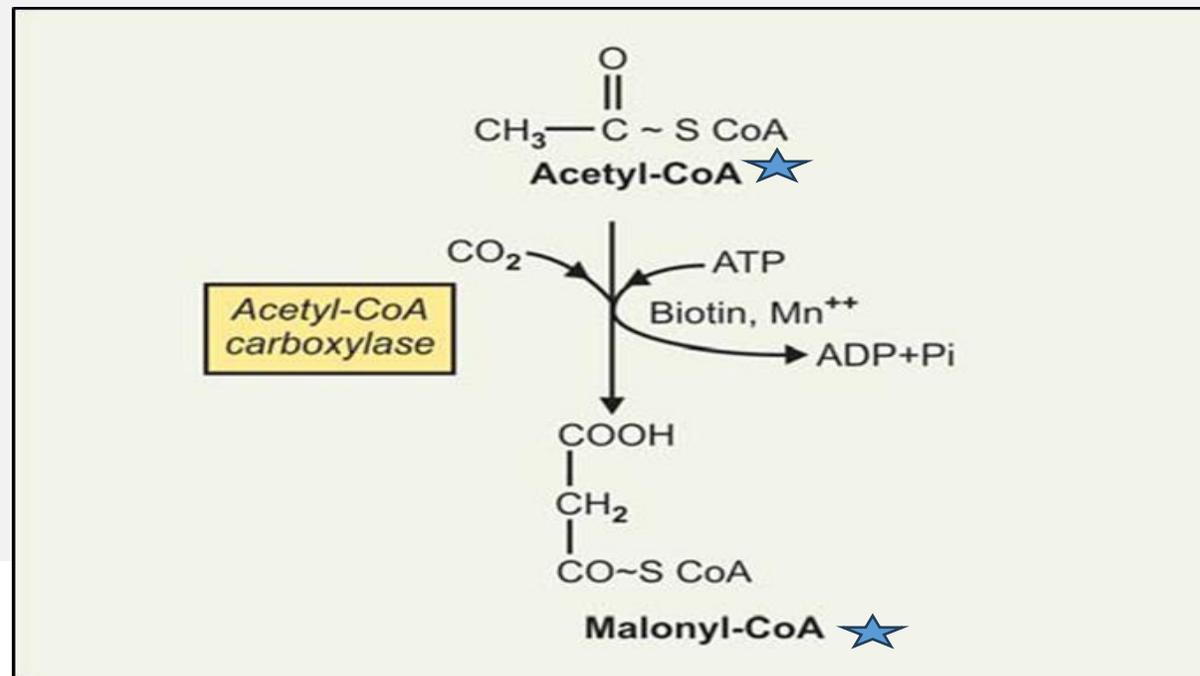
- **2 subunits** (dimers), lie in **head to tail** configuration.
- Each subunit contains all **7 enzymes** of FA synthase & **2 SH groups**



4. Steps of Extra-mitochondrial synthesis of palmitic acid (16 carbon FA)

1. Synthesis of malonyl CoA (3C):

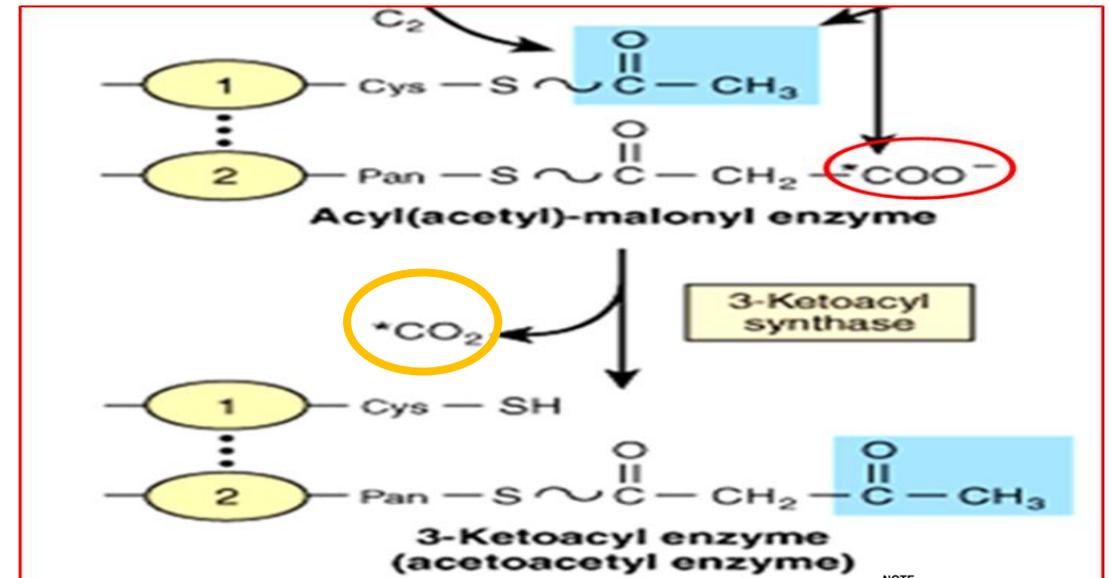
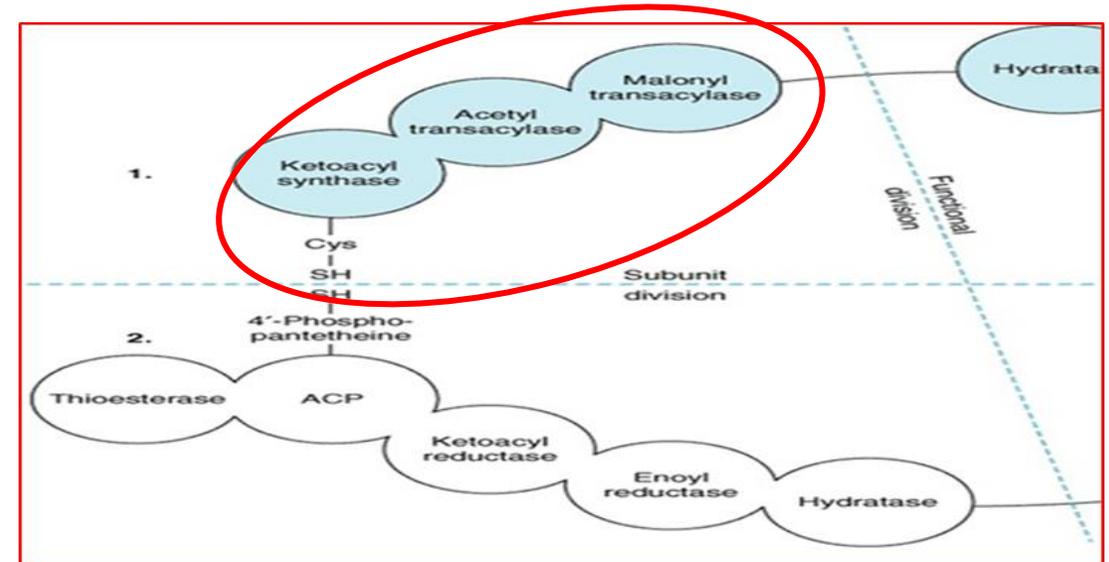
- It is synthesized from acetyl CoA by **acetyl CoA carboxylase**
- In presence of **ATP, biotin and bicarbonate** as a source of CO₂.



2. Synthesis of palmitate by the fatty acid synthase complex:

The first 3 steps involves 3 enzyme actions:

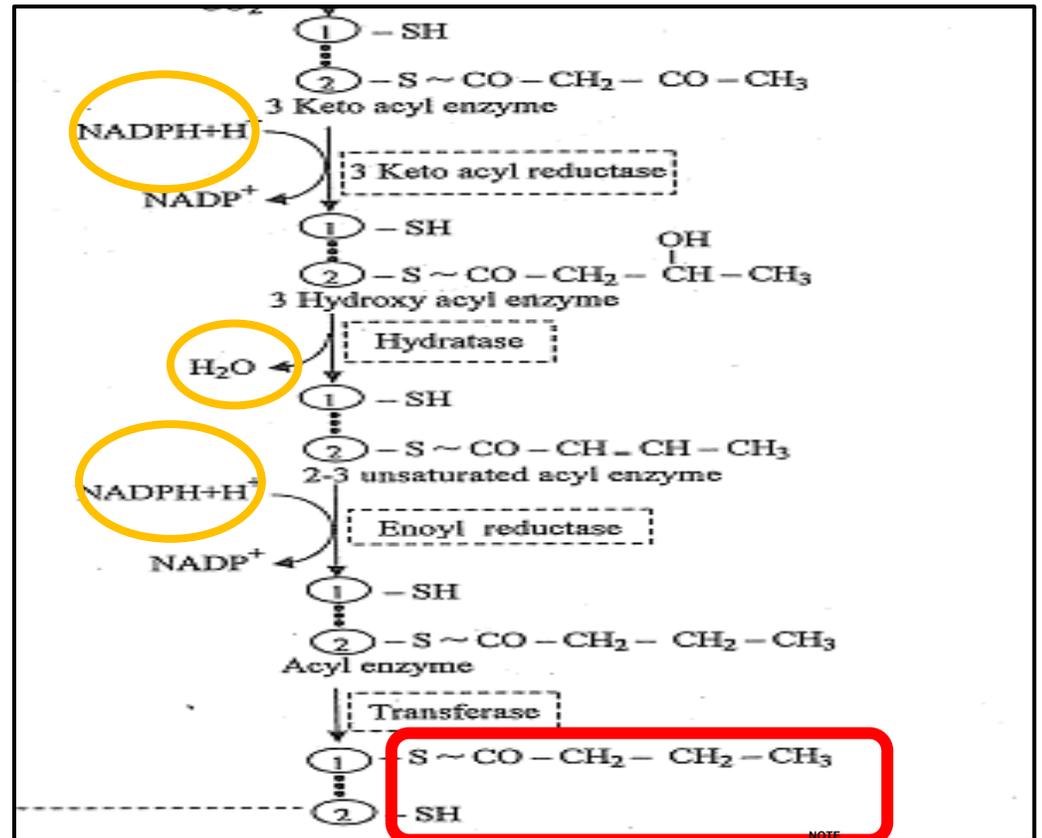
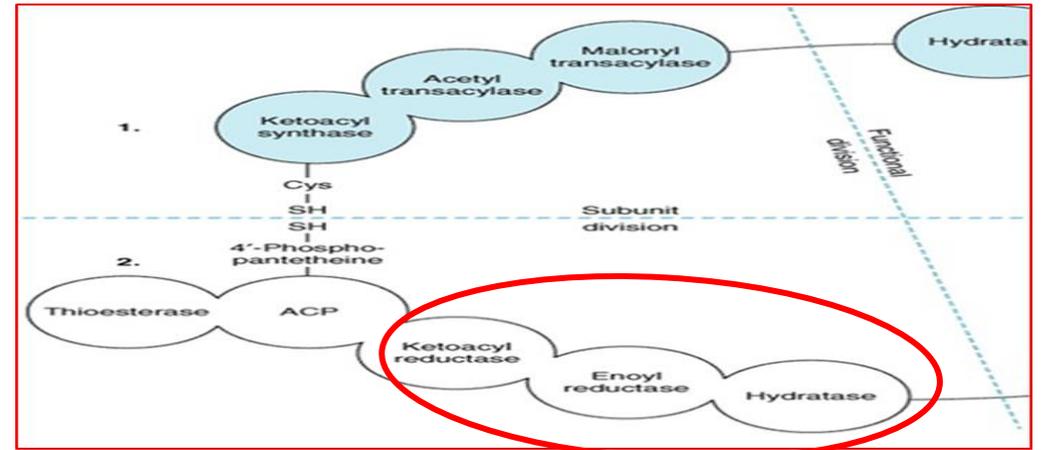
- 1st step is addition of an acetyl CoA to SH group at position 1
- 2nd step is addition of a malonyl CoA to SH group at position 2
- 3rd step is condensation of the acetyl CoA in position 1 with the malonyl CoA in position 2 with the release of **one CO₂**.



- A next 4 steps process involves 3 enzyme actions:

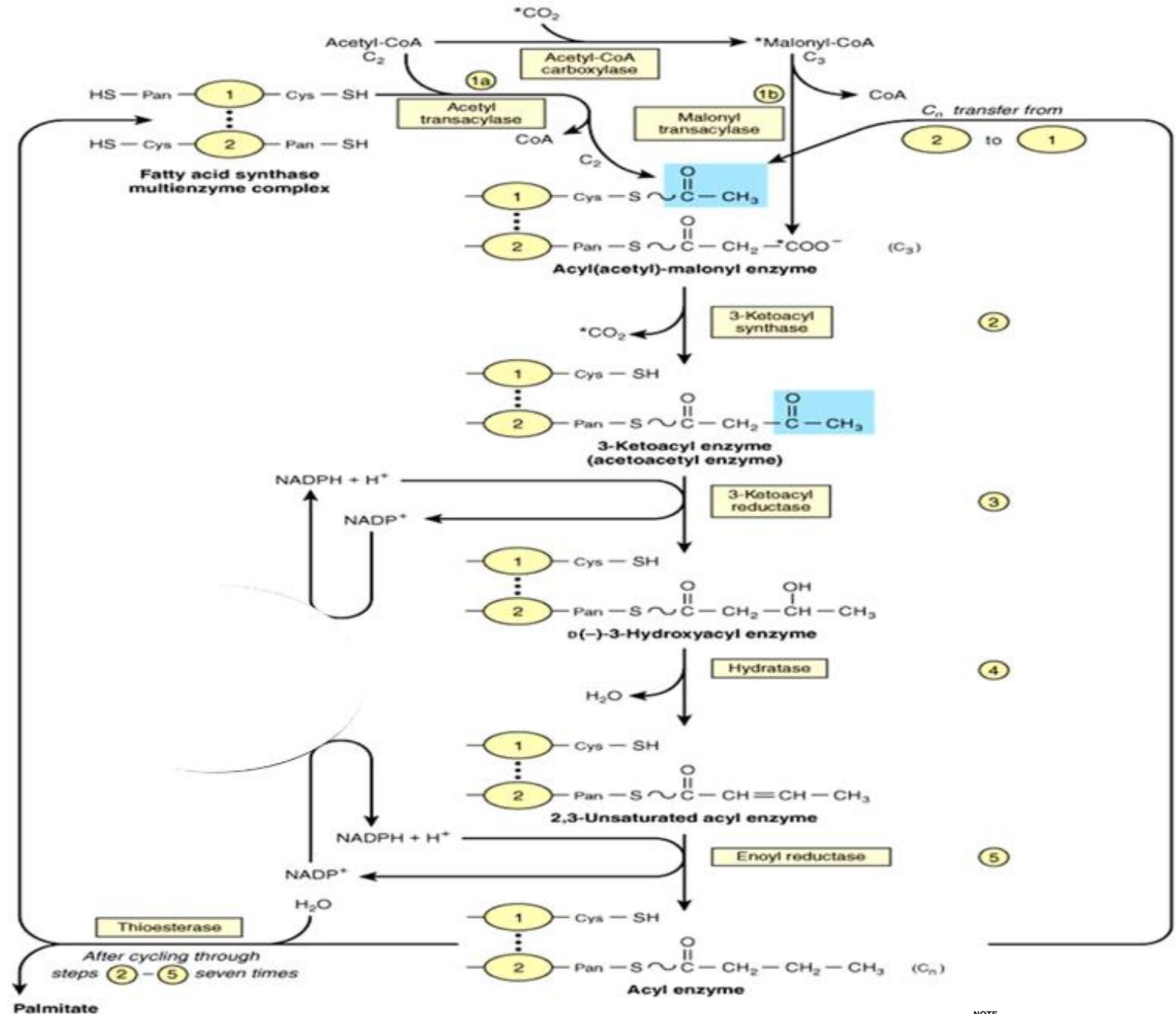
1. Reduction in the presence of **NADPH+**.
2. Dehydration by release of one H₂O.
3. Reduction in the presence of **NADPH+**.
4. Transfer from position 2 to 1.

- Now position 2 is free to accept a new malonyl CoA.



- The process is **repeated 6** more times.

- At the end of FA synthesis, **thioesterase** hydrolyzes the thioester bond to release palmitate from the enzyme complex.



❖ The overall reaction for palmitic acid synthesis:

7 molecules of malonyl CoA is added to **1 molecule of acetyl CoA** in 7 cycles by FA synthase enzyme complex to form **palmitic acid**



FA synthase complex



5. Main product of FA synthesis : free Palmitic acid (16 C)

❖ Fates of palmitic acid

Free palmitic acid must be activated to **palmitoyl CoA** (by **thiokinase= acyl CoA synthetase**) to be used for:

- 1. Esterification** with glycerol (to form TG)**** or with cholesterol (to form cholesterol esters).
- 2. Chain elongation:** For synthesis of longer chain F.A.
- 3. Desaturation:** Synthesis of unsaturated FA.
- 4. Sphingosine formation:** palmitoyl CoA + Serine

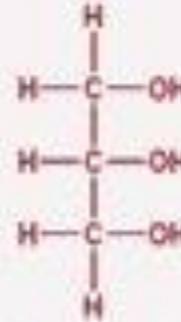
C. Synthesis of Triglycerides (esterification of active glycerol and active FA)

1- Synthesis of glycerol-3-phosphate

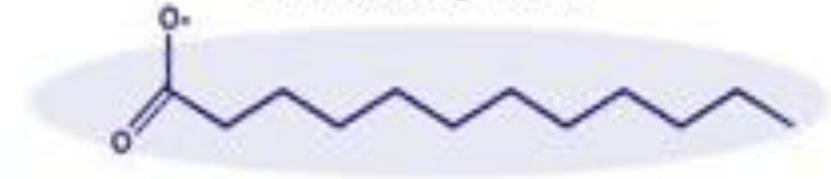
2- Synthesis of fatty acyl CoA

3- Synthesis of TG

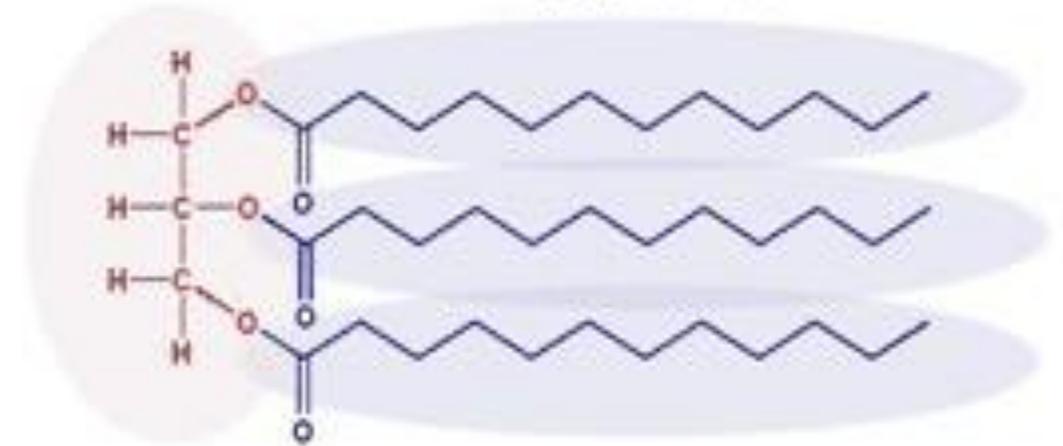
Glycerol



Free fatty acid



Triglyceride



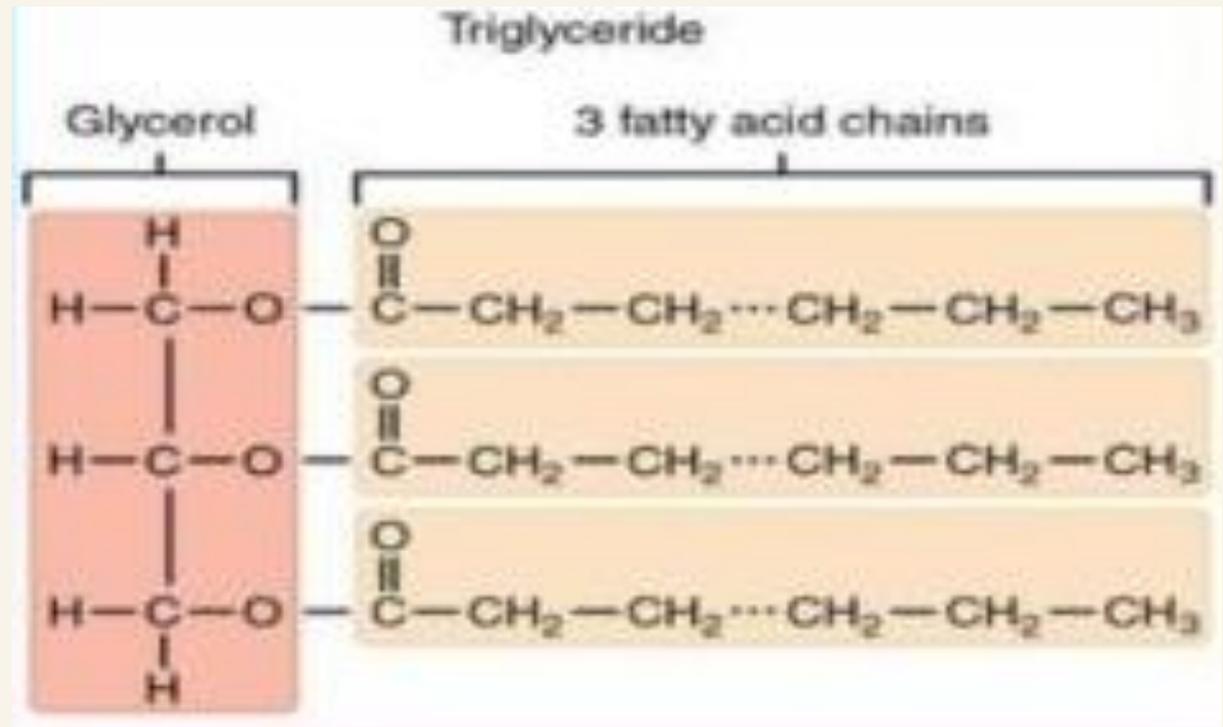
Synthesis of Triglycerides

➤ The active FA then esterified to active glycerol forming triacylglycerol (TAG)

➤ TAG is formed of:

- 1 glycerol
- 3 FAs

Attached by **ester bonds**



➤ TGs then stored **Mainly in adipocyte** to be ready for mobilization (utilization) if the body needs energy

REGULATION OF ADIPOSE TISSUE METABOLISM

6. Regulation of adipose tissue metabolism

Dietary, metabolic and hormonal factors regulate adipose tissue metabolism :

1- Dietary factor:

- Excess glucose (↑ carbohydrate meal) → ↑↑ lipogenesis
- Fasting → ↑↑ lipolysis

2- Metabolic factor

- Uncontrolled Diabetes mellitus → ↑↑ lipolysis

3- Hormonal factor regulation

I. Rate limiting enzyme

- ❖ Rate limiting enzyme of **lipogenesis** is **Acetyl-CoA carboxylase** which present in 2 forms
 - **Active form**: Dephosphorylated
 - **Inactive form** Phosphorylated
- ❖ Rate limiting enzyme of **lipolysis** is **Hormone sensitive lipase** which present in 2 forms
 - **Active form**: Phosphorylated
 - **Inactive form**: Dephosphorylated

3- Hormonal factor regulation

II. During fed state

- ❖ ↑ Blood glucose → release of **insulin** → **dephosphorylation** of the enzymes:
 - **Dephosphorylate HSL** → inactivate it → ↓↓ **lipolysis**
 - **Dephosphorylate acetyl CoA carboxylase** → ↑↑ **lipogenesis**

III. During fasting

- ❖ ↓ Blood glucose → release of **anti-insulin hormones** (glucagon & adrenalin) → **activate adenyl cyclase** enzyme → **Phosphorylation** of the enzymes:
 - **Phosphorylate HSL** → activate it → ↑↑ **lipolysis**
 - **Phosphorylate acetyl CoA carboxylase** → ↓↓ **lipogenesis**

QUESTION 1

- Which of the following cofactors or their derivatives must be present for the conversion of acetyl CoA to malonyl CoA in extra-mitochondrial fatty acid synthesis?
 - (A) Biotin
 - (B) FAD
 - (C) FMN
 - (D) ACP
 - (E) NAD^+

QUESTION 3

Fatty acid synthase enzyme complex involves which of these vitamins:

- a) Vit B1
- b) Vit B3
- c) Vit B5
- d) Vit B10
- e) Vit B12

QUESTION 2

Acetyl CoA required for extra mitochondrial fatty acid synthesis is produced mainly by:

- (A) Pyruvate dehydrogenase complex
- (B) Pyruvate carboxylase
- (C) Citrate lyase
- (D) Thiolase
- (E) Carnitine-acyl transferase

Life
isn't about
finding yourself.

...

Life
is about
creating yourself.



References

