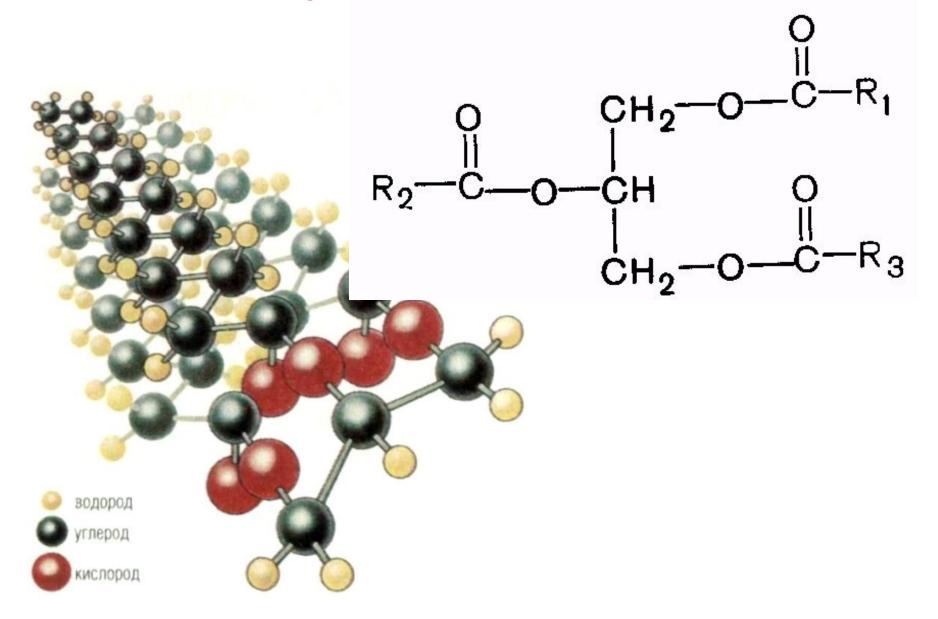
LIPID METABOLISM



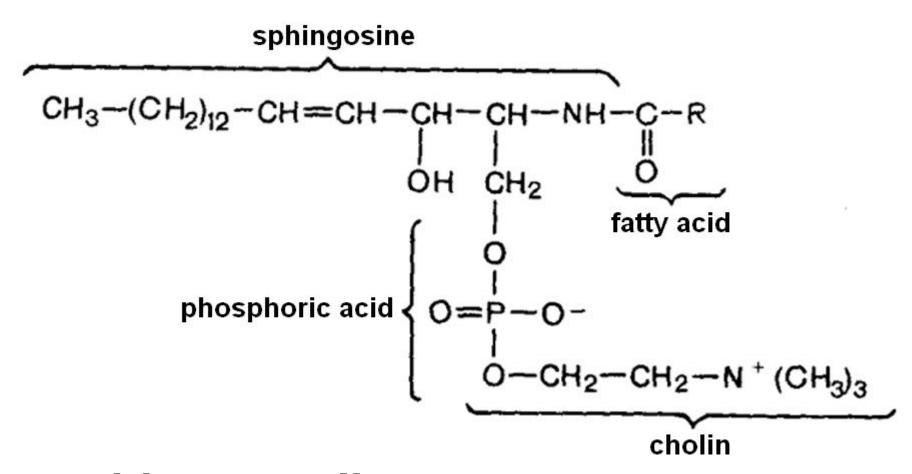
Fats (Triacylglycerols)



Phospholipids

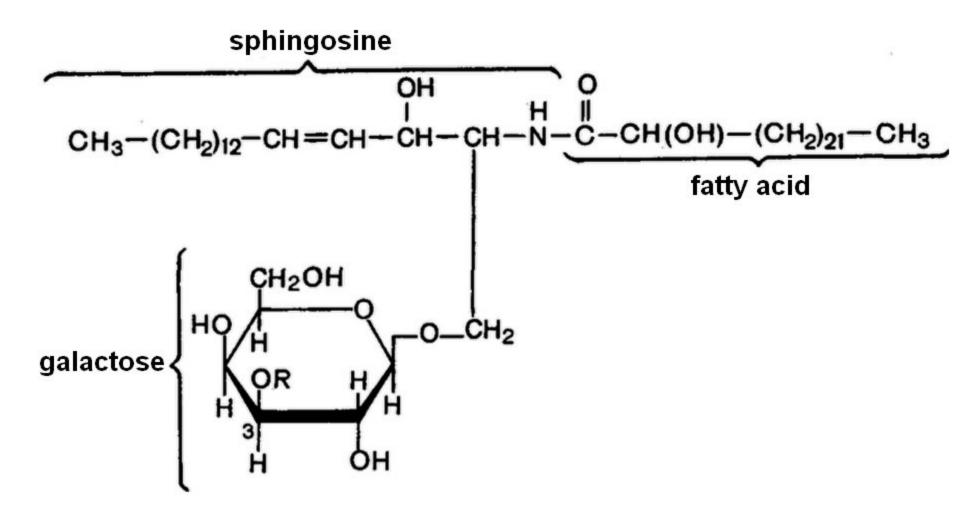
acetylcholine

Sphingophospholipids

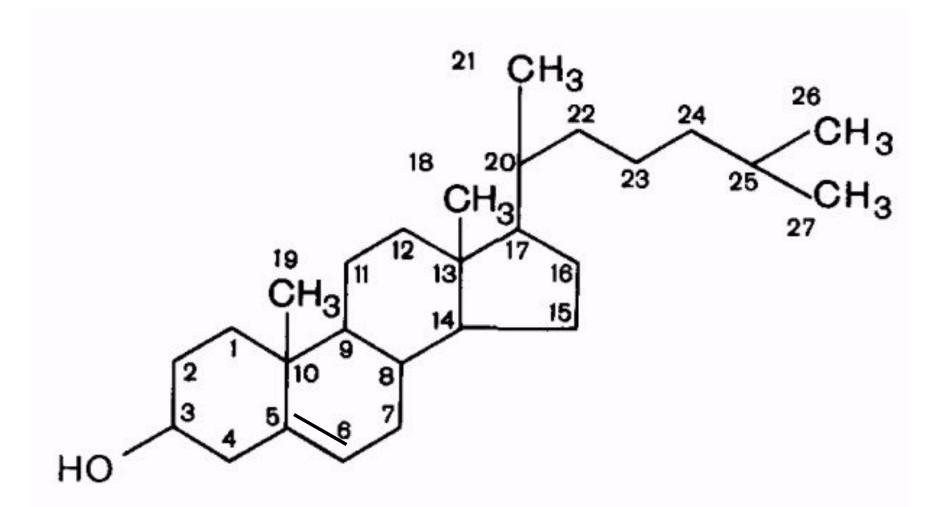


sphingomyelin

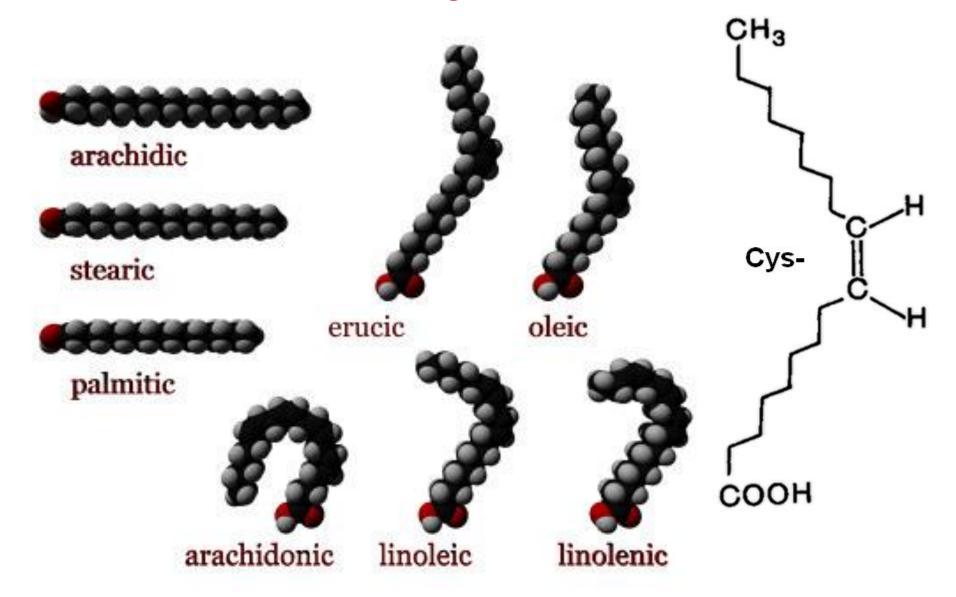
Glycolipids



Cholesterol



Fatty acids

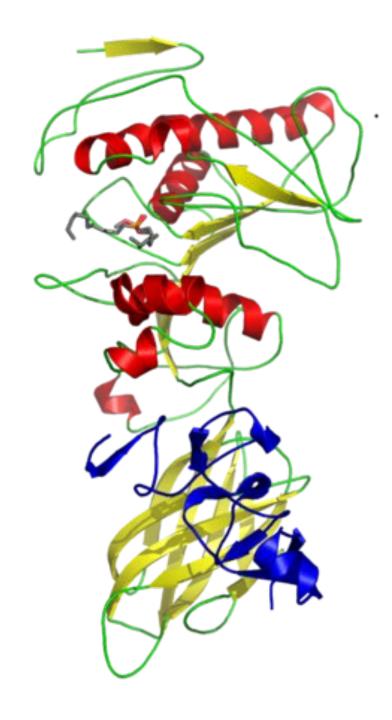


Bile acids

(b) Sodium glycocholate (a bile salt)

Lipid functions

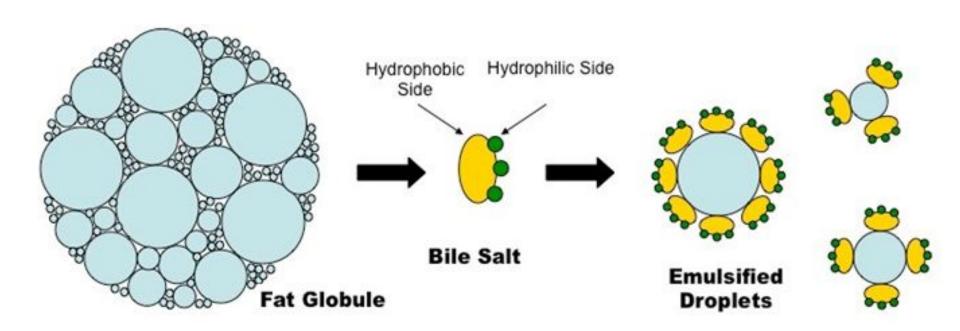
- Storage form of energy
- Supply essential fatty acids
- Structural components of cell membranes
- Electrical insulation
- Protect body from cold
- Mechanical protection of internal organs
- Metabolic regulators (hormones)
- Help transport fat soluble vitamins



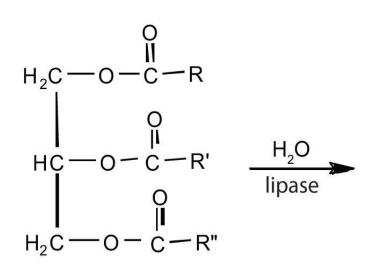
Human pancreatic lipase (activation by colipase)

Colipase is colored in blue

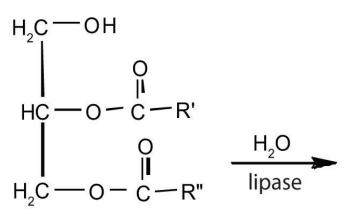
Emulsification



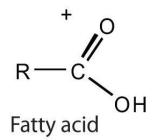
Breakdown of fats

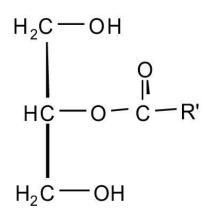


Triglyceride

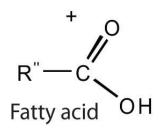


Diglyceride

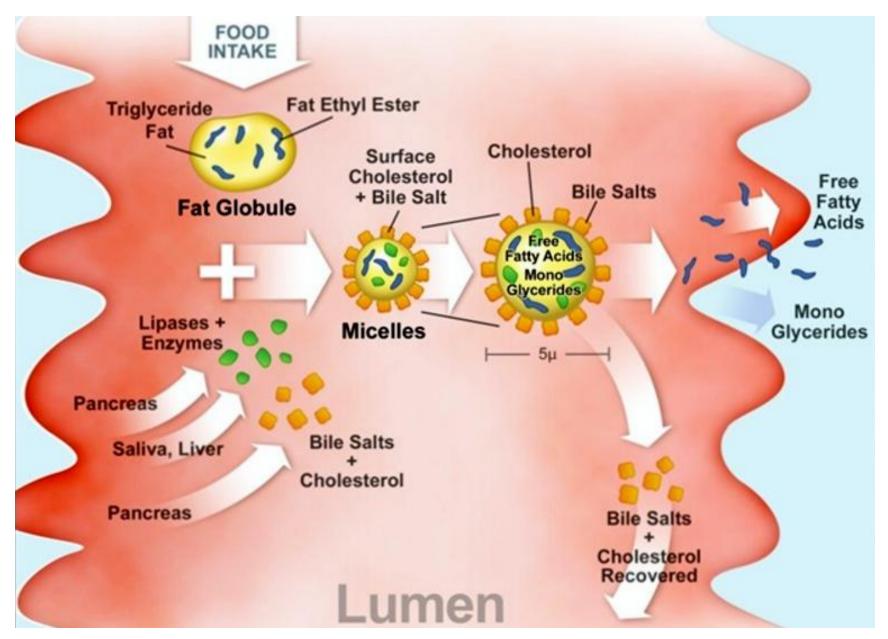




Monoglyceride



Absorbtion



Steatorrhea

Increased excretion of neutral fat

Increased excretion of fatty acids

Pancreatic exocrine insufficiency

Intestinal pathology

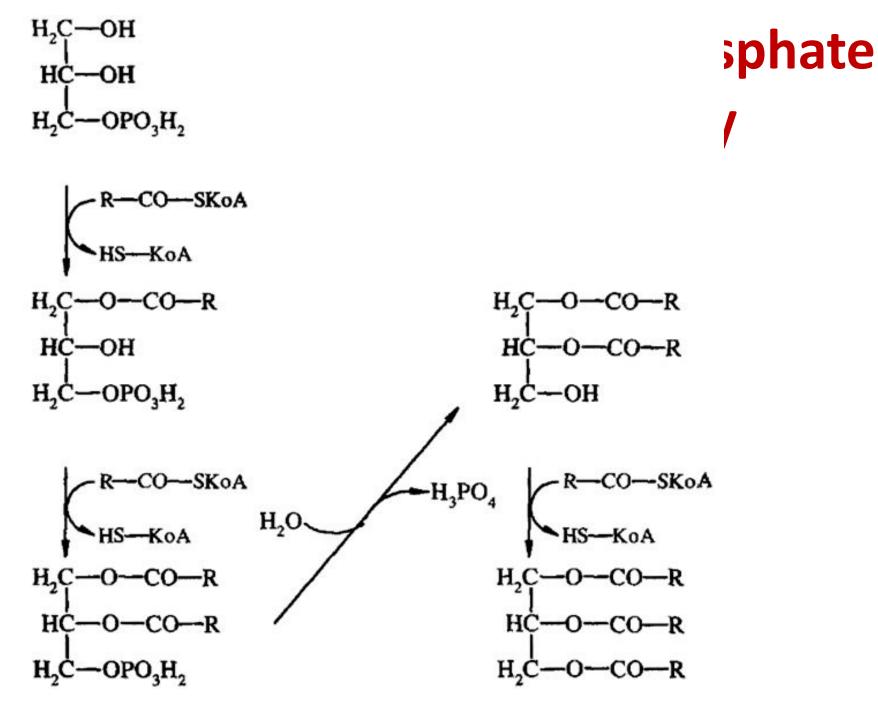
Lipid resynthesis in cells of intestinal wall epithelium β-monoglyceride pathway

β-monoglyceride+R-CO-SCoA →

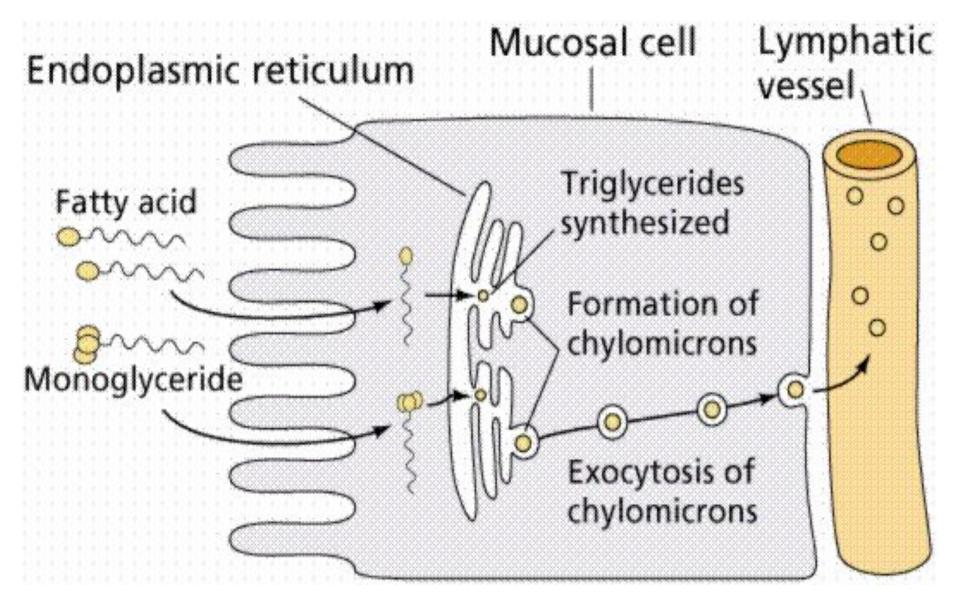
→ diglyceride + HS-CoA

diglyceride+R-CO-SCoA→

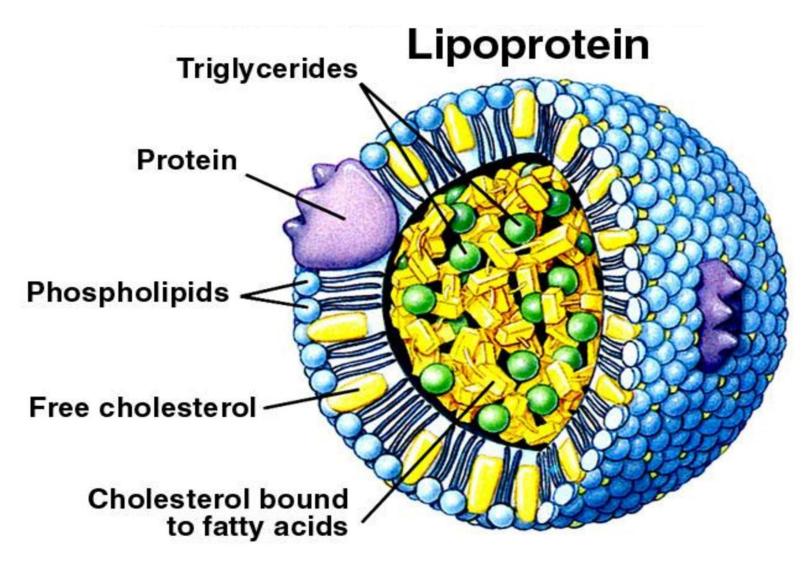
→ triglyceride +HS-CoA



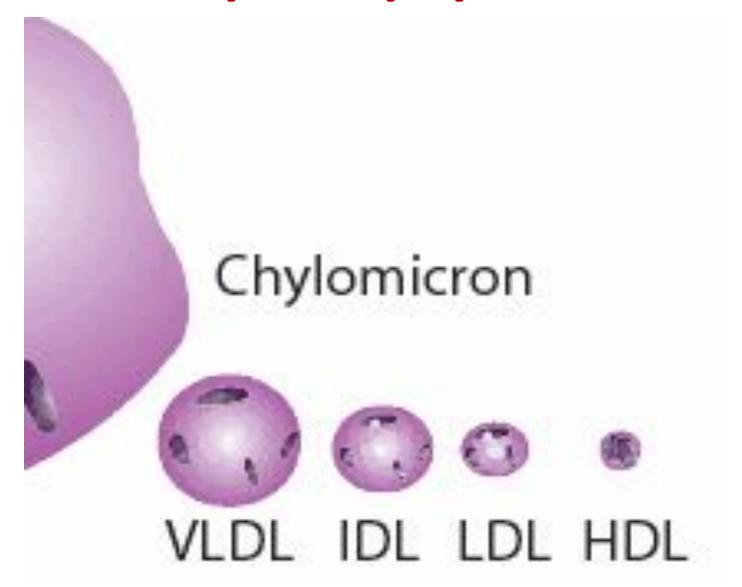
Formation of chylomicrons



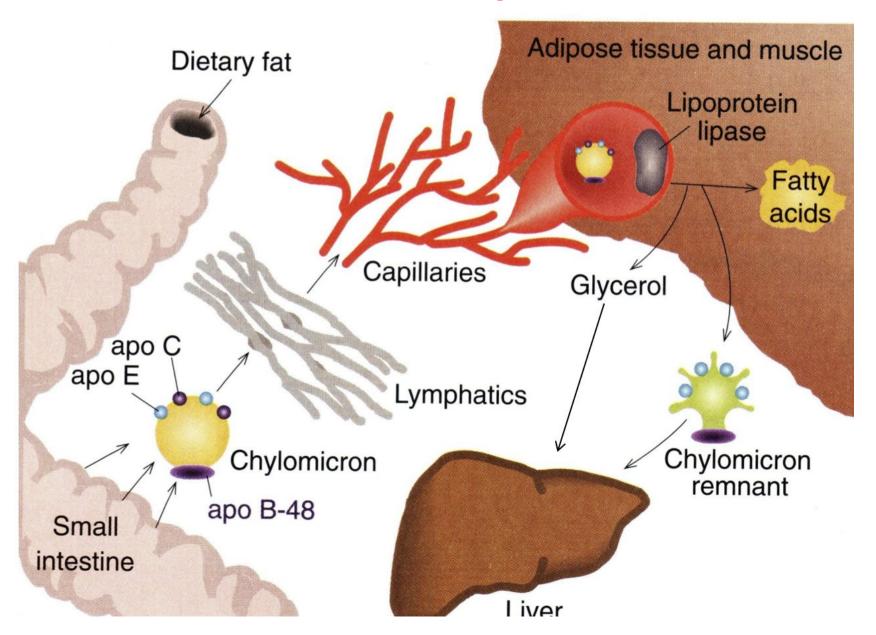
Lipoprotein structure



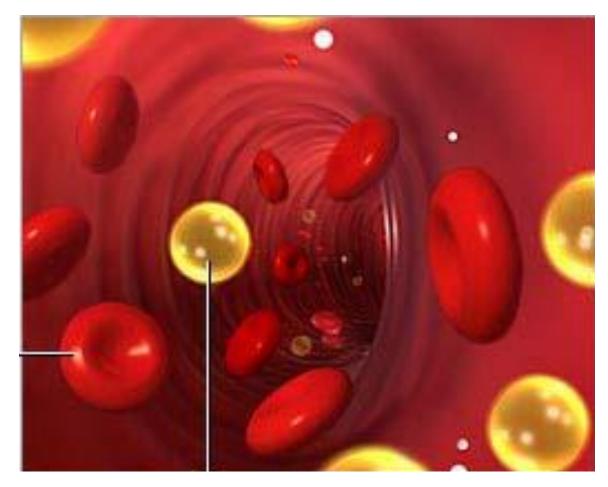
Transport lipoproteins



Fat transport

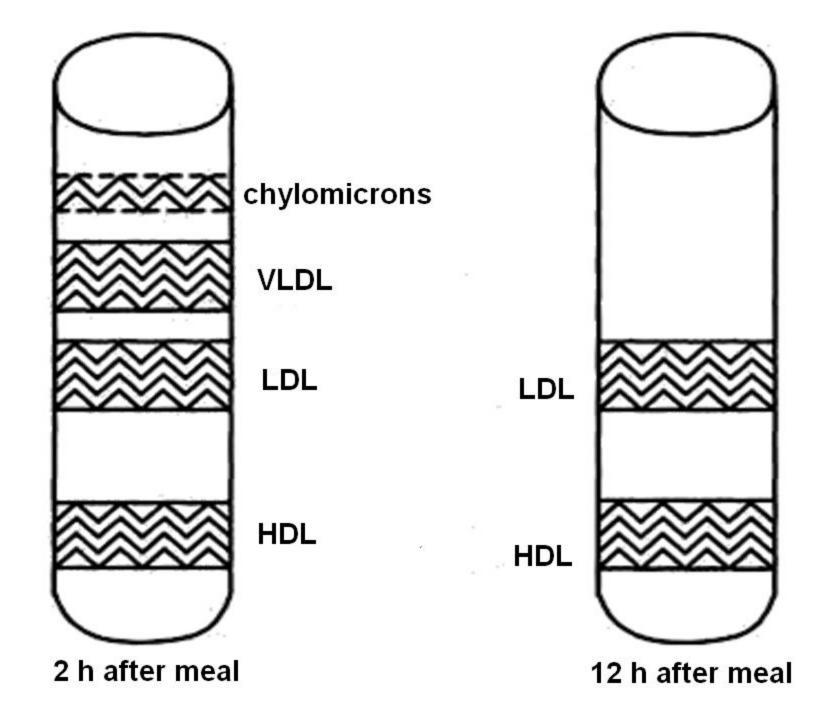


Artery section



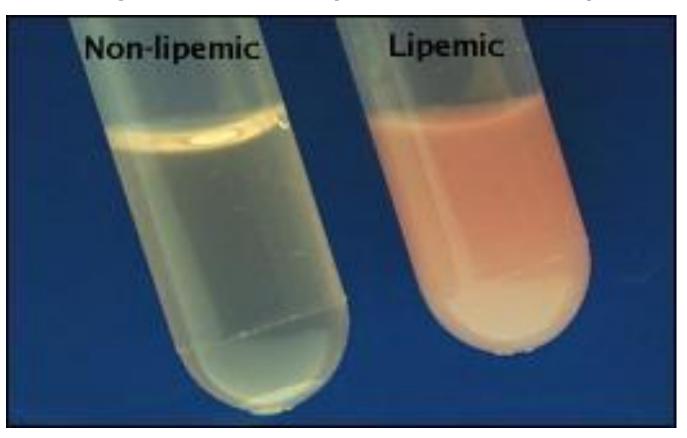
RBC

Chylomicron



Dislipoproteinemia (hyperlipoproteinemia) Type I

- □ Reduced activity of lipoprotein lipase (LPL)
- □ Deficiency of apoCII (LPL activator)



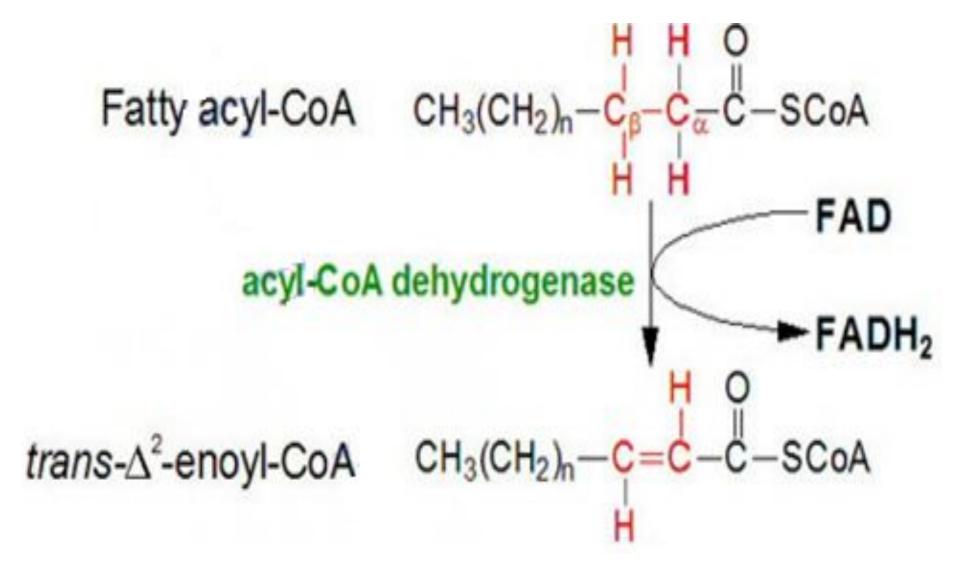
Tissue lipase adipocyte activation **Proteinkinase ATP** inactive Glucagon cAMP **Adrenalin Proteinkinase** active blood **TAG-lipase TAG-lipase** inactive active **Glycerol Glycerol TAG Oxidation Fatty Fatty** in tissues acids acids

β-Oxidation of fatty acids

carnitine acyl transferase

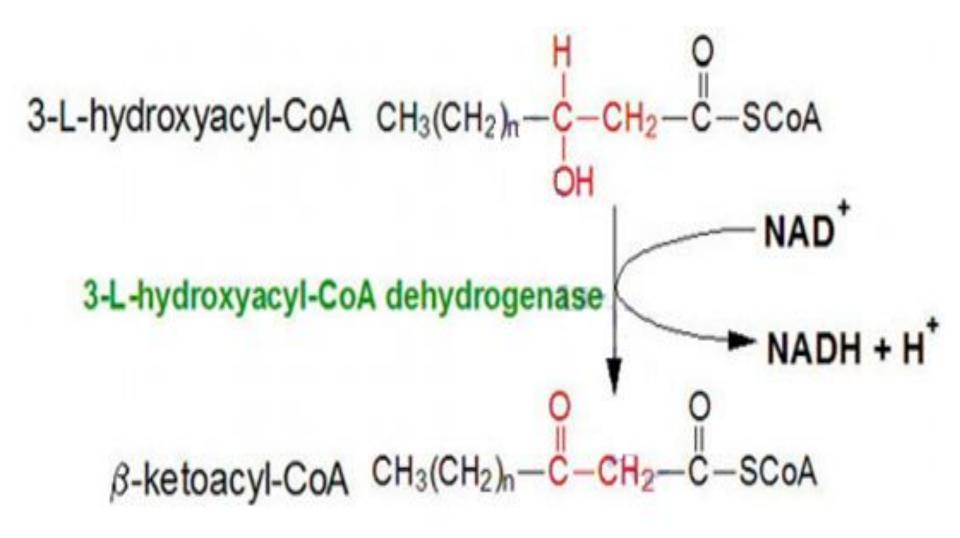
Breakdown of acyl carnitin in mitochondria

Dehydrogenation



Hydration

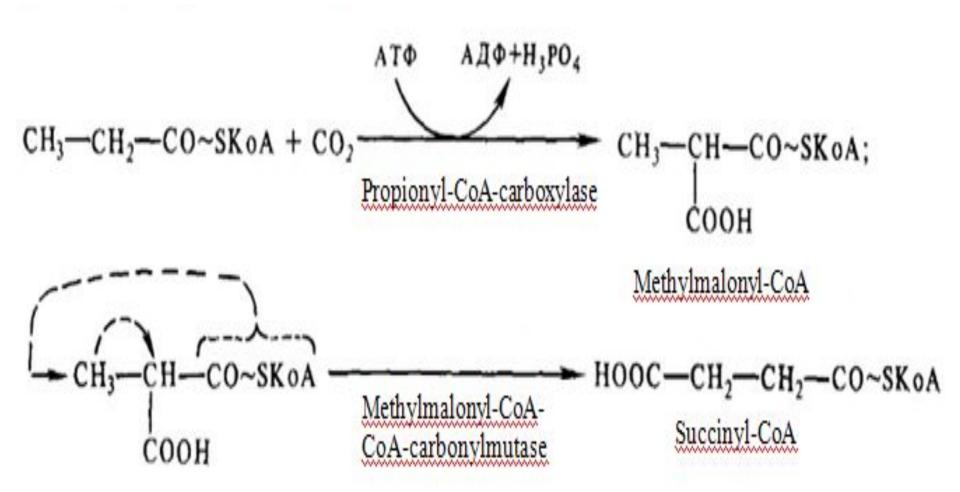
Dehydrogenation

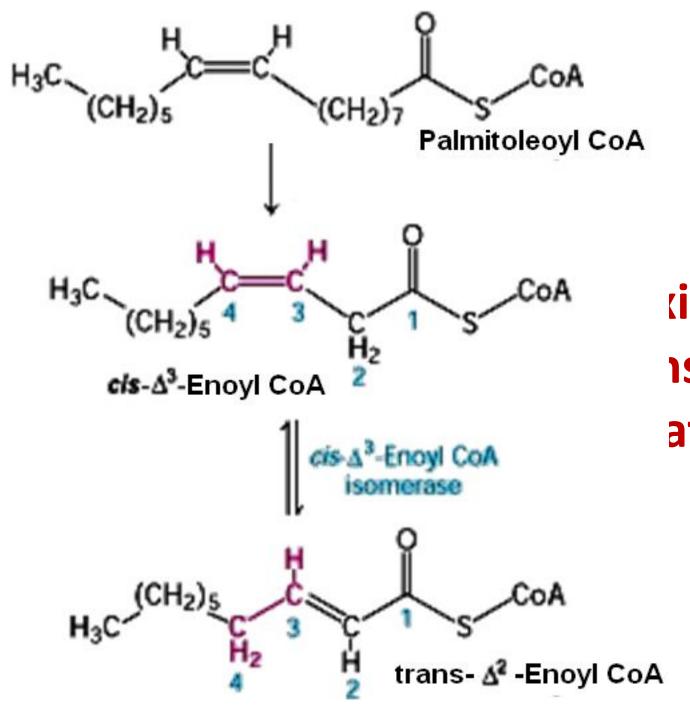


Thiolase reaction

$$\beta$$
-ketoacyl-CoA CH₃(CH₂)_n-C-CH₂-C-SCoA
$$\beta$$
-ketothiolase HSCoA
$$CH_3(CH_2)_n-C-SCoA + CH_3-C-SCoA$$
Fatty acyl-CoA Acetyl-CoA

Propionyl-CoA metabolism





ridadion of saturated atty acids

α-Oxidation (nervous tissue, > 20 C)

$$CH_{3}-(CH_{3})_{n}-CH_{2}\left[\overset{\bigcirc}{C}-O^{-}\right]\longrightarrow CH_{3}-(CH_{2})_{n}-\overset{\bigcirc}{CH}\left[\overset{\bigcirc}{C}-O^{-}\right]\longrightarrow CH_{3}-(CH_{2})_{n}-\overset{\bigcirc}{C}H_{3}-(CH_{2})_{n}-\overset{\bigcirc}{C}-O^{-}+\overset{\bigcirc}{[CO_{2}]}$$

$$CH_{3}-(CH_{2})_{n}-\overset{\bigcirc}{C}H\left[\overset{\bigcirc}{C}-O^{-}\right]\longrightarrow CH_{3}-(CH_{2})_{n}-\overset{\bigcirc}{C}-O^{-}+\overset{\bigcirc}{[CO_{2}]}$$

ω-Oxidation (in pathology)

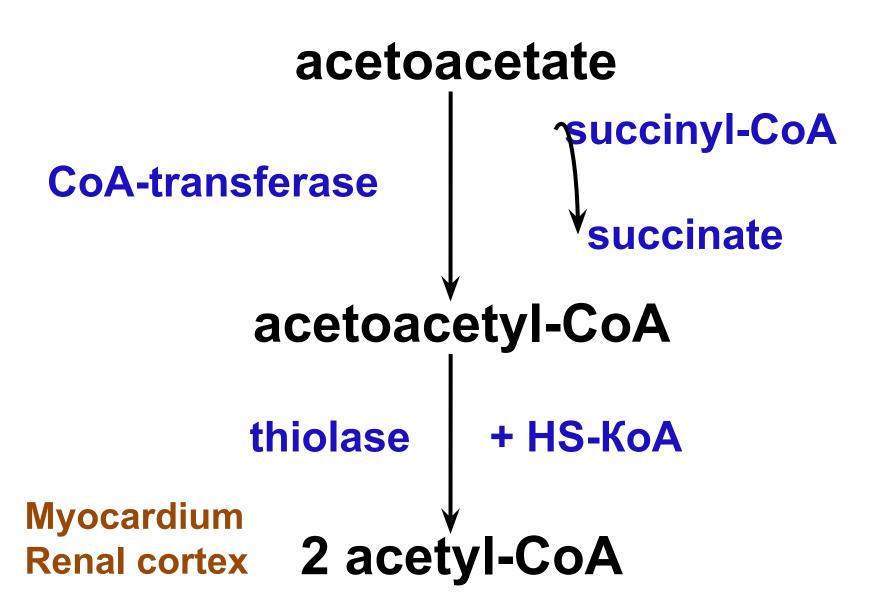
Violations of fatty acid oxidation

- Hereditary defects of carnitine acyl transferase I or enzymes of carnitine synthesis
- Genetic defect of dehydrogenase of fatty acids with medium chain

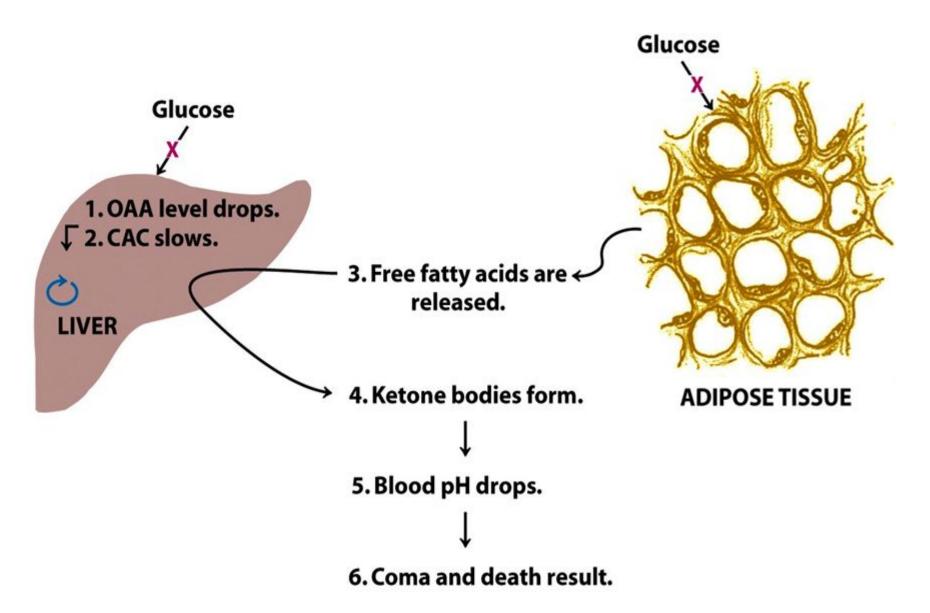
Synthesis of ketone bodies

Liver!
$$2 \text{ CH}_3\text{-CO-SCoA}$$
 $C\text{H}_3\text{-CO-CH}_2\text{-CO-S-CoA}$
 $+ \text{CH}_3\text{-CO-S-CoA}$
 $+ \text{CH}_3\text{-CO-S-CoA}$
 $+ \text{CH}_3\text{-CO-S-CoA}$
 $+ \text{CO-CH}_2\text{-COH}$
 $+ \text{CO-CH}_2\text{-CO-S-CoA}$
 $+ \text{CH}_3\text{-CO-S-CoA}$
 $+ \text{CH}_3\text{-CO-CH}_2\text{-COOH}$
 $+ \text{NADH} + \text{H}^+$
 $- \text{NAD}^+$
 $- \text{CO}_2$
 $+ \text{CH}_3\text{-CO-CH}_3\text{-CO-CH}_3$

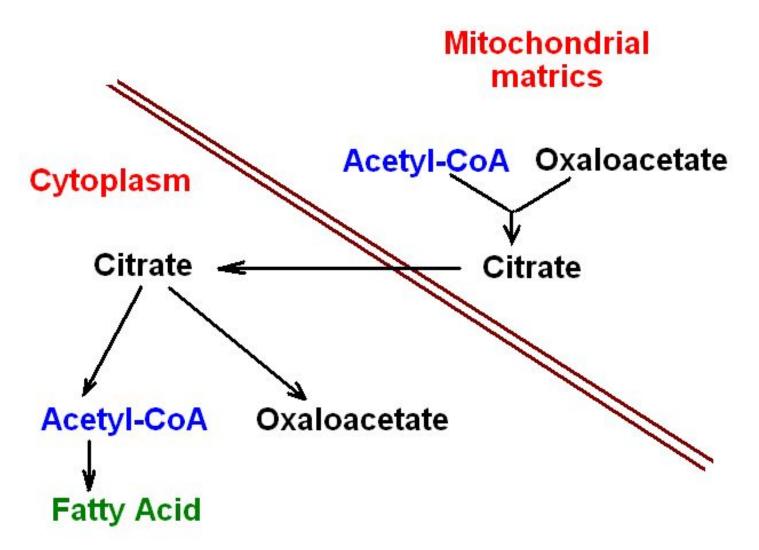
Oxidation of ketone bodies



Mechanism of ketosis



Lipogenesis Acetyl-CoA transport



Lipogenesis

 CO_2 + ATP + biotin-enzyme \rightarrow

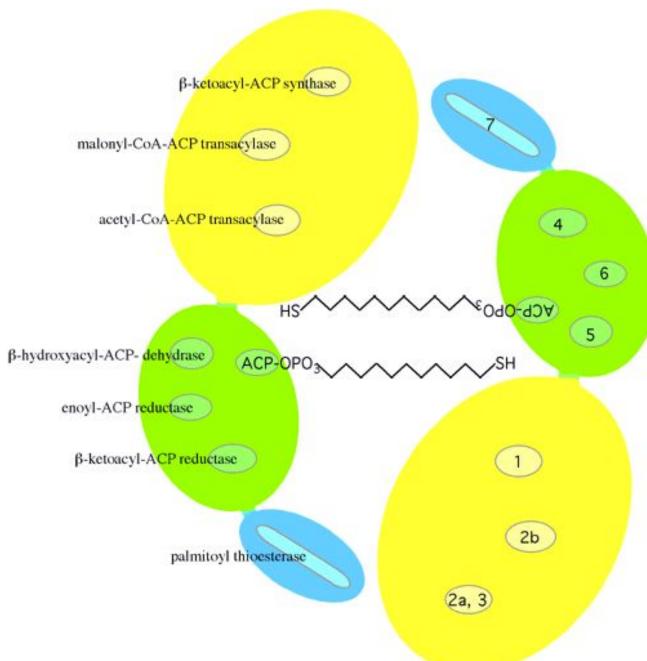
→ carboxybiotin-enzyme + ADP + P_i

carboxybiotin-enzyme + CH_3 -CO- $SKoA \rightarrow$

→ HOOC-CH₂-CO-S-KoA + biotin-enzyme

malonyl-CoA

Rate-limiting!



Fatty acid synthase structure

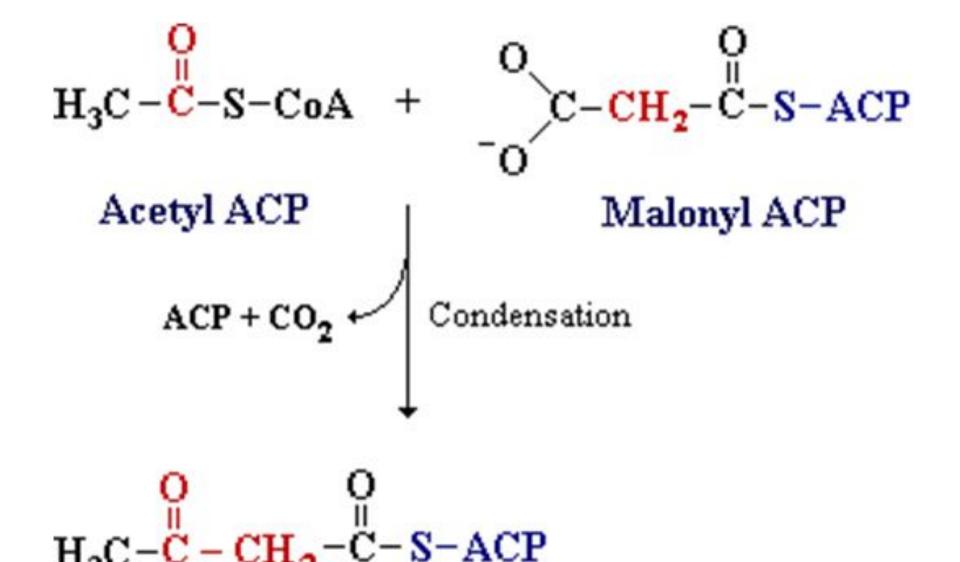
acetyl transacylase

Acetyl CoA + ACP --- Acetyl-ACP + HSCoA

malonyl transacylase

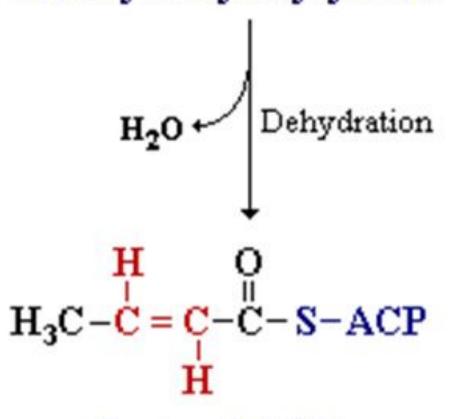
Malonyl CoA + ACP Malonyl-ACP + HSCoA

(ACP = acyl carrier protein)

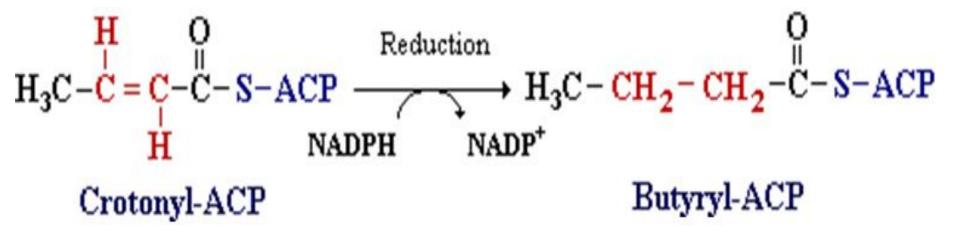


Acetoacetyl-ACP

D-3-Hydroxybutyryl-ACP



Crotonyl-ACP

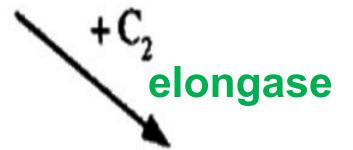


Formation of other fatty acids

Palmitic acid 16:0



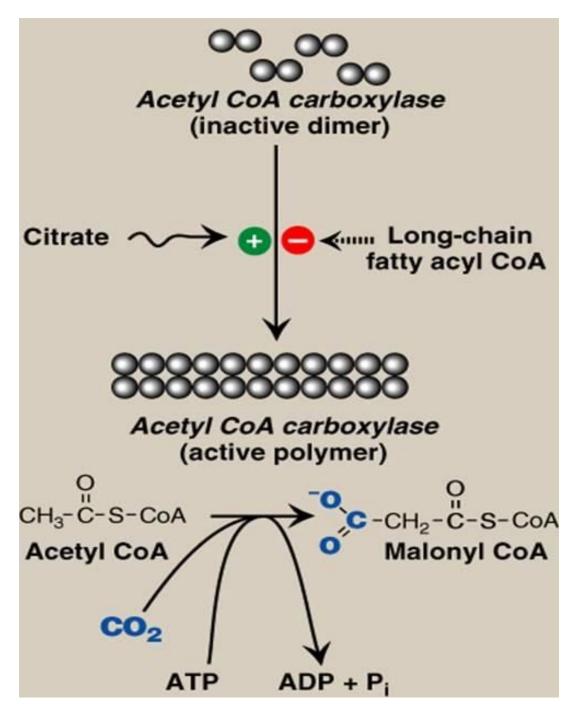
Palmitoleic Acid 16:1 (9)



Stearic acid 18:0



Oleic acid 18:1(9)



Regulation of acetyl-CoA carboxylase activity

Regulation of acetyl-CoA carboxylase activity

