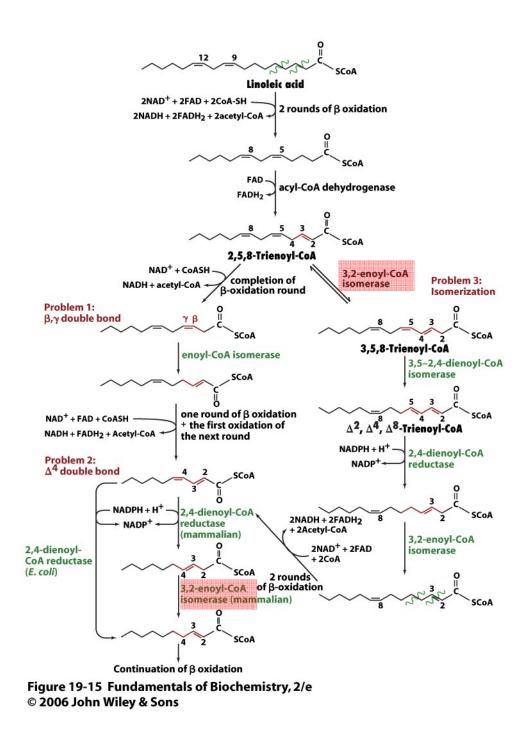
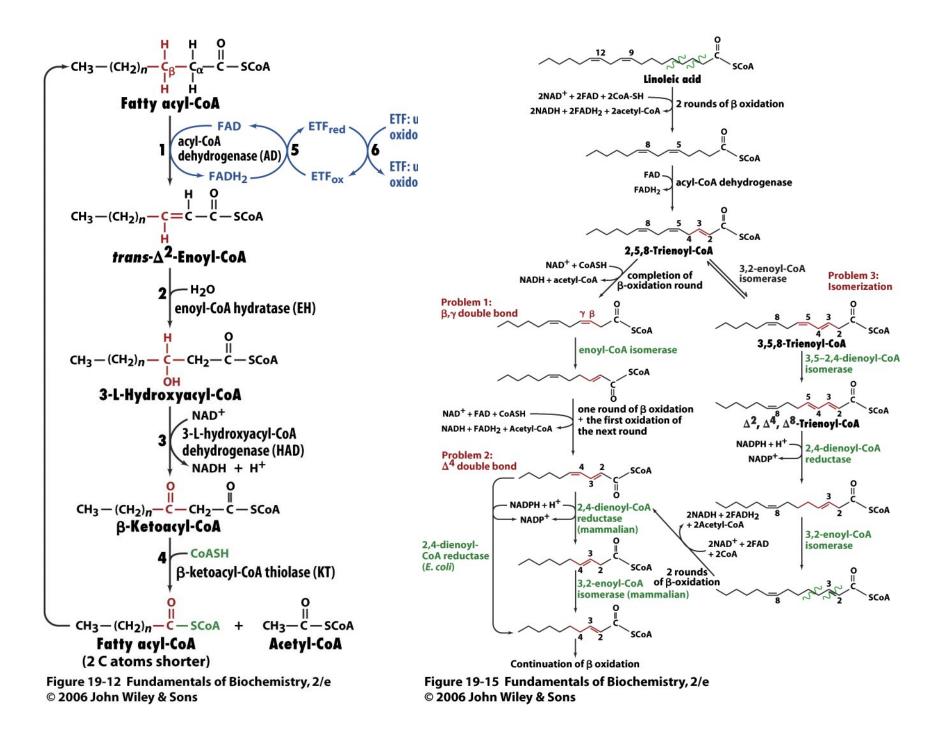
Chapter 7-II: Lipid Metabolism

Oxidation of unsaturated fatty acids

Double bonds One: C9 Two or three: C12, C15

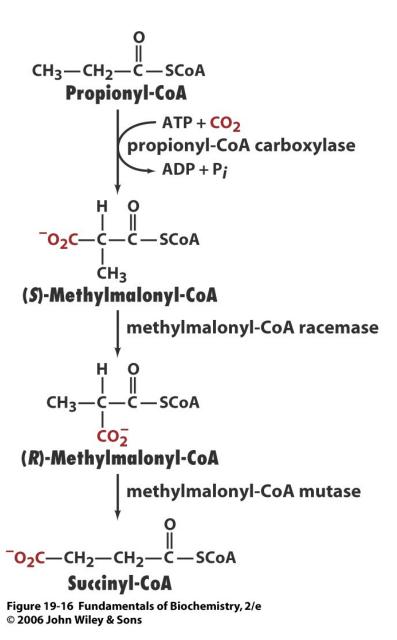
Confront with three problems Enoyl-CoA isomerase 2,4-dienoyl-CoA reductase 3,2-enoyl-CoA isomerase 3,5-2,4-dienoyl-CoA isomerase

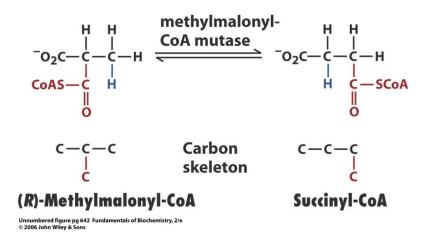




Oxidation of odd-chain fatty acids

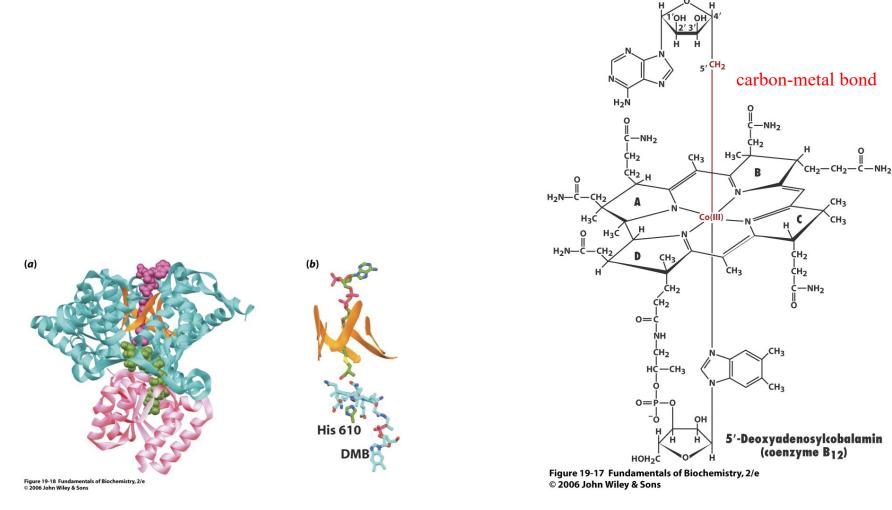
The end product is propionyl-CoA Converted to succinyl-CoA Entry into the citric acid cycle





Methylmalonyl-CoA mutase

Prosthetic group: 5'-deoxyadenosylcobalamine (AdoCbl, coenzyme B12)



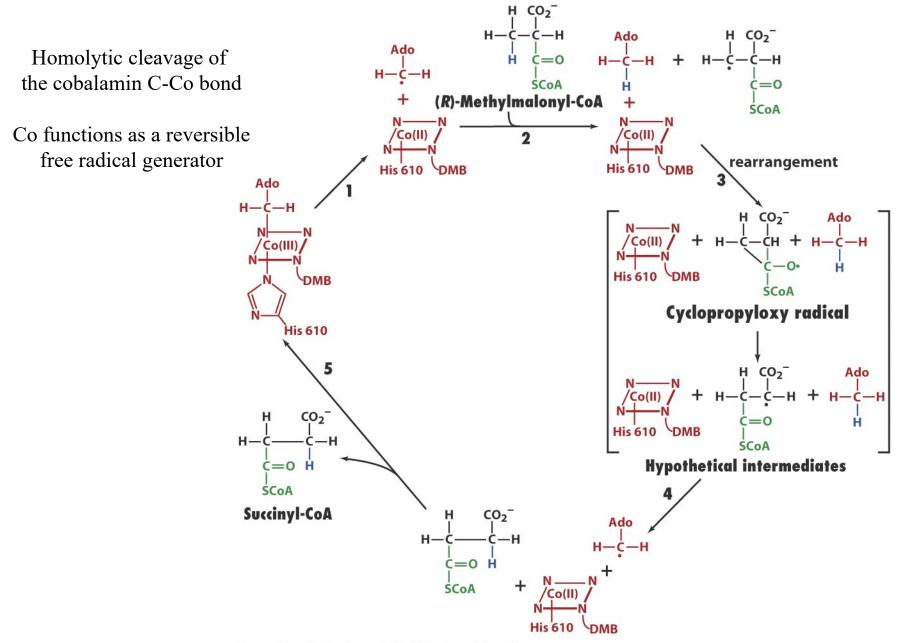
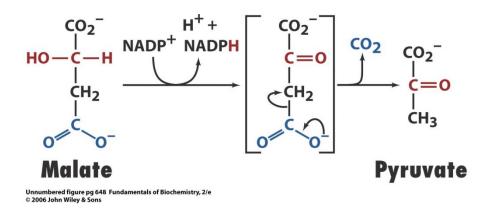


Figure 19-19 Fundamentals of Biochemistry, 2/e © 2006 John Wiley & Sons

Succinyl-CoA to malate Transport to the cytosol Oxidative decarboxylation to pyruvate and CO2



Peroxisomal β oxidation

Animal cells: very long chains or branched chains Plant cells: exclusively in the peroxisomes and glyoxysomes

NADH transport electron to O_2 to generate H_2O_2 Animal cells: very long chain fatty acids (>22 chains) are shortened and transported to mito Three enzymes

Acyl-CoA oxidase Peroxisomal enoyl-CoA hydratase and 3-L-hydroxyacyl-CoA dehydrogenase Peroxisomal thiolase

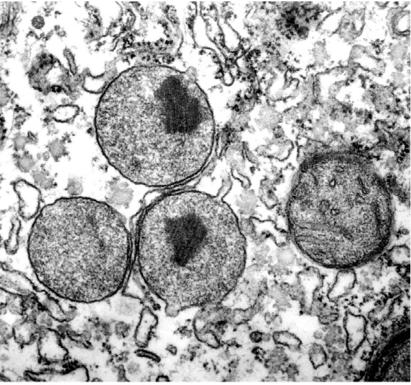
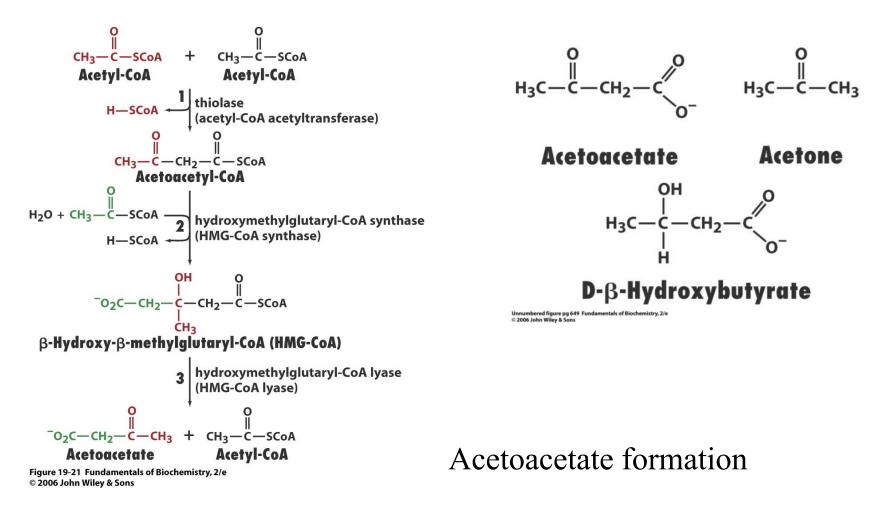


Figure 19-20 Fundamentals of Biochemistry, 2/e

Ketone bodies

Ketogenesis: acetyl-CoA to acetoacetate or D-β-hydroxybutyrate Important metabolic fuels for heart & skeletal muscle During starvation the brain depends on ketone bodies



Ketone bodies to acetyl-CoA

ketosis

$$\begin{array}{c} CH_{3} \\ CH_{2} \\ CO_{2}^{-} \end{array} \qquad \begin{array}{c} H^{+} + \\ P^{-}hydroxybutyrate} \\ CH_{2} \\ CO_{2}^{-} \end{array} \qquad \begin{array}{c} H^{+} + \\ P^{-}hydroxybutyrate} \\ CH_{2} \\ CO_{2}^{-} \end{array} \qquad \begin{array}{c} D^{-}P^{-}Hydroxybutyrate} \\ CH_{2} \\ CO_{2}^{-} \end{array} \qquad \begin{array}{c} D^{-}P^{-}Hydroxybutyrate} \\ D^{-}P^{-}Hydroxybutyrate} \\ CH_{3} - C^{-}CH_{2} - CO_{2}^{-} \\ H \\ D^{-}P^{-}Hydroxybutyrate} \\ 0 \\ CH_{3} - C^{-}CH_{2} - CO_{2}^{-} \\ H \\ P^{-}P^{-}Hydroxybutyrate} \\ CH_{3} - C^{-}CH_{2} - CO_{2}^{-} \\ Acetoacetate \\ 0 \\ CH_{3} - C^{-}CH_{2} - CO_{2}^{-} \\ Acetoacetate \\ 0 \\ CH_{3} - C^{-}CH_{2} - CO_{2}^{-} \\ Acetoacetate \\ 0 \\ CH_{3} - C^{-}CH_{2} - CO_{2}^{-} \\ Succinyl-COA \\ COA \\ CH_{3} - C^{-}CH_{2} - CC_{3} \\ CCC - CH_{2} - CH_{2} - CO_{2}^{-} \\ Succinate \\ 0 \\ CH_{3} - C^{-}C-SCOA \\ Acetoacetyl-COA \\ H^{-}SCOA \\ H^{-}SCOA \\ H^{-}H^{-}SCOA \\ H^{-}H^{-}SCOA \\ H^{-}H^{-}SCOA \\ CH_{3} - C^{-}SCOA \\$$

Fatty acid biosynthesis

Reverse of β -oxidation process

In humans, fatty acids are predominantly formed in the liver and lactating mammary glands, and to a lesser extent, the adipose tissue.

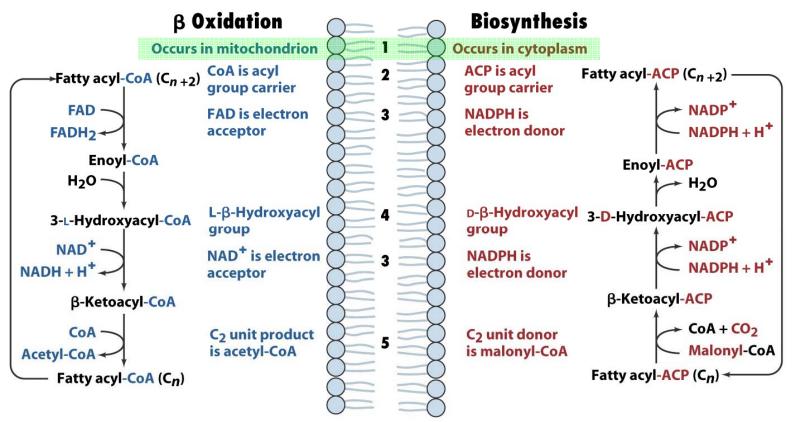


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Acetyl group transfer from mito to cytosol

tricarboxylate transport system ATP-citrate lyase

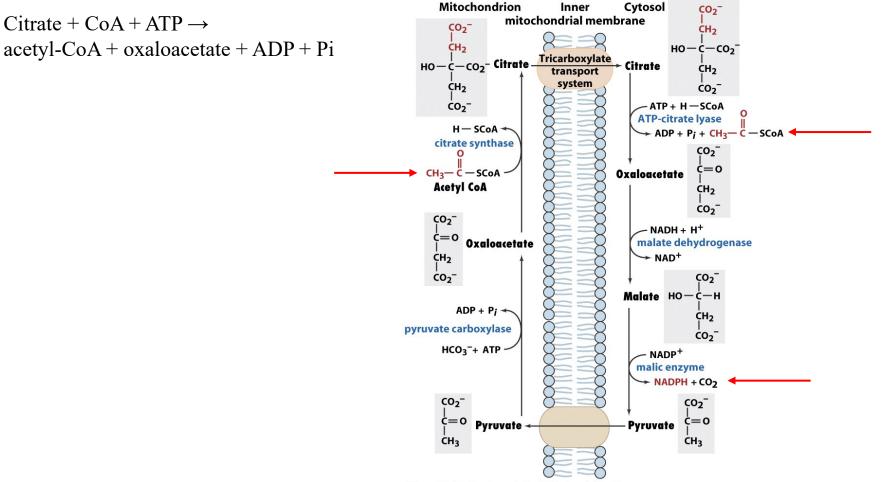
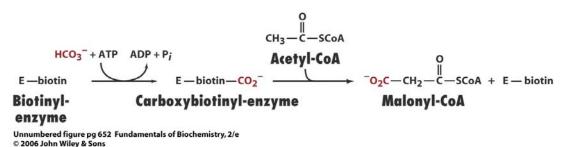


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Acetyl-CoA carboxylase (ACC) The first committed step of fatty acid synthesis & r.d.s. Allosteric and covalent regulation CO₂ activation and carboxylation



Mammalian ACC: two isoforms

adipose tissue, α -ACC; heart muscle, β -ACC; liver, both Heart muscle does not synthesize fatty acids What is the function of β -ACC?

Malonyl-CoA strongly inhibits the mito import of fatty acyl-CoA

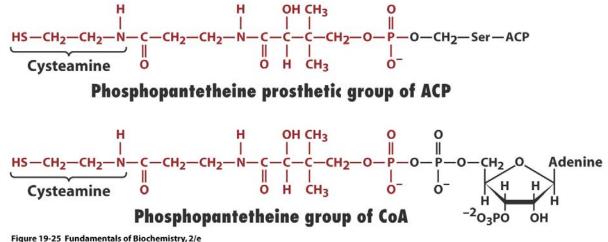
E. coli ACC

regulated by guanine nucleotides why? Fatty acid synthesis is coordinated with cell growth

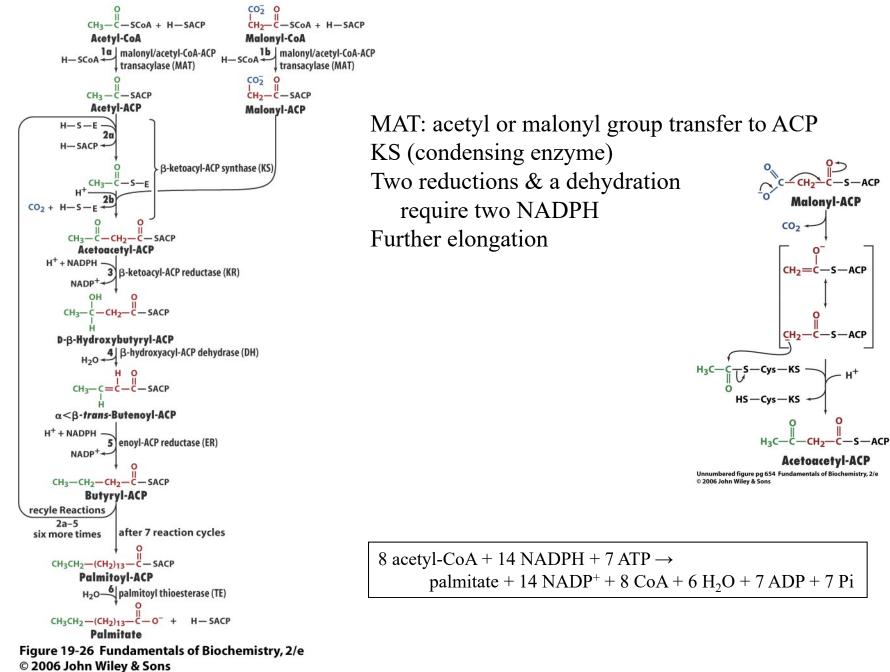
Fatty acid synthase

E. coli by individual enzymes
Plant: in chloroplast by individual enzymes
Yeast: cytosolic 2500-kD multifunctional enzyme α6β6
Animal: 534-kD consisting of two identical polypeptide chains

Acyl-carrier protein (ACP) in *E. coli* 10-kD polypeptide in animal a part of the multifunctional complex



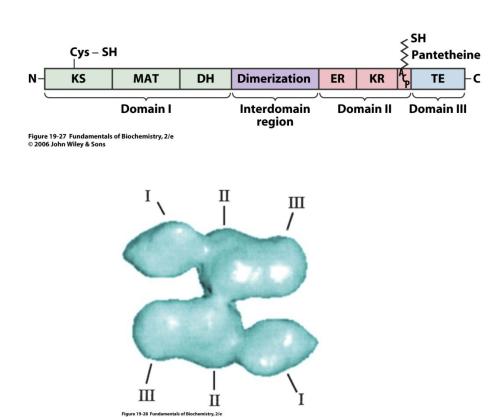
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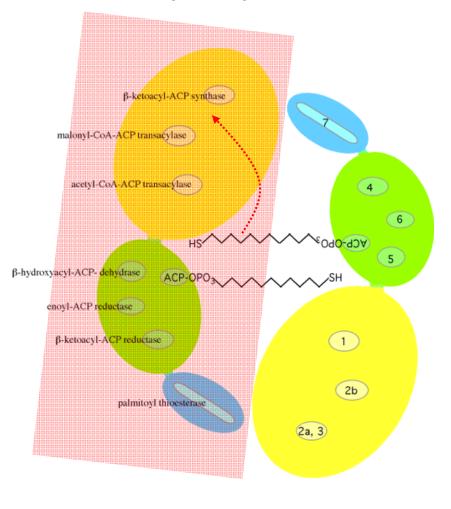
-ACP

Animal fatty acid synthase

Dimers operate in concert: head to tail 7 reactions by 6 discrete active sites (two by MAT)



Mammalian Fatty Acid Synthase



Malignant tissues: high levels of fatty acid synthase An inhibitor of fatty acid synthesis: possible anticancer agent

Triclosan (5-chloro-2-(2,4-dichlorophenoxy)phenol) antibacterial agent inhibits enoyl-ACP reductase emergence of resistant strains





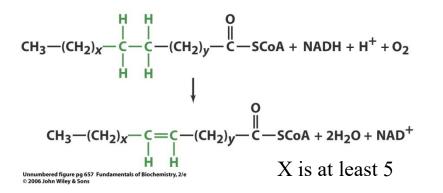
ox 19-3 figure 2 Fundamentals of Biochemistry, 2/e

Elongases and desaturases

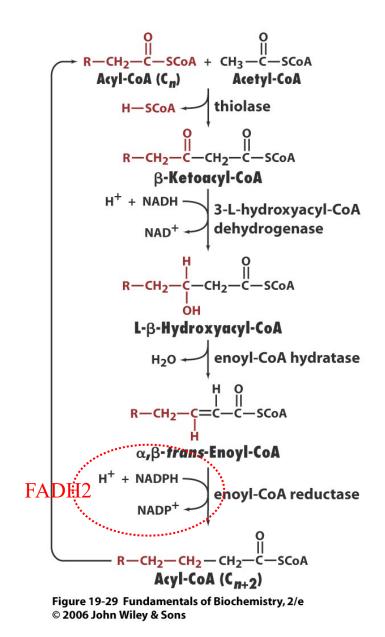
Elongases: mito & ER but in different mechanisms more than C16 in mito: reversal of fatty acid oxidation in ER: successive addition of malonyl-CoA

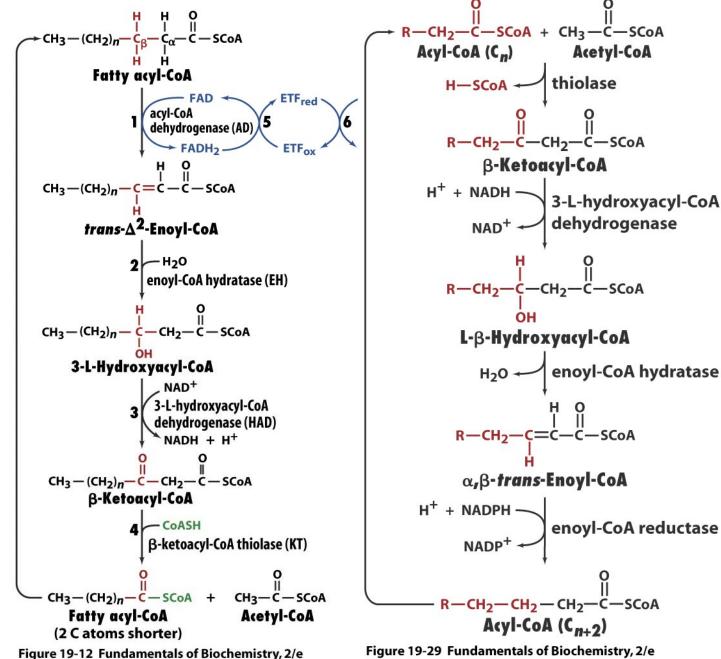
Terminal desaturases

4 enzymes of broad chain-length specificities Δ^9 -, Δ^6 -, Δ^5 -, Δ^4 - fatty acyl-CoA desaturases



Essential fatty acid: linoleic acid (9,12-octadecadienoic acid) A double bond at C6 from methyl end

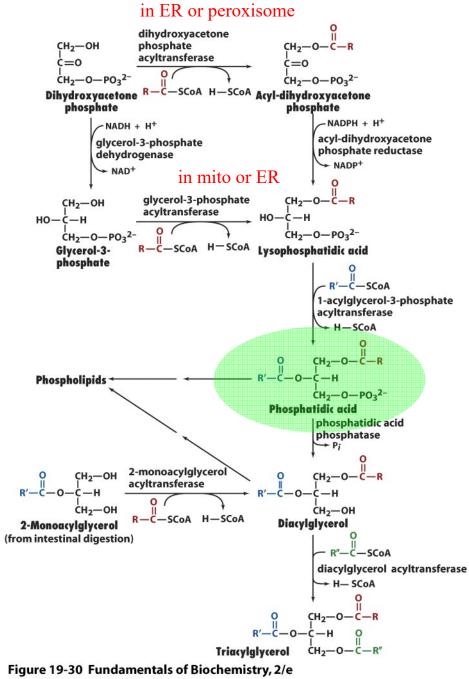




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Synthesis of triacylglycerols

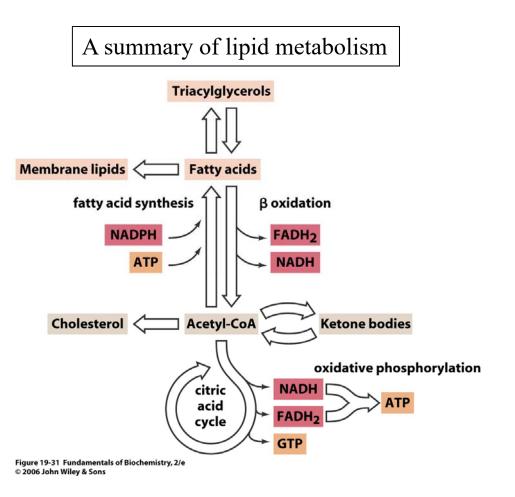




Glyceroneogenesis

important for triacylglycerol biosynthesis

dihydroxyacetone phosphate and glycerol-3-phosphate from glycolysis oxaloacetate via gluconeogenesis (important in times of starvation)



Regulation of Fatty acid metabolism

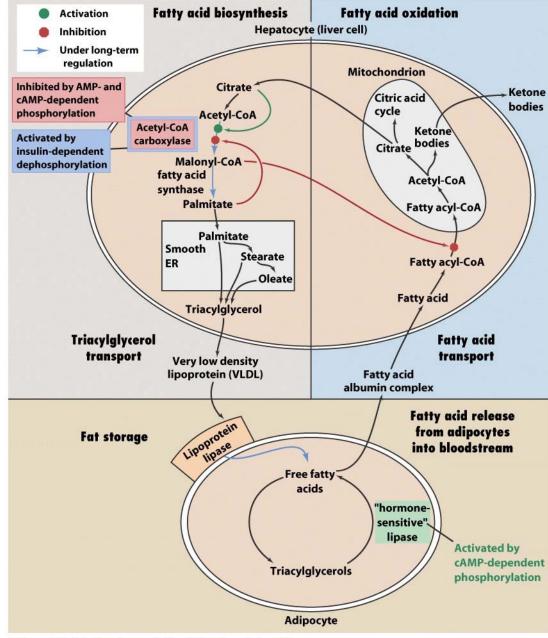


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Lipase

Lipoprotein lipase (LPL)

Lipoprotein lipase is found in vascular endothelium. It is activated by insulin, ACTH, TSH, glucagon and thyroid hormone. Its activity is enhanced by heparin. As discussed above, lipoprotein lipase hydrolyzes CM and VLDL to free fatty acids and glycerol and VLDL-remnants, respectively. Apolipoprotein C is essential for activation of LPL.

Hepatic lipase

This enzyme hydrolyzes surface phospholipids on lipoproteins and is responsible for converting VLDL to LDL.

Hormone sensitive lipase

This enzyme is responsible for lipolysis (mobilization of triglycerides from adipose tissue to yield free fatty acids and glycerol). The enzyme is stimulated by catecholamines, growth hormone, thyroxine, corticosteroids and prostaglandins. It is inhibited by insulin. Fatty acids are transported to the liver (free or albumin-bound), where they are taken up and used for energy (beta oxidation), combined with triglycerides to form VLDL or incorporated into ketones. Therefore, lipolysis will increase VLDL production.

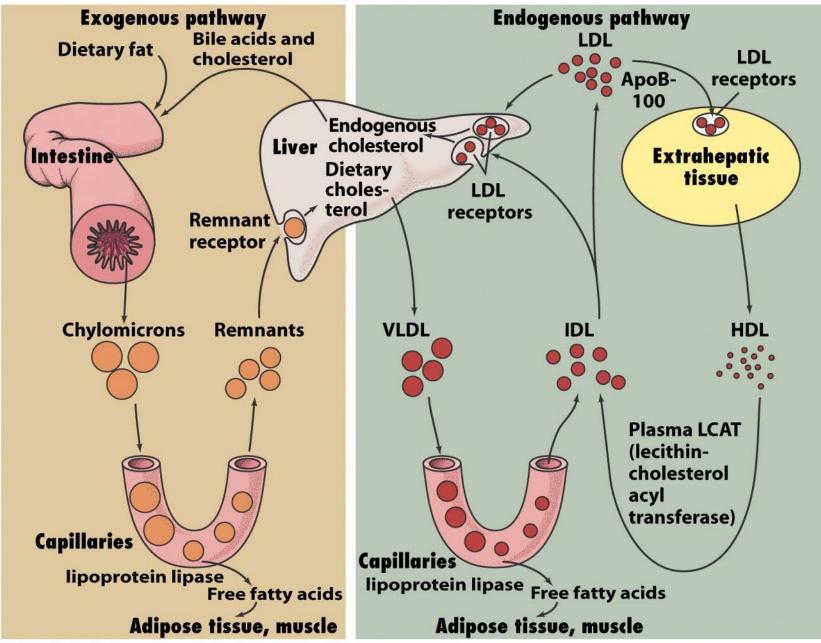
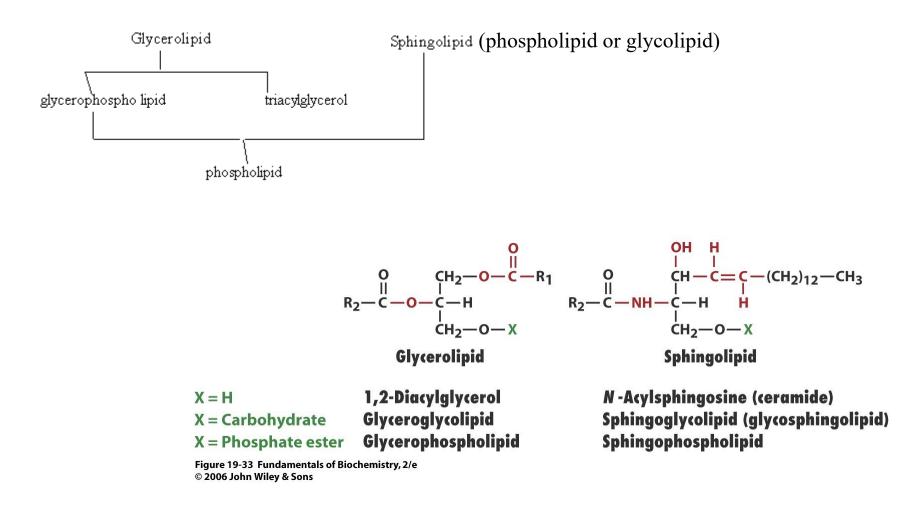


Figure 19-7 Fundamentals of Biochemistry, 2/e © 2006 John Wiley & Sons

Synthesis of other lipids

Membrane lipids and signal molecules

Synthesis in membranes of the cytosolic side of ER & then transport to their destinations Gylcerolipids & sphingolipids



Synthesis of glycerophospholipids

C1: saturated

C2: unsaturated

C3: phosphoester group: ethanolamine, choline, serine, etc

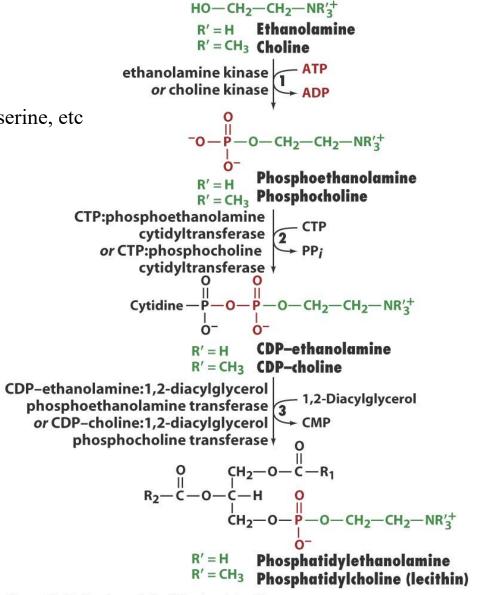
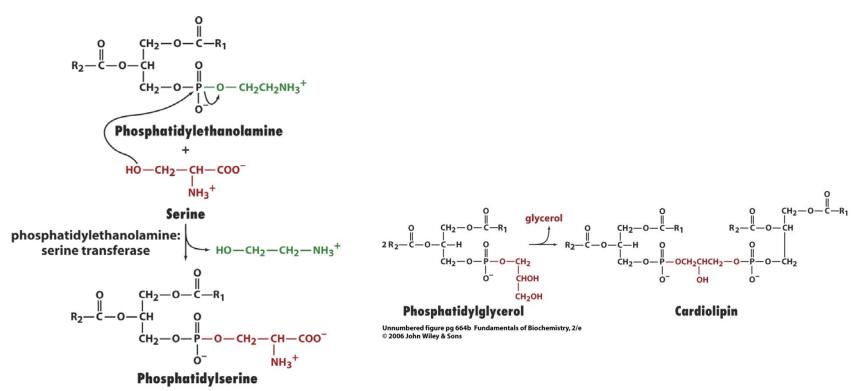
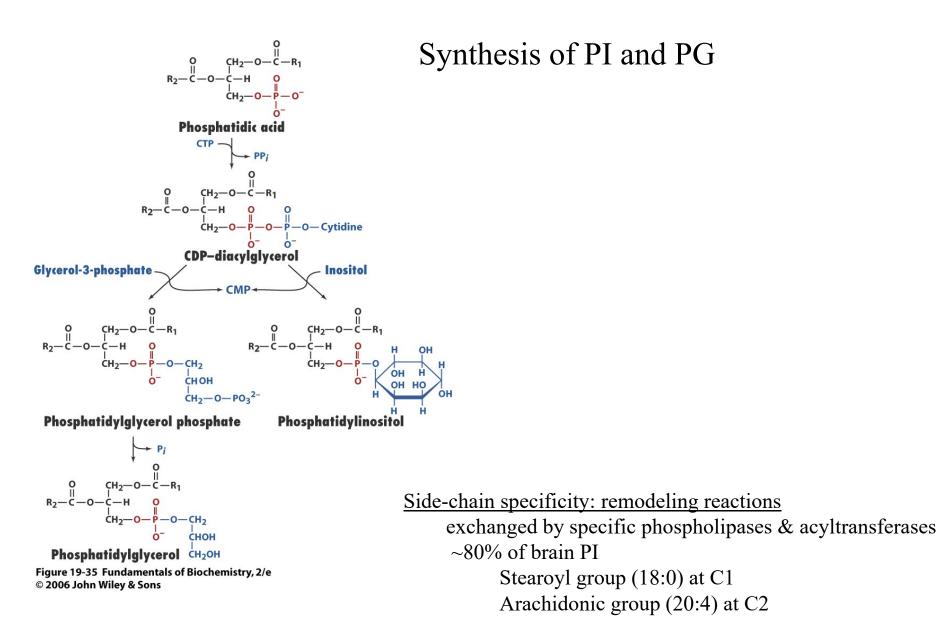


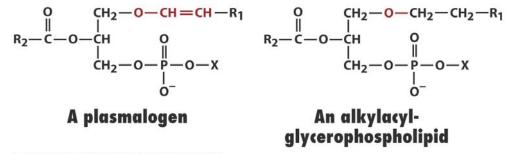
Figure 19-34 Fundamentals of Biochemistry, 2/e © 2006 John Wiley & Sons



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~40% of lung PC Palmitoyl groups (16:0) at C1& C2 Biosynthesis of plasmalogen & alkylacylglycerophospholipids C1 in ether linkage variable abundance among species and tissues abundant in nervous, immune, and cardiovascular system



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Biosynthesis of sphingolipids

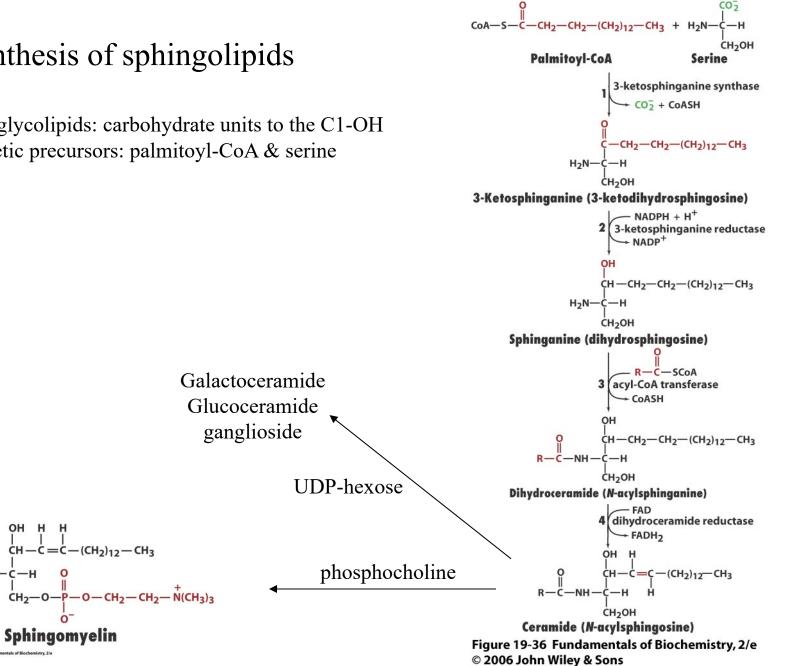
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R - C - NH - C - H

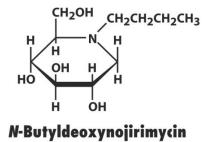
Sphingomyelin

Most are glycolipids: carbohydrate units to the C1-OH Biosynthetic precursors: palmitoyl-CoA & serine



Sphingolipid degradation and Lipid storage diseases

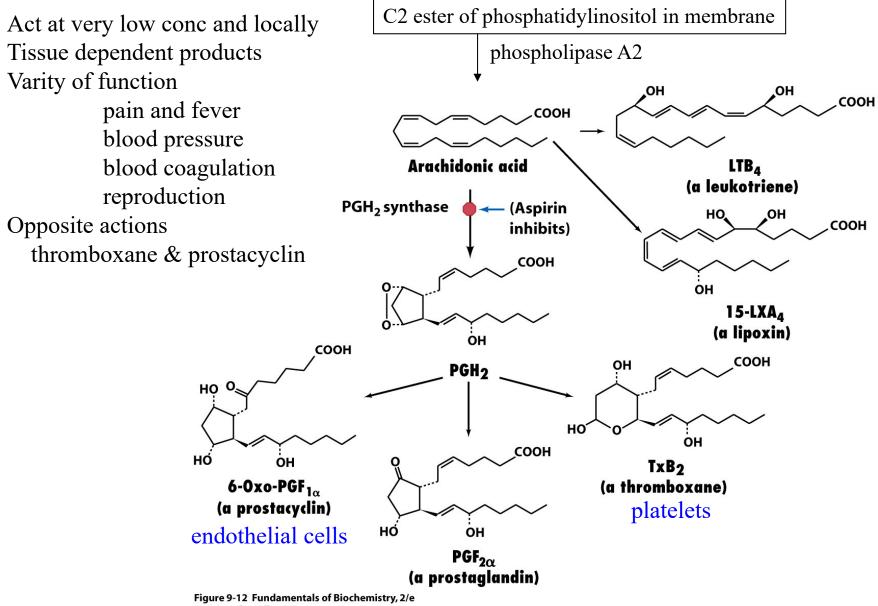




NANA $Gal \xrightarrow{\beta} GalNAc \xrightarrow{\beta} Gal \xrightarrow{\beta} Glc \xrightarrow{\beta} Cer$ **Ganglioside GM1** $G_{M1} \beta$ -galactosidase **G_{M1}** gangliosidosis $GaINAc \stackrel{\beta}{-} GaI \stackrel{\beta}{-} GIc \stackrel{\beta}{-} Cer$ Ganglioside G_{M2} hexosaminidase A Tay-Sachs disease ➤ GalNAc $GaINAc \xrightarrow{\beta} GaI \xrightarrow{\alpha} GaI \xrightarrow{\beta} GIc \xrightarrow{\beta} Cer$ NANA Globoside $Gal \stackrel{\beta}{-} Glc \stackrel{\beta}{-} Cer$ hexosaminidase A and B **Ganglioside GM3** Sandhoff's disease ganglioside → GalNAc neuraminidase ► NANA Gal $Gal \xrightarrow{\alpha} Gal \xrightarrow{\beta} Glc \xrightarrow{\beta} Cer$ $Gal \xrightarrow{\beta} Glc \xrightarrow{\beta} Cer$ Lactosyl ceramide α -galactosidase A Trihexosylceramide Fabry's disease β-galactosidase $^{-}O_{3}S - Gal - \frac{\beta}{2}Cer$ Glc^{-B}Cer **Sulfatide** Glucocerebroside glucocerebrosidase arylsulfatase A Gaucher's disease Metachromatic + Glc ► SO4²⁻ leukodystrophy phosphocholine Gal Cer-phospho-Ceramide P Cer Galcholine sphingomyelinase galactocerebrosidase Galactocerebroside (sphingomyelin) Niemann-Pick **Krabbe's disease** ceramidase disease Farber's -> sphingosine lipogranulomatosis Fatty acid

Box 19-4 figure 1 Fundamentals of Biochemistry, 2/e © 2006 John Wiley & Sons

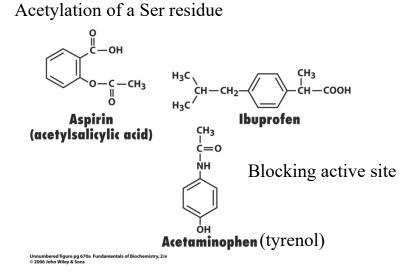
Eicosanoids from arachidonic acid (p 247, Fig. 9-12)

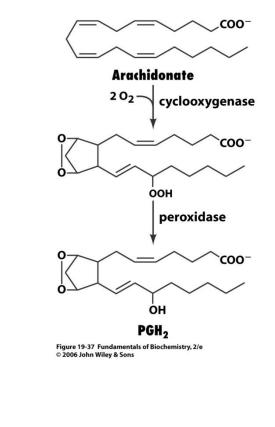


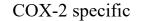
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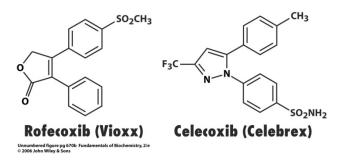
Prostaglandins

Prostaglandin H2 synthase (COX) Cyclooxygenase & peroxidase Two isoforms: COX-1 & COX-2 COX-1: constitutive expression in most tissue COX-2: certain tissue in response to inflammatory stimuli









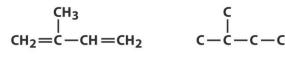
Finding of COX-3: a target of acetaminophen? Poor binding of acetaminophen to COX-1 & -2

Cholesterol metabolism

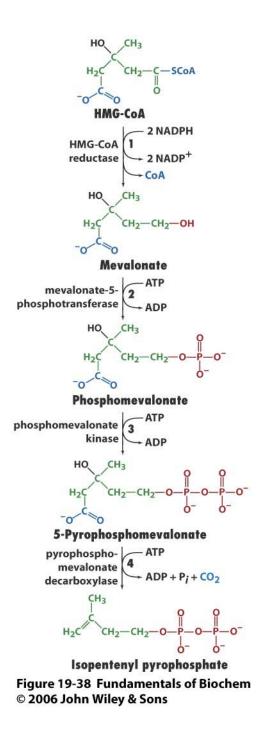
<u>Biosynthesis</u> HMG-CoA synthesis in cytosol: thiolase & HMG-CoA synthase (in mitochondria for ketone bodies)

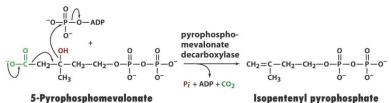
Mevalonate (C6): by HMG-CoA reductase (ER membrane protein) the rate limiting step

Isopentenyl pyrophosphate (C5)



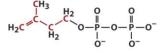
Isoprene An isoprene unit (2-methyl-1,3-butadiene)





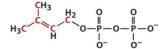
5-Pyrophosphomevalonate

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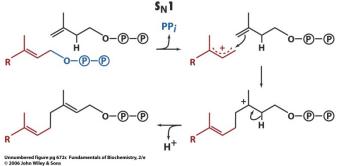
Isopentenyl pyrophosphate

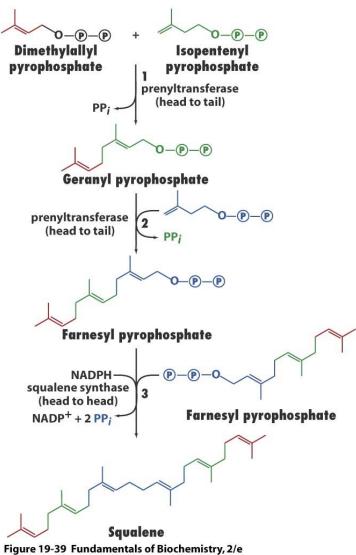
isopentenyl pyrophosphate isomerase



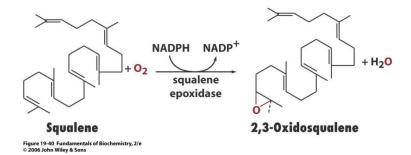
Dimethylallyl pyrophosphate Unnumbered figure pg 672b Fund © 2006 John Wiley & Sons amentals of Biochemistry, 2/e







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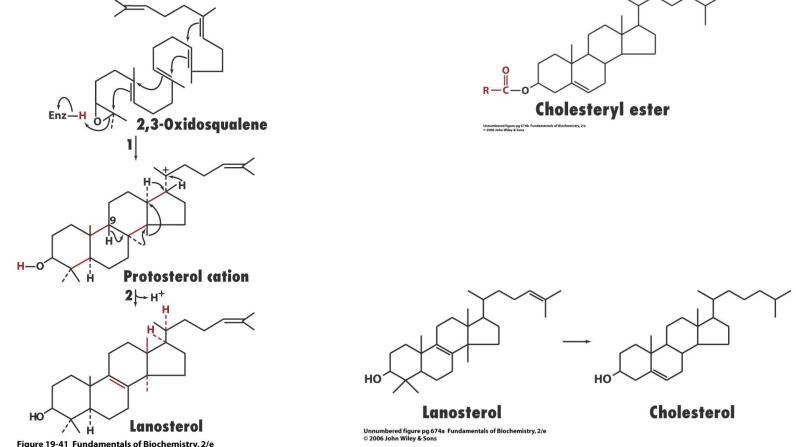


Figure 19-41 Fundamentals of Biochemistry, 2/e © 2006 John Wiley & Sons

Regulation of cholesterol synthesis

The main regulation: HMG-CoA reductase

<u>Short-term regulation</u> Competitive inhibition Allosteric effects Covalent modification: phosphorylation by AMPK

Reductase ATP kinase ÔН Phosphoprotein phosphatase Reductase kinase kinase ADP Reductase H,O kinase ADP ATP INACTIVE HMG-CoA HMG-CoA reductase reductase ÓН cholesterol Phosphoprotein phosphatase ĥо PPI-1 (a) ADP H₂O Phosphoprotein phosphatase cAMP ATI P PPI-1 (b) óн copyright 1998 S.Marchesini

AMPK http://www.indstate.edu/thcme/mwking/ampk.html

http://www.med.unibs.it/~marchesi/cholest.html

Long-term control: gene expression

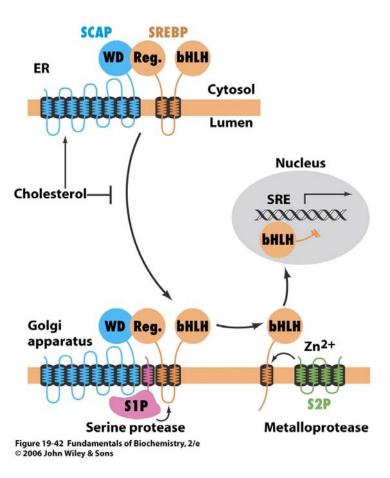
the primary regulation

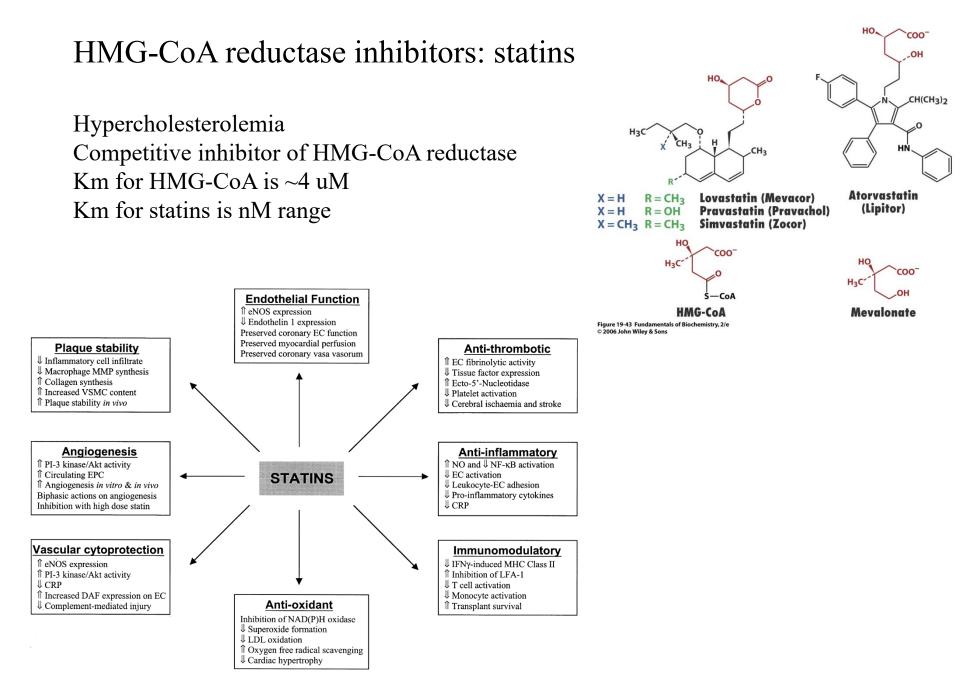
Increased as much as 200-fold along with >20 other genes for synthesis and uptake

Sterol regulatory element (SRE) SREBP: regulatory & bHLH domains SCAP: SREBP cleavage-activating protein sterol-sensing domain & WD repeat

Activation procedure

Low cholesterol in ER SCAP conformation change Transport to golgi apparatus via membranous vesicles Site-1 protease Site-2 protease bHLH binding to SRE



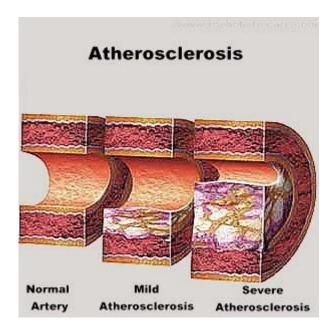


http://www.clinsci.org/cs/105/0251/cs1050251f03.htm?resolution=HIGH

Cholesterol transport and atherosclerosis

Cellular cholesterol concentration depends on the rate of cholesterol synthesis the ability of cell to absorb cholesterol from circulating lipoproteins

High LDL is a strong risk factor for cardiovascular disease Accumulation of lipid in vessel walls: Atherosclerosis Myocardial infarction (heart attack) Stroke (brain)



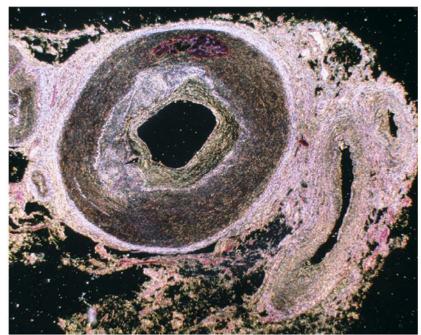
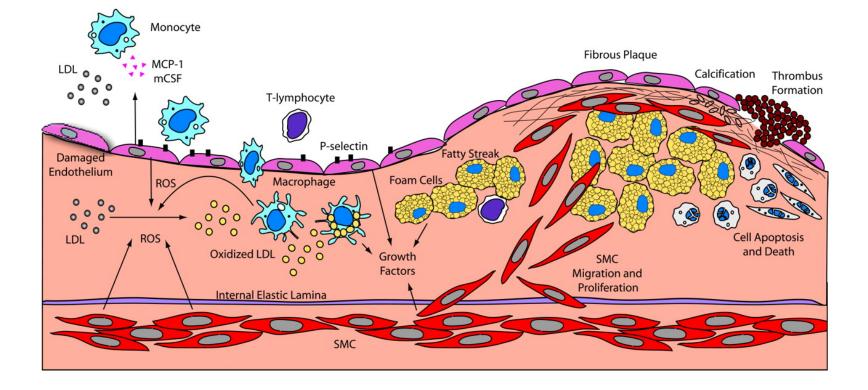


Figure 19-44 Fundamentals of Biochemistry, 2/e



Role of the LDL receptors

familial hypercholesterolemia (FH) Long-term ingestion of a high-fat/high-cholesterol diet

Cholesterol efflux from cells

LDL receptor: FH ABCA1 (ATP-cassette binding protein A1): Tangier disease no HDL synthesis accumulation of cholesteryl ester in macrophages develop atherosclerosis

