

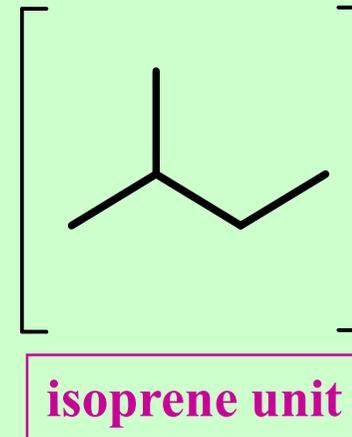
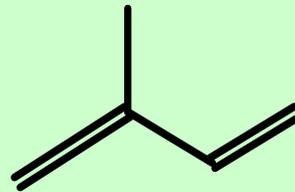
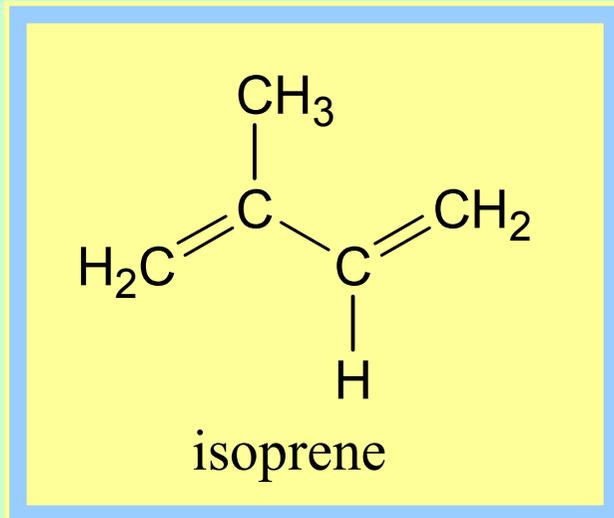
**Metabolism of
cholesterol and other
isoprenoids
(bile acids,
steroid hormones,
vitamins)**

Željka Vukelić
zeljka.vukelic@mef.hr

Dienes

(with condensed, conjugated, or isolated double bonds)

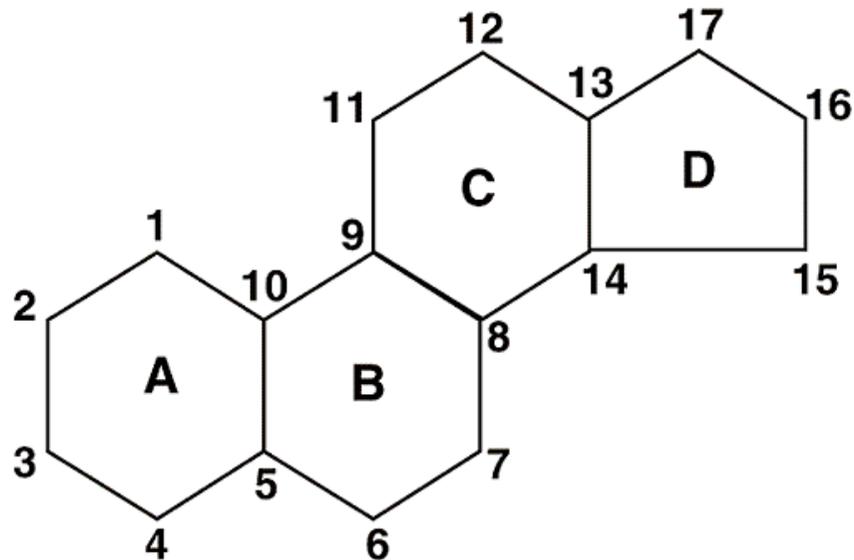
isoprene (2-methyl-1,3-butadiene)



For the synthesis of cholesterol, steroids, some vitamins...

- In biosynthetic reactions, acetyl-CoA is converted to the isoprene units

GONANE



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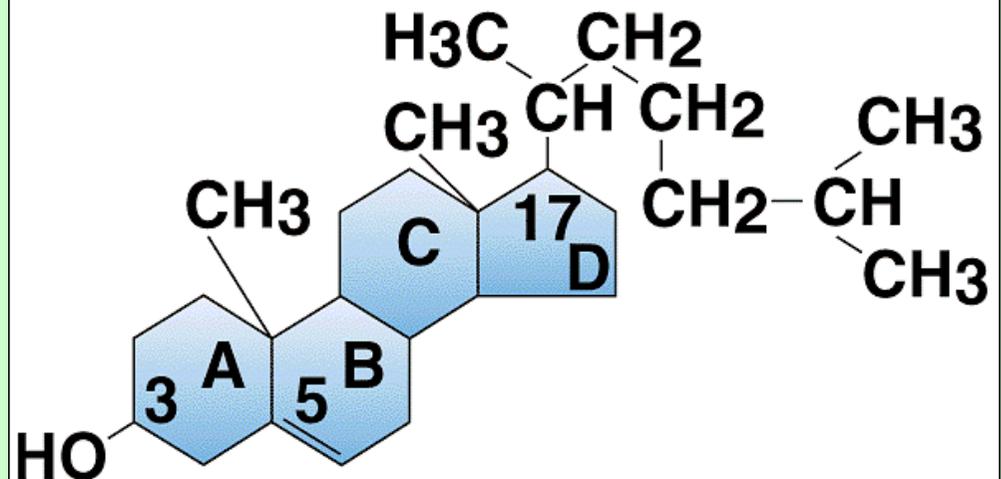
Tetracyclic system:
3 cyclohexane- and
1 cyclopentane ring

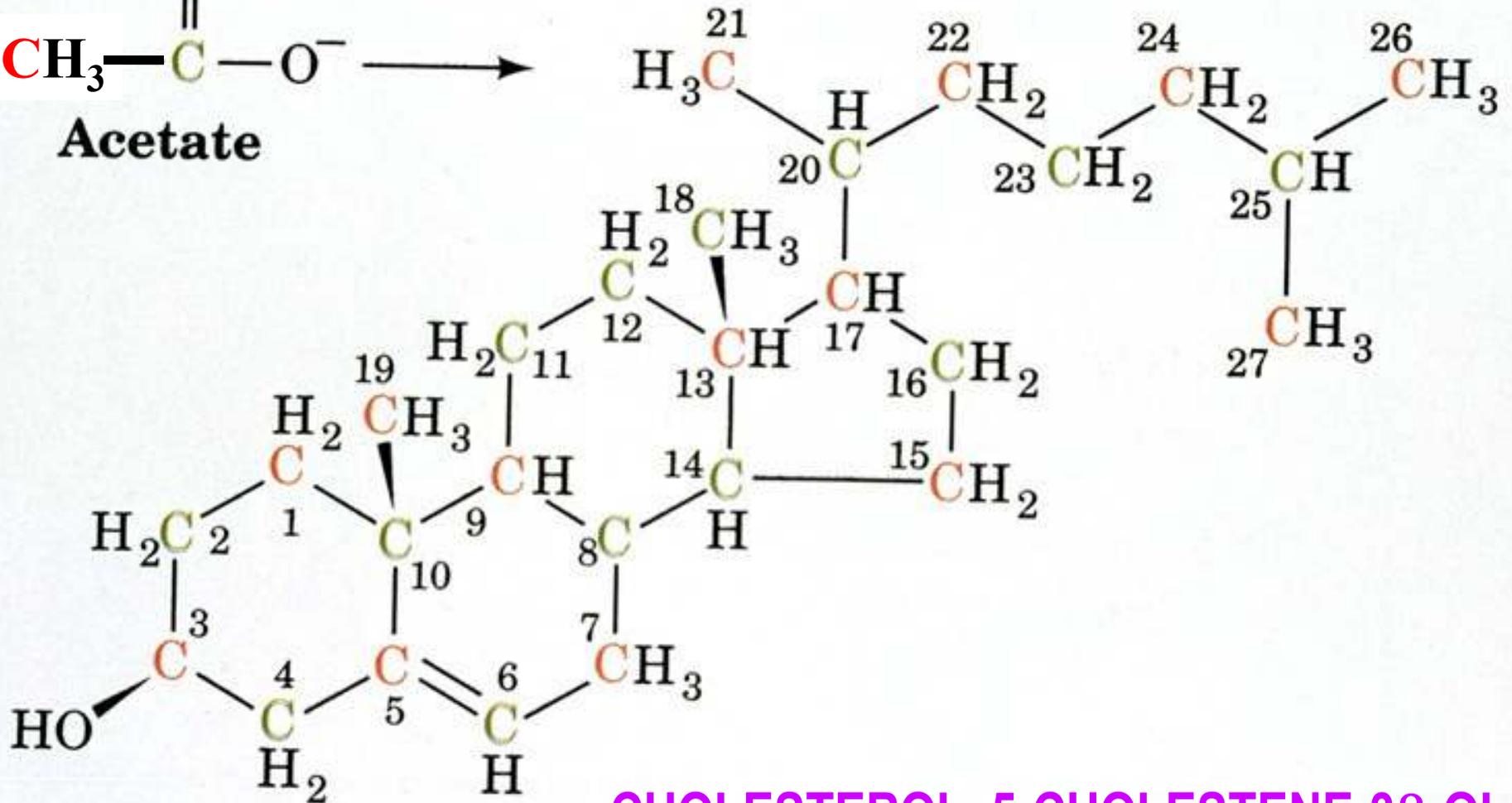
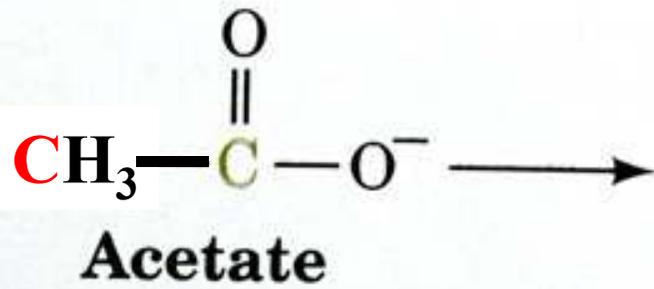
CHOLESTEROL

Steroid skeleton:

5- α -gonan

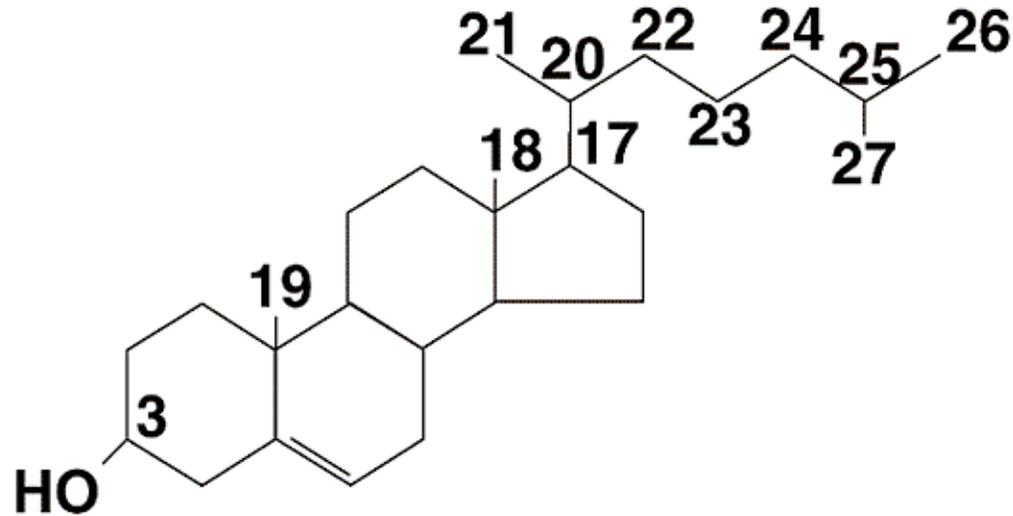
(α -configuration at C-5)





CHOLESTEROL, 5-CHOLESTENE-3β-OL

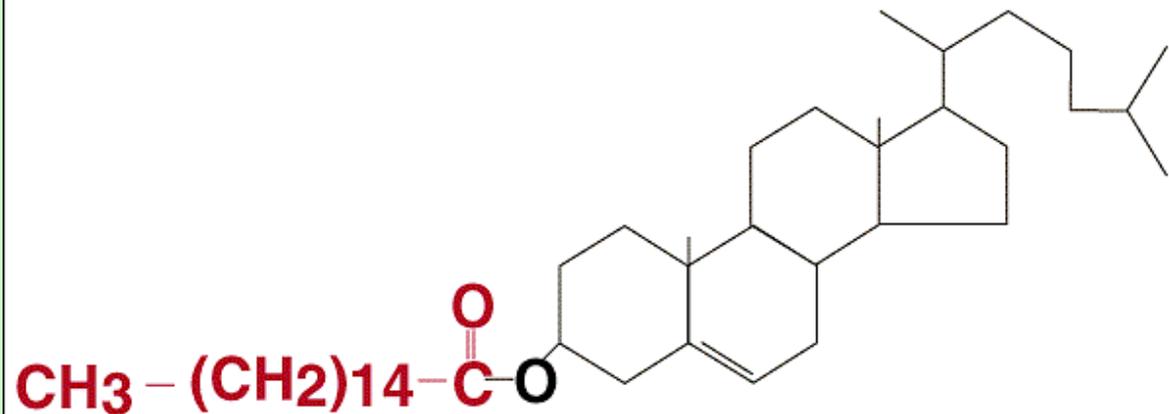
CHOLESTEROL

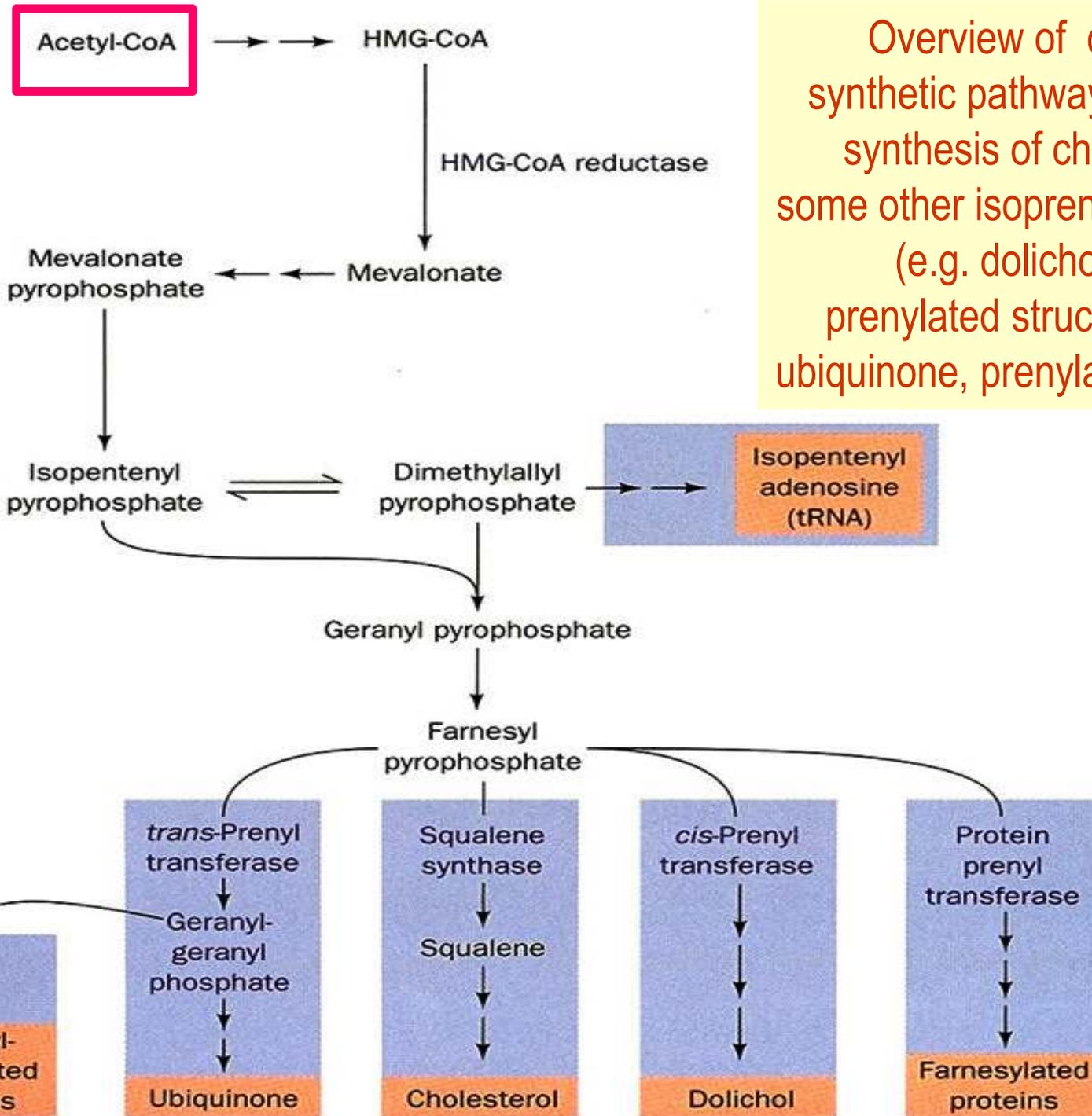


Acyl-CoA-
cholesterol-
acyl-transferase
(ACAT)
- Catalyzes
esterification of
cholesterol in LIVER

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CHOLESTEROL-(PALMITOYL) ESTER





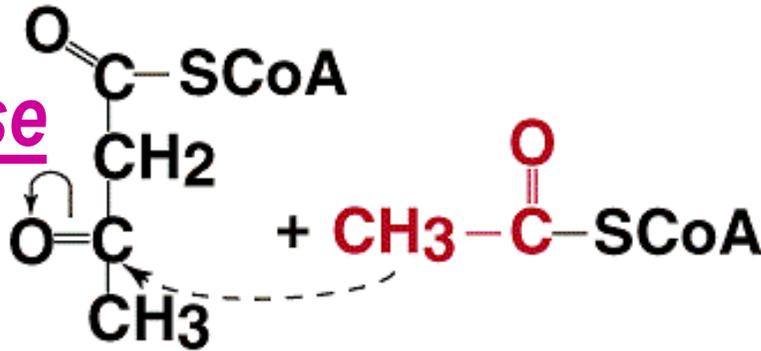
Overview of de novo synthetic pathway leading to synthesis of cholesterol, some other isoprenoid structures (e.g. dolichol) and prenylated structures (e.g. ubiquinone, prenylated proteins)

CHOLESTEROL BIOSYNTHESIS

2 acetyl-CoA

thiolase

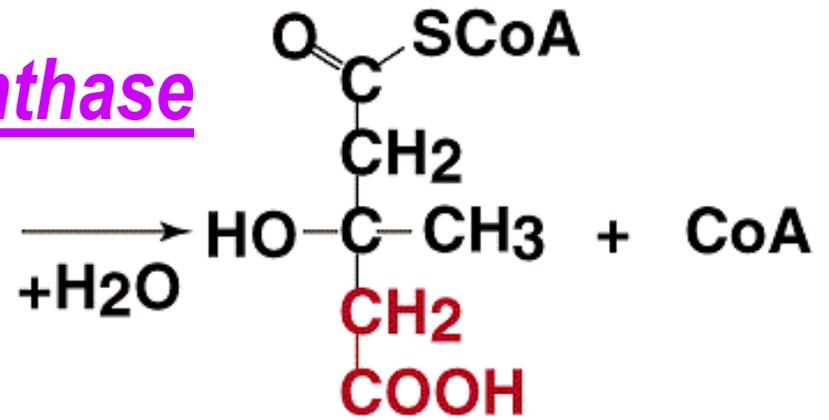
CoA-SH



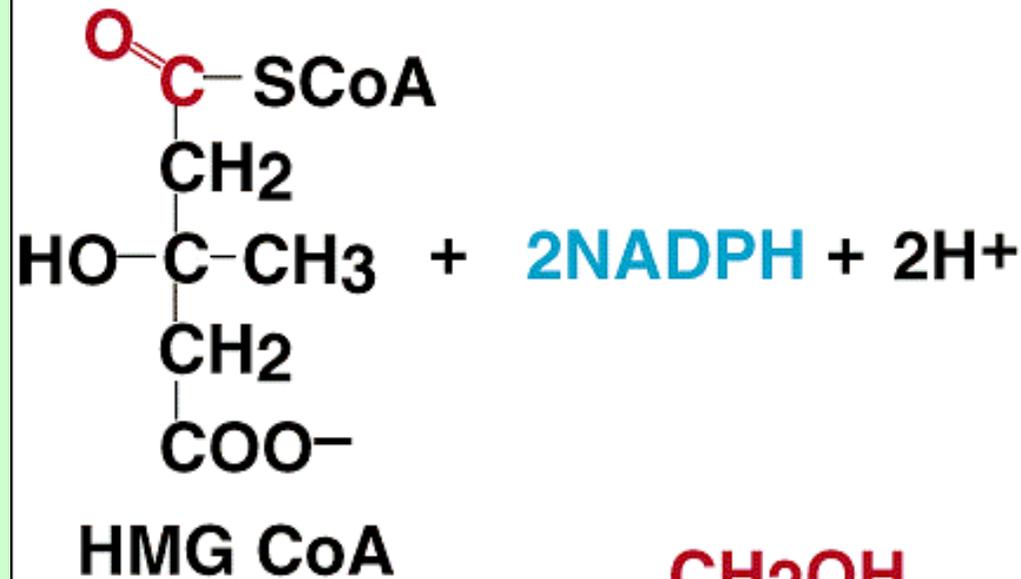
Acetoacetyl CoA

Acetyl CoA

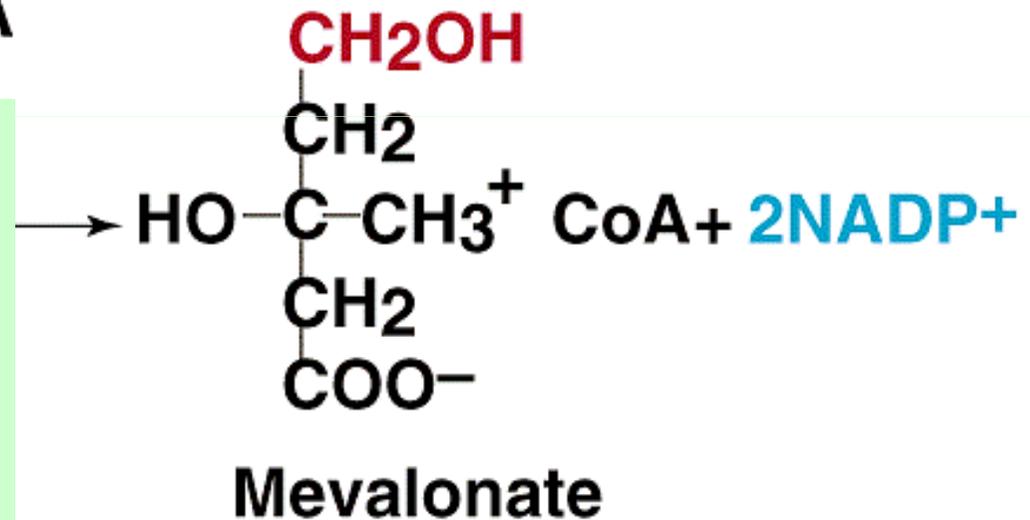
HMG-CoA synthase
(cytosolic)

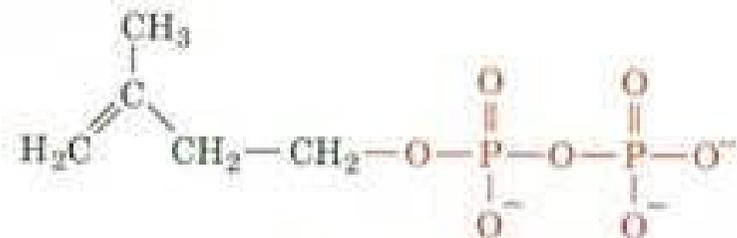
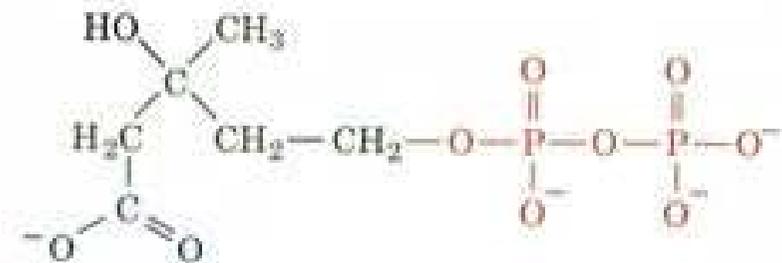
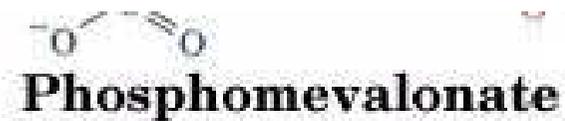
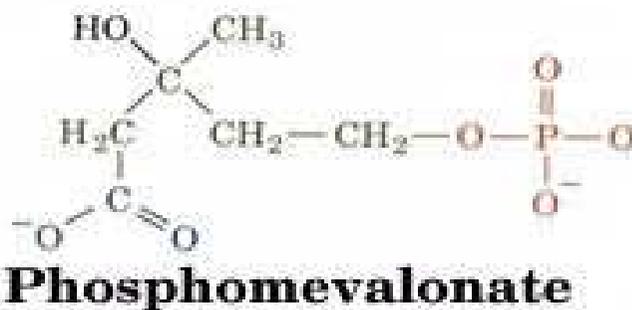
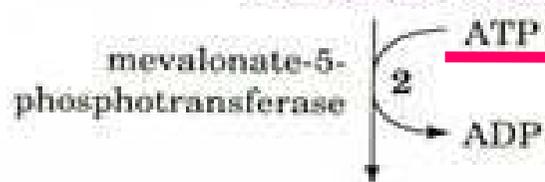
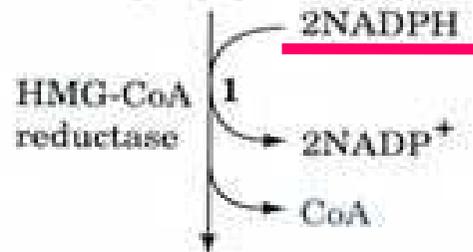
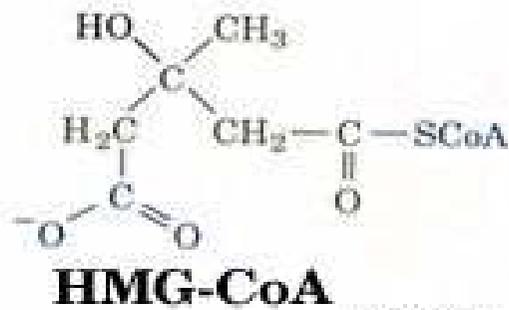


HMG CoA

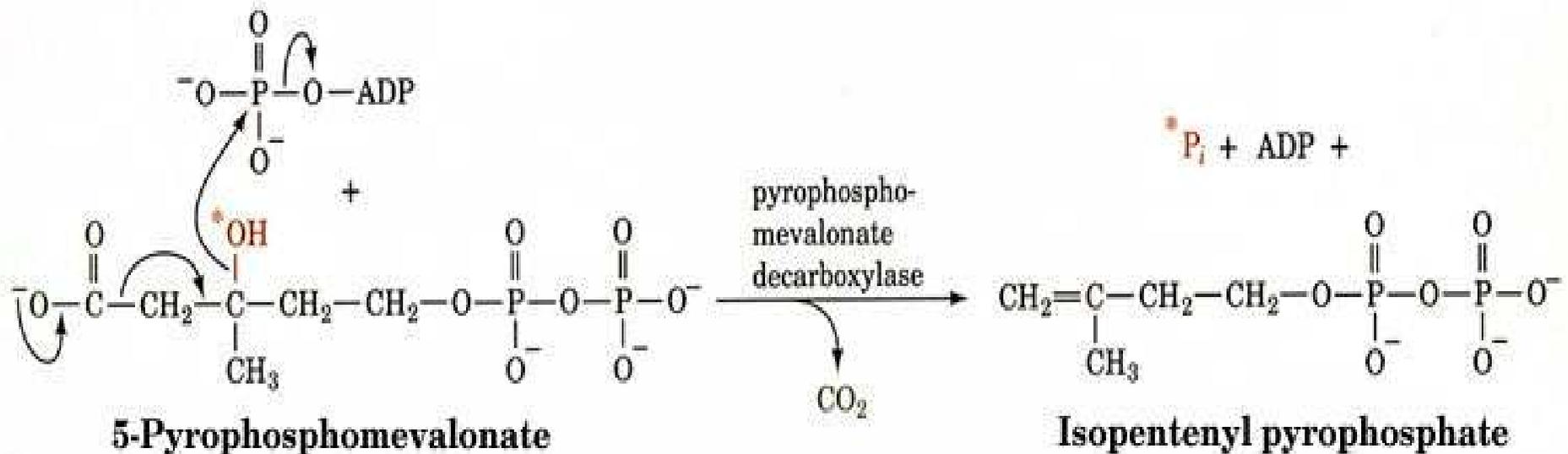


HMG-CoA reductase
*(integral protein of
smooth ER)*
- major regulatory site
of the biosynthesis



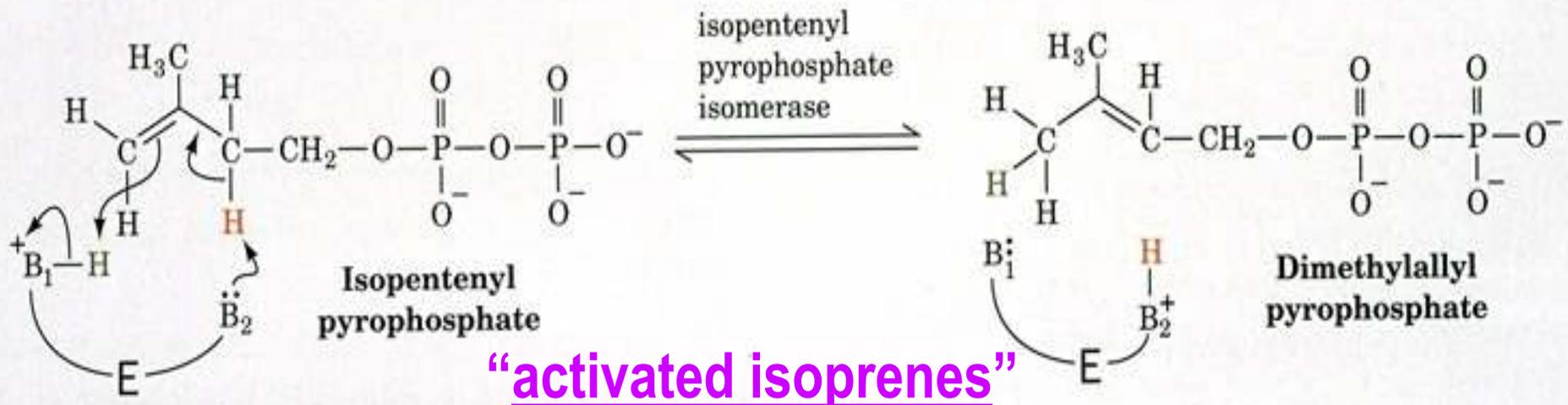


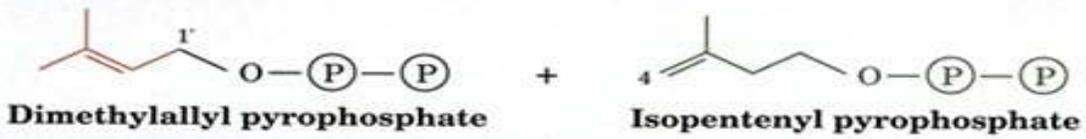
The enzyme catalyzes the ATP-dependent dehydration-decarboxylation reaction:



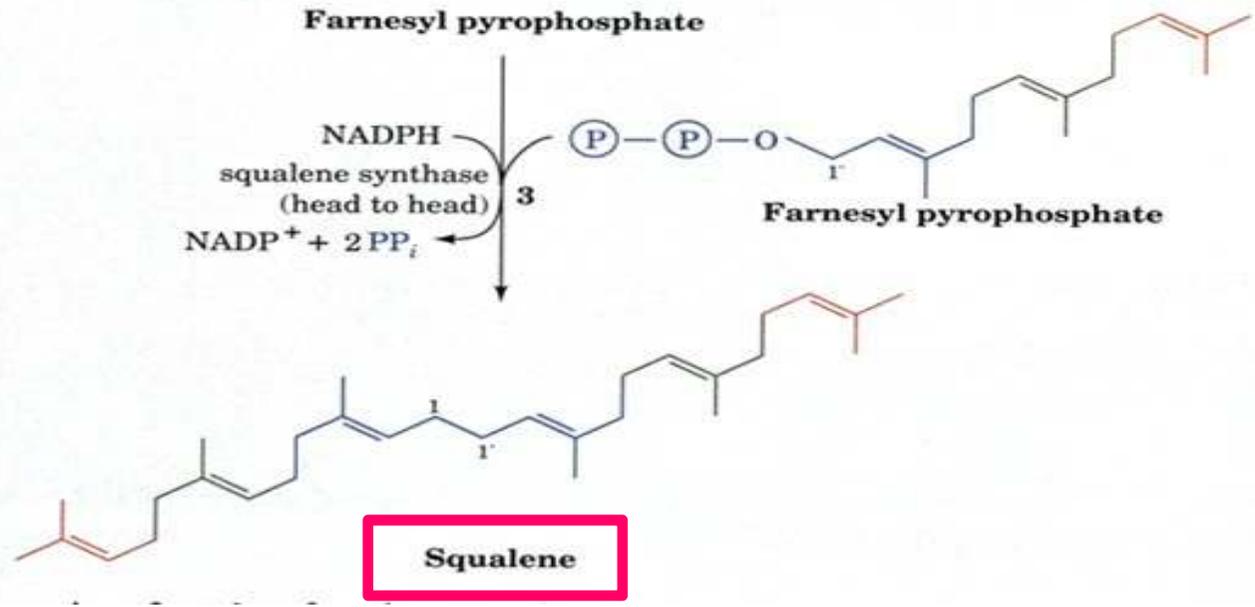
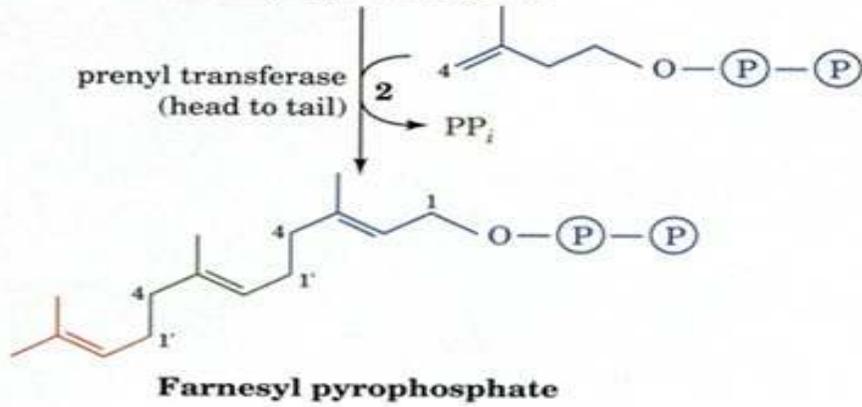
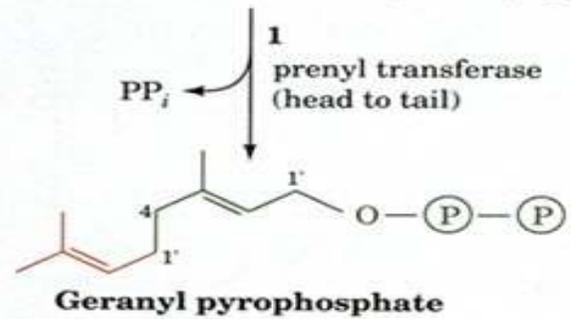
INTERCONVERSION:

The isomerase catalyzes the protonation/deprotonation reaction





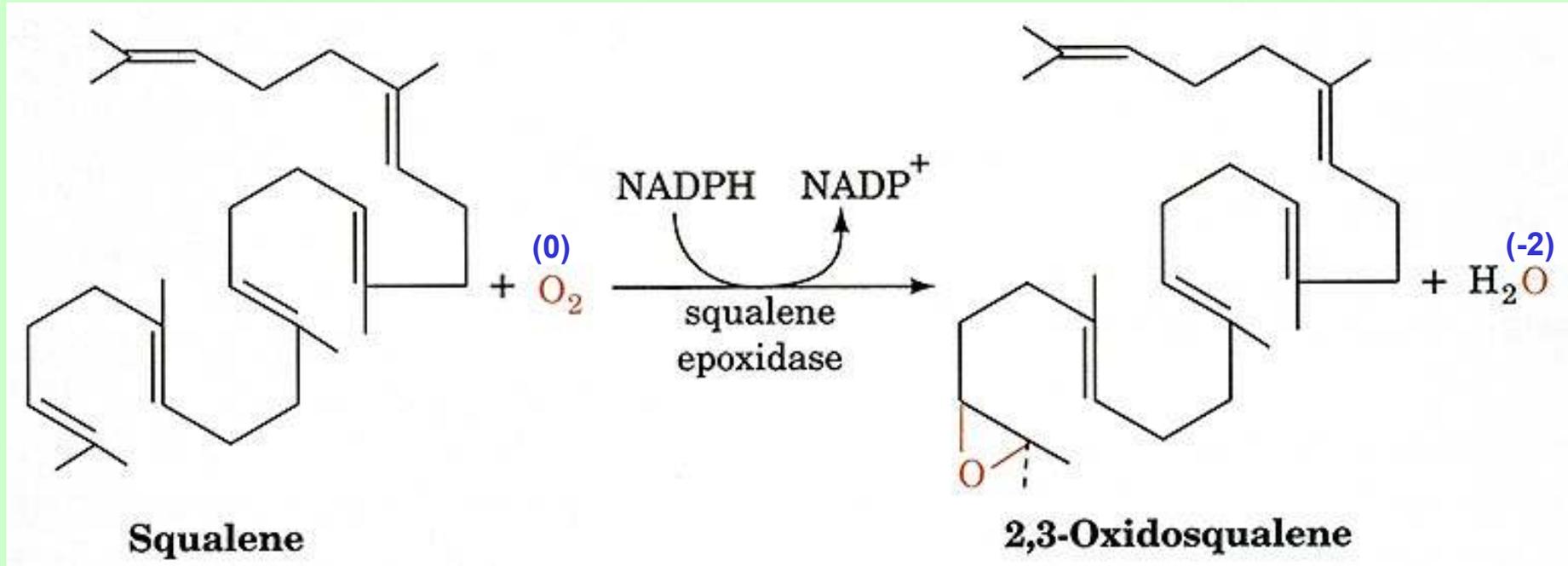
“activated isoprenes”



2 head-to-tail
condensation
reactions (1,2)

The
head-to-head
condensation
reaction (3)

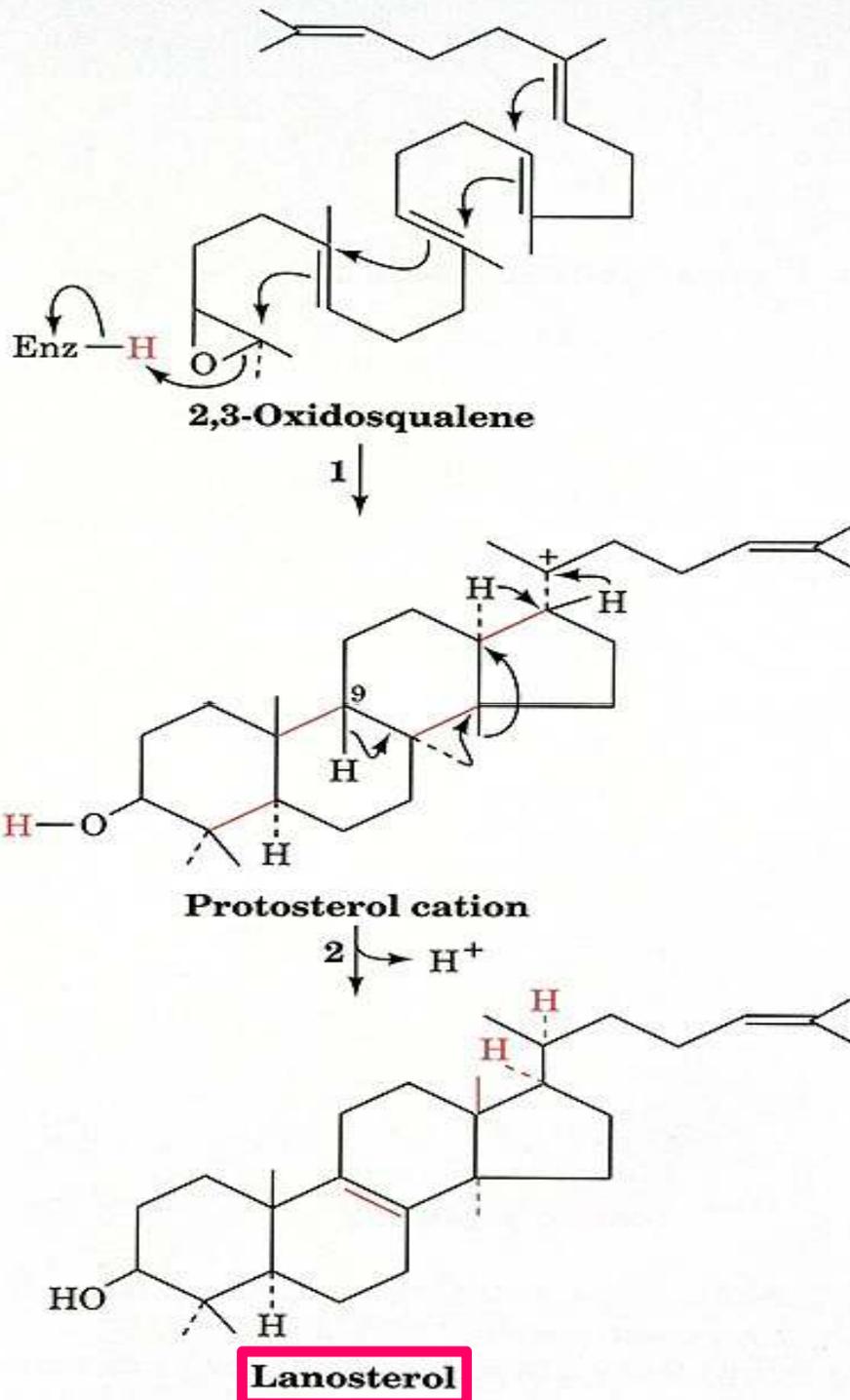
SQUALENE is converted to LANOSTEROL by CYCLISATION...

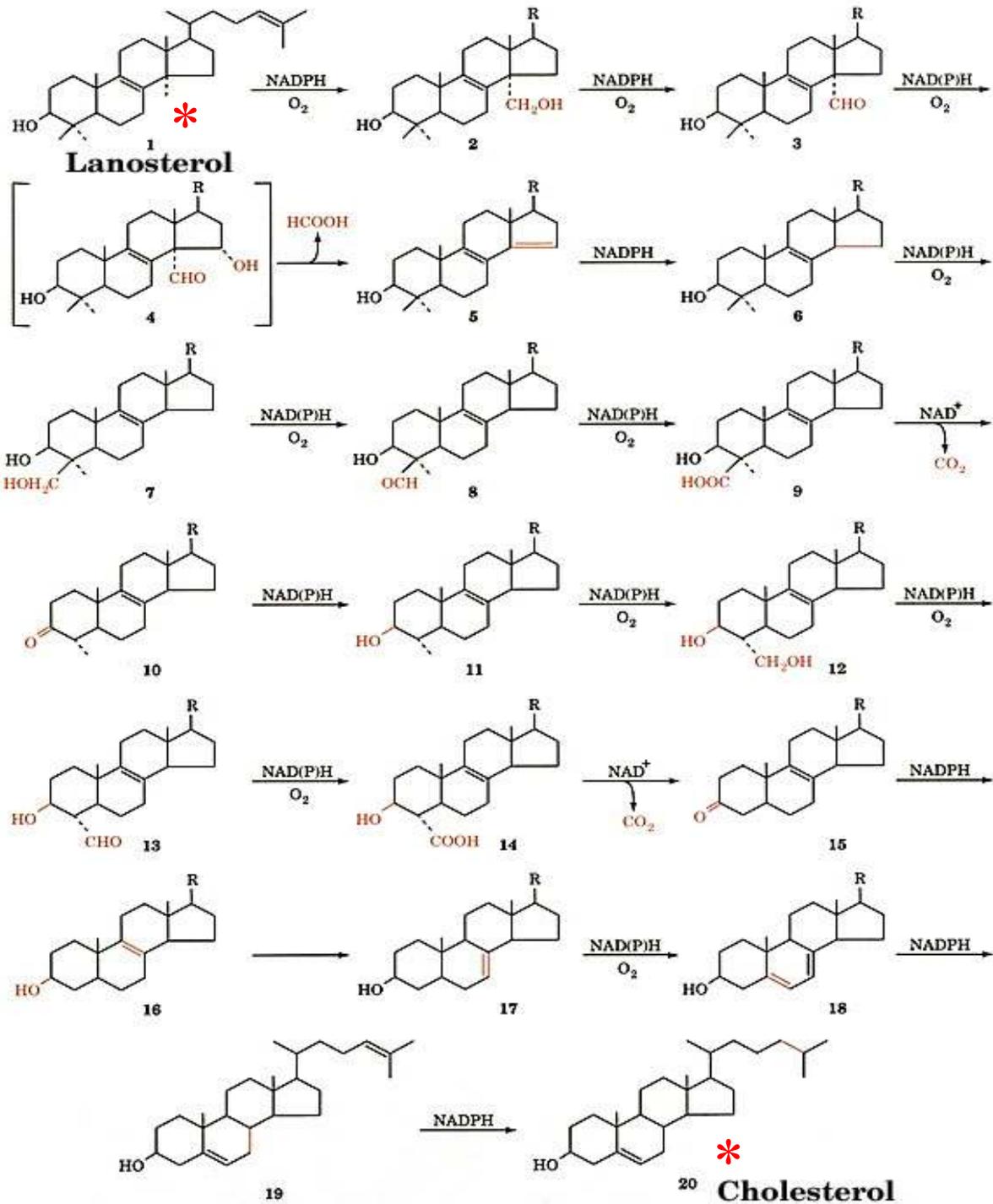


The

SQUALENE-OXIDOCYCLASE REACTION

1. PROTONATION of "O"
(lack of electrons in the center)
2. ELIMINATION of PROTON
from C9; formation of "=" bond





19 REACTIONS OF LANOSTEROL to CHOLESTEROL CONVERSION.

Enzymes incorporated in
the ER membrane

Oxidation and
demethylation (3 groups)

For all reactions:
**NADPH and O_2
required!!!**

*... the reactions are
just for information...*

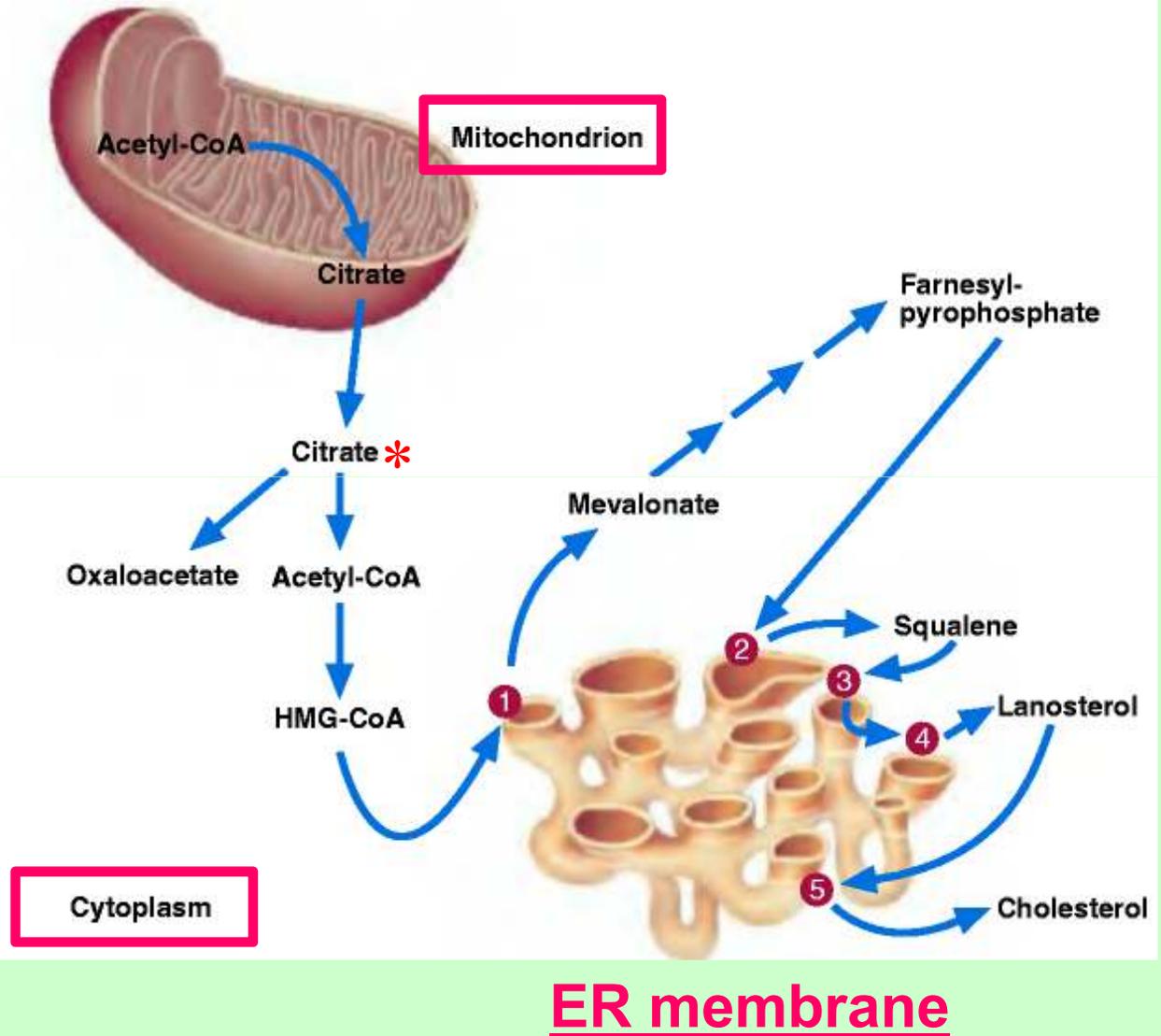
Intracellular localization of cholesterol biosynthesis

FIGURE 12.23

Cholesterol Synthesis.

Several reactions occur in the cytoplasm, but most enzymes involved in cholesterol biosynthesis occur within ER membrane. The enzymes are indicated by the following numbers: 1 = HMG-CoA reductase, 2 = Squalene synthase, 3 = Squalene monooxygenase, 4 = 2,3-Oxidosqualene lanosterol cyclase, 5 = Enzymes catalyzing 20 separate reactions. Note that squalene and lanosterol are acted upon by ER membrane enzymes while they are bound to carrier proteins in the cytoplasm.

*** Citrate carries acetyl units from mitochondrion into cytosol!!!**



REGULATION of cholesterol biosynthesis

Regulation of cholesterol formation balances synthesis with dietary uptake and energy state!!!

Short-term regulation by:

1) AMP-Dependent Protein Kinase

2) Hormones (insulin, glucagon)

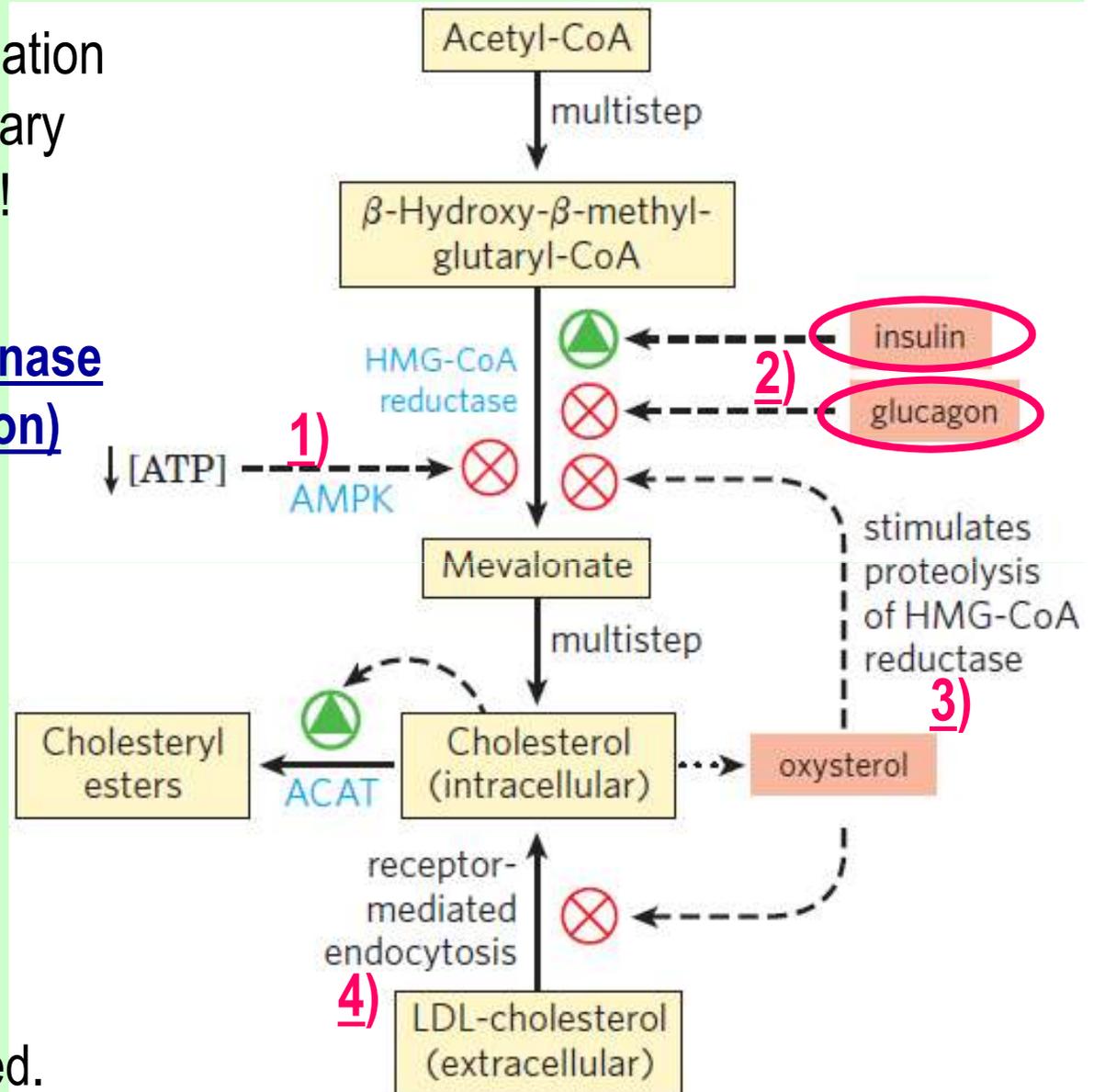
Long-term regulation:

3) by proteolysis

(the enzyme degradation)

4) at the transcription level

(when extracellular cholesterol is taken up by the cells, the genes for synthesis of HMG-Co A reductase and of LDL-receptor are not activated.



Short-term regulation:

1) HMG-CoA Reductase is inactivated (inhibited) by phosphorylation, catalyzed by AMP-Dependent Protein Kinase, AMPK, (also regulates fatty acid synthesis and catabolism).

This kinase is active when cellular [AMP] is high, corresponding to when ATP is low.

Thus, when cellular [ATP] is low, energy is not expended in synthesizing cholesterol.

2) **Insulin** promotes dephosphorylation (activation) of HMG-CoA reductase; **glucagon** promotes its phosphorylation (inactivation);

Long-term regulation is by varied **formation** and **degradation** of HMG-CoA reductase and other enzymes of the pathway for synthesis of cholesterol.

3) Regulated proteolysis of **HMG-CoA Reductase**:

Degradation of HMG-CoA Reductase is stimulated by cholesterol, oxidized derivatives of cholesterol (oxysterol or 24(S)-hydroxycholesterol), mevalonate, farnesol (dephosphorylated farnesyl pyrophosphate), and probably bile acids.

HMG-CoA Reductase includes a transmembrane sterol-sensing domain that has a role in activating degradation of the enzyme *via* the proteasome.

4) Regulated transcription:

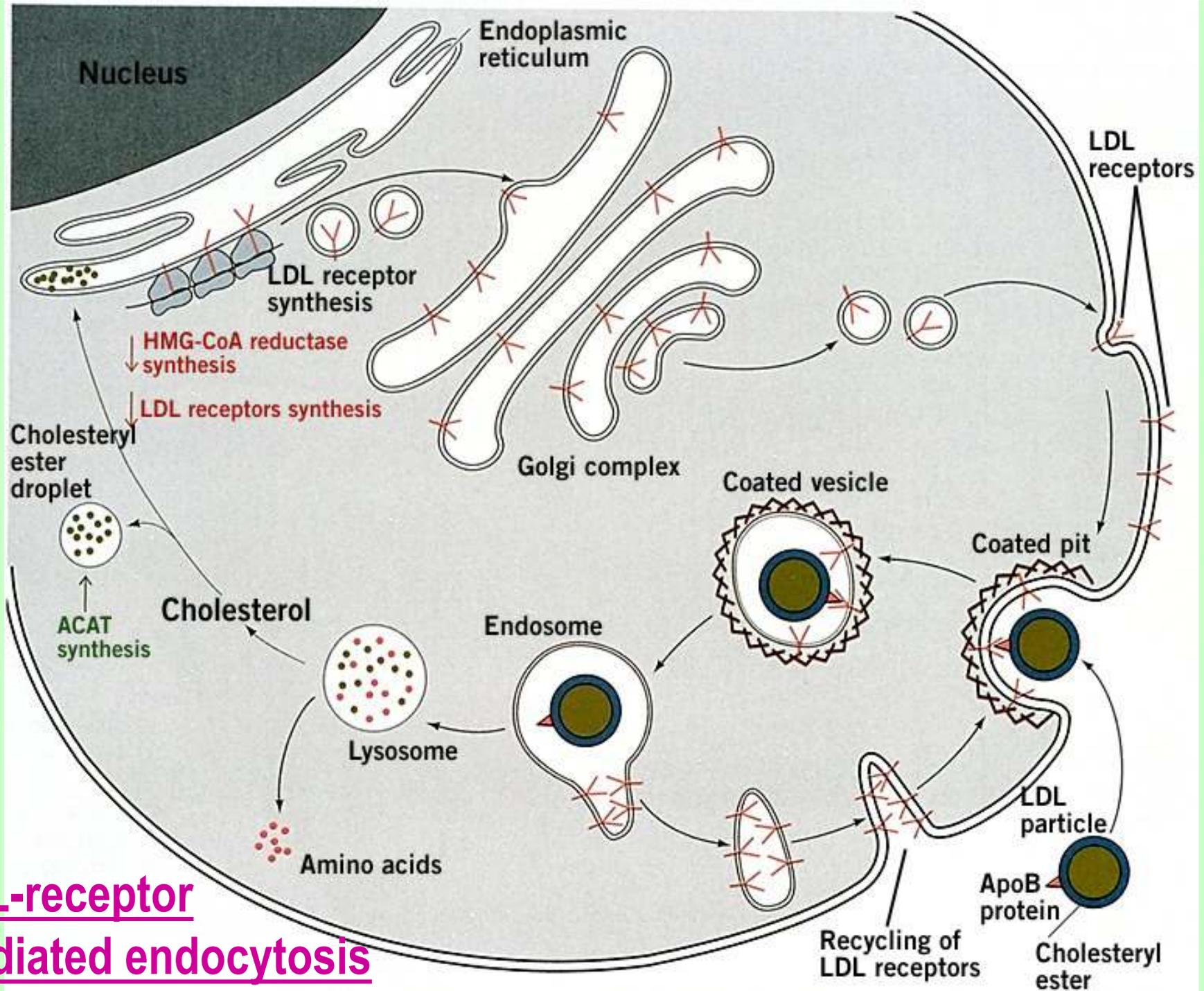
A family of transcription factors designated **SREBP** (sterol regulatory element binding proteins) regulate synthesis of cholesterol and fatty acids.

Of these, **SREBP-2** mainly **regulates cholesterol synthesis**.

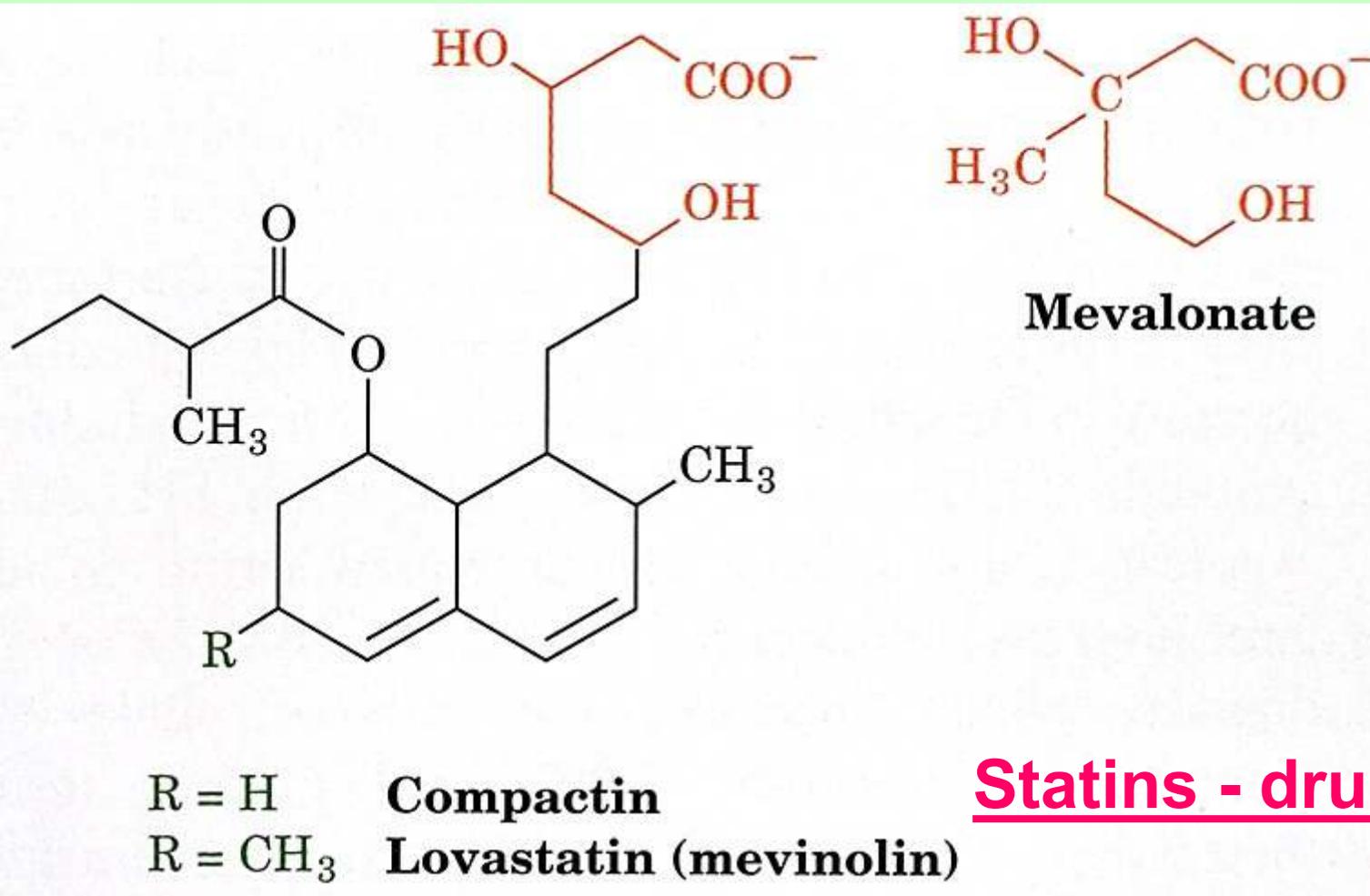
(SREBP-1c mainly regulates fatty acid synthesis.)

When sterol concentrations are **low**, **SREBP-2** is **released** by cleavage of a membrane-bound precursor protein.

SREBP-2 **activates transcription** of genes for **HMG-CoA Reductase** and other enzymes of the pathway for cholesterol synthesis.



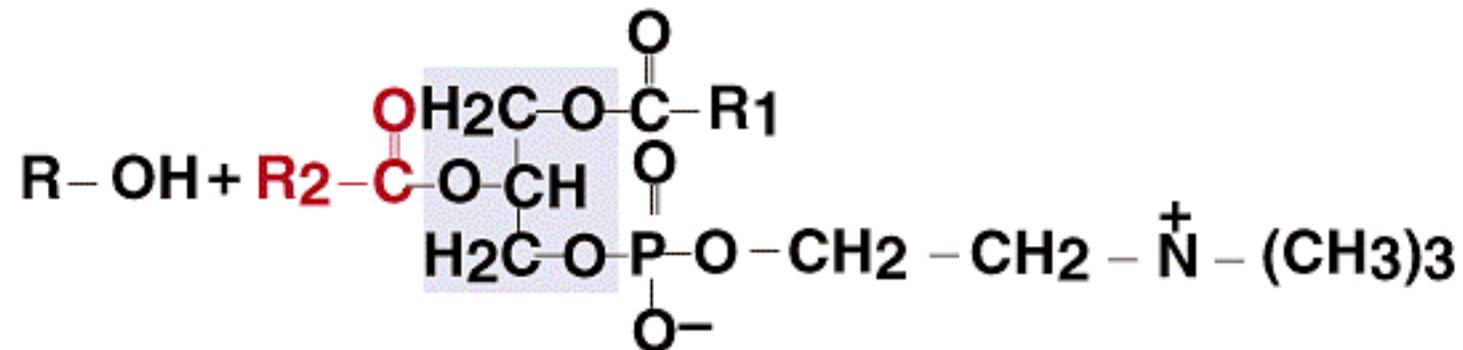
LDL-receptor mediated endocytosis



INHIBITORS of HMG-CoA reductase
(competitive inhibitors due to
similarity with mevalonate).

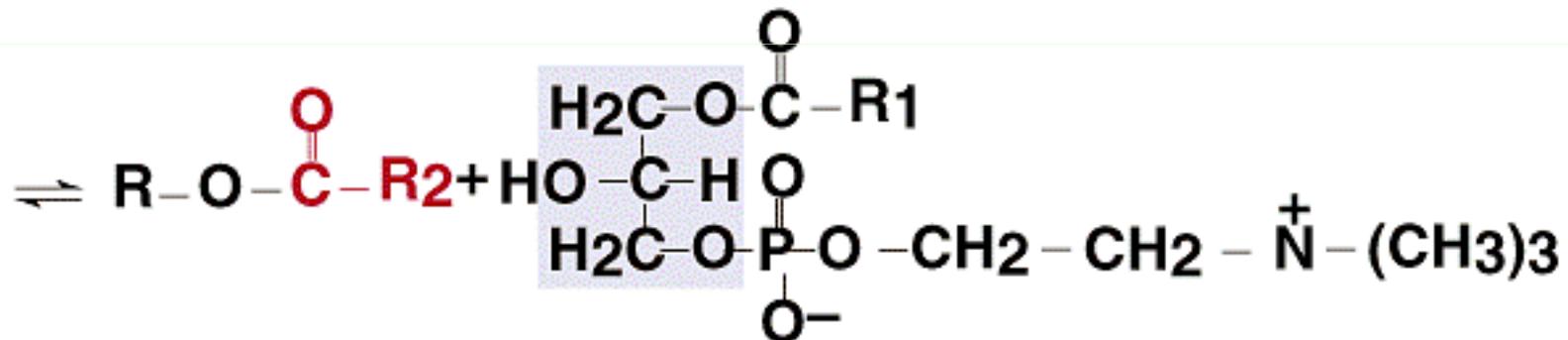
Cholesterol

Phosphatidylcholine

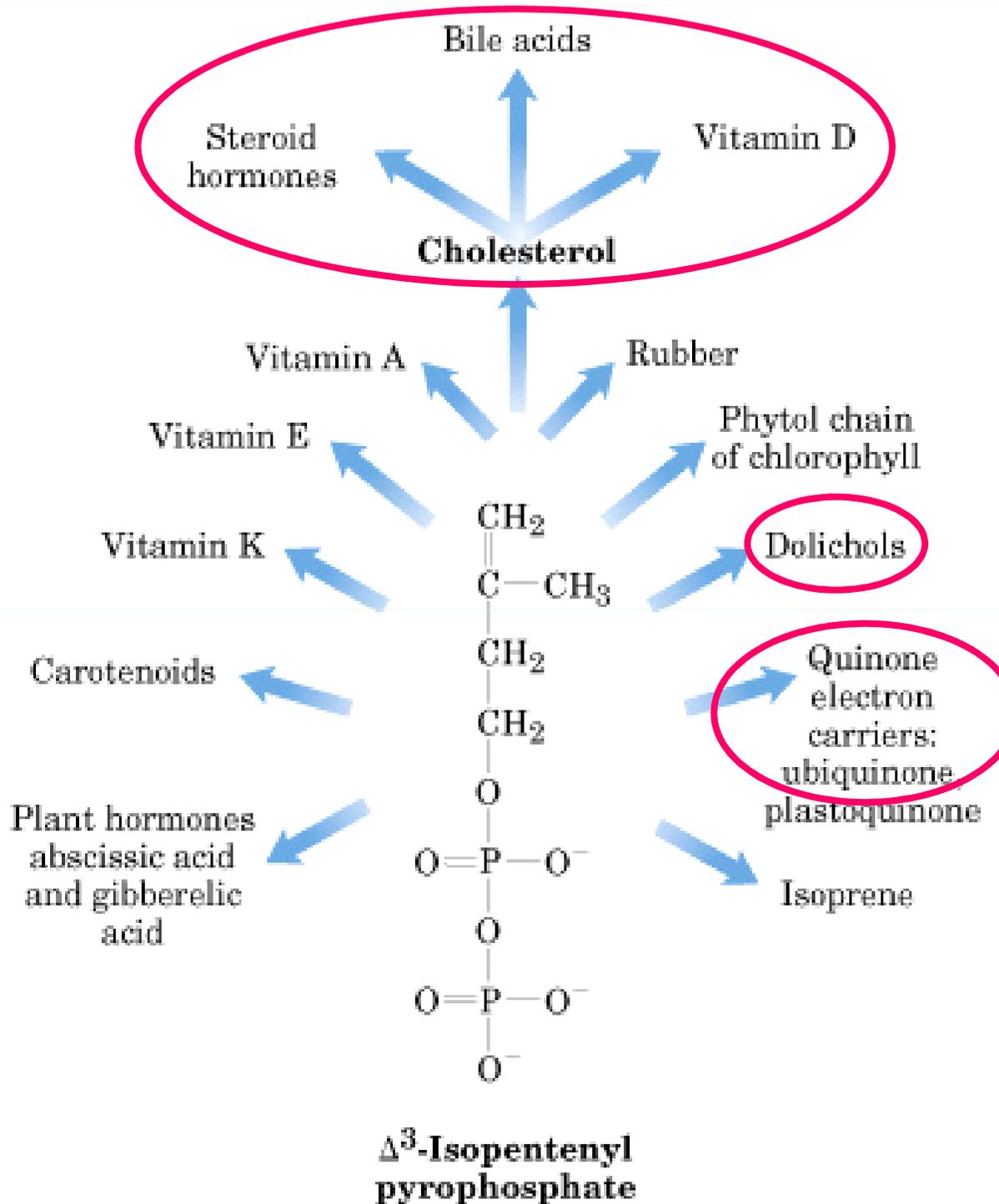


Cholesterol ester

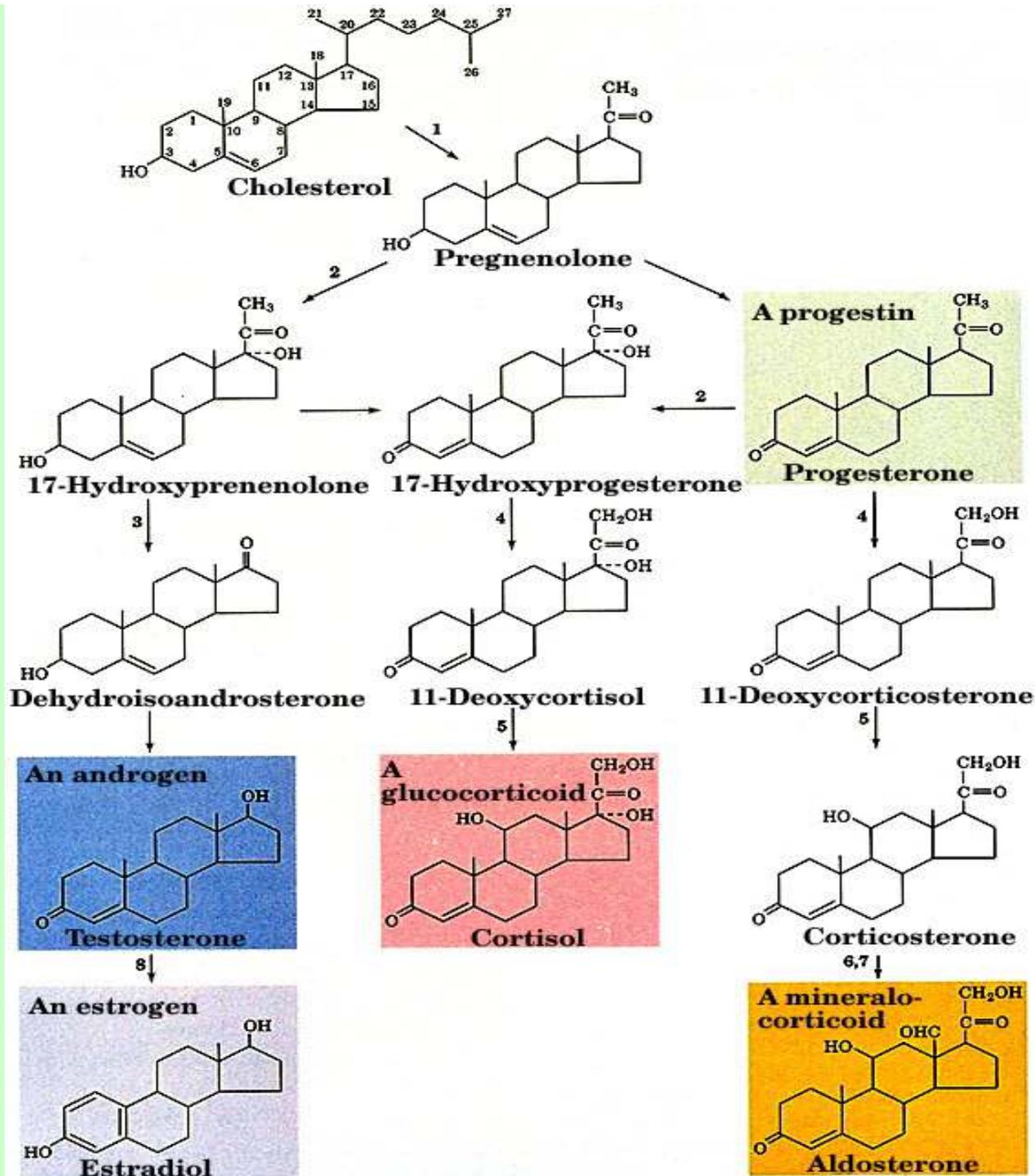
Lysophosphatidylcholine



Conversion of cholesterol into cholesterol-esters by the **LECITHIN:CHOLESTEROL ACYL TRANSFERASE (LCAT)**, enzyme in plasma bound to HDL (synthesized by the liver, mostly) - the cholesterol-esters diffuse into the core of HDL.

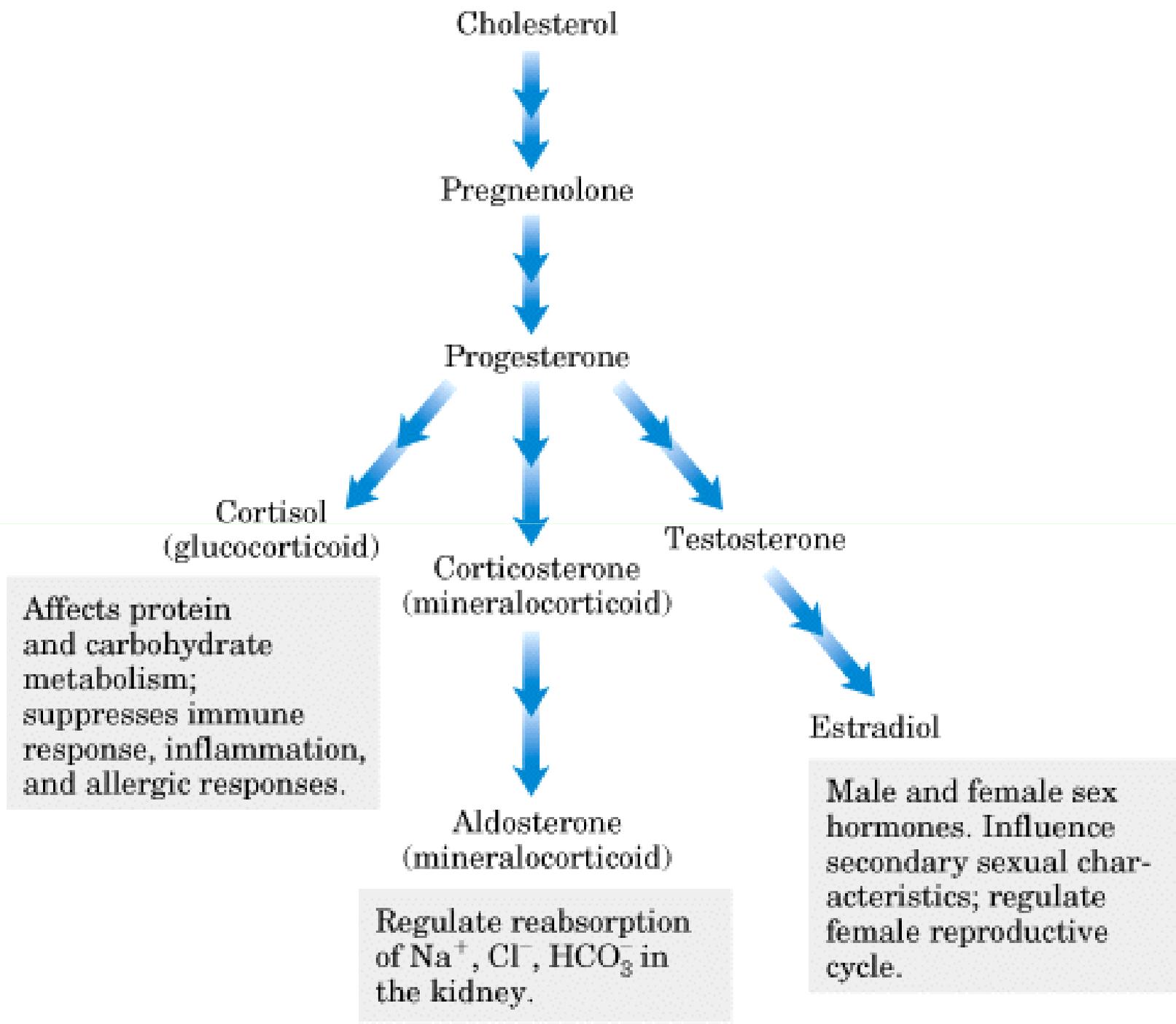


In humans...

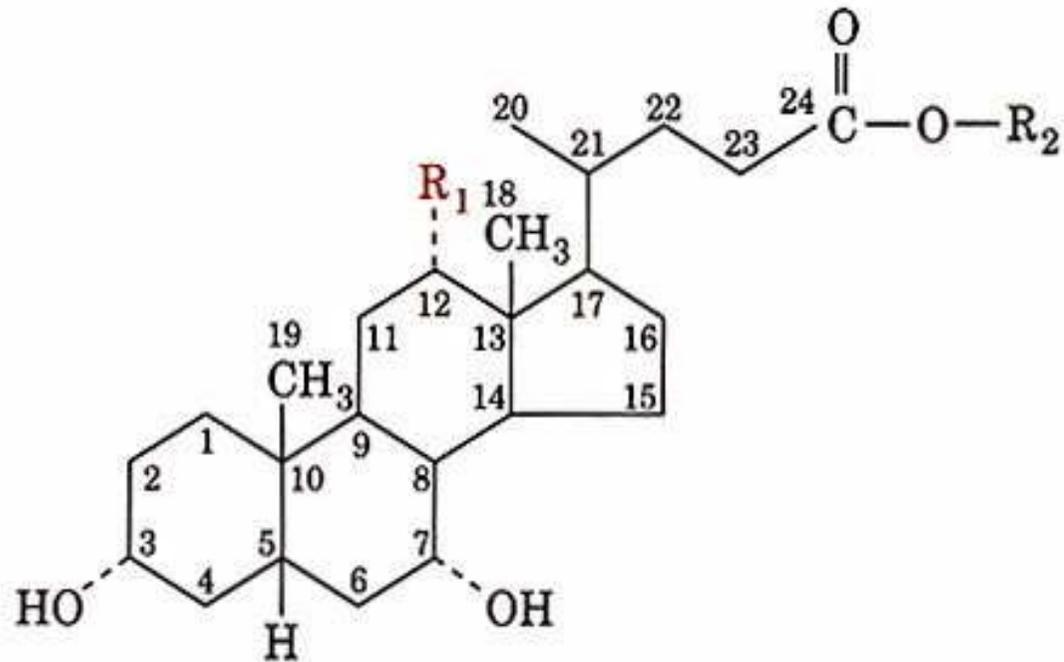


HORMONES, synthesis

1. Enzyme for the side chain cleavage
2. Steroid-C17-hydroxylase
3. Steroid-C17,C20-lyase
4. Steroid-C21-hydroxylase
5. Steroid-11 β -hydroxylase
6. Steroid-C18-hydroxylase
7. 18-hydroxysteroid-oxidase
8. aromathase



Bile acids



$R_1 = \text{OH}$

$R_1 = \text{H}$

$R_2 = \text{H}$

$R_2 = \text{NH} - \text{CH}_2 - \text{COOH}$

$R_2 = \text{NH} - \text{CH}_2 - \text{CH}_2 - \text{SO}_3\text{H}$

Cholic acid

Glycocholic acid

Taurocholic acid

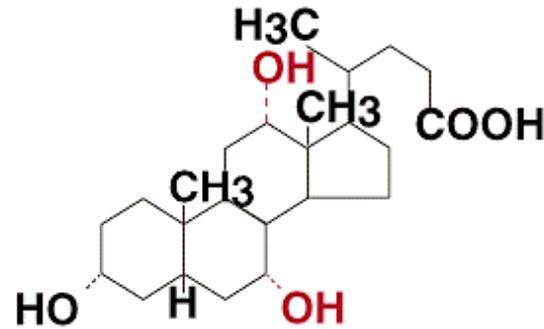
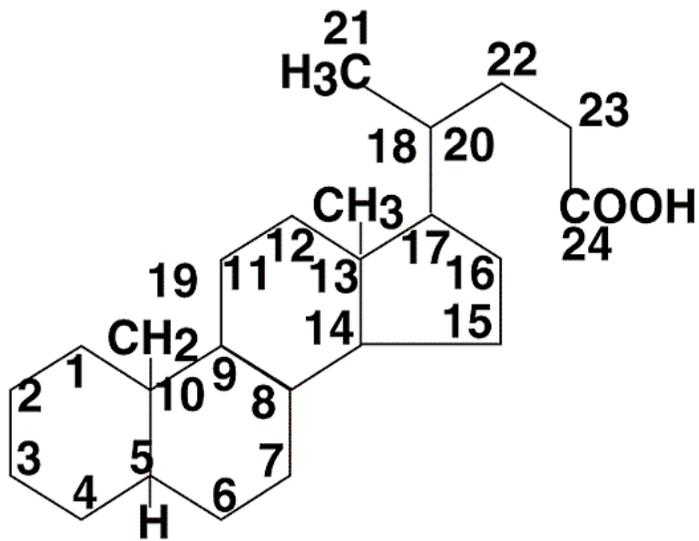
Chenodeoxycholic acid

Glychenodeoxycholic acid

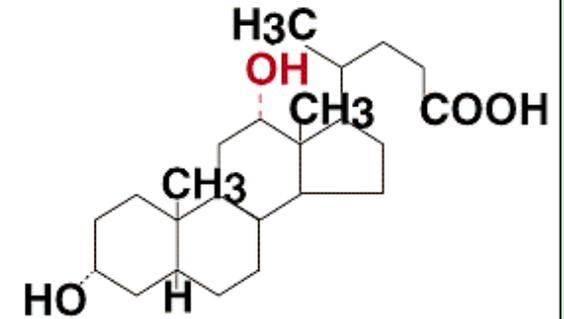
Taurochenodeoxycholic acid

Bile acids

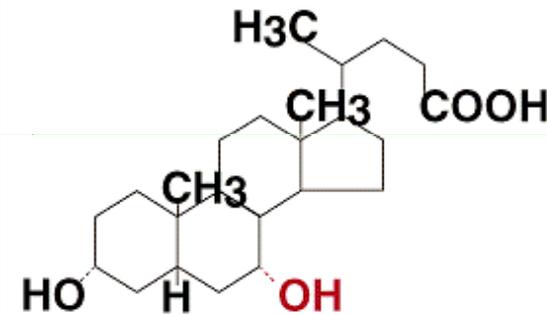
Cholanic acid



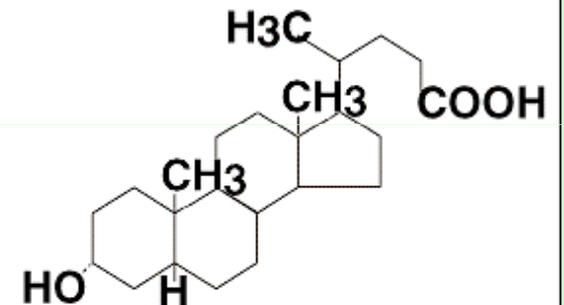
Cholic acid



Deoxycholic acid



Chenodeoxycholic acid



Lithocholic acid

CHOLANIC ACID DERIVATIVES

The primary bile acids are synthesized in the LIVER from cholesterol. The first synthesized are cholic acid (found in the largest amount) and chenodeoxycholic acid.

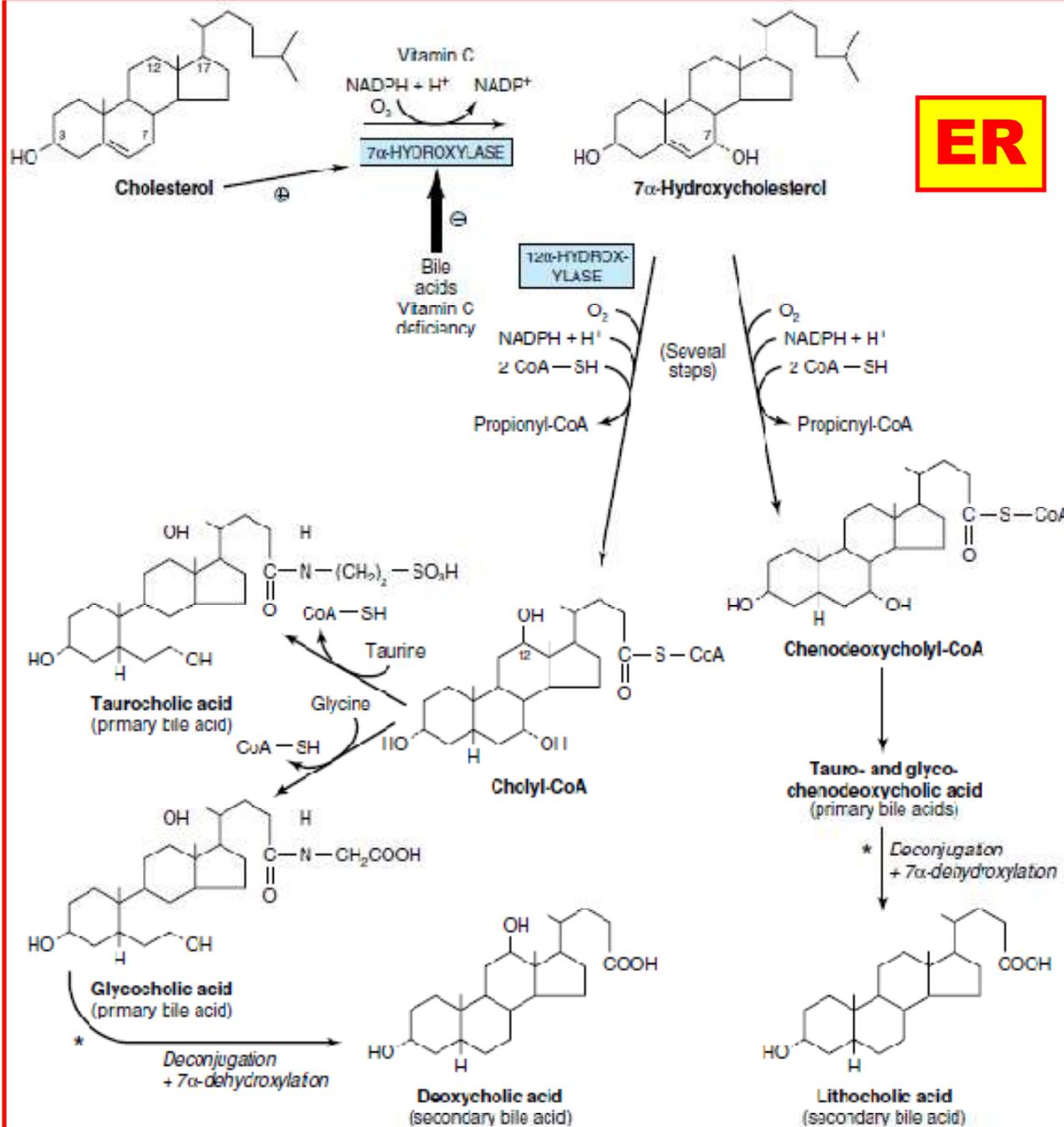


Figure 26-7. Biosynthesis and degradation of bile acids. A second pathway in mitochondria involves hydroxylation of cholesterol by sterol 27-hydroxylase. Asterisk: Catalyzed by microbial enzymes.

The 7 α -hydroxylation of cholesterol is the first and principal regulatory step in the biosynthesis of bile acids catalyzed by 7-hydroxylase, a microsomal enzyme. (A typical monooxygenase, it requires oxygen, NADPH, and cytochrome P450.)

Subsequent hydroxylation steps are also catalyzed by monooxygenases.

The pathway of bile acid biosynthesis divides early into one subpathway leading to **choly-CoA**, characterized by an extra α -OH group on position 12, and another pathway leading to **chenodeoxycholy-CoA**.

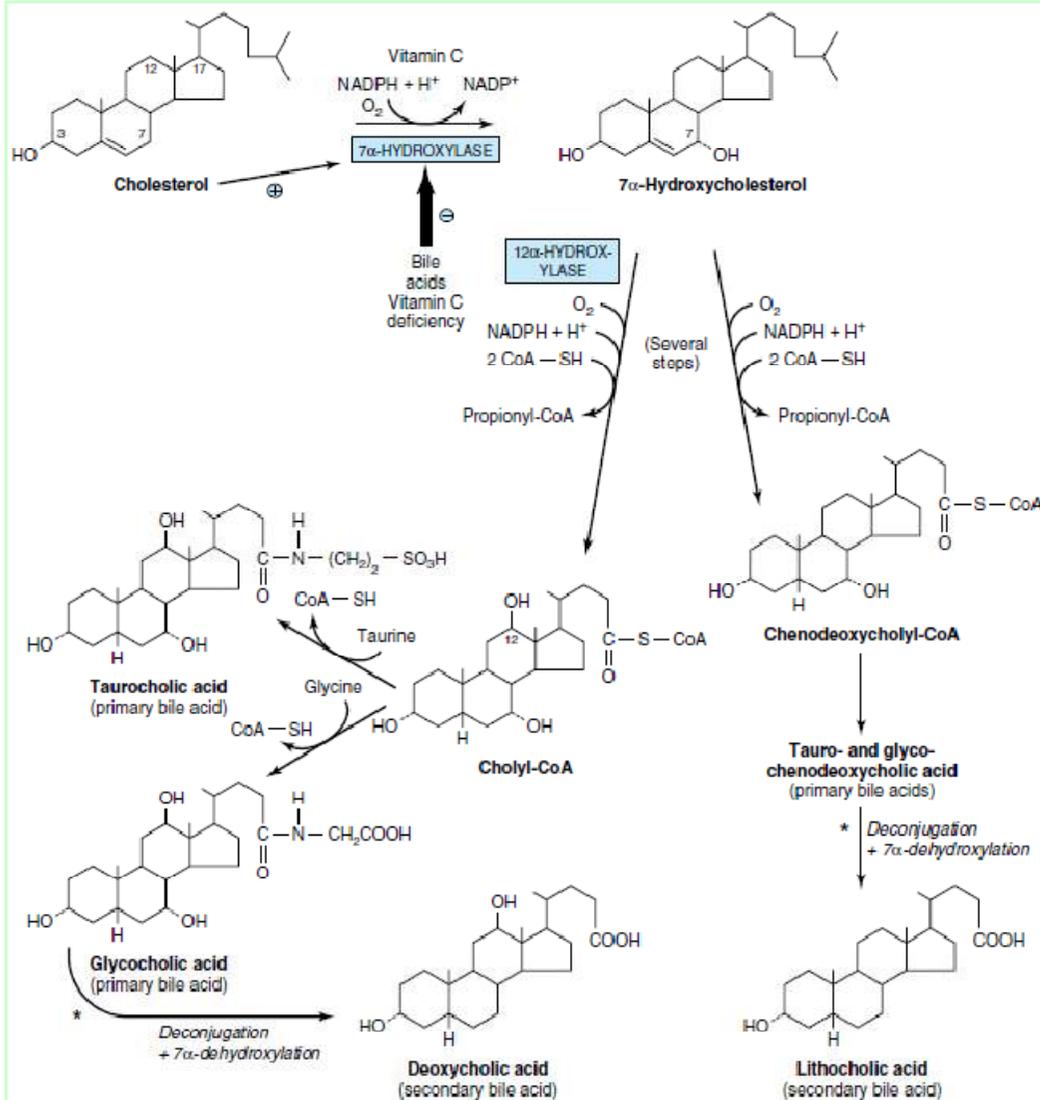


Figure 26-7. Biosynthesis and degradation of bile acids. A second pathway in mitochondria involves hydroxylation of cholesterol by sterol 27-hydroxylase. Asterisk: Catalyzed by microbial enzymes.

A second pathway in mitochondria involving the **27-hydroxylation** of cholesterol by **sterol 27-hydroxylase** as the first step is responsible for a significant proportion of the primary bile acids synthesized.

The primary bile acids **enter the bile** as **glycine or taurine conjugates**.

Conjugation takes place **in peroxisomes**.

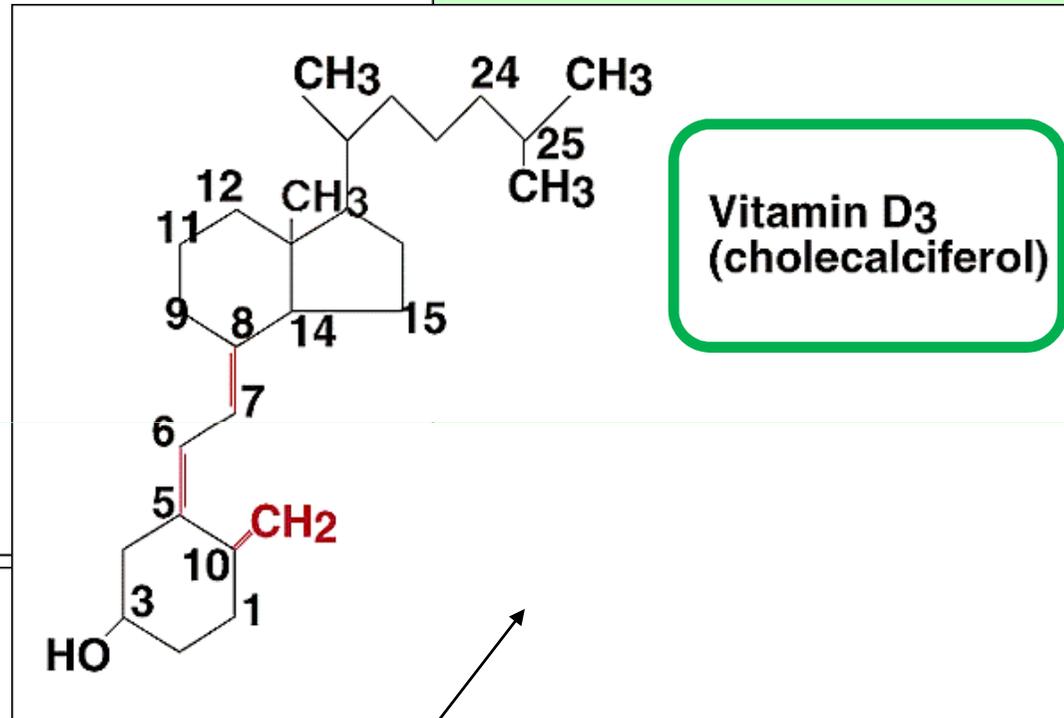
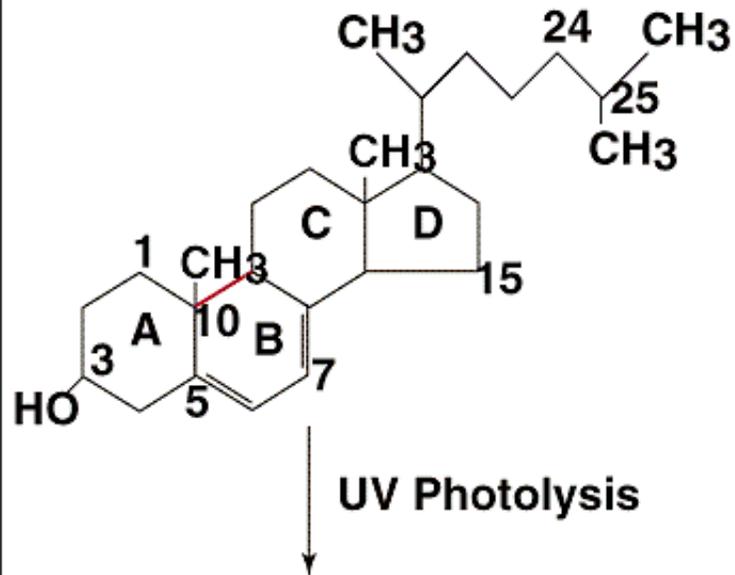
In humans, the ratio of the glycine to the taurine conjugates is normally 3:1.

(In the alkaline bile, the bile acids and their conjugates are assumed to be in a salt form — hence the term “bile salts.”)

A portion of the primary bile acids **in the intestine** is subjected to further changes by **the activity of the intestinal bacteria**. These include **deconjugation** and **7α-dehydroxylation**, which produce the **secondary bile acids**, deoxycholic acid and lithocholic acid.

(in skin)

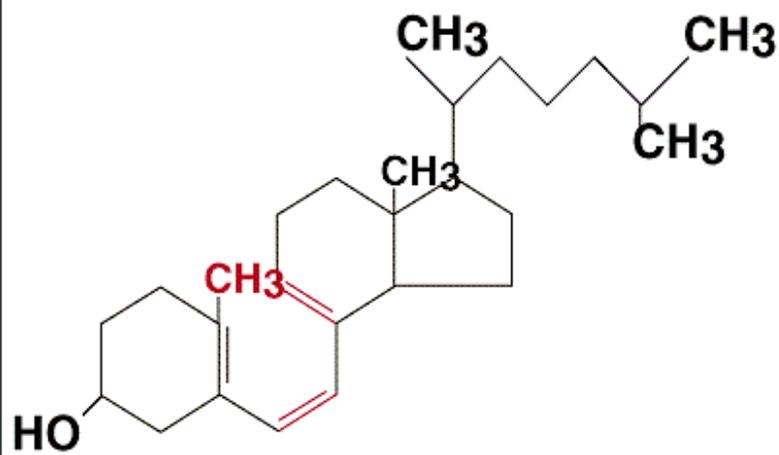
7-Dehydrocholesterol



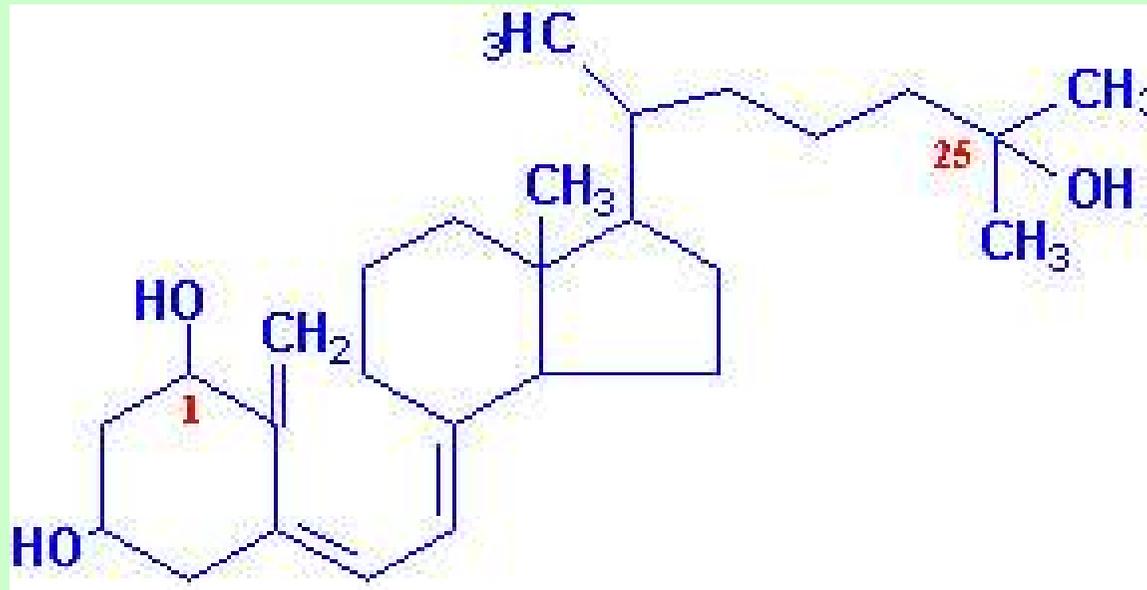
Vitamin D₃
(cholecalciferol)

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Previtamin D₃



..... Hydroxylation at C25 and C1:



1,25-dihydroxyvitamin D₃ (calcitriol; hormon)

- Hydroxylation at C25: in liver
- Hydroxylation at C1: in kidney, bones and placenta

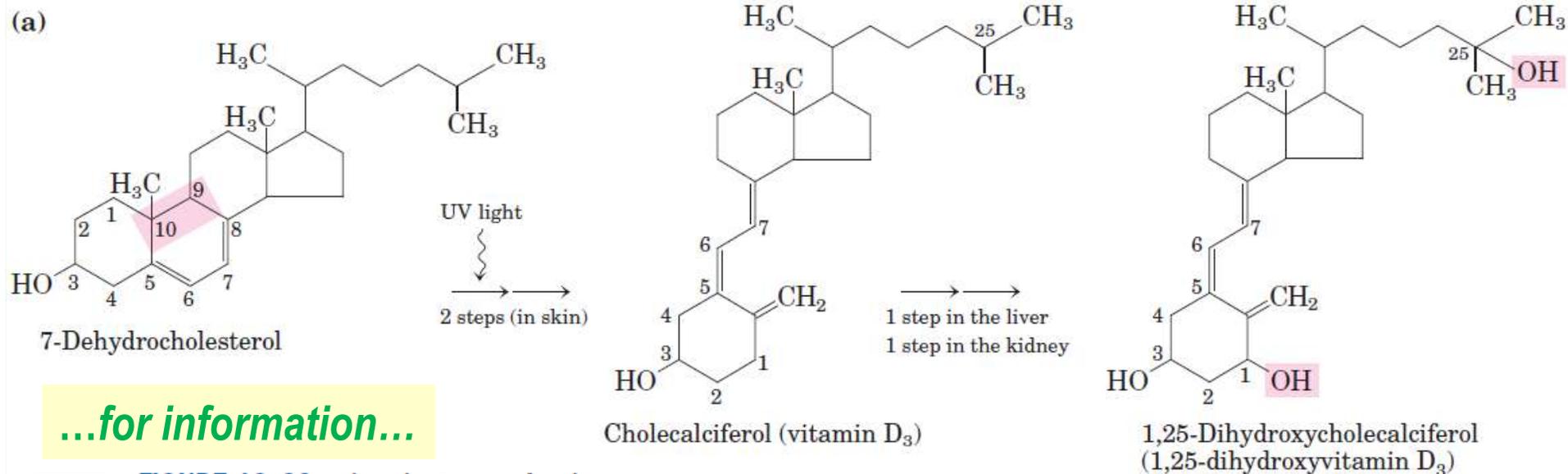


FIGURE 10-20 Vitamin D₃ production and metabolism. (a) Cholecalciferol

(vitamin D₃) is produced in the skin by UV irradiation of 7-dehydrocholesterol, which breaks the bond shaded pink. In the liver, a hydroxyl group is added at C-25 (pink); in the kidney, a second hydroxylation at C-1 (pink) produces the active hormone, 1,25-dihydroxycholecalciferol. This hormone regulates the metabolism of Ca²⁺ in kidney, intestine, and bone.

(b) Dietary vitamin D prevents rickets, a disease once common in cold climates where heavy clothing blocks the UV component of sunlight necessary for the production of vitamin D₃ in skin. On the left is a 2½-year-old boy with severe rickets; on the right, the same boy at age 5, after 14 months of vitamin D therapy.

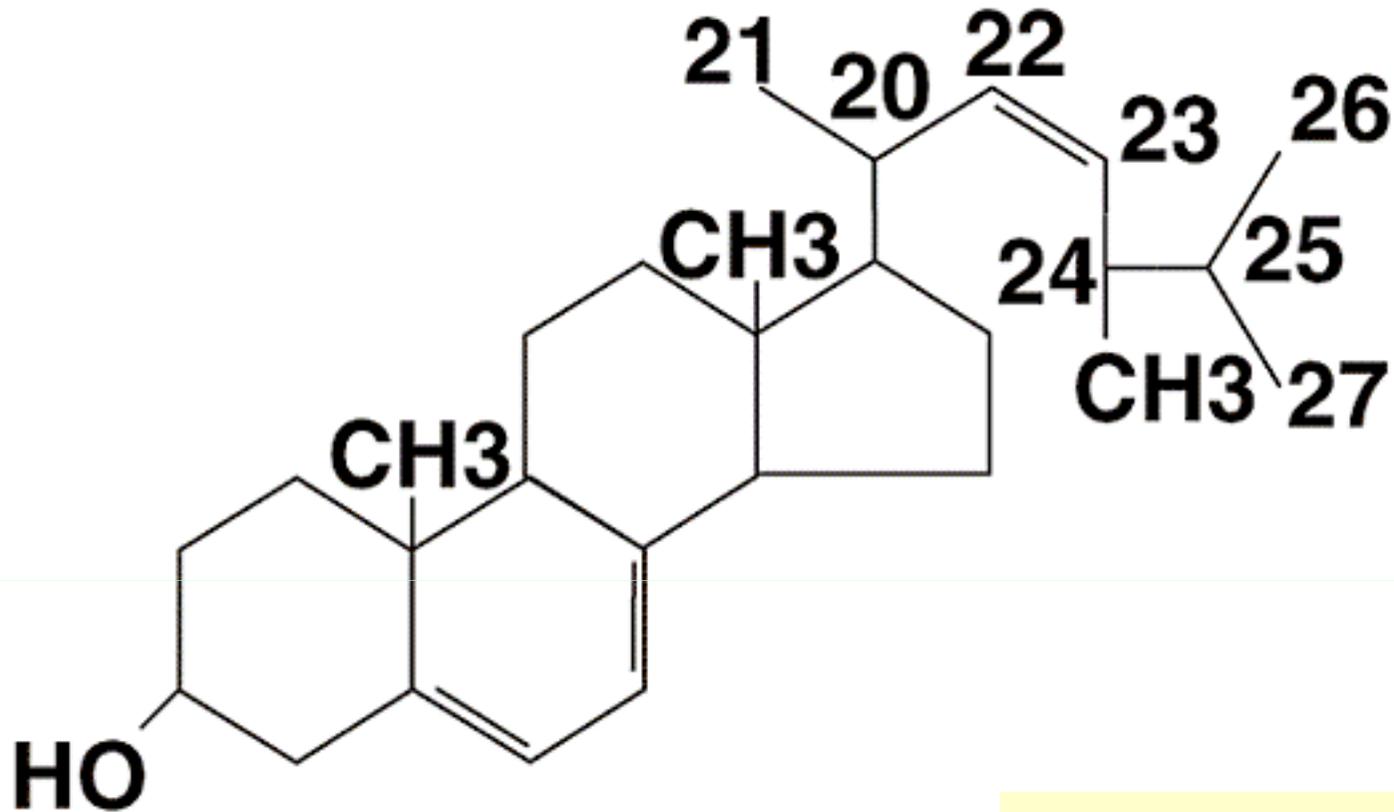


Before vitamin D treatment



After 14 months of vitamin D treatment

(b)

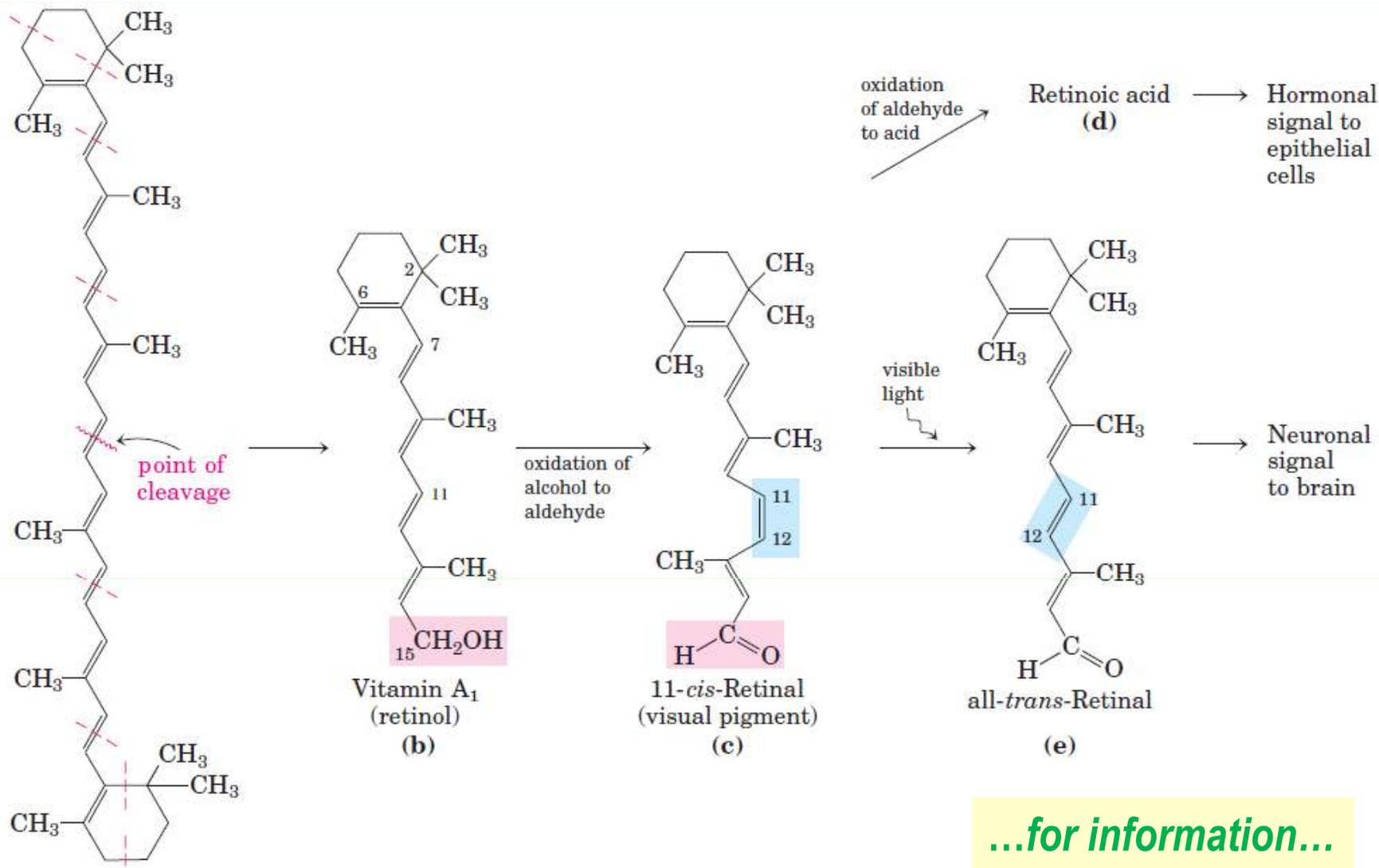


...for information...

ERGOSTEROL (PLANT STEROL)

Precursor of VITAMIN D₂ (ergocalciferol):

- transformation occurs in skin by UV-LIGHT



...for information...

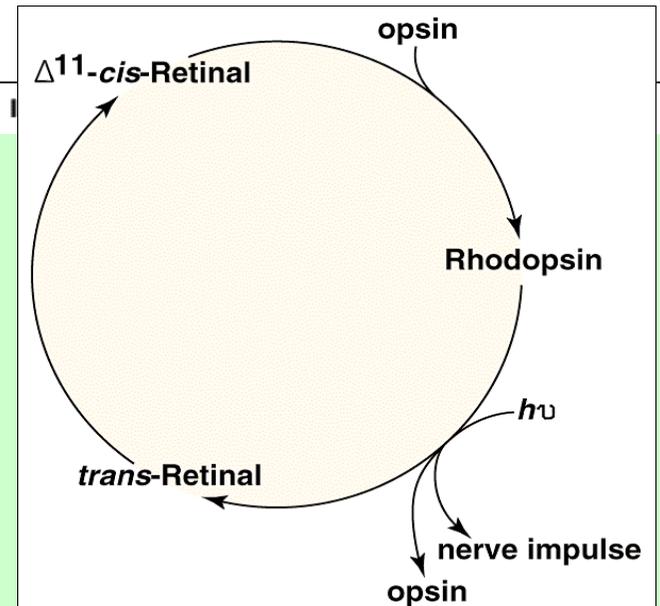
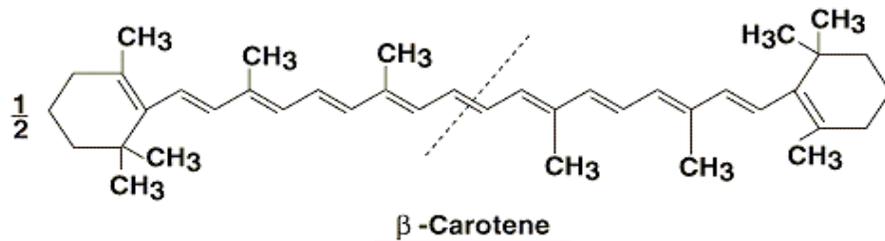
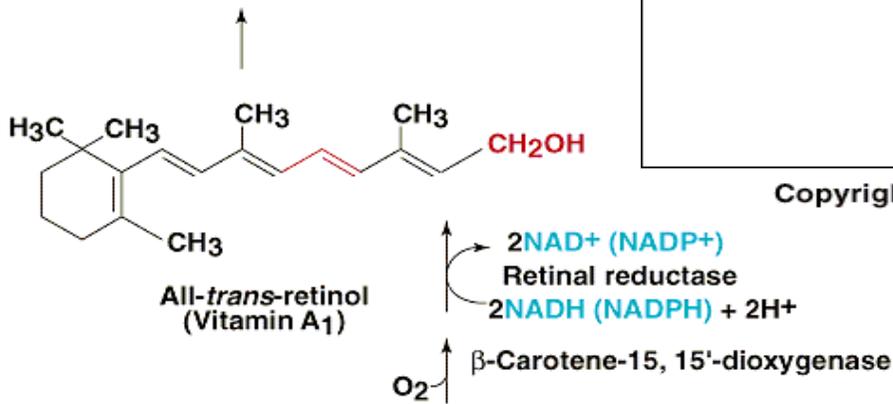
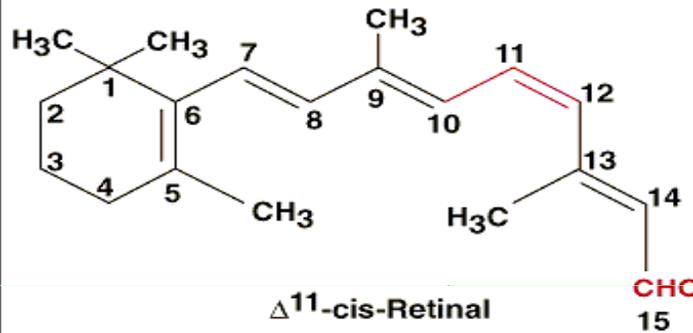
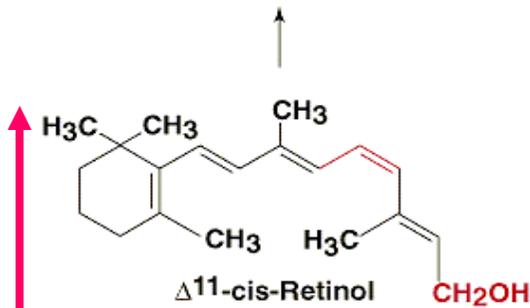
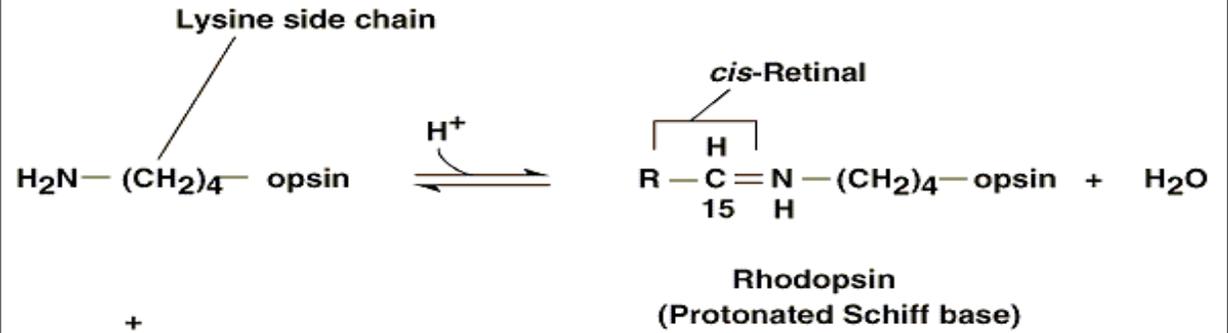


FIGURE 10-21 Vitamin A₁ and its precursor and derivatives.

(a) β -Carotene is the precursor of vitamin A₁. Isoprene structural units are set off by dashed red lines. Cleavage of β -carotene yields two molecules of vitamin A₁ (retinol) (b). Oxidation at C-15 converts retinol to the aldehyde, retinal (c), and further oxidation produces retinoic acid (d), a hormone that regulates gene expression. Retinal combines with the protein opsin to form rhodopsin (not shown), a visual pigment widespread in nature. In the dark, retinal of rhodopsin

is in the 11-*cis* form (c). When a rhodopsin molecule is excited by visible light, the 11-*cis*-retinal undergoes a series of photochemical reactions that convert it to all-*trans*-retinal (e), forcing a change in the shape of the entire rhodopsin molecule. This transformation in the rod cell of the vertebrate retina sends an electrical signal to the brain that is the basis of visual transduction, a topic we address in more detail in Chapter 12.

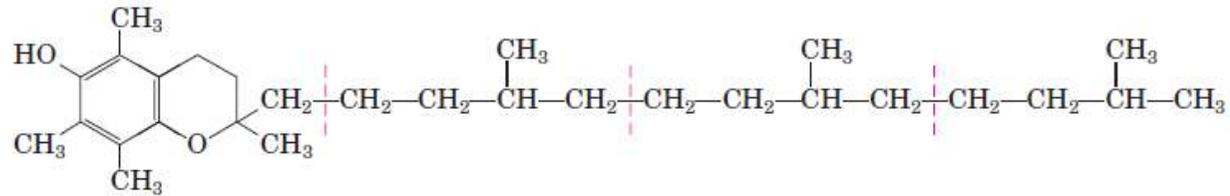
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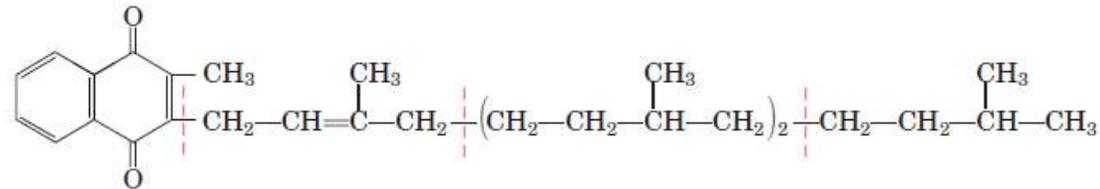
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(a)
Vitamin E: an antioxidant



(b)
Vitamin K₁: a blood-clotting cofactor (phylloquinone)



(c)
Warfarin: a blood anticoagulant

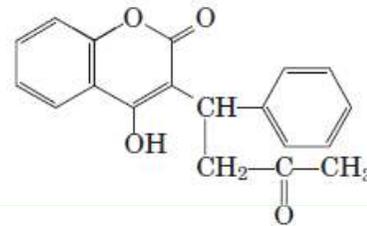
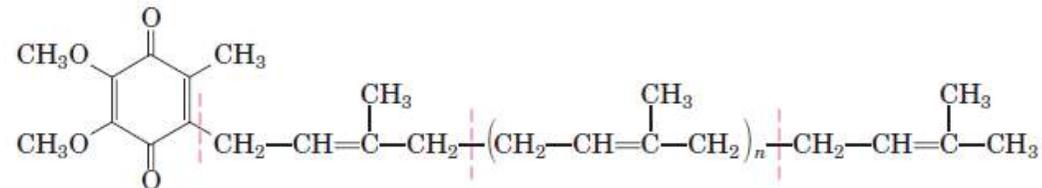
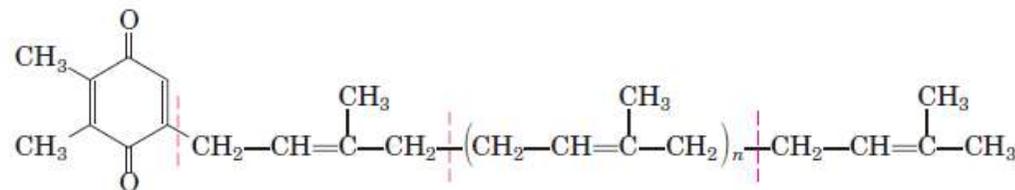


FIGURE 10-22 Some other biologically active isoprenoid compounds or derivatives. Isoprene structural units are set off by dashed red lines. In most mammalian tissues, ubiquinone (also called coenzyme Q) has 10 isoprene units. Dolichols of animals have 17 to 21 isoprene units (85 to 105 carbon atoms), bacterial dolichols have 11, and those of plants and fungi have 14 to 24.

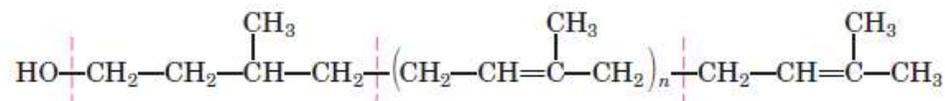
(d)
Ubiquinone: a mitochondrial electron carrier (coenzyme Q)
($n = 4$ to 8)



(e)
Plastoquinone: a chloroplast electron carrier ($n = 4$ to 8)



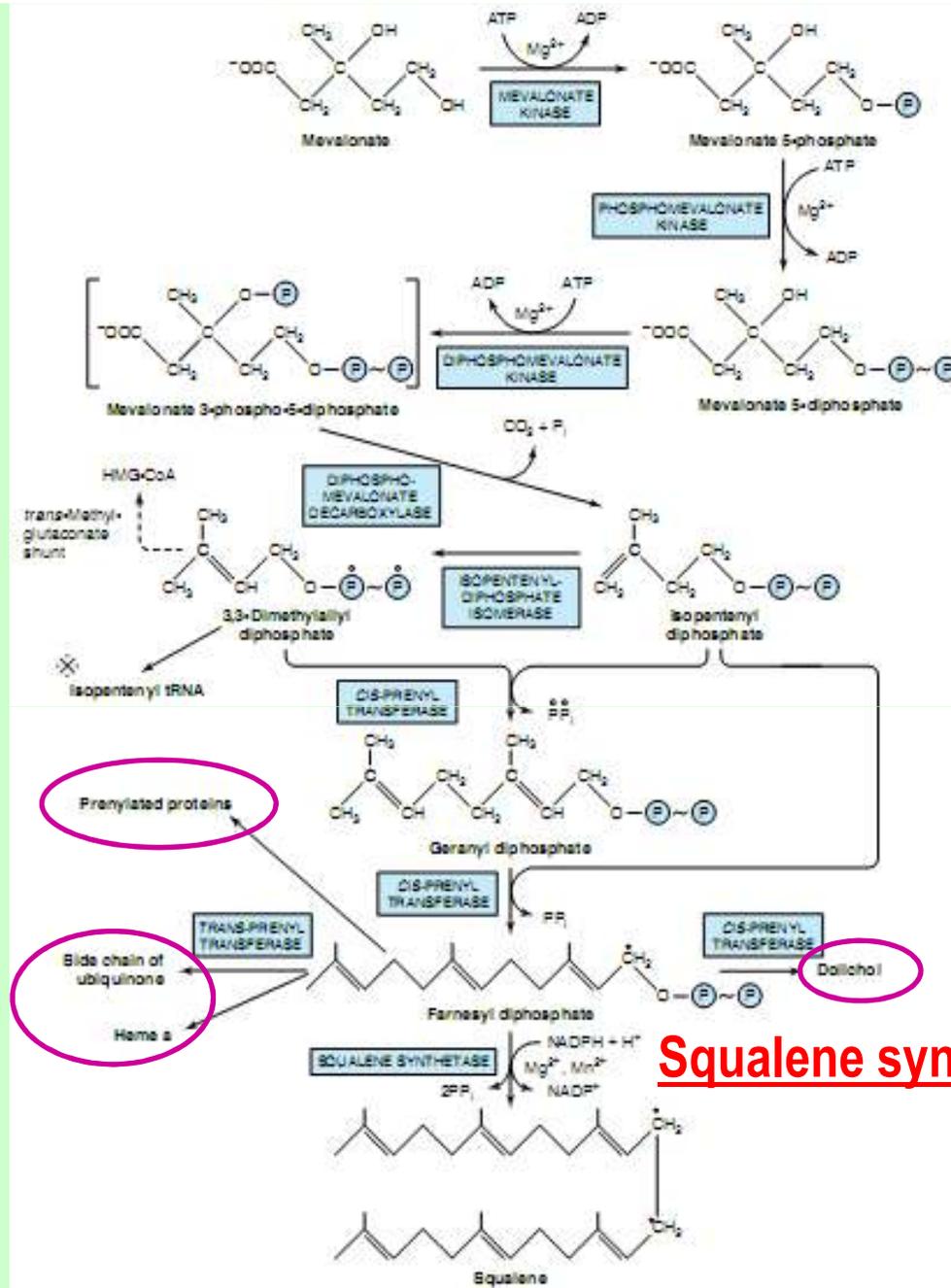
(f)
Dolichol: a sugar carrier
($n = 9$ to 22)



...for
information...

Farnesyl diphosphate:
precursor for synthesis of
 some other
 isoprenoid structures
 (e.g. dolichol)
 and
 prenylated structures
 (e.g. ubiquinone,
 prenylated proteins)

...for information...



Squalene synthase

Figure 26-2. Biosynthesis of squalene, ubiquinone, dolichol, and other polyisoprene derivatives. (HMG, 3-hydroxy-3-methylglutaryl; x, cytoskeleton.) A farnesyl residue is present in heme a of cytochrome oxidase. The carbon marked with asterisk becomes C₁₁ or C₁₂ in squalene. Squalene synthetase is a microsomal enzyme; all other enzymes indicated are soluble cytosolic proteins, and some are found in peroxisomes.

Outline of Cholesterol Synthesis

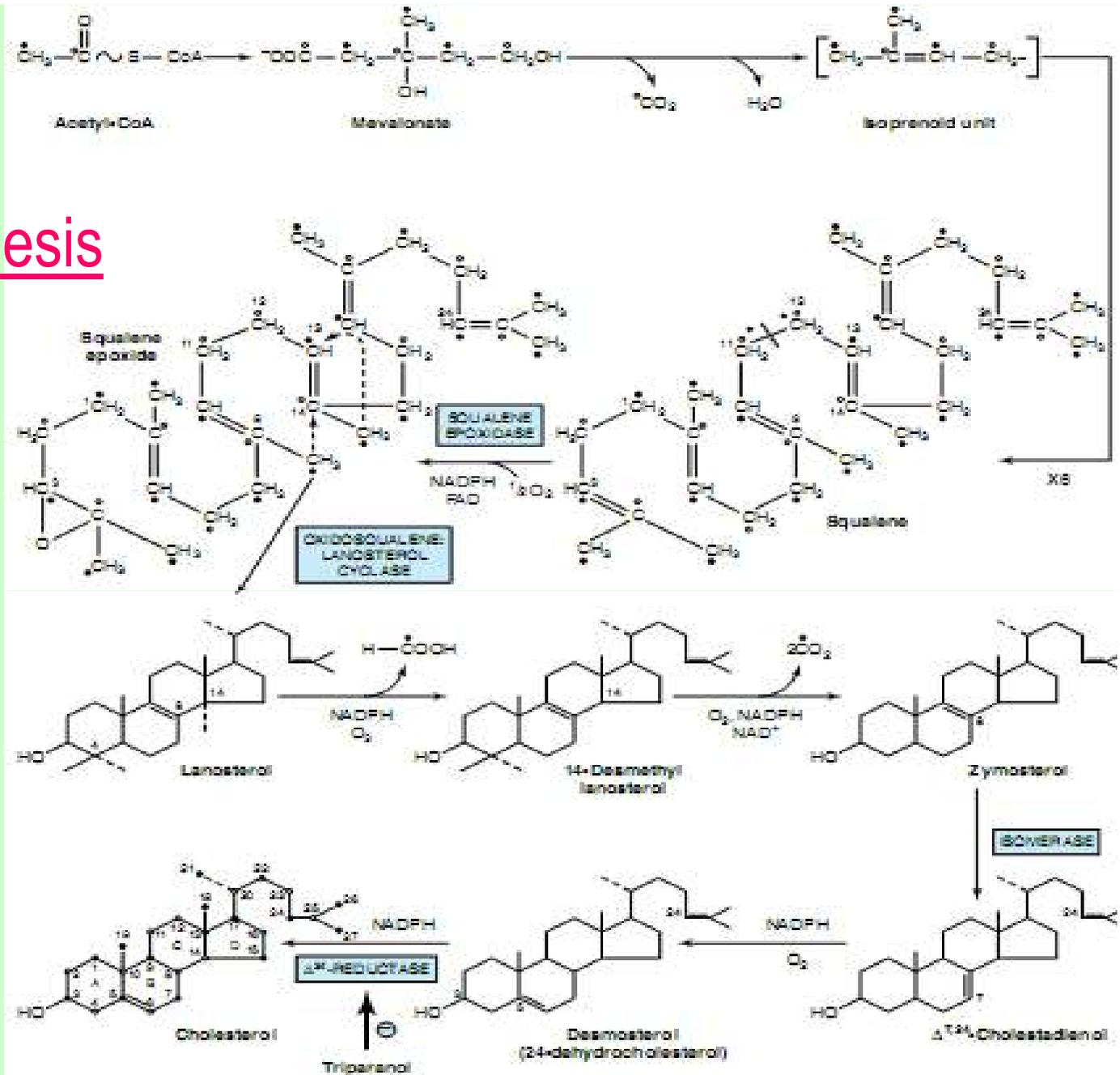


Figure 26-3. Biosynthesis of cholesterol. The numbered positions are those of the steroid nucleus and the open and solid circles indicate the fate of each of the carbons in the acetyl moiety of acetyl-CoA. Asterisks: Refer to labeling of squalene in Figure 26-2.

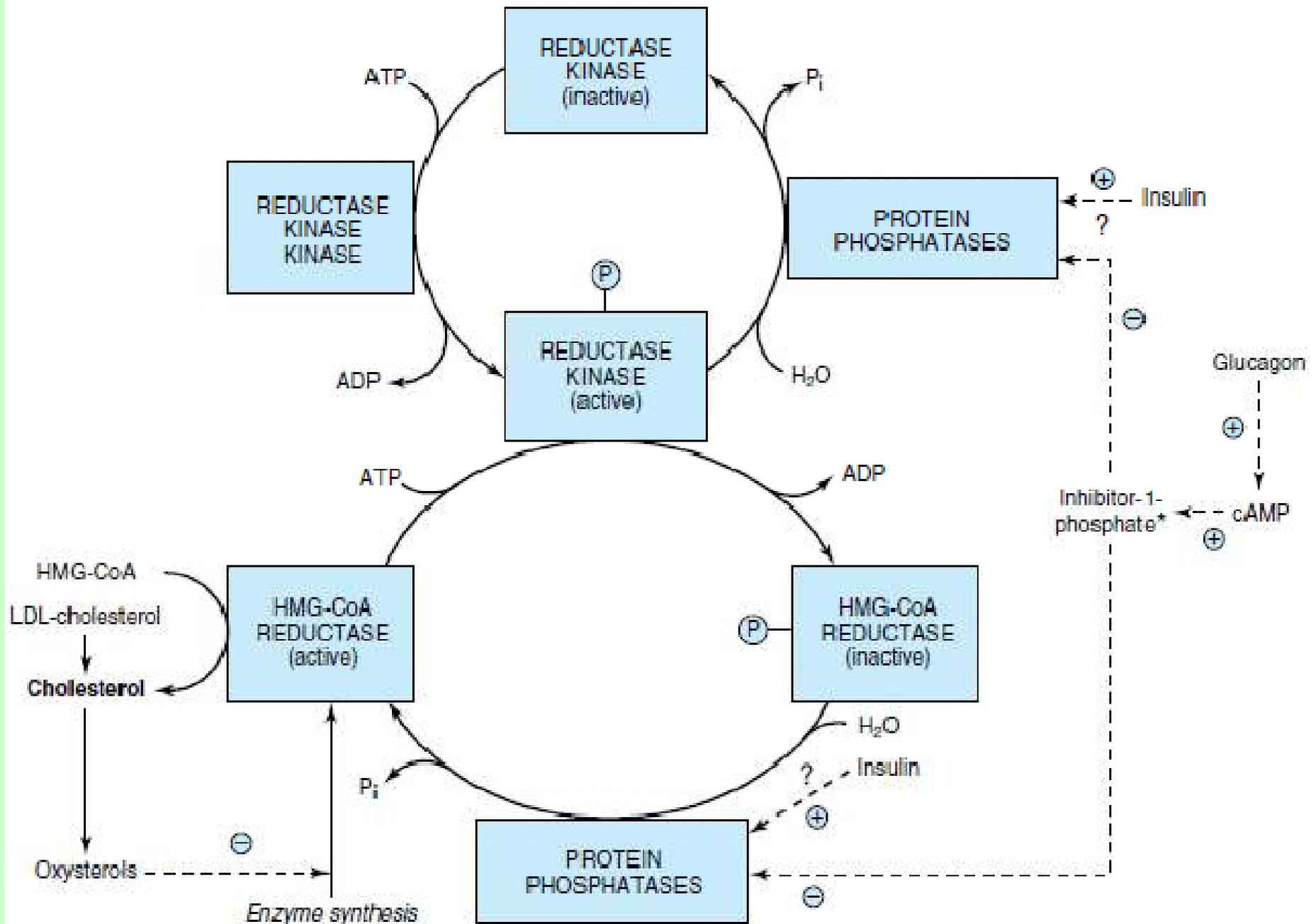


Figure 26-4. Possible mechanisms in the regulation of cholesterol synthesis by HMG-CoA reductase. Insulin has a dominant role compared with glucagon. Asterisk: See Figure 18-6.

HOMEWORK: Questions to be answered

1. Represent by structural formulas the reaction step of cholesterol biosynthesis catalyzed by the key regulatory enzyme of the pathway! Name the key enzyme!
2. Briefly explain the levels and ways of regulation of the key regulatory enzyme of cholesterol biosynthesis!
3. Which metabolically important biomolecules are synthesized from the cholesterol as precursor molecule?