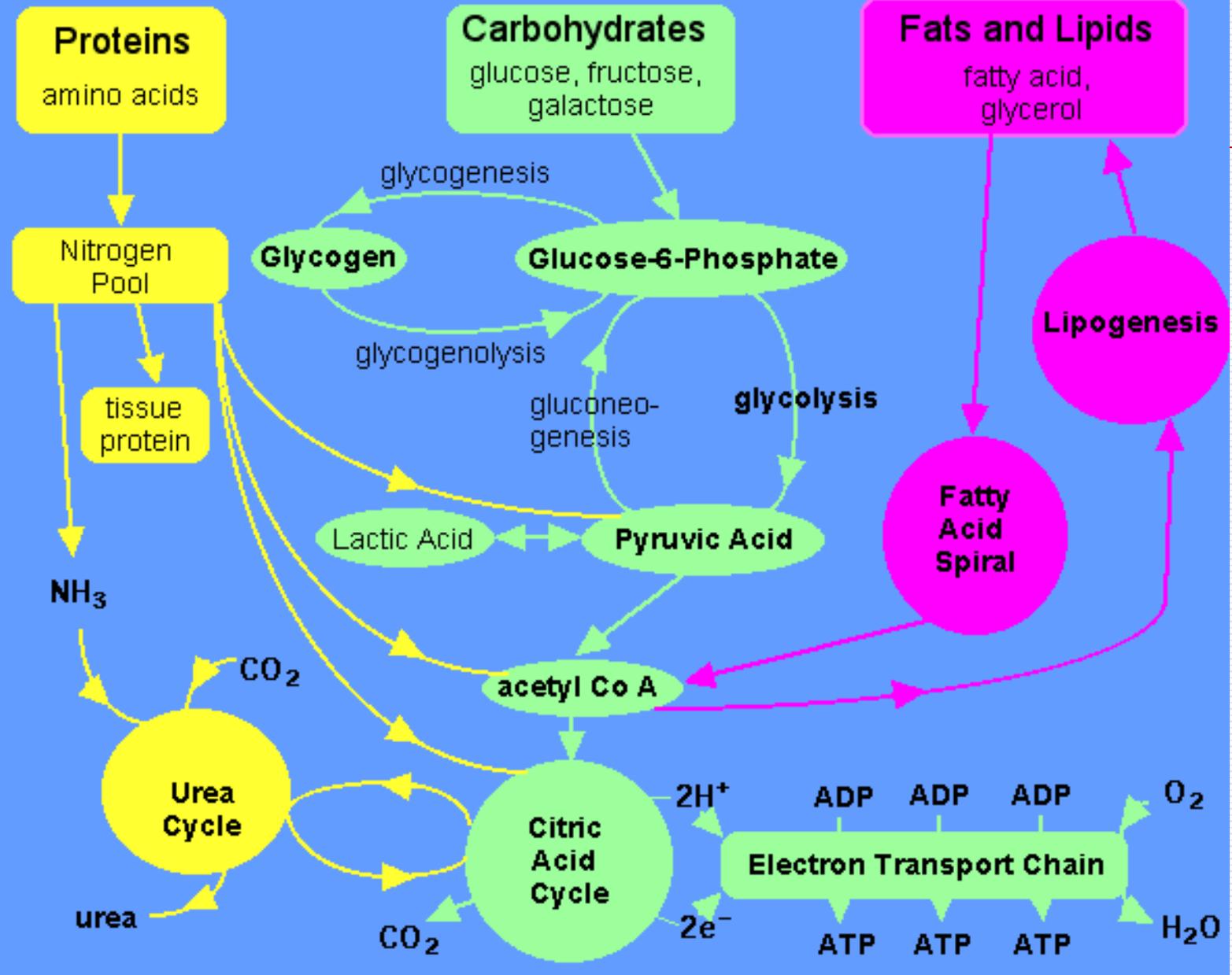


Lipids Metabolism

VLA 2019

2. 4. 2019

Metabolism Summary



Metabolism stages

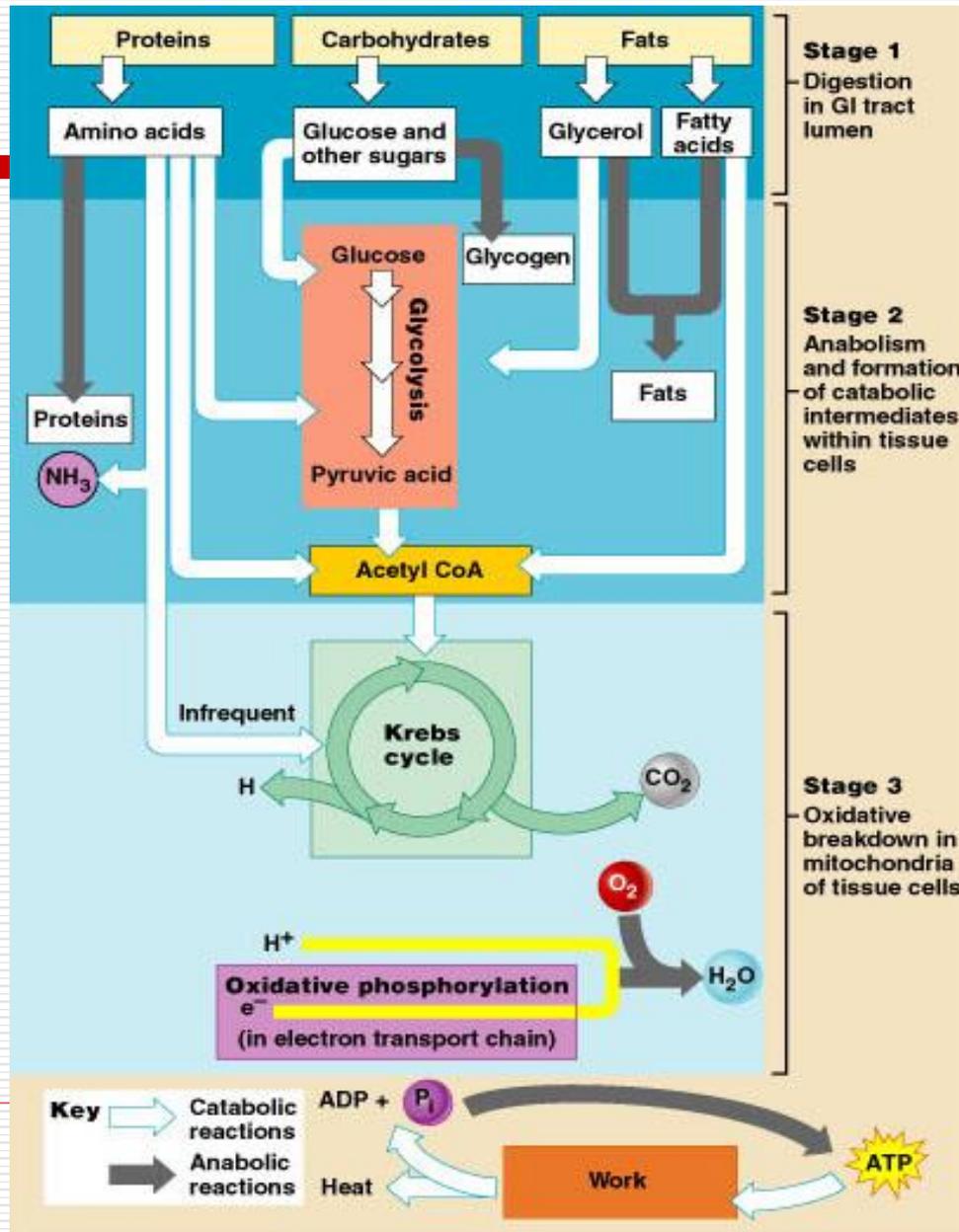
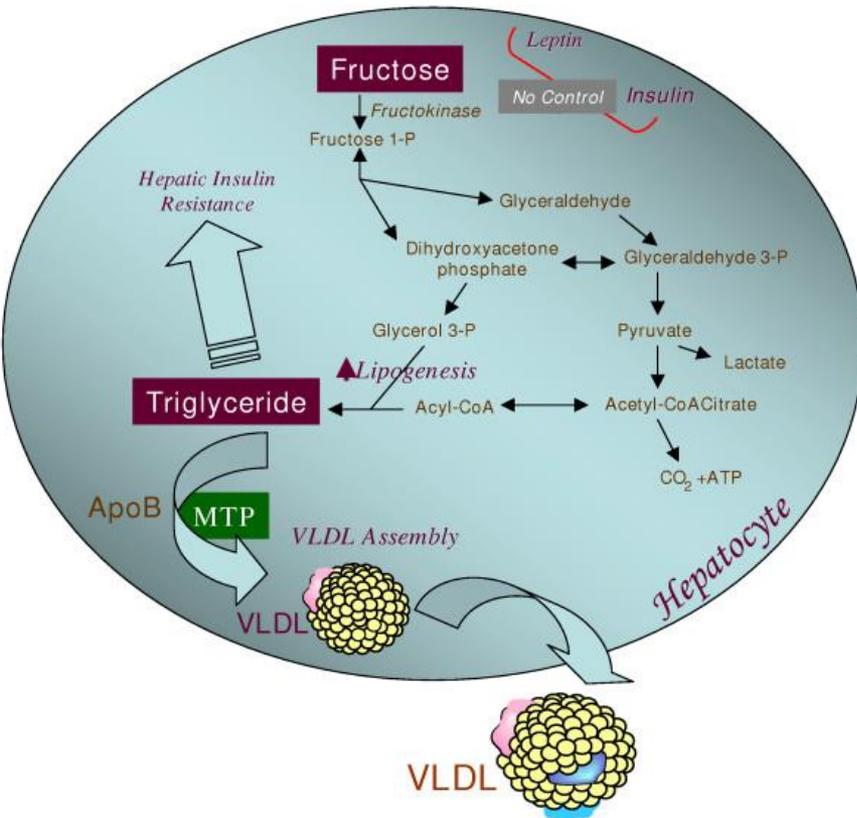


Figure 24.3

Complete oxidation – energy expenditure zisk

- Fatty acids: 9 kcal/g
 - Sugars: 4 kcal/g
 - Proteins: 4 kcal/g
-



Hepatic fructose metabolism: A

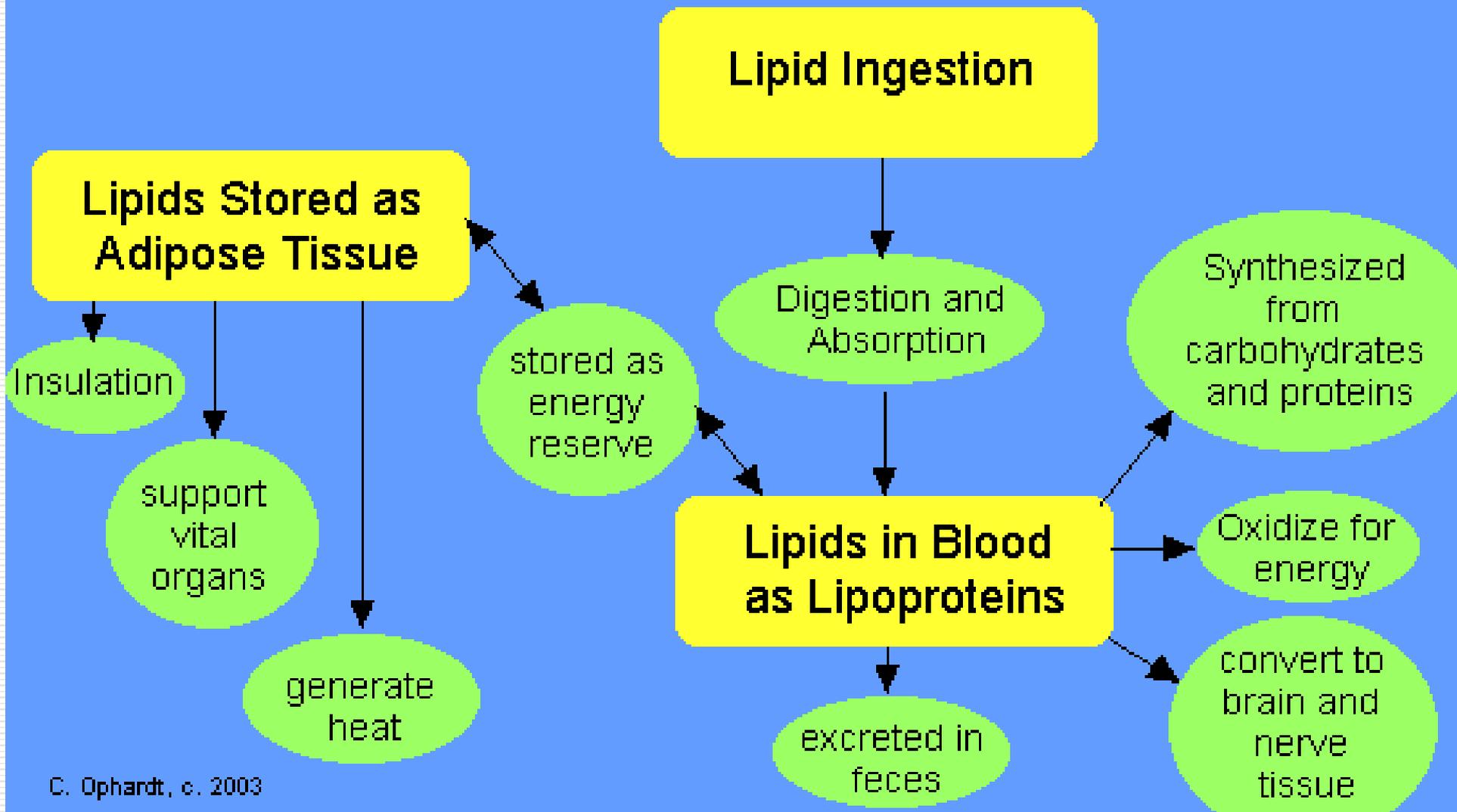
highly lipogenic pathway. Fructose is readily absorbed from the diet and rapidly metabolized principally in the liver.

Fructose can provide carbon atoms for both the glycerol and the acyl portions of triglyceride. **Fructose is thus a highly**

efficient inducer of *de novo* lipogenesis.

High concentrations of fructose can serve as a relatively unregulated source of acetyl CoA. In contrast to glucose, dietary fructose does NOT stimulate insulin or leptin (which are both important regulators of energy intake and body adiposity). Stimulated triglyceride synthesis is likely to lead to hepatic accumulation of triglyceride, which has been shown to reduce hepatic insulin sensitivity, as well as increased formation of VLDL particles due to higher substrate availability, increased apoB stability, and higher **MTP (microsomal triglyceride transfer protein)**, the critical factor in VLDL assembly.

Lipid Function and Metabolism Summary

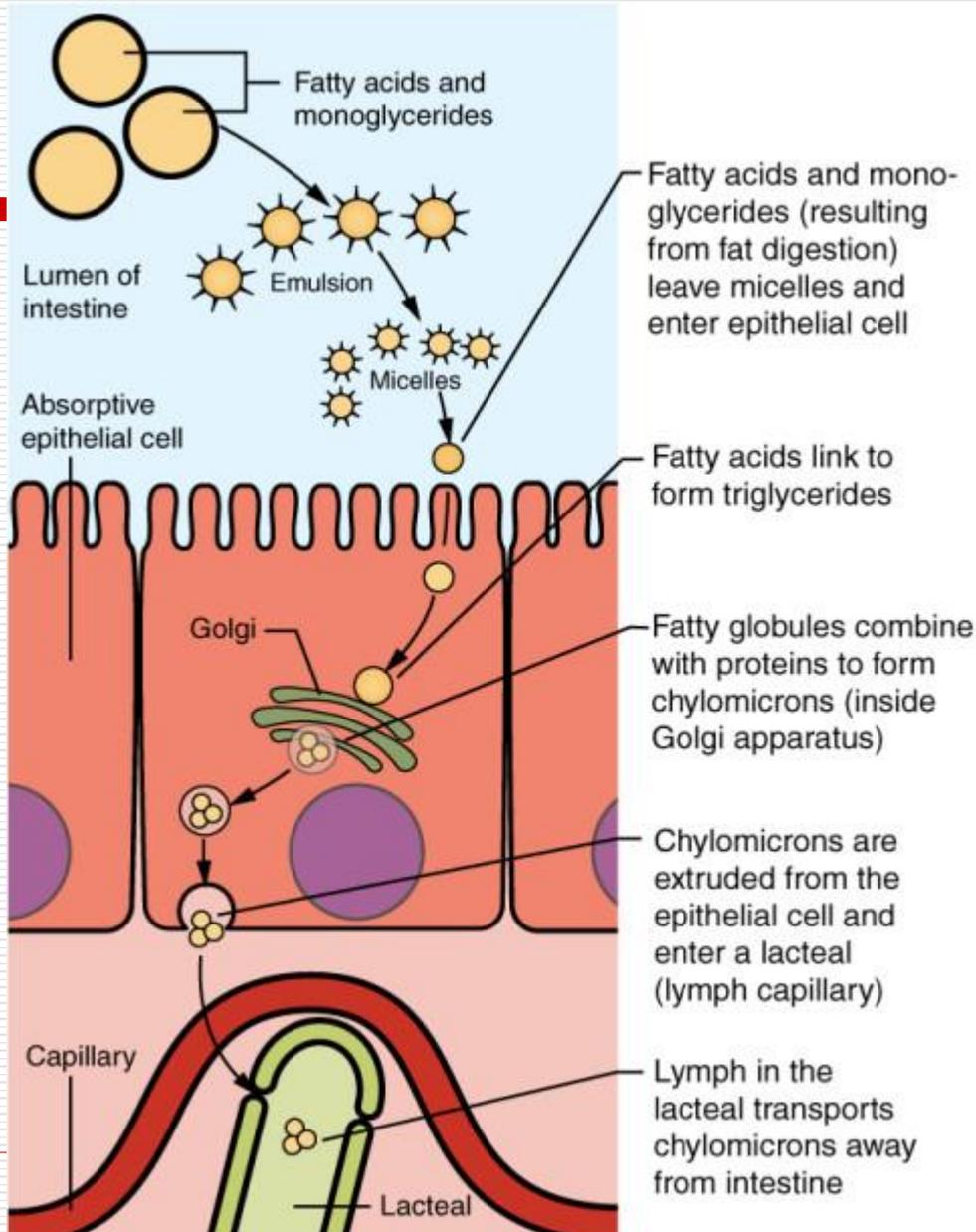


Lipids as a energy reserve

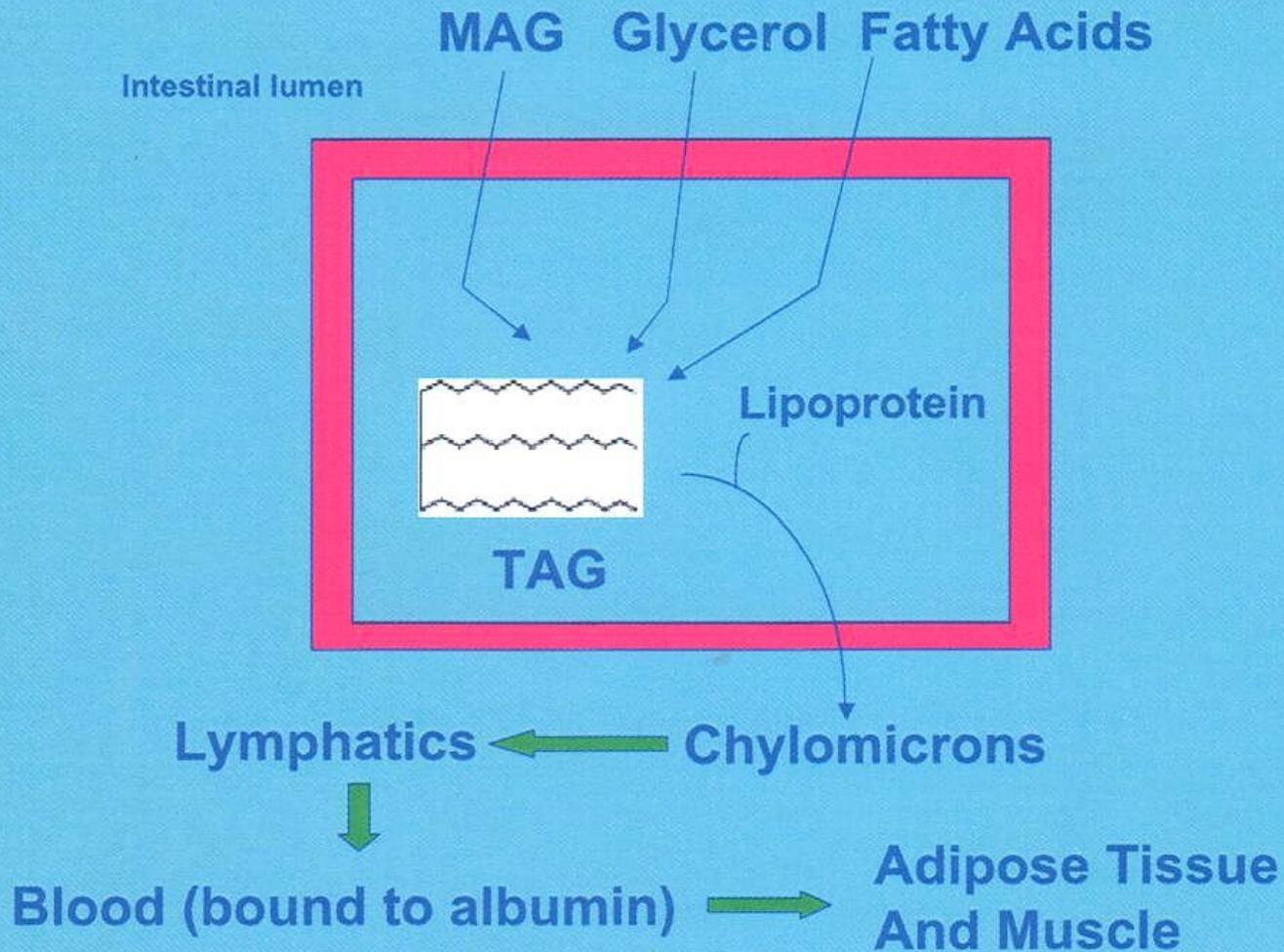
- ❑ Most of body energy is formed by oxidation of sugars and lipids.
 - ❑ Sugars: quick source of energy
 - ❑ Lipids: energy reserve
 - ❑ Energy reserve of lipids is much higher compared to glycogen reserve
-

Lipids metabolism

- Most of lipids metabolism products is transported to lymph as chylomicrons.
 - Lipids in chylomicrons are hydrolysed by plasma enzymes and absorbed by cells.
 - For energy formation only neutral lipids are oxidized
 - Lipids catabolism includes two distinct pathways:
 - Glycerol pathway
 - Pathway of fatty acids
-



Epithelial Cell (Intestinal Wall)



Fat Storage

- **Mainly as triacylglycerols (triglycerides) in adipose cells**
- **Constitute 84% of stored energy**
 - Protein - 15%
 - Carbohydrate (glucose or glycogen) - <1%

Processing of Lipid Reserves: Overview

1. **Lipid Mobilization:**
In adipose tissue TAGs hydrolyzed to fatty acids plus glycerol
2. **Transport of Fatty Acids in Blood To Tissues**
3. **Activation of Fatty Acids as CoA Ester**
4. **Transport into Mitochondria**
5. **Metabolism to Acetyl CoA**

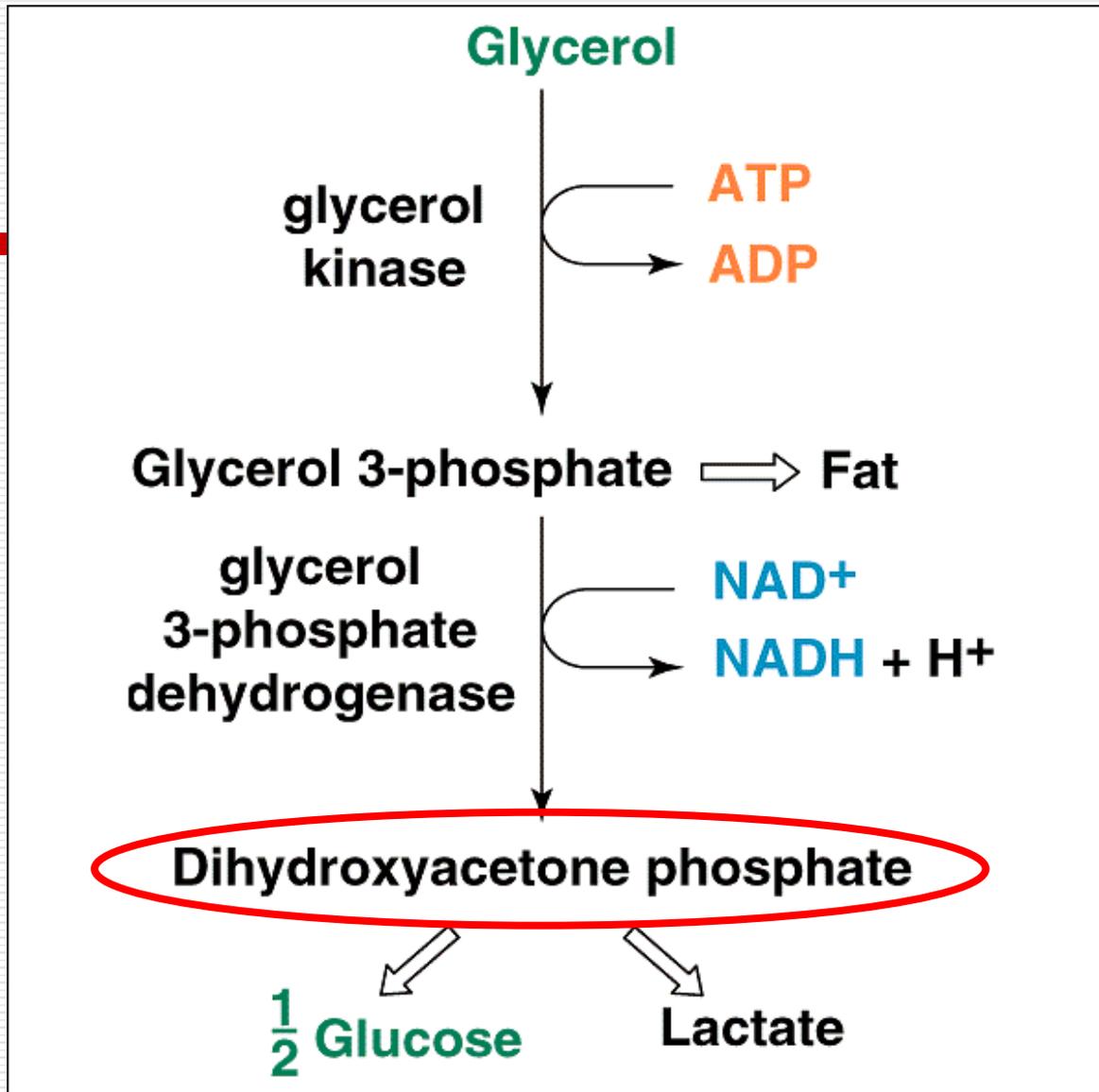
Subcutaneous [adipocyte](#) tissue (SCAT) and visceral adipocyte tissue (VAT)

- When compared to SCAT, VAT is more vascular, cellular and innervated. It contains a greater number of [immune cells](#) and large adipocytes when compared to that of SCAT. There are also more androgen and [glucocorticoid receptors](#) in VAT than in SCAT. Adipocytes in VAT are not only more metabolically active but also are more sensitive to [lipolysis](#) than adipocytes in SCAT. VAT also generates higher amounts of [free fatty acids](#), has an enhanced uptake of glucose and a higher sensitivity to [adrenergic stimulation](#), while SCAT is more involved in absorbing circulating free fatty acids and [triglycerides](#) (TGs).
- Overall, VAT accumulation increases the risk of [metabolic disorders](#), such as T2D, hypertension, [hyperlipidemia](#) and [atherosclerosis](#)

Due to fructose-induced inflammation (increased infiltration of adipose tissue by macrophages) there is also an increase in 11 β -hydroxysteroid dehydrogenase-1. This leads to an increase in intracellular [cortisol](#) in subcutaneous [adipocyte](#) tissue (essentially making them insulin resistant) causing less [fatty acids](#) to enter the subcutaneous adipocyte while more are expelled for storage into visceral depots and in and around organs, such as the liver, [skeletal muscle](#), heart, and pancreas, further disrupting metabolic processes and impairing organ function.

TOFI

- ❑ Thus, overconsumption of added sugars promotes “thin on the outside, fat on the inside” (TOFI).
- ❑ The term TOFI is used to describe lean individuals with an increased fat deposition within their abdomen (visceral adiposity).
- ❑ Subjects with TOFI have a [body mass index](#) < 25 kg/m² with an increase in risk factors associated with the [metabolic syndrome](#). This phenotype is a subtype of “metabolically-obese but normal-weight”.
- ❑ The prevalence of TOFI is uncertain but it is estimated that 14% of men and 12% of women have TOFI.
- ❑ TOFI can be diagnosed by [MRI](#) or [CT scan](#) which can help in differentiating fat (bright white) and other tissues (dark).



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Lipolysis

Hormone
(Adrenalin, Glucagon, ACTH)

Receptor (7TM)

Activates
Adenylyl
Cyclase
ATP → c-AMP

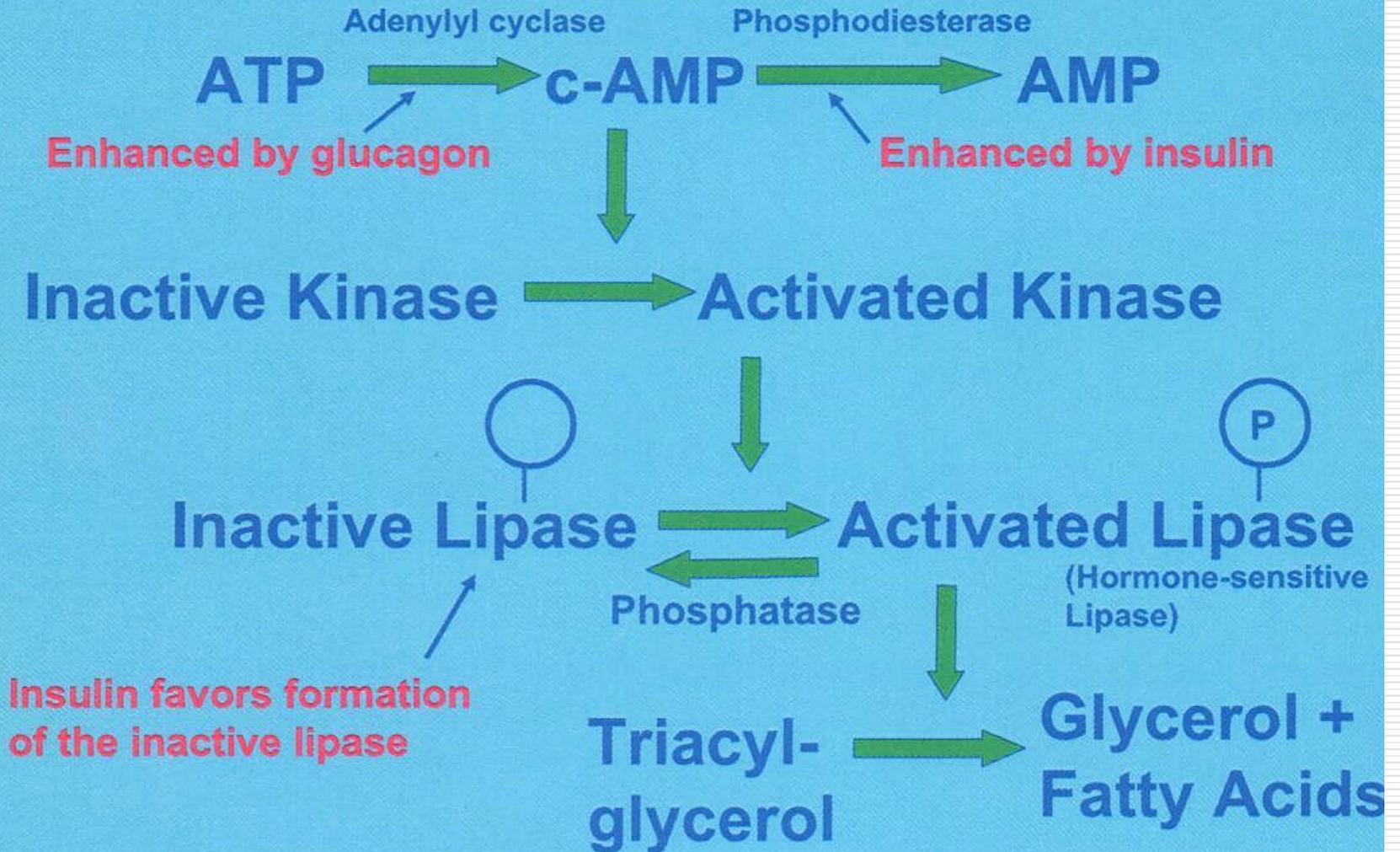
Activates lipase

Triacylglycerols → Glycerol +
Fatty acids

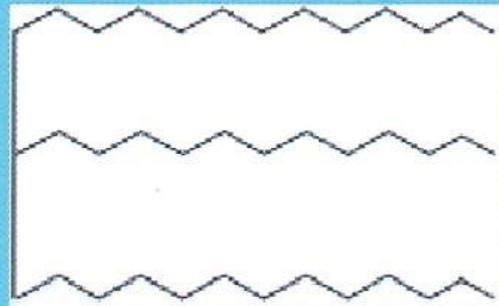
Blood

Adipose Cell

Insulin
blocks this
step



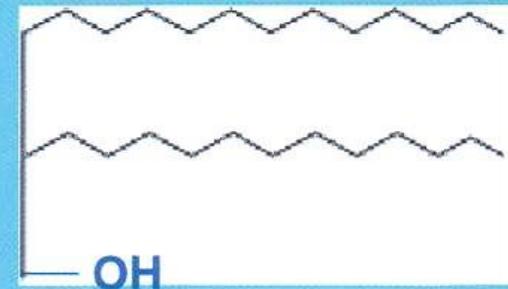
Acylglycerol Lipases



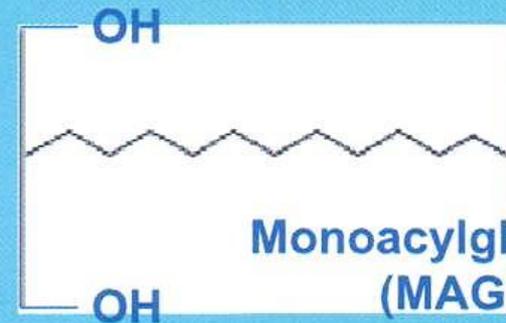
Triacylglycerol (TAG)

Triacylglycerol
Lipase

Diacylglycerol (DAG)

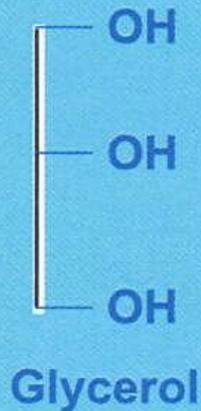


Diacylglycerol
Lipase



Monoacylglycerol
(MAG)

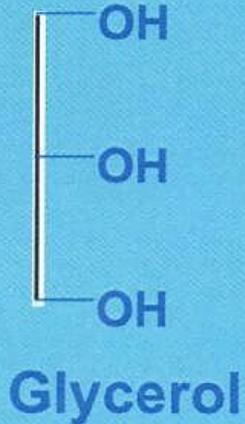
Monoacylglycerol
Lipase



Glycerol

Fate of Glycerol

In Liver:



Dihydroxyacetone
Phosphate

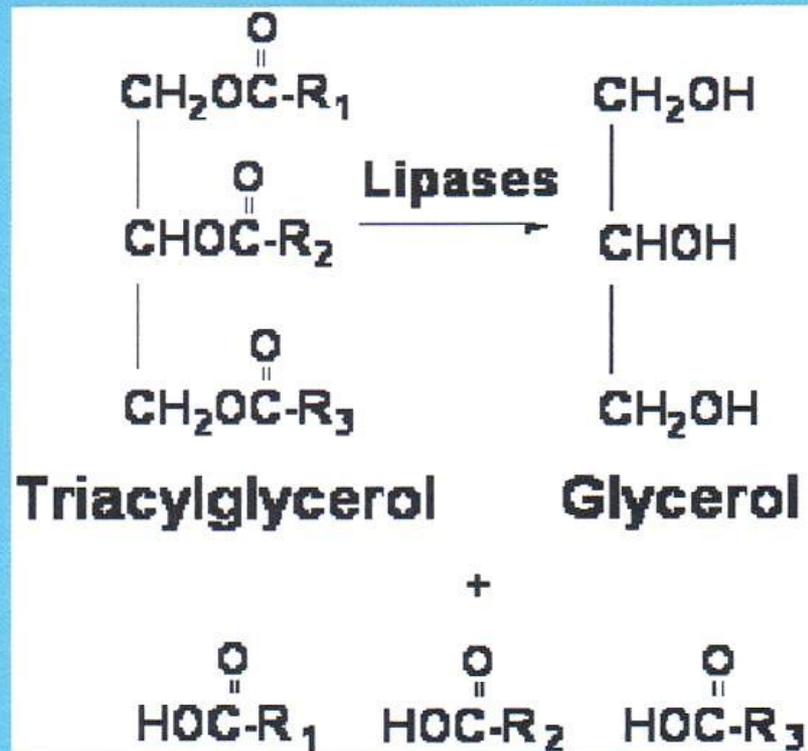
Glycolysis

Pyruvate

Gluconeogenesis

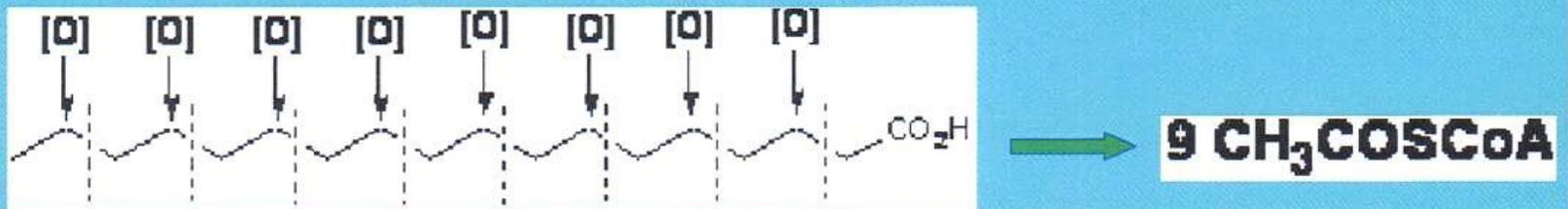
Glucose

Release of Fatty Acids from Triacylglycerols



Beta Oxidation

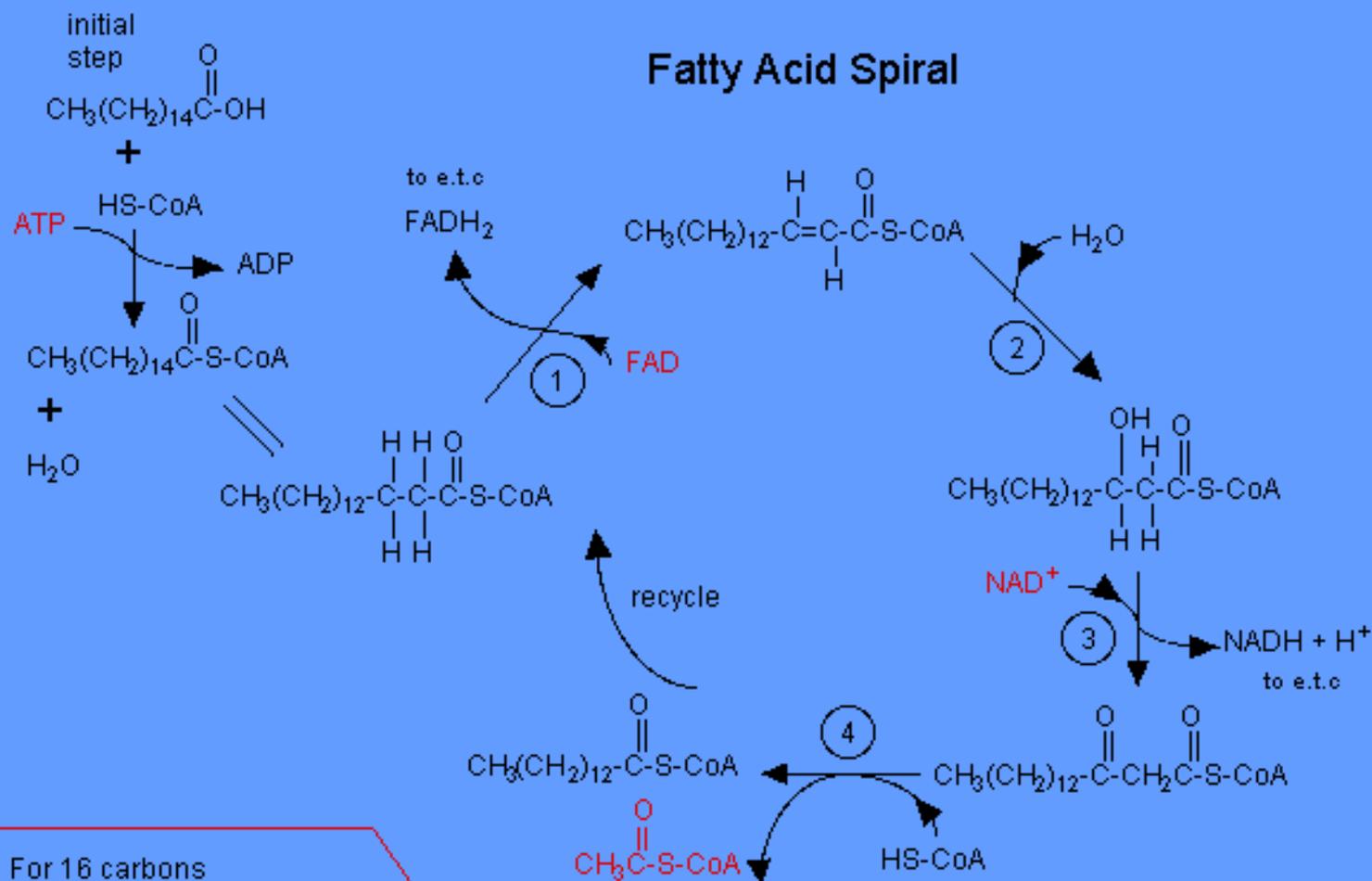
- Cleavage of fatty acids to acetate in tissues
- Occurs in mitochondria



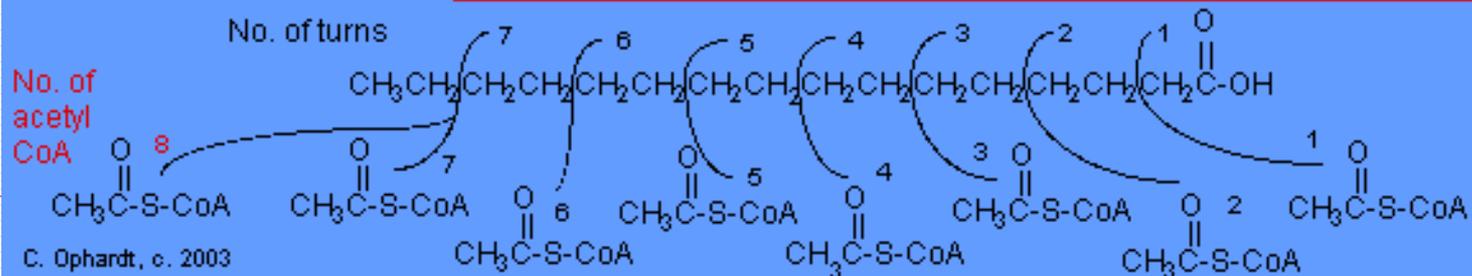
Steps in Beta Oxidation

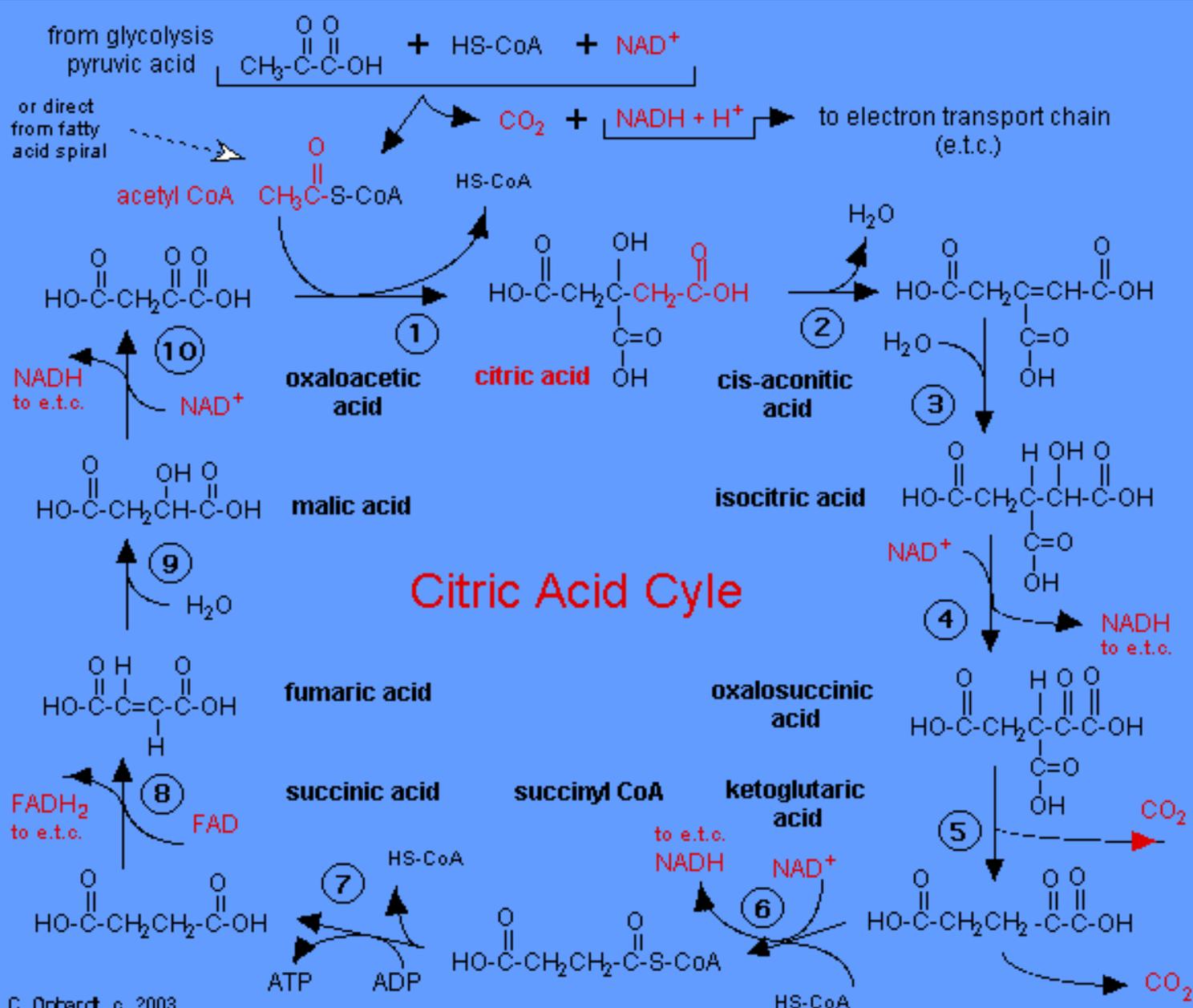
- **Fatty Acid Activation by Esterification with CoASH**
- **Membrane Transport of Fatty Acyl CoA Esters**
- **Carbon Backbone Reaction Sequence**
 - Dehydrogenation
 - Hydration
 - Dehydrogenation
 - Carbon-Carbon Cleavage (Thiolase Reaction)

Fatty Acid Spiral



For 16 carbons



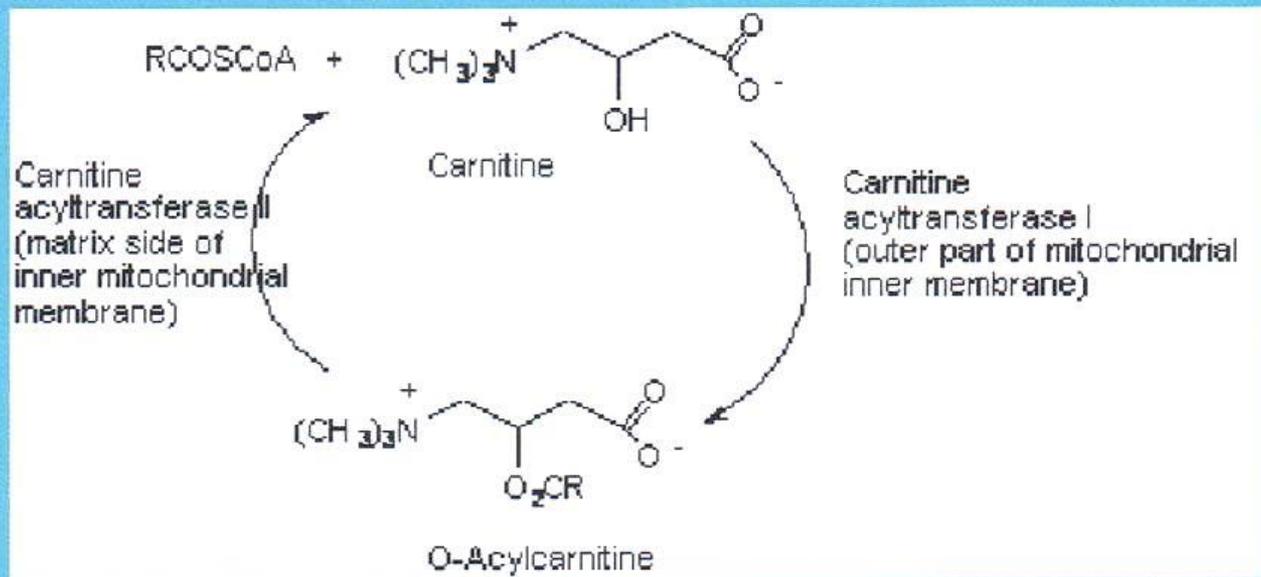


Citric Acid Cycle

Acetyl CoA

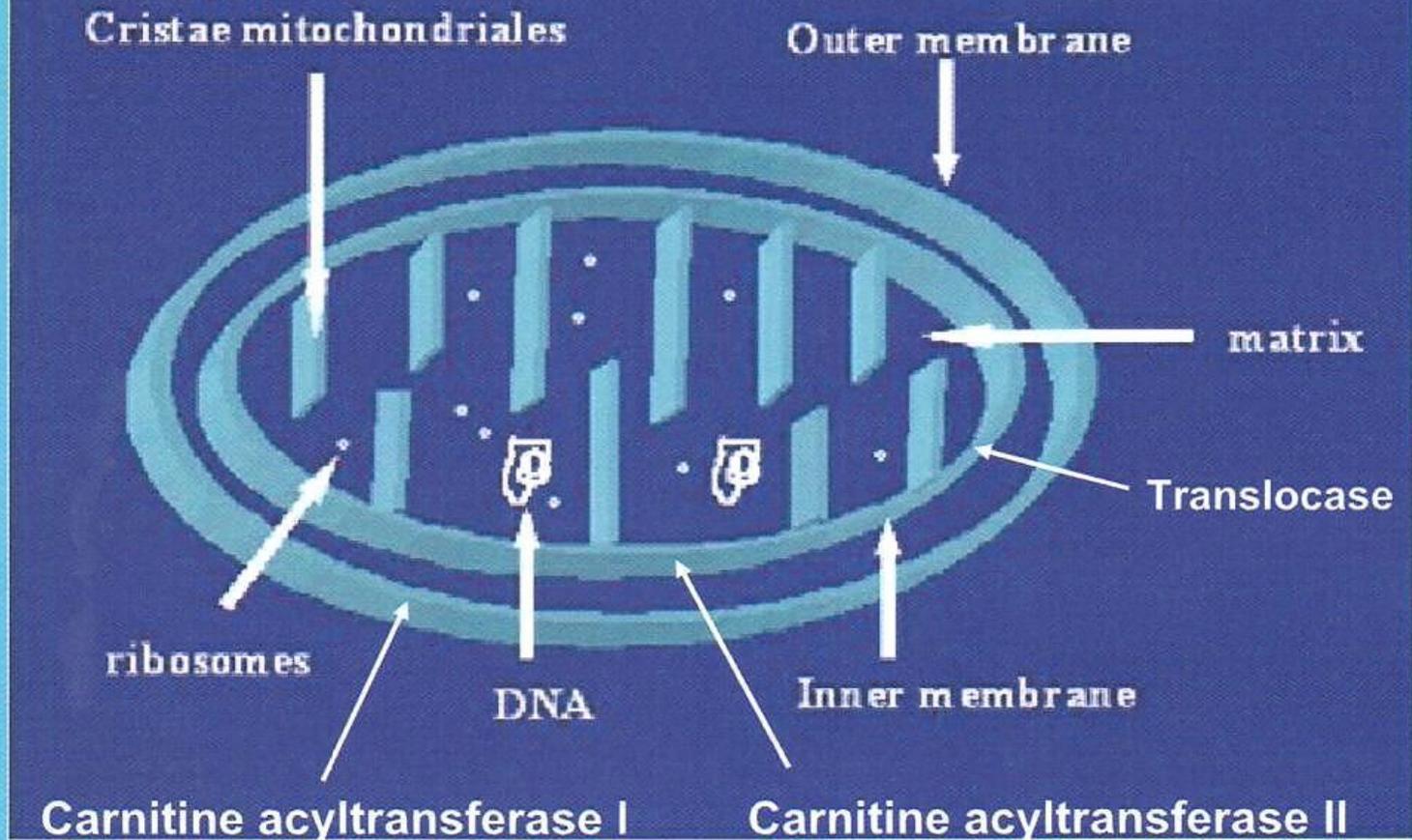
- Under aerobic conditions the end product of glycolysis is pyruvic acid. The next step is the formation of **acetyl coenzyme A** (acetyl CoA) - this step is technically not a part of the citric acid cycle, but is shown on the diagram on the top left.
 - Acetyl CoA, whether from glycolysis or the fatty acid spiral, is the initiator of the citric acid cycle. In carbohydrate metabolism, acetyl CoA is the link between glycolysis and the citric acid cycle.
 - The initiating step of the citric acid cycle occurs when a four carbon compound (oxaloacetic acid) condenses with acetyl CoA (2 carbons) to form citric acid (6 carbons).
 - The whole purpose of a "turn" of the citric acid cycle is to produce two carbon dioxide molecules. This general oxidation reaction is accompanied by the loss of hydrogen and electrons at four specific places. These oxidations are connected to the electron transport chain where many ATP are produced.
-

Membrane Transport of Fatty Acyl CoA Esters



↑
Transported across inner mitochondrial membrane by translocase

Mitochondrial Compartments



Ketone Bodies As Energy Sources

In liver

β -Hydroxybutyrate \longleftrightarrow Acetoacetate

Acetoacetate is major energy source in cardiac muscle and renal cortex; also in brain in starvation and diabetes

Succinyl CoA

β -Ketoacyl CoA transferase

Not found in liver

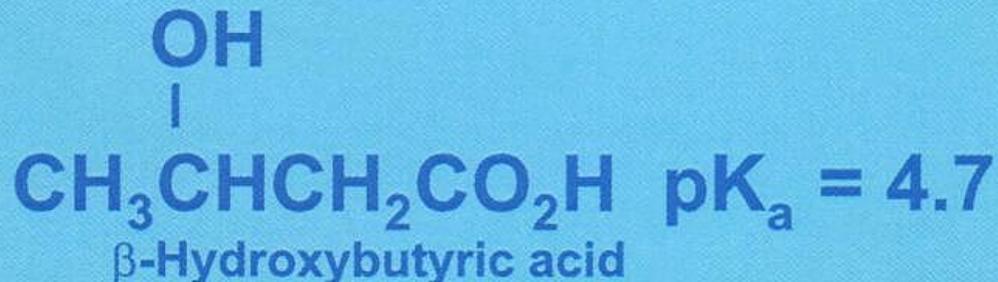
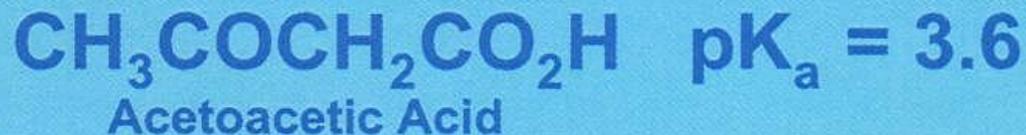
2 Acetyl CoA $\xrightleftharpoons{\text{Thiolase}}$ Acetoacetyl CoA

Succinate

Combines with oxaloacetate

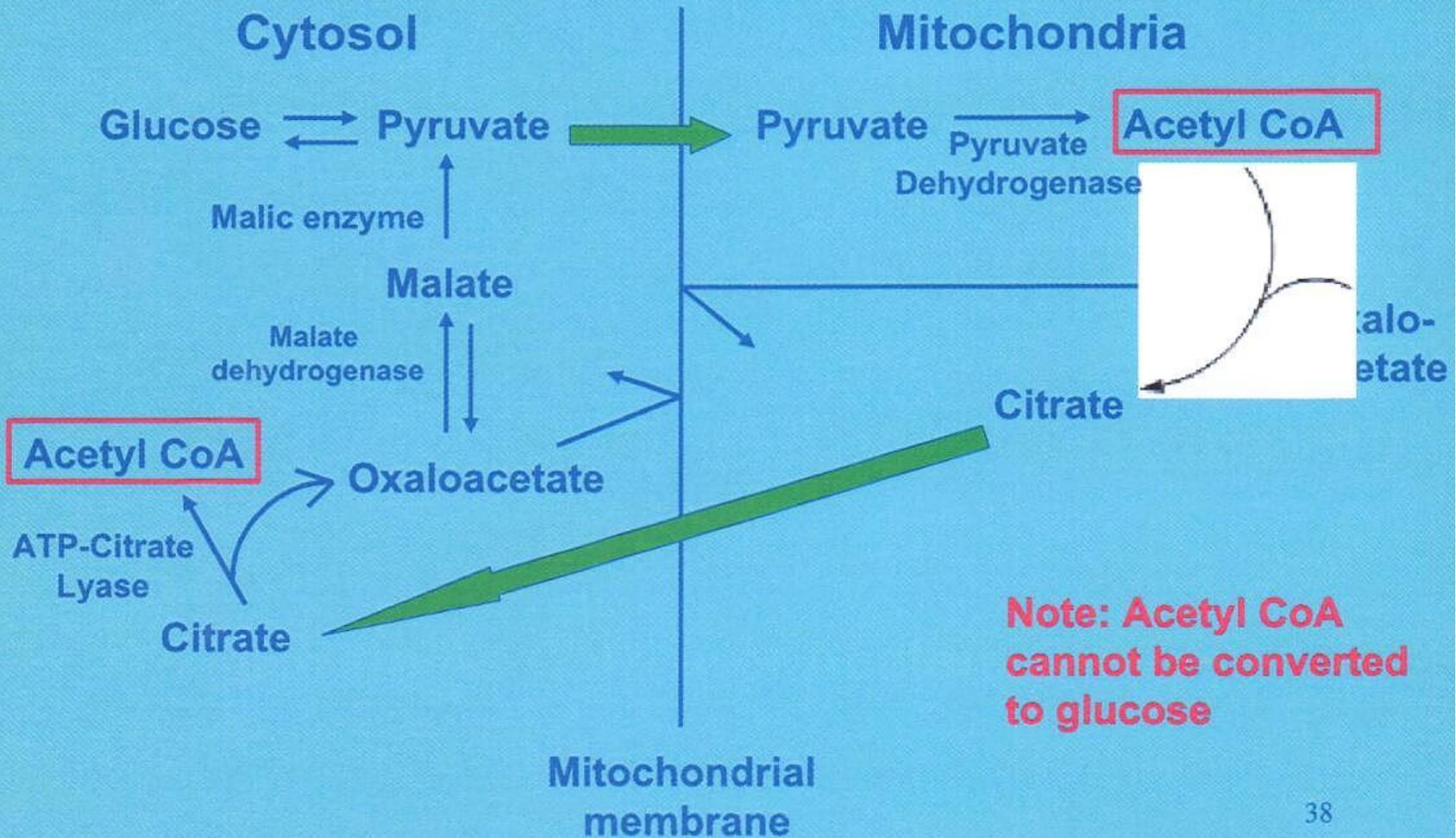
TCA Cycle

Metabolic Acidosis in Untreated Diabetes Mellitus



↑ Concentration of acetoacetic acid can result in metabolic acidosis (pH 7.1) → ↓ affinity of Hb for O₂.

Citrate As Carrier of Acetate Groups



Glycolysis

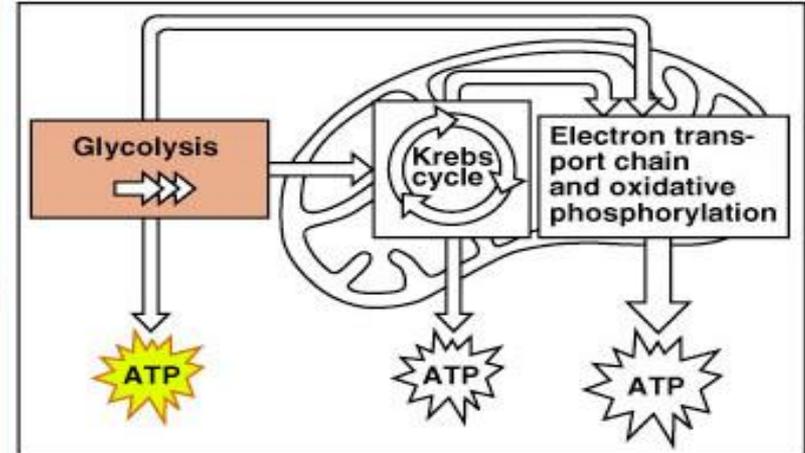
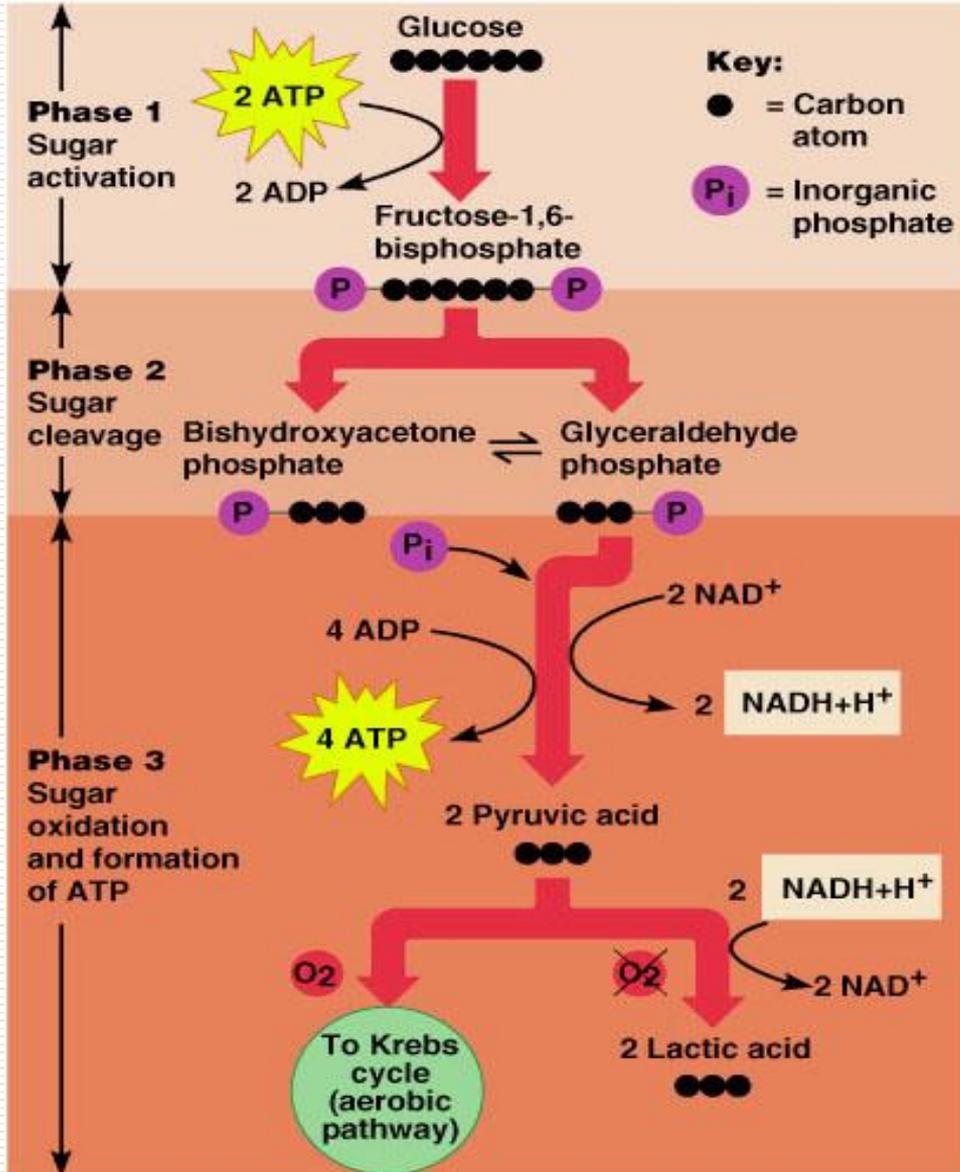
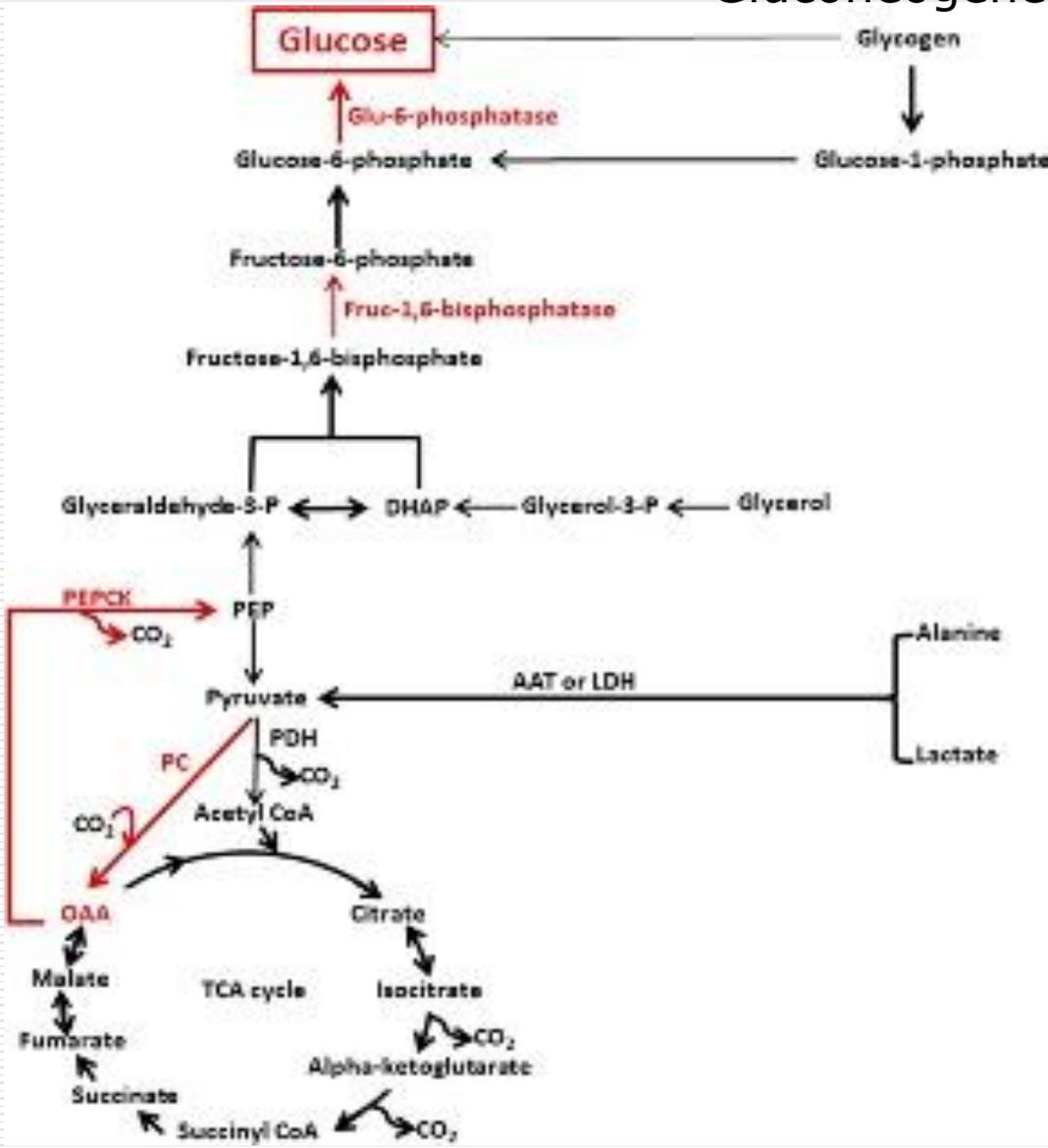


Figure 24.6

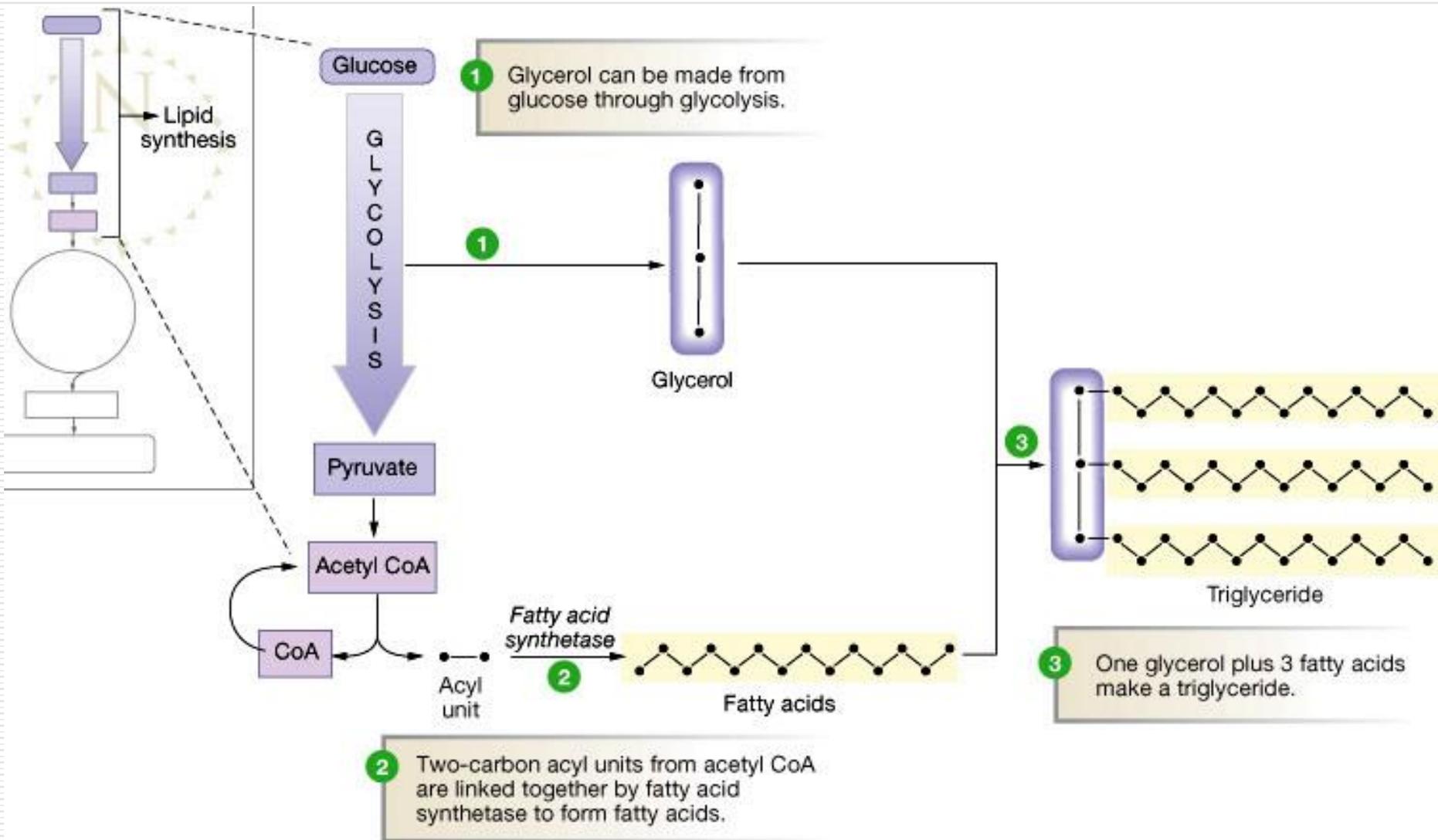
Gluconeogenesis



Major enzymes and substrates involved in the regulation of gluconeogenesis. Red arrows and text represent the major enzymes and pathways involved in the regulation of gluconeogenesis. Direct glucose release from glycogen via the debranching enzyme accounts for <10% of the total glucose made via gluconeogenesis.

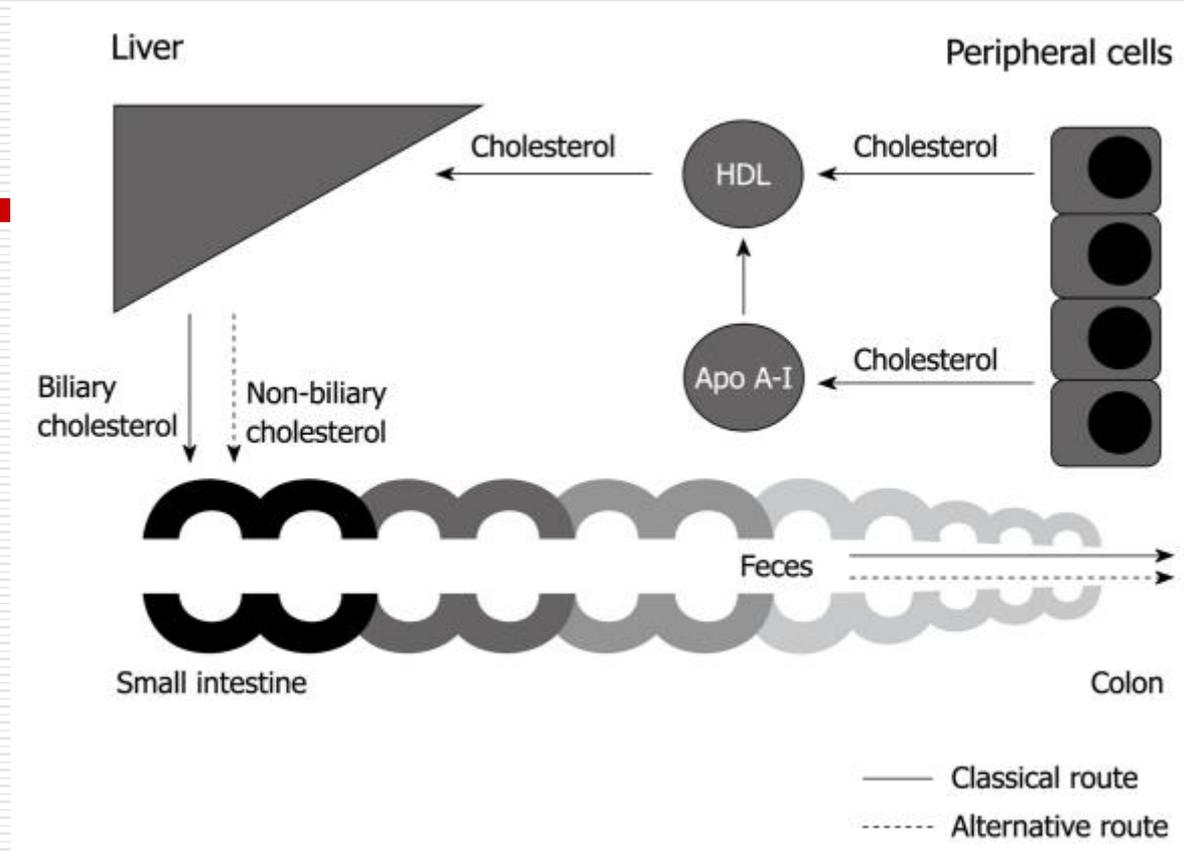
AAT, alanine aminotransferase;
 fruc-1,6-bisphosphatase, fructose-1,6-bisphosphatase;
 glu-6-phosphatase, glucose-6-phosphatase;
 glyceraldehyde-3-P, glyceraldehyde-3-phosphate;
 glycerol-3-P, glycerol-3-phosphate;
 LDH, lactate dehydrogenase;
 OAA, oxaloacetate;
 PC, pyruvate carboxylase;
 PDH, pyruvate dehydrogenase
 phosphoenolpyruvate carboxykinase (PEPCK)

Lipids Synthesis

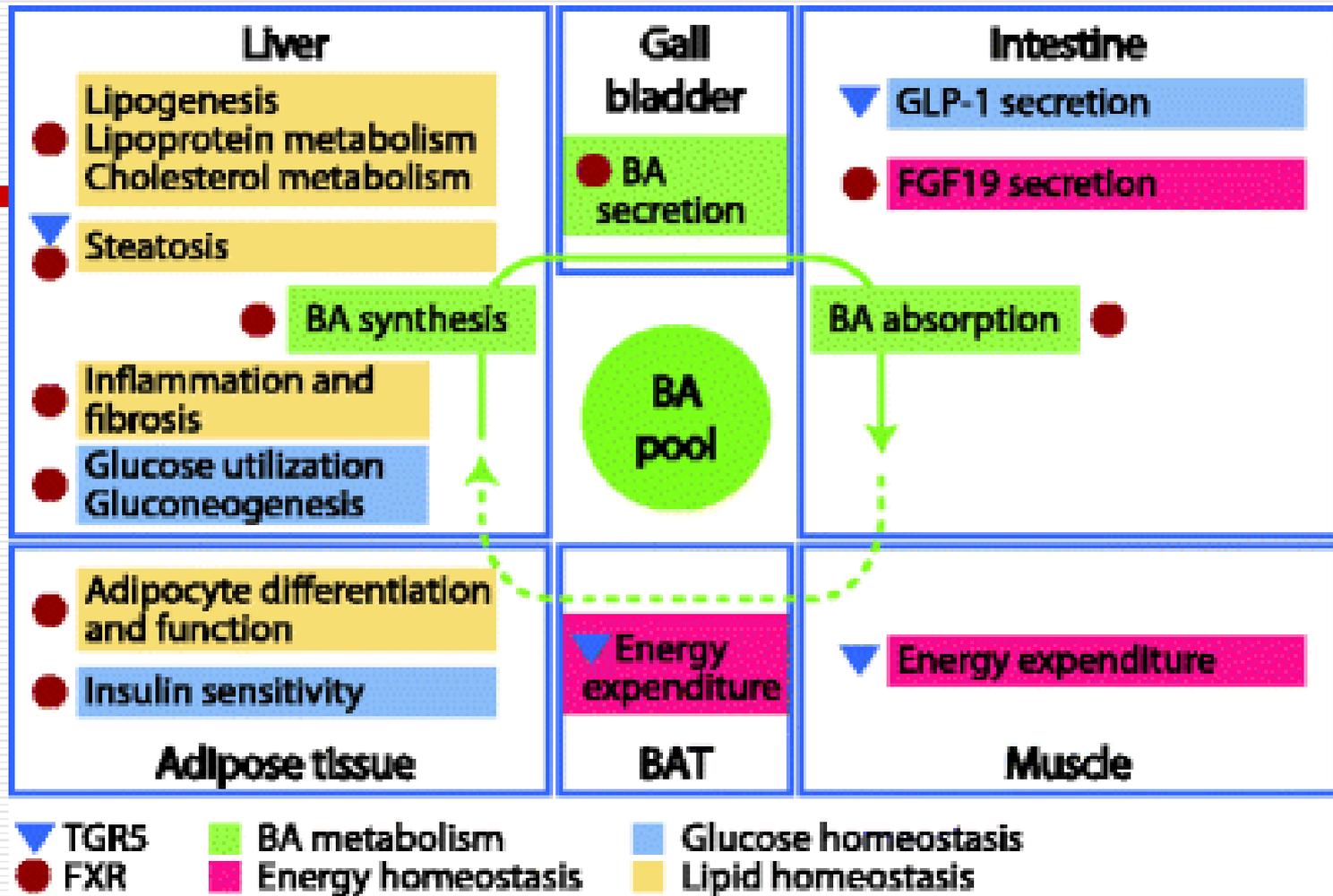


Functions of human plasma lipoproteins

Lipoprotein class	Origin	Function
Chylomicrons	Intestine	Transport lipids from intestine to liver and tissues
Very low density (VLDL)	Liver	Transport lipid from tissues to liver
Intermediate density (IDL)	VLDL	Precursor of LDL
High density (HDL 2 and 3)	Intestine	Remove cholesterol from tissues

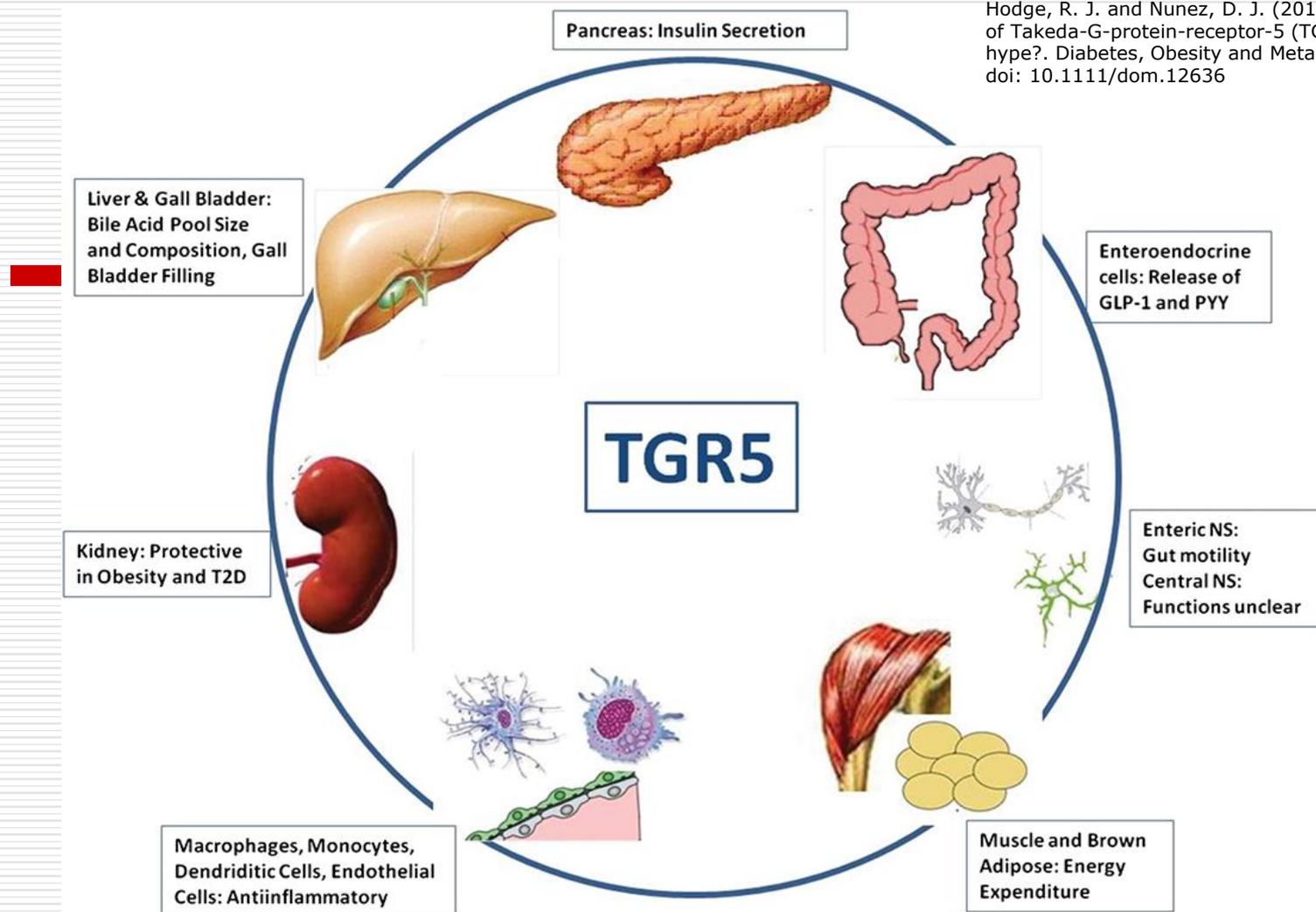


Cholesterol – the ways of excretion.
 Biliary and non-biliary cholesterol.



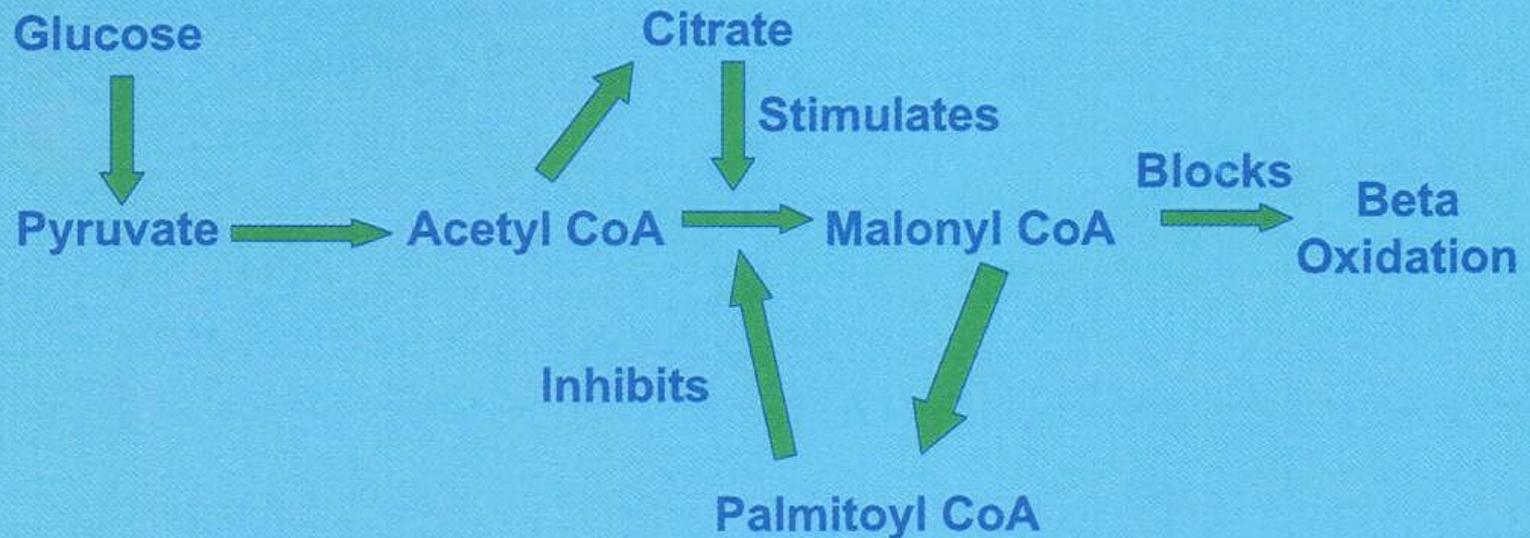
Functions of bile acids (BA) in regulation of BA, energy, glucose and lipid metabolism via farnesoid X receptor (FXR) and TGR5-mediated signaling pathways. BAT—brown adipose tissue; FGF—fibroblast growth factor; GLP-1—glucagon-like peptide 1

Hodge, R. J. and Nunez, D. J. (2016), Therapeutic potential of Takeda-G-protein-receptor-5 (TGR5) agonists. Hope or hype?. Diabetes, Obesity and Metabolism. doi: 10.1111/dom.12636



A wide range of Takeda-G-protein-receptor-5 (TGR5) effects. A variety of downstream effects has spawned intense interest in the therapeutic potential of TGR5 agonists for the treatment of metabolic and inflammatory diseases. GLP-1, glucagon-like peptide-1; NS, nervous system; PYY, peptide tyrosine tyrosine; T2D, type 2 diabetes.

Metabolite Regulation of Fatty Acid Synthesis and Breakdown



Tay-Sachs Disease

GM₂ (a ganglioside):

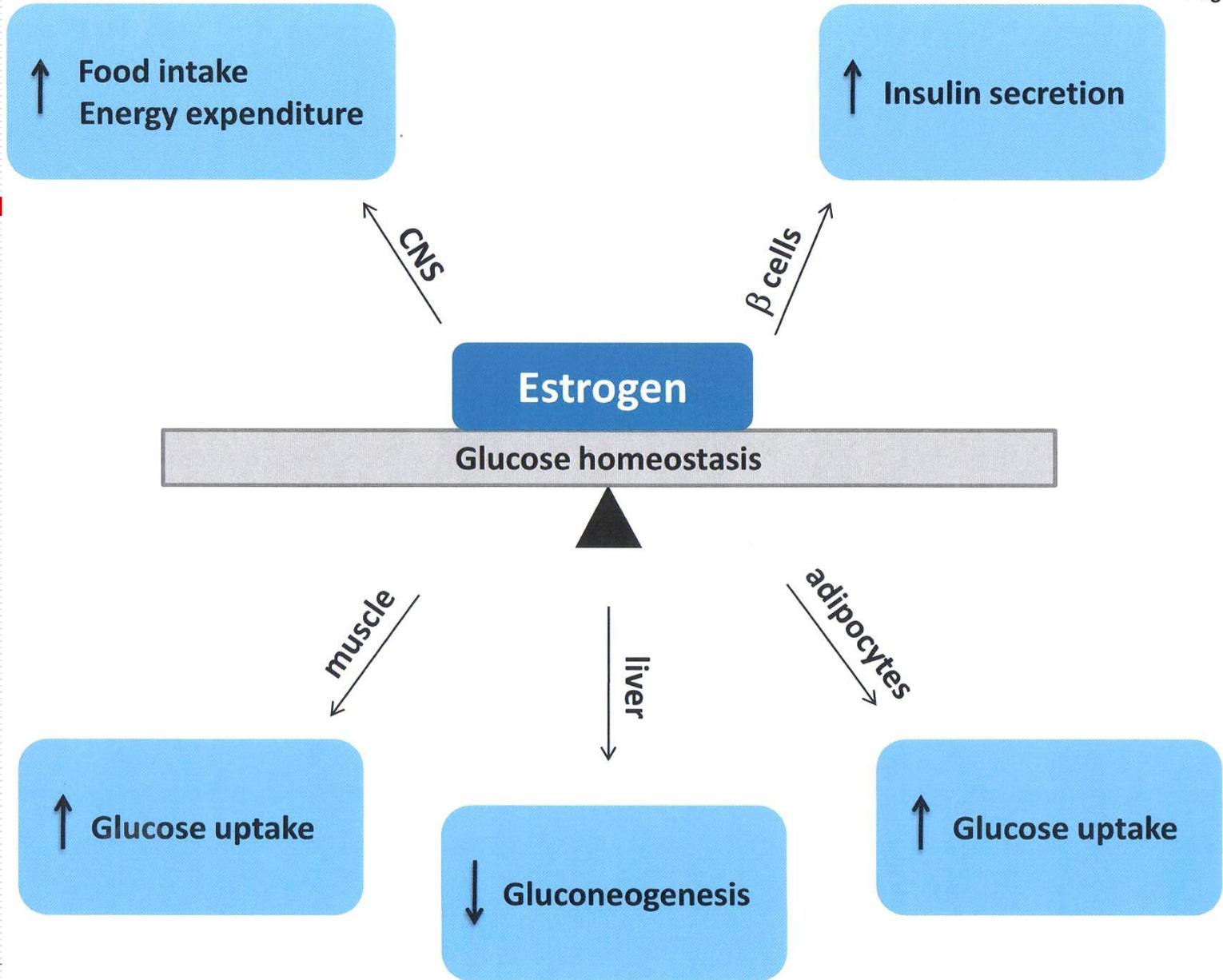
Ceramide - O - Glucose - Galactose - N-Acetylgalactose

↑
**Hexoseaminidase A
catalyzes cleavage of this
glycoside linkage**

Autosomal recessive disorder characterized by deficiency
of hexoseaminidase A; accumulation of gangliosides in brain
Most prevalent in Jews from Eastern Europe

For further information see:

http://www.marchofdimes.com/professionals/681_1227.asp



Estrogen deficiency



INSULIN RESISTANCE



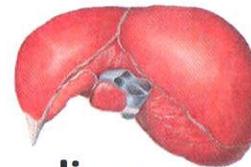
pancreatic β cells

Impaired insulin secretion



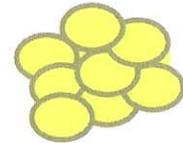
muscle

Impaired glucose uptake



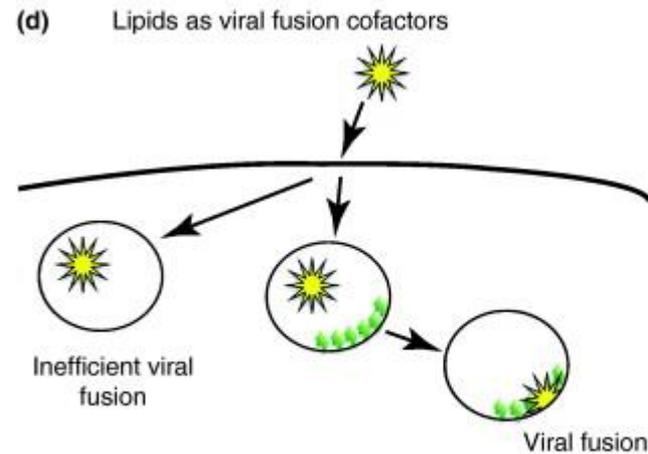
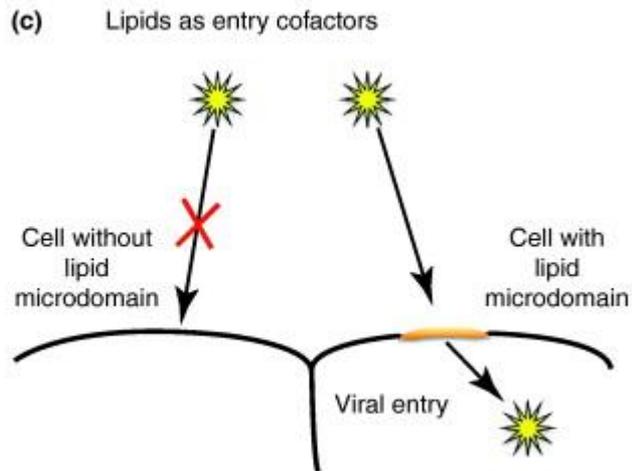
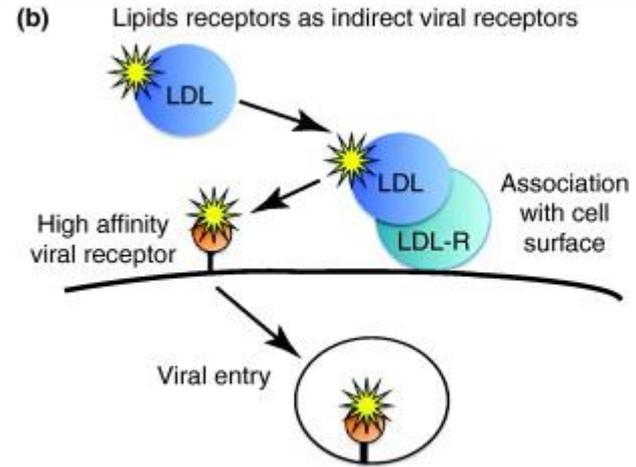
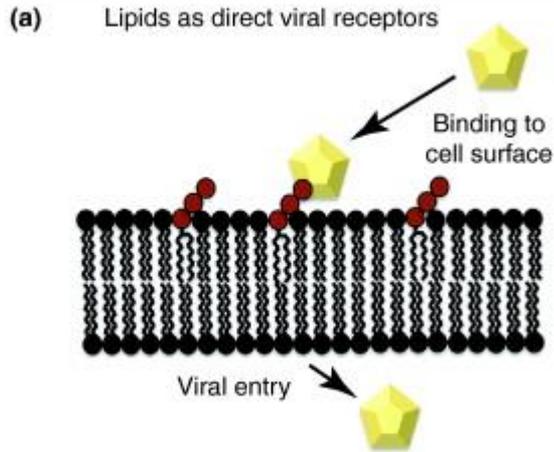
liver

Increased gluconeogenesis
Increased lipogenesis
TG accumulation
Increased VLDL production
Decreased insulin clearance



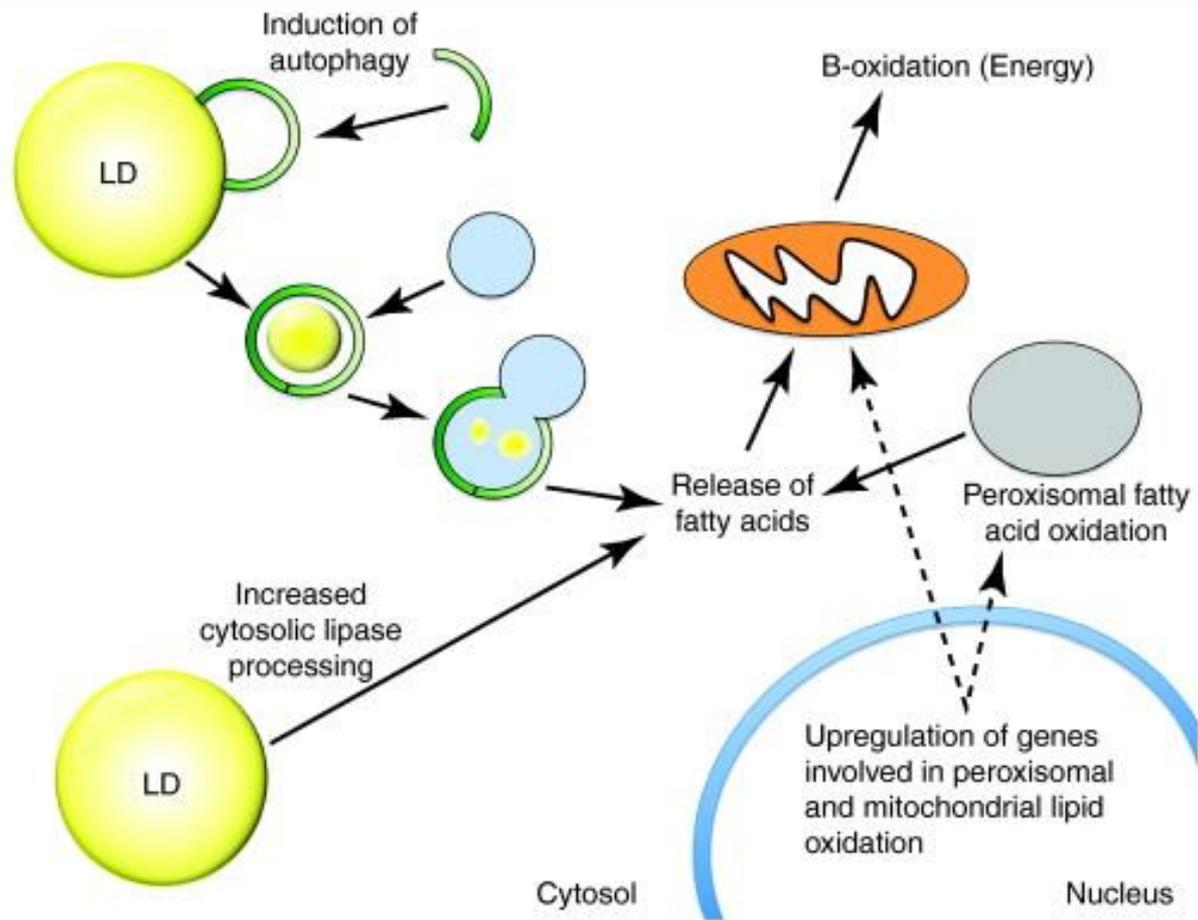
adipose tissue

Increased lipolysis
Increased adipocyte size
Inflammation



Lipid droplets

- Storage neutral lipids, i.e. triacylglycerols (TAG) and sterol esters (SE), are stored in the form of lipid droplets (LDs) in almost all eukaryotic cells.
 - LDs are dynamic subcellular organelles that not only govern the storage and turnover of lipids, but also function in membrane and lipid trafficking, protein storage and degradation, and even in the replication of hepatitis C virus.
 - All LDs comprise a core of storage neutral lipids which are wrapped by a monolayer of phospholipids with proteins embedded. LDs are believed to originate from the endoplasmic reticulum (ER), although the exact mechanism underlying their biogenesis remains to be determined.
-



Lipodystrophies

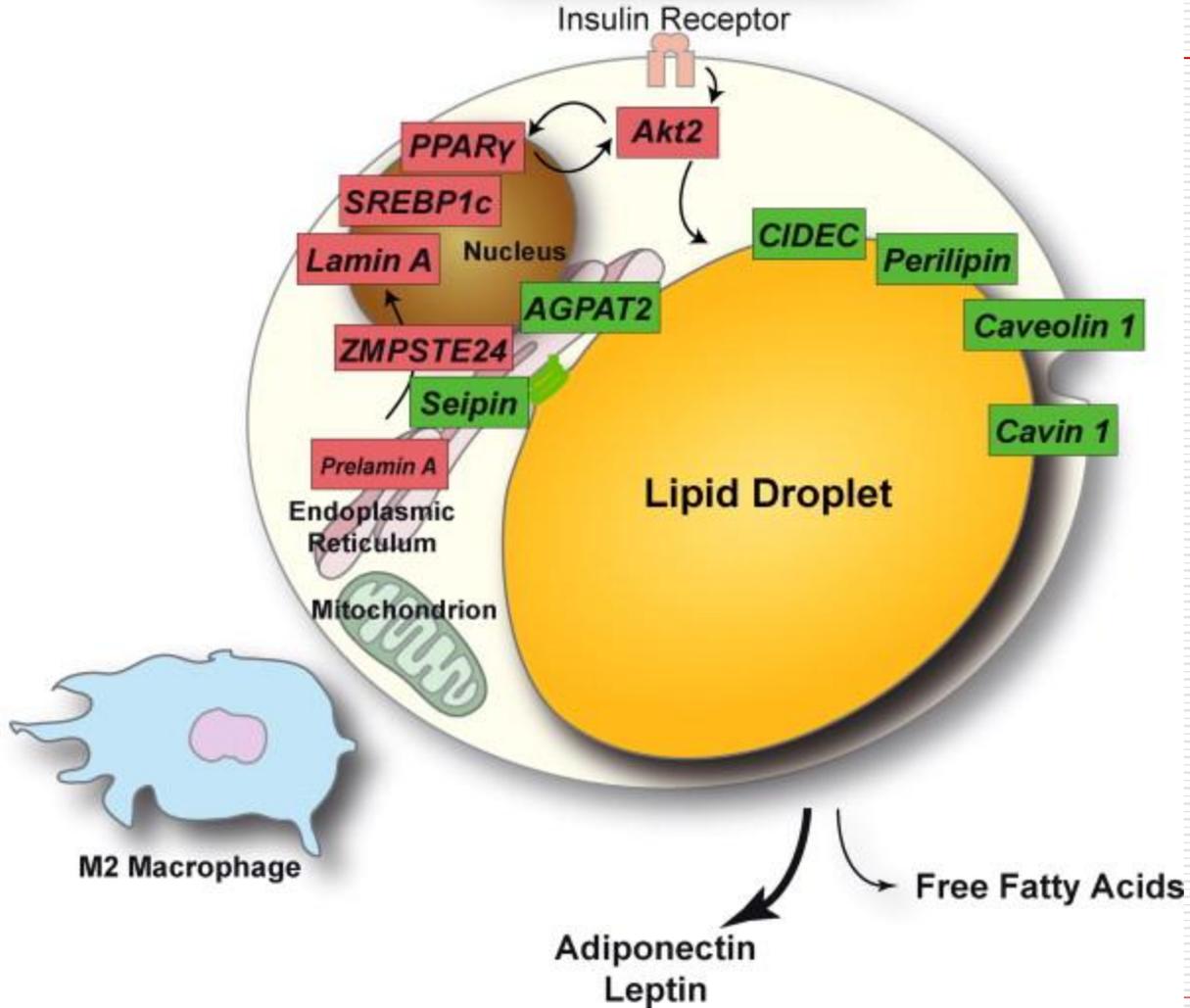
- ❑ Heterogenic group of diseases defined as **localised or generalised loss of body fat**.
 - ❑ If localised, usually related with **fat hypertrophy in other side of the body**.
 - ❑ Usually associated with sever metabolic changes including insulin resistance, dyslipidemia and glucose intolerance.

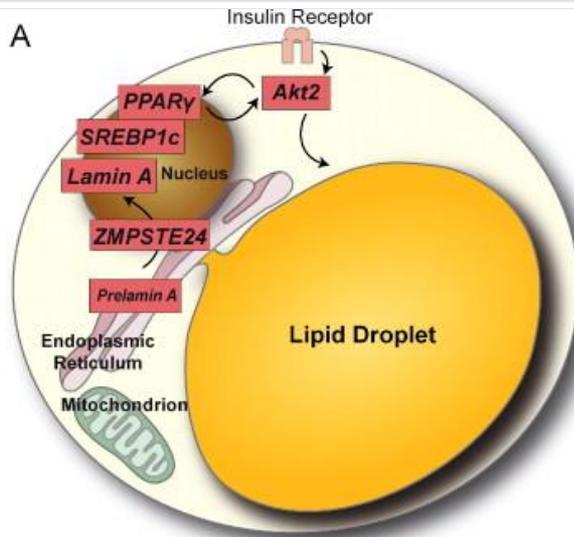
 - ❑ Different phenotypes:
 - ❑ **Familiar parcial lipodystrophy**, type Dunnigan (FPLD): fat reduction on the lower part of the body, hypertrophy on the upper part
 - ❑ **Barraquer-Simons** syndrome – reverse phenotype, milder metabolic changes

 - ❑ Problems on the level of:
 - ❑ adipogenesis, insulin sensitivity, TAGs storage, lipid droplets formation, oxidative stress and fat remodellation.
-

A

Healthy Adipose Tissue

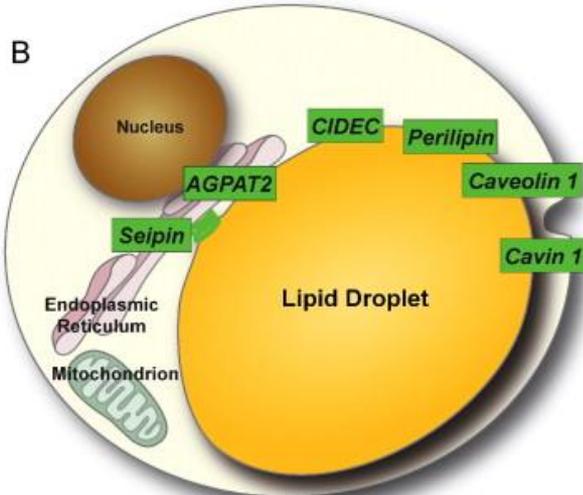




Adipogenesis and Insulin Signaling

Cellular targets altered by mutations in lipodystrophies

A: proteins taking part in **adipogenesis** at the level of nuclear DNA and in insulin signal transduction pathway

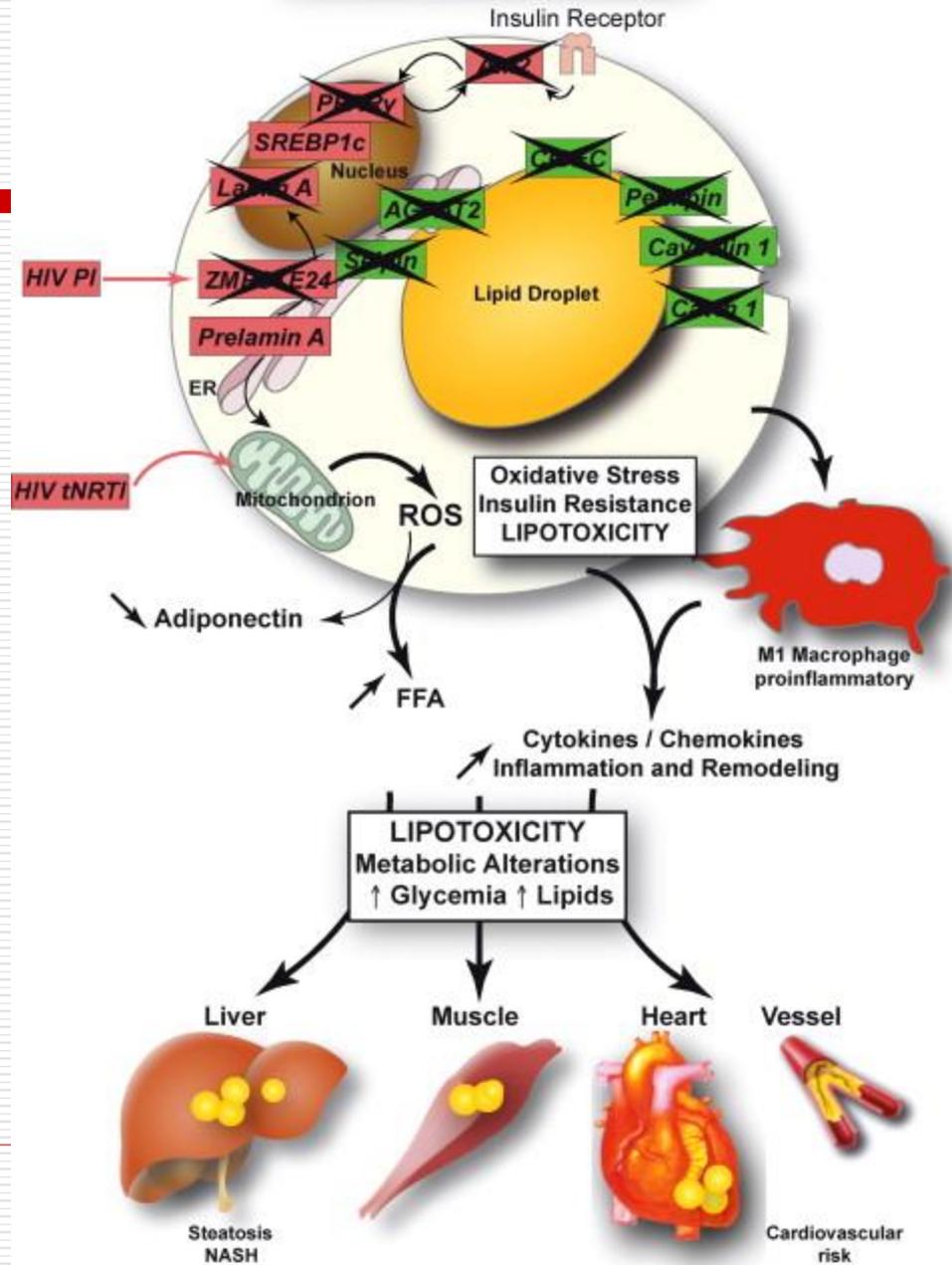


Lipid Storage

B: proteins of endoplasmic reticulum and lipid droplets during **fat storage**

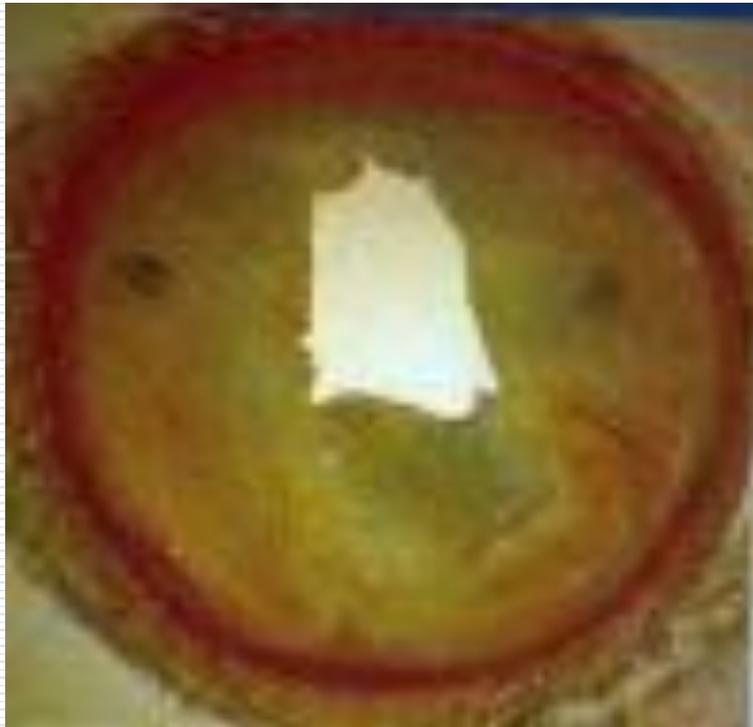
B

Lipodystrophic Adipose Tissue



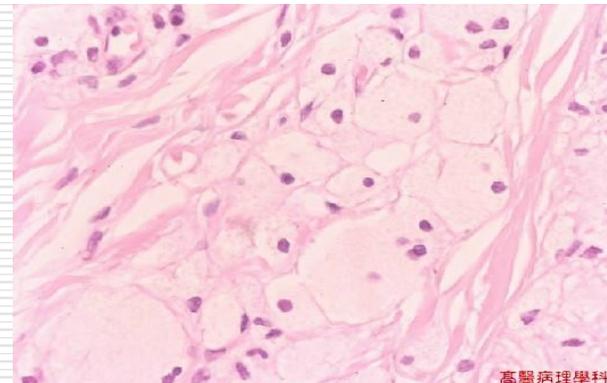
Hyperlipidemia Signs

- Atheroma- plaques in blood vessels



Hyperlipidemia signs

- Xanthoma- plaques or nodules composed of lipid-laden histiocytes (foamy cells) in the skin, especially the eyelids



Tendinous Xanthoma

Xanthoma deposits in tendon, commonly the Achilles



Corneal arcus

- Lipid deposit in cornea

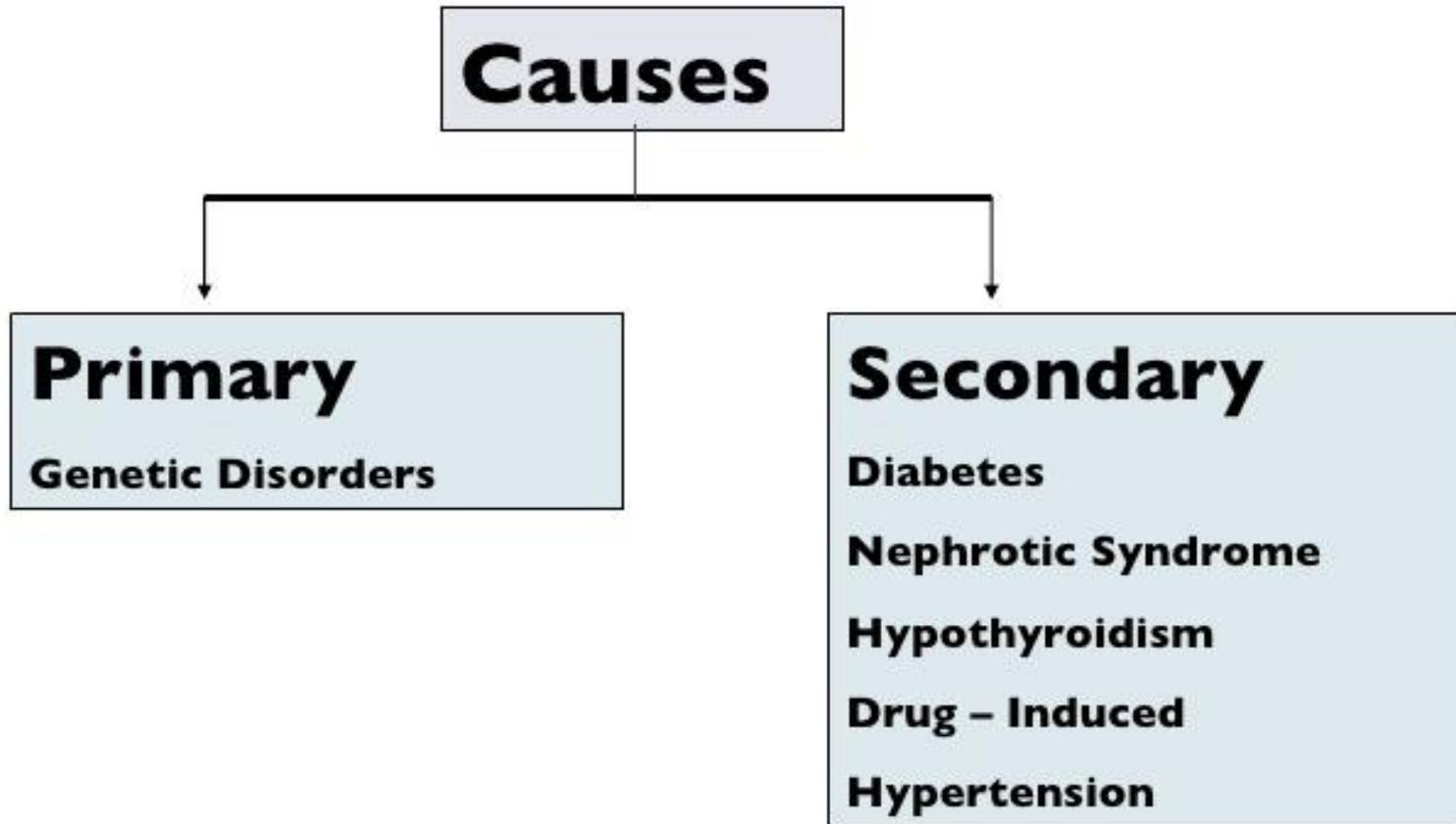


Dyslipidemia

- Disorder of Lipid & Lipoprotein Metabolism
- A common form of Dyslipidemia is characterized by three lipid abnormalities:
 - Elevated triglycerides,
 - Elevated LDL and
 - Reduced HDL cholesterol.
- Important Modifiable Risk Factor for CAD



Dyslipidemia



Secondary causes of Dyslipidemia

Selected Causes of Secondary Dyslipidemia

Increased LDL cholesterol level

Diabetes mellitus

Hypothyroidism

Nephrotic syndrome

Obstructive liver disease

Drugs

Anabolic steroids

Progestins

Beta-adrenergic blockers (without
intrinsic sympathomimetic action)

Thiazides

Increased triglyceride level

Alcoholism

Diabetes mellitus

Hypothyroidism

Obesity

Renal insufficiency

Drugs

Beta-adrenergic blockers (without
intrinsic sympathomimetic action)

Bile acid-binding resins

Estrogens

Ticlopidine (

Decreased HDL cholesterol level

Cigarette smoking

Diabetes mellitus

Hypertriglyceridemia

Menopause

Obesity

Puberty (in males)

Uremia

Drugs

Anabolic steroids

Beta-adrenergic blockers (without
intrinsic sympathomimetic action)

Progestins

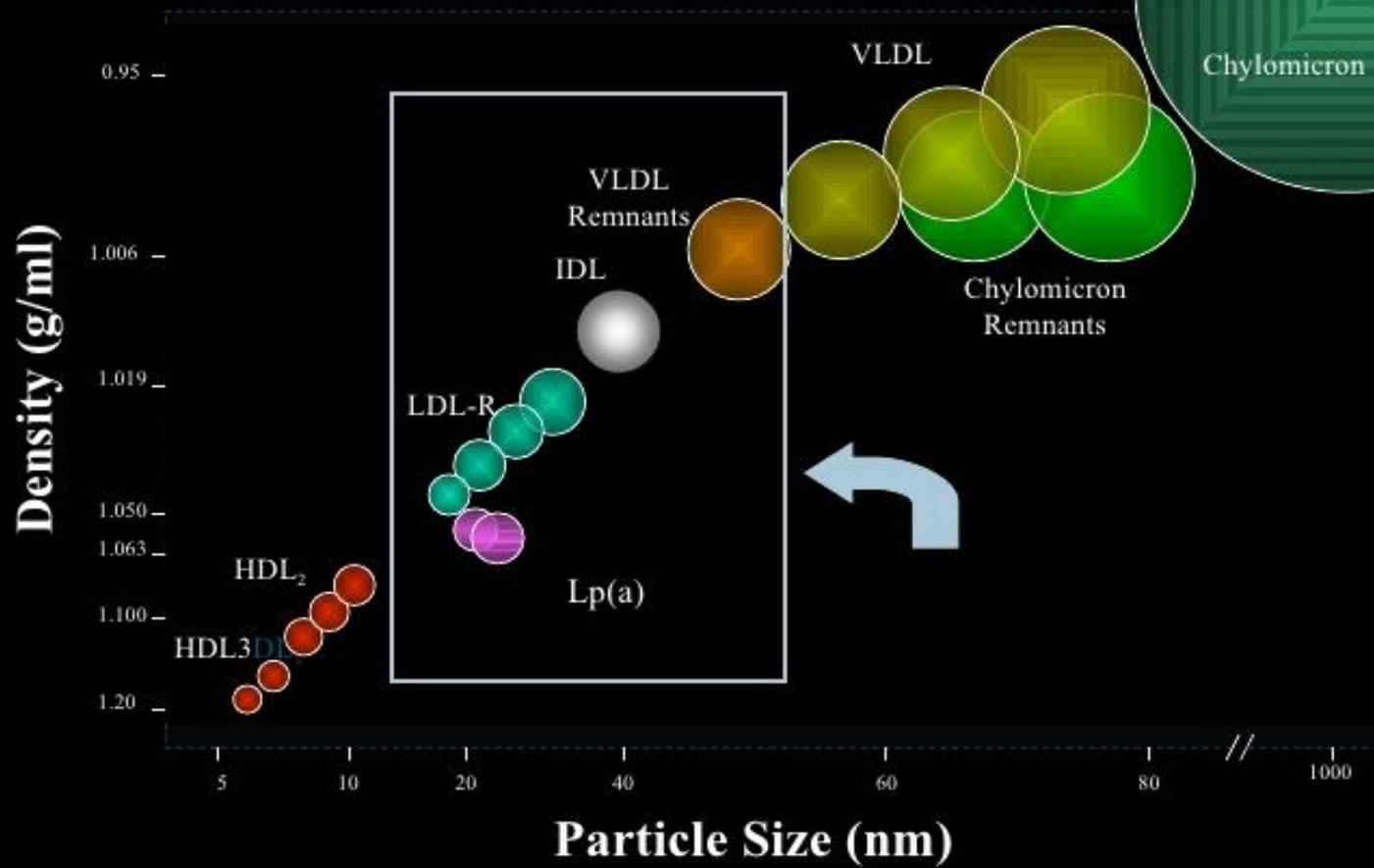
LDL=low-density lipoprotein; HDL=high-density lipoprotein.

Adapted with permission from Schaefer EJ. Diagnosis and management of lipoprotein disorders. In: Rifkind BM, ed. Drug

PRIMARY DYSLIPIDEMIA (Fredrickson's Classification)

Type	Primary Lipid Elevation	Lipoprotein	Occurrence
I	TG	Chylomicrons	Rare
II _a	C	LDL	Common
II _b	C, TG	LDL, VLDL	Most Common
III	C, TG	IDL	Rare
IV	TG	VLDL	Common
V	TG	VLDL, Chylomicrons	Rare

Lipoprotein Particles



Genetic Causes of Dyslipidemia

- ◆ **Type I – Familial Hyperchylomicronemia**

 - Fasting triglycerides > 1000 mg/dl

 - Defect in lipoprotein lipase or apo CII

 - Not necessarily at increased risk of CAD

- ◆ **Type II - Familial Hypercholesterolemia (type II)**

 - LDL-C > 95th percentile for age and gender

 - CAD in men by 3rd or 4th decade

 - Defect in LDL receptor

 - Autosomal dominant inheritance

 - Prevalence 1:500

- ◆ **Familial Defective apo B 100**

 - Defective apo B alters LDLr handling

 - Previously undetectable from FH

Genetic Causes of Dyslipidemia

Type III – Hyperlipoproteinemia

Increased TC, VLDL, decreased HDL; Increased VLDL: TG

Defect in apo E results in increased concentration of remnant particles

Rare

Type IV – Familial Hypertriglyceridemia

Increased TC (due to VLDL), TG, decreased LDL, HDL

Results from hepatic overproduction of VLDL

Prevalence 1:100 – 1:50; Association with CAD not as strong as FH

Heterogeneous inheritance

Very sensitive to diet and EtOH

Type V

Increase in chylomicrons and VLDL

Rare

Genetic Causes of Dyslipidemia

Familial Combined Hyperlipidemia

Increased TC, LDL and/or triglycerides; decreased HDL

Most common genetic dyslipidemia: prevalence 1:50

Heterogenous inheritance

Accounts for 10-20% of patients with premature CAD

Defects in HDL Metabolism

Most often low HDL is secondary to other dyslipidemia

Not all associated with increased CAD risk (e.g. apo A_I_{Milano})

Tangier's Disease

CETP defects result in increased HDL

Thank you for your attention

