



METABOLISM

2025-2024

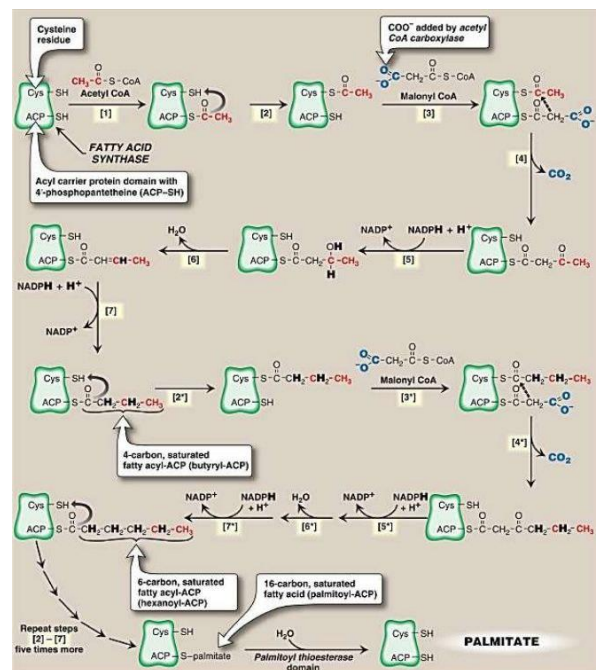
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Fatty acids synthesis

- At low energy state, high ADP level leads for the activation of TCA cycle to generate energy
- FAs are synthesized in the *high energy state*, which causes:
 - Inhibition of TCA cycle, especially isocitrate dehydrogenase enzyme, causing the accumulation of citrate which is transported from the mitochondria into the cytosol by citrate transporter and cleaved into *oxaloacetate* and *acetyl CoA* by *ATP citrate lyase*
 - Acetyl CoA is then used in the synthesis of FAs
- FAs synthesis occurs in the *cytosol* of *liver* and *adipose tissues*, by Fatty acid synthase (FAS)
- FAS is a multi-functional homo-dimeric enzyme
 - Each monomer has 6 enzymatic domains and 1 binding domain (ACP)
- **ACP** (Acyl-carrier protein) carries acyl groups and contains phosphopantetheine
 - *Phosphopantetheine* is a derivative of pantothenic acid (*vitamin B5*) with a terminal thiol group

FAs synthesis steps:

- *Acetyl CoA* is added to the **ACP** domain
- Acetyl is transferred into the **Cys** residue
- *Malonyl CoA* is added to the **ACP** domain
- **Condensation** reaction producing *4-carbon molecule* on the **ACP** and decarboxylation of carboxyl group in malonyl CoA, *released as CO₂*
- **Reduction** (requires **NADPH**) which converts the carbonyl group into hydroxyl
- **Dehydration** which involves the removal of H₂O
- **Reduction** (requires **NADPH**) removes double bond
- Addition of more malonyl CoA, and repeating the same steps for each *malonyl CoA* added increasing the length of the FA by *2 carbons each turn*



- *Palmitoyl thioesterase* domain of the FAS **cleaves** (releases) the palmitate from the ACP domain
- Synthesis of FAs in the *liver* and *adipose* tissue terminates by reaching 16 C molecule called **palmitate**
 - In the lactating *mammary* gland, the reaction terminates early producing **SCFAs** and **MCFAs**
- The source of substances used in FA synthesis:
 - **Acetyl CoA**: from *pyruvate oxidation*
 - **NADPH**: from *PPP* and conversion of *malate into pyruvate*
 - **NADH** (important for the conversion of OAA into malate then into pyruvate): from *glycolysis*
 - **Malonyl CoA**: *Carboxylation of acetyl CoA* by ACC

- Carboxylation of acetyl CoA produces Malonyl CoA, where CO₂ is brought as a ***bicarbonate (HCO₃⁻)***
 - This enzyme is catalyzed by ***acetyl CoA carboxylase (ACC)***, and it is ATP dependent
 - It requires ***vitamin B7*** as a coenzyme which is covalently bound to lysine in the ACC
 - It is the ***rate limiting step*** of FA synthesis
- ACC expression (production) requires the presence of 2 transcription factors:
 - Carbohydrate response element– binding protein (***ChREBP***)
 - Sterol regulatory element–binding protein-1c (***SREBP-1c***)
- Short term regulation of ACC activity:
 - Activated by ***citrate, phosphatase*** (dephosphorylation) and ***insulin***
 - Inhibited by ***palmitoyl CoA, AMP, glucagon*** and ***epinephrine***
 - ✓ AMP, glucagon and epinephrin inhibit ACC by activating ***AMPK*** (phosphorylation) which phosphorylates ACC
- Long term regulation of ACC activity:
 - ***PKA & AMPK*** phosphorylates ChREBP preventing its entry to the nucleus (reduces ACC synthesis)
 - ***Insulin*** activates SREBP-1c (increases ACC synthesis)
- ***Metformin (Glucophage)*** is a drug given for ***pre-diabetic patients*** which:
 - ***Lowers plasma TAGs*** by activating AMPK which inhibits ACC and FA synthase expression
 - ***lowers blood glucose*** by increasing AMP-mediated glucose uptake in the muscles
- For the production of 1 palmitate (16:0), it requires:
 - 1 acetyl CoA, 7 Malonyl CoA, 14 NADPH, 14 H⁺
 - 8 Acetyl CoA, 7 CO₂, 7 ATP, 14 NADPH, 14 H⁺
- Further ***elongation*** of FAs occurs in:
 - ***SER***: where the 2-carbon donor is ***malonyl CoA*** and the source of electrons is ***NADPH***
 - ***Mitochondria***: the 2-carbon donor is ***acetyl CoA***, and the source of electrons is ***NADH & NADPH***
 - ✓ ***Brain*** has enzymes allowing it to produce VLCFA
- Chain desaturation is done by ***fatty acyl CoA desaturases*** which act on LCFAs in the ***SER***
 - The electron acceptor is ***O₂*** and ***Cyt b5*** (which have FAD-linked reductase)
 - Donor of electrons: ***NADH***
- The ***first double bond*** is inserted between carbons 9 and 10, producing ***oleic acid (18:1)***, and small amounts of ***palmitoleic acid (16:1)*** by Δ⁹ desaturase
- Humans have carbon Δ⁹, Δ⁶, Δ⁵ and Δ⁴ ***desaturases***
- We cannot introduce double bonds from carbon 10 to the ω end of the chain, so they are essential
 - The polyunsaturated ***ω-6 linoleic acid*** and ***ω-3 linolenic acid*** are ***essential***

AMPK: AMP-activated
Protein Kinase

- Notes
 - SCFAs and MCFAs usually do not require a carrier in plasma, but LCFAs *requires albumin*
 - Glucose can be used to produce fats, but fats **can't** produce glucose
 - ACC2 Inhibitors can be used as wight loss agents
 - CoA also contains Vitamin B5

Glyceroneogenesis

- Glycerol production from *non-glucose precursors* such as lactate, alanine and aspartate
- It occurs in the adipose tissue and liver, and after synthesis glycerol leaves adipose to the liver
- It *regulates FA levels* in the blood
- Failure in regulating glyceroneogenesis may lead to *Type 2 diabetes*
- Steps:
 - Precursor is converted into pyruvate which is converted into OAA by pyruvate carboxylase (PC), which is converted into PEP (phosphoenolpyruvate) by PEP-CK-C (PEP carboxykinase C)
 - PEP is converted into G3P then DHAP which is converted into glycerol 3-phosphate

TAGs synthesis

- Synthesis of TAGs in the liver and adipose tissue involves:
 - Glycerol 3-phosphate synthesis, by 2 mechanisms:
 - ✓ *Conversion of DHAP* (from glycolysis) into glycerol 3-phosphate
 - ✓ *Phosphorylation of glycerol* into glycerol 3-phosphate by glycerol kinase
 - Liver can use both mechanisms; adipose tissue can only use the first one
 - Glucose enters the adipose tissue by GLUT4 transporter
 - Activation of fatty acids by adding CoA, by:
 - ✓ *Adding AMP* from ATP into FA producing acyl-adenylate and PP_i by Thiokinase
 - ✓ *AMP is replaced by CoA* producing Acyl CoA
 - Synthesis of triacylglycerol
 - ✓ Acyltransferase adds a FA-CoA to glycerol 3-phosphate on C1
 - ✓ Acyltransferase adds a FA-CoA to lysophosphatidic acid on C2
 - ✓ Phosphatase removes phosphate group from phosphatidic acid
 - ✓ Acyltransferase adds a FA-CoA to diacylglycerol on C3 forming TAG
- TAGs in the **adipose** tissue are produced to be *stored* as lipid droplets
- TAGs in the **liver** are produced mainly to be *packaged in VLDL* and a small portion for storage

- In the **intestines**, TAGs are synthesized by **MAG pathway**, which involves:
 - Monoacylglycerol enters mucosal cells
 - 2 FAs are activated into FA-CoA, to be added into the monoglyceride producing TAG
 - TAGs are then packaged into **chylomicrons** into the lymph

Degradation of FAs and TAGs

- Lipids represent a **major storage site of energy** (long term storage), especially TAGs
 - Lipids are hydrophobic (stored without water), producing 9 Kcal per gram
 - Carbohydrates are hydrophilic (which attracts water), producing 4 Kcal per gram
 - ✓ **Glucose is the major circulating fuel**
- The release of FAs from TAGs:
 - **Adipose triglyceride lipase (ATGL)** releases the **1st** FA of TAG, producing diglyceride
 - **Hormone-sensitive lipase (HSL)** releases the **2nd** FA producing monoglyceride
 - **Monoacylglyceride lipase (MGL)** releases the **3rd** FA producing glycerol
- In well-fed state, **insulin** inhibits ATGL synthesis and activates phosphatase which inhibits HSL
 - **Perilipin** protein that coats fat droplets, act as an emulsifier and blocks HSL
- In fasting state, **Glucagon** and **epinephrine** increases cAMP which activates PKA that phosphorylates perilipin releasing (disassembles) it and activating HSL
 - **Phosphodiesterase inhibitors** activate the HSL

β-Oxidation of FAs

- It is the process of cleaving FAs on the **β-C**, producing **2-C fragments (Acetyl CoA)**
 - It occurs in the **mitochondrial matrix**
- SCFAs and MCFAs can enter the mitochondrial matrix **easily**
- LCFAs are shuttled from the cytosol into the mitochondrial matrix by the **carnitine shuttle mechanism**
- Carnitine shuttle mechanism
 - FAs are **activated** by attaching CoA via **Acyl CoA synthetase** (**outer** membrane), requiring **ATP**
 - Fatty acyl CoA passes through the outer mitochondrial membrane
 - **CPT I** (carnitine palmitoyl transferase I) in the **outer** membrane converts Fatty acyl CoA & carnitine into fatty **acyl-carnitine** and **CoA** and it is inhibited by **malonyl CoA**
 - Fatty acyl carnitine is passed through the inner membrane by a translocase into the matrix
 - **CPT II** (**inner** membrane) reform **fatty acyl-CoA**, and a translocase passes **carnitine** out

- Carnitine is originally synthesized from 2 amino acids L-methionine and L-lysine
 - The cardiac and skeletal **muscles** contain 97% of carnitine in the body
 - Muscles don't have ACC1 (no FA synthesis) but have mitochondrial ACC2 (regulate degradation)
- Sources of carnitine are meat and synthesis in the kidney and liver
- Carnitine deficiencies:
 - **Primary carnitine deficiency**
 - ✓ *Defects in membrane transporter*: No uptake of carnitine by muscles and the kidneys (excreted)
 - ✓ Treated by carnitine supplementation
 - **Secondary carnitine deficiency**
 - ✓ Taking *valproic acid (antiseizure)*, decreases renal reabsorption
 - ✓ *Liver diseases* cause decreased carnitine synthesis
 - ✓ *Defective fatty acid oxidation*, causes acyl-carnitines accumulation and its excretion into urine
 - ✓ *CPT-I deficiency*: affects the **liver**, causing severe **hypoglycemia, coma, and death**
 - ✓ *CPT-II deficiency*: affects the **liver, cardiac muscle, and skeletal muscle**
 - Avoidance of fasting
 - A diet with high carbohydrates, low fat and supplemented with medium-chain TAG
- β -oxidation of FAs:
 - *Oxidation*: fatty acyl CoA is oxidized by **Acyl CoA dehydrogenase** which reduces FAD into **FADH₂**
 - ✓ This enzyme has 4 isozymes SCAD, MCAD, LCAD, VLCAD
 - *Hydration*: addition of water by hydratase
 - *Oxidation*: by **hydroxyacyl-CoA dehydrogenase**, reducing NAD⁺ into **NADH**
 - *Thiolytic cleavage*: by **thiolase** via breaking the bond between alpha and beta carbon
 - ✓ Inhibited by **Acetyl CoA**
- The number of cycles of β -oxidation and FA synthesis = **$(n/2) - 1$**
- Number of acetyl CoA produced during β -oxidation = **$n/2$**
 - NADH = 2.5 ATP, FADH₂ = 1.5 ATP, Acetyl CoA = 12 ATP
 - **Net** ATP yield of oxidation of FA = the net ATP yield of oxidation of fatty acyl CoA **- 2**
 - Not all acetyl CoA are used to produce ATP, some are used in gluconeogenesis and ketogenesis
- Medium chain fatty-acyl CoA dehydrogenase (**MCAD**) *deficiency* is an autosomal-recessive disorder
 - Most common inborn error of β -oxidation (1:14,000 births worldwide)
 - Higher incidence among Caucasians of Northern European descent
 - Decreased ability to oxidize MCFAs (lack of energy) and severe hypoglycemia and hypoketonemia
 - Treatment: *avoidance of fasting* by regular and frequent meals and snacks and diet high in carbohydrates and low in fat

Oxidation of ODD FA

- Odd FAs undergo ***β-oxidation*** until reaching the last 3-C fragment which is called ***propionyl CoA***
 - Propionyl CoA is carboxylated by ***propionyl CoA carboxylase*** which requires ***ATP*** and ***CO₂*** in the form of ***HCO₃⁻*** and carried by biotin (***Vit. B7***) forming ***D-methylmalonyl CoA***
 - D-methylmalonyl CoA is converted into ***L-methylmalonyl CoA*** by ***epimerase***
 - L-methylmalonyl CoA is converted into ***Succinyl CoA*** by ***methylmalonyl CoA mutase*** which requires ***vitamin B12*** as a coenzyme
 - ✓ Deficiency in this enzyme or vitamin B12 causes neurological manifestations (developmental delay, seizures or intellectual disabilities) and metabolic acidosis
 - ✓ Succinyl CoA enters the TCA cycle

Oxidation of unsaturated FA

- **Monounsaturated** FAs are isomerized where the double bond is transferred between α and β carbons
 - It is done by ***isomerase*** and it causes the skip of the first step of β -oxidation leading to the loss of electrons (***loss of FADH₂***)
- Oxidation of **polyunsaturated** FAs such as linoleic acid (18:2) requires an NADPH dependent ***2,4-dienoyl CoA reductase*** in addition to the ***isomerase***
 - It causes the loss of electrons (***loss of FADH₂, NADPH***)

Peroxisomal β -oxidation

- Oxidation of ***VLCFAs into LCFAs*** then return to the mitochondrial β -oxidation
 - It is very similar to the mitochondrial β -oxidation, but it uses ***acyl CoA oxidase*** instead of dehydrogenase in the first step
 - ✓ FADH₂ is oxidized forming hydrogen peroxide (***H₂O₂***) causing the ***loss of electrons***
- ***Zellweger syndrome*** (peroxisomal biogenesis disorder) and ***X-linked adrenoleukodystrophy*** (dysfunctional transport VLCFA) causes VLCFAs accumulation

Peroxisomal α -oxidation

- Occurs in the ***phytanic acid metabolism***
 - Phytanic acid is a branched fatty acid produced from the breakdown of ***chlorophyll***
 - Cleaves the bond between the ***1st*** and ***α carbons*** ***producing CO₂***
- Steps:
 - Phytanic acid is ***activated*** by CoA (requires ***ATP***) then it is transported into the peroxisomes
 - It is hydroxylated by phytanoyl CoA α -hydroxylase (***PhyH***) which has ***Fe⁺²*** as a prosthetic group

➤ 2-hydroxy phytanoyl-CoA lyase releases formyl-CoA which is then converted into formic acid and then CO_2 is released

• When fully degraded, it generates 4 byproducts: acetyl-CoA, 2-methyl-propionyl-CoA, formyl-CoA, propionyl-CoA

• *Refsum disease* is an autosomal-recessive disorder caused by a deficiency of peroxisomal PhyH

SER ω -oxidation

• ω -Oxidation is a minor pathway of the SER where FAs is converted into *dicarboxylic* acids by forming carboxyl group on the ω carbon

• It is upregulated in certain conditions such as MCAD deficiency

Sources of energy

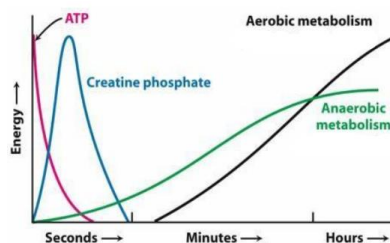
• Sources of energy during an exercise:

➤ For the first 5 seconds, our reservoir of ATP is consumed

➤ For the first 30 seconds, creatinine phosphate is used to replenish ATP reservoir by creatinine kinase

➤ Aerobic and anaerobic metabolism starts:

- ✓ Anaerobic can't continue for a long time because of the accumulation of lactic acid
- ✓ Aerobic metabolism works after a while and persist for long periods



• For *short distance runners*, they rely on *ATP* and *creatinine phosphate*

• For *marathon runners*, they mostly rely on *aerobic metabolism* so must attain adequate oxygen

VARIABLE	SYNTHESIS	DEGRADATION
Greatest flux through pathway	After carbohydrate-rich meal	In starvation
Hormonal state favoring pathway	High insulin/glucagon ratio	Low insulin/glucagon ratio
Major tissue site	Primarily liver	Muscle, liver
Subcellular location	Cytosol	Primarily mitochondria
Carriers of acyl/acetyl groups between mitochondria and cytosol	Citrate (mitochondria to cytosol)	Carnitine (cytosol to mitochondria)
Phosphopantetheine-containing active carriers	Acyl carrier protein domain, coenzyme A	Coenzyme A
Oxidation/reduction coenzymes	NADPH (reduction)	NAD ⁺ , FAD (oxidation)
Two-carbon donor/product	Malonyl CoA: donor of one acetyl group	Acetyl CoA: product of β -oxidation
Activator	Citrate	—
Inhibitor	Palmitoyl CoA (inhibits acetyl CoA carboxylase)	Malonyl CoA (inhibits carnitine palmitoyltransferase-I)
Product of pathway	Palmitate	Acetyl CoA
Repetitive four-step process	Condensation, reduction, dehydration, reduction	Dehydrogenation, hydration, dehydrogenation, thiolysis

	carbohydrates	lipids
Stored as...?	Starch - plants Glycogen - animals	Fats & oils (plants) Fat (animals)
Long/short term storage?	Starch: long-term Glycogen: short-term	Long term
Ease of digestion/ release of energy?	Easy to release energy	Harder to release energy (needs more oxygen)
Energy per gram?	17kJ/g	38kJ/g
Solubility in water? (and consequence)	Soluble	Not soluble
Use of oxygen in metabolism? (and consequence)	Needs less oxygen, useful for high-demand activity	Needs more oxygen, less efficient to release energy



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