# Regulation of Lipid Synthesis

This unit will cover the synthesis of lipids including cholesterol, fatty acid and triglyceride synthesis. For more details on these topics, refer to Chapters 28 and 29 in Biochemistry: A Short Course available in reserve<sup>1</sup>.

<sup>1</sup> John L Tymockzko, Jeremy M Berg, and Lubert Stryer. *Biochemistry: A Short Course.* W.H. Freeman and Co, New York, NY, 3rd edition, 2015

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# Learning Objectives

- Identify the functions of cholesterol within our body
- Understand how fatty acids are synthesized in the body
- Describe the initial highly regulated step of FA synthesis
- Explain the tissue-specific energetic costs of triglyceride synthesis.
- Understand the reactions resulting in triacylglycerol synthesis resulting in 3 fatty acids esterified to glycerol
- Describe the breakdown of triacylglycerol to glycerol plus fatty acids and the fates of these products
- Describe fatty acid breakdown (beta-oxidation)
- Determine the amount of energy produced by fatty acid breakdown in comparison to glucose
- Understand when ketogenesis occurs, what organs ketones are a fuel source for and consequences of overactive ketogenesis
- Describe how triglyceride synthesis is regulated in the liver, and how liver triglycerides are transported to other tissues

# *Synthesis of Triglycerides from Fatty Acids*

Fatty acids are obtained from the diet, or can be made from excess glucose/amino acids<sup>2</sup>. The esterification of these fatty acids within cells is very important because free fatty acids can be toxic to a cell once they build up. Most tissues are able to store excess fatty acids to some degree but the major sites of synthesis are adipose tissue, and

Generally cells have to decide between three general fates for the fatty acids that arrive inside the cell<sup>3</sup>:

Without energy demand: esterify with glycerol as triglycerides

With energy demand, but with OAA availability: oxidize fatty acid to Acetyl-CoA and use that for fuel in the TCA cycle to generate ATP.

With energy demand but insufficient OAA: oxidize to Acetyl-CoA and convert that to ketone bodies for release (primarily in the liver<sup>4</sup>.

We will discuss the regulation of  $\beta$ -oxidation, which is the main process by which fatty acids become Acetyl-CoA in the next lecture. When fatty acid synthesis is active, the elevations in malonyl-CoA<sup>5</sup>

<sup>&</sup>lt;sup>2</sup> A process known as de novo lipogene-

<sup>&</sup>lt;sup>3</sup> Though in growing and dividing cells, the fraction of fatty acids that become phospholipids becomes much more relevant.

<sup>&</sup>lt;sup>4</sup> Two main reasons why this is mainly in the liver, the first is that there is typically less cataplerosis in nonhepatic tissues because there is less gluconeogenesis. The second is that the enzymes of ketogenesis are at higher levels in the liver.

<sup>5</sup> Described below

will generally suppress  $\beta$ -oxidation via inhibition of Carnitine Palmitoyltransferase I<sup>6</sup>. This ensures that fatty acid synthesis and oxidation do not occur simultaneously.

TRIGLYCERIDE SYNTHESIS INVOLVES THE SEQUENTIAL ADDITION OF FATTY ACIDS TO A GLYCEROL BACKBONE. This involves enzymatic reactions starting with phosphorylated glycyerol and an activated fatty acid. To prepare fatty acids for esterification, an fatty acid<sup>7</sup> is conjugated to coenzyme A by Acyl-CoA Synthetase:

$$FattyAcid + CoA + ATP \rightarrow AcylCoA + AMP + PPi$$
 (1)

Note that this reaction consumes two high ATP equivalents for each fatty acid to be added. This fatty acid is now in the activated form and is available to be conjugated to phosphorylated glycerol by the following four sequential reactions:

$$AcylCoA + Glycerol3P \rightarrow LysophosphatidicAcid$$
 (2)

$$AcylCoA + LysophosphatidicAcid \rightarrow PhosphatidicAcid$$
 (3)

$$PhosphatidicAcid \rightarrow Diacylglycerol + Pi$$
 (4)

$$AcylCoA + Diacylglycerol \rightarrow Triacylglycerol$$
 (5)

The phosphatidic acid generated in step 3 is the precursor FOR MOST PHOSPHOLIPIDS. Enzymes can convert phosphatidic acid to phosphatidylserine, or phosphatidylinositol. Phosphatidylserine is the substrate for the generation of phosphatidylethanolamine, which in turn generates phosphatidylcholine<sup>8</sup> For more details about phospholipid synthesis see Kent [1995]. These structural lipids are particularly important during development, and are targets to reduce the growth of cancer cells, which require generation of substantial membranes.

FATTY ACIDS ARE ADDED IN A SPECIFIC MANNER. Generally at the sn2 position is an unsaturated fatty acid, while there are saturated fats at the sn1 position. The sn3 position seems to have much less specificity in mammalian triglyceride formation [Brockerhoff, 1971]. This is relevant for phospholipids that form membranes, as these will tend to have one saturated and one unsaturated fatty acid at each of the sn1 and sn2 positions<sup>9</sup>.

<sup>&</sup>lt;sup>6</sup> Which is the rate limiting step of fatty acid breakdown.

<sup>&</sup>lt;sup>7</sup> Also known as an acyl group

<sup>&</sup>lt;sup>8</sup> This is the reaction catalyzed by PEMT, the enzyme that has variants which conver choline dependence.

<sup>&</sup>lt;sup>9</sup> The sn<sub>3</sub> position is where the phosphate group is located.

## Glycerol-3-Phosphate Supply for Triglycerides

While each triglyceride molecule needs three fatty acids, they also require a phosphorylated glycerol backbone for reaction 2. The source of this glycerol varies between tissues. In the liver, glycerol can be generated by phosphorylation of glycerol via Glycerol Kinase at a cost of an ATP:

$$Glycerol + ATP \rightarrow Glycerol3P$$
 (6)

Adipocytes on the other hand have low levels of glycerol kinase activity. Therefore when fatty acids are presented to adipocytes they require glycerol to be made from glucose. The glycolytic intermediate dihydroxyacetone phosphate (DHAP) is generated by Aldolase<sup>10</sup> and can be converted into glycerol.

$$DHAP + NADH \rightleftharpoons Glycerol3P + NAD^{+} \tag{7}$$

This reaction comes at a cost of one NADH molecule, or 2.5 ATP equivalents. The lack of glycerol kinase in adipocytes is thought to be an adaptation to prevent the futile release and then re-esterification of fatty acids, wheras after lipolysis in adipocytes both glycerol and fatty acids can go to the liver, so the liver always has sufficient glycerol to prevent fatty acid accumulation. Nonetheless, because of this requirement, a substantial amount of the glucose used in adipocytes is used to generate glycerol backbones. In the absence of glucose, gluconeogenic precursors such as lactate and pyruvate can be converted to glycerol in adipocytes, a process known as glyceroneogenesis<sup>11</sup>.

THE ENERGY COSTS OF TRIGLYCERIDE SYNTHESIS, based on these pathways is substantial. Three acyl chains must activated at a cost of 2 ATP equivalents each for 6 ATP in total. This is tissue independent. In the liver, one more ATP to activate glycerol so a total 7 ATP are required to form one triglyceride. In adipose tissue the cost is more because you use 2 x ATP in the preparation phase of glycolysis and another NADH to form Glycerol-3-phosphate in reaction 7. This means the cost in adipose tissue is 10.5 ATP, so it is not only more energy demanding, but also consumes some of the glucose that would be used to generate those ATP molecules.

#### Regulation of Triglyceride Esterification Enzymes

Both in the liver and in adipose triglyceride synthesis will increase in response to insulin. The mechanisms of this include both shortterm and long term regulation. Recall first, that in adipocytes glucose 10 Two steps after PFK1

<sup>11</sup> This is equivalent to gluconeoegenesis, up to the DHAP step, where reaction 7 takes precedence over the Aldolase reaction.

uptake is increased by insulin via stimulation of GLUT4 translocation. Insulin can acutely activate GPAT<sup>12</sup> and Lipin <sup>13</sup>. Together the increased glucose and fatty acid flux, combined with more activity of these triglyceride synthethizing enzymes will result in efficient triglyceride storage in adipose tissue after a meal. Simulataneous to these effects, as we will discuss in the lipid transport lecture, insulin also suppresses triglyceride breakdown, a process known as *lipolysis*. For more insights into how insulin regulates triglyceride synthesis, see the recent review by Coleman and Mashek [2011].

CHRONIC REGULATION OF TRIGLYCERIDE SYNTHESIS IS TRAN-SCRIPTIONAL and is regulated by the two transcription factors SREBP1c<sup>14</sup> and PPAR $\gamma^{15}$  Both of these nuclear hormone receptors increase the number of the key triglyceride synthesis enzymes, including GPAT, Lipin and AGPAT<sup>16</sup>. The regulation of SREBP1c is quite complicated but involves insulin- and mTORC1-dependent signals (reviewed in Bakan and Laplante [2012]). In this way both insulin and nutritional status can promote the efficiency of triglyceride storage.

# *De Novo Fatty Acid Synthesis*

Triglycerides can use fatty acids derived from the diet, or that are generated endogenously. Certain fatty acids (the  $\omega_3$  and  $\omega_6$  derived fatty acids) are entirely dependent on nutritional inputs, but others are generated endogenously. This can vary substantially, with individuals on a high carbohydrate diet generating many of their fatty acids de novo, with those on a low carbohydrate diet relying almost entirely on dietary intake. All tissues can perform this function<sup>17</sup>, but the major sites are liver and adipose tissue.

#### Fatty Acid Synthesis from Glucose and Ketogenic Amino Acids

Fatty acid biosynthesis occurs in the endoplasmic reticulum, which presents the first problem in terms of substrates. The initial input, Acetyl-CoA is generated from its precursors<sup>18</sup> in the mitochondria, and there are no Acetyl-CoA transporters. Therefore Acetyl-CoA must first be converted to Citrate via Citrate Synthase<sup>19</sup>:

$$AcetylCoA + OAA + H_2O \rightarrow Citrate + CoA$$
 (8)

Citrate transporters do exist in the mitochondria, so citrate is removed and then reconverted to Acetyl-CoA via an enzyme called

- 12 The enzyme that catalyzes reaction 2.
- 13 The enzyme that catalyzes reaction 4, this is mediated by activation of mTORC1 [Harris et al., 2007]

- 14 Sterol response element binding protein 1c, which has a similar isoform SREBP2, that plays a key role in cholesterol synthesis, more on this in the next section.
- 15 This stands for Peroxisome proliferator-activated receptor gamma, a nuclear hormone receptor which promotes both triglyceride synthesis and new adipocyte formation. It is the target of the anti-diabetic drugs of the thiazolidinedione family.
- 16 Which catalyzes reaction 2.

- <sup>17</sup> As it may be important for phospholipid synthesis
- <sup>18</sup> Hypothetically from pyruvate via Pyruvate Dehydrogenase, from ketogenic amino acids or as the product of  $\beta$ -oxidation, though the latter is unlikely as the cell will not typically break down fatty acids and then immediately resynthesize then.
- 19 This is the same Citrate Synthase that is the first step of the TCA cycle.

ATP-Citrate Lyase via the following reaction:

$$Citrate + ATP + CoA \rightarrow AcetylCoA + OAA + ADP + Pi$$
 (9)

Once the Acetyl-CoA is in the cytoplasm<sup>20</sup>, the next and ratelimiting step is catalyzed by an enzyme called Acetyl-CoA Carboxylase or ACC. This enzyme catalyzes the following reaction:

20 As we will discuss later, the Oxaloacetate from ATP-Citrate Lyase activity in the cytoplasm is also important for generating reducing agents. This is described in reaction 12 below.

$$AcetylCoA + ATP + HCO_3 \rightarrow MalonylCoA + ADP + Pi + H^+$$
 (10)

Malonyl-CoA and Acetyl-CoA are both the substrates of Fatty Acid Synthase, a multifunctional enzyme that catalyzes the next several reactions. Acetyl-CoA serves as both a primerAs in an initial substrate and a source of carbon units<sup>21</sup>. The first round generates a four carbon fatty acid condensing Acetyl-CoA and Malonyl-CoA and using two NADPH<sup>22</sup> molecules. After that, each round of the reaction consumes two NADPH and another Malonyl-CoA, resulting in the sequential addition of carbons two at a time. This ends when the acyl chain is 16 carbons long, releasing Palmitate, a C16:0 fatty acid. In sum the overall reaction for a single palmitate is:

<sup>21</sup> Through MalonylCoA generated by ACC

<sup>22</sup> Not NADH, remember this from the pentose phosphate shunt?

$$8AcetylCoA + 7ATP + 14NADPH \rightarrow Palmitate + 14NADP^+ + 7ADP + 8CoA + 7Pi + 6H_2O$$
(11)

THE NADPH REQUIREMENTS OF FATTY ACID SYNTHESIS ARE HIGH. Each palmitate requires 14 NADPH molecules along with a substantial amount of ATP. The NADPH comes from two sources the pentose phosphate shunt<sup>23</sup>, and the activity of Malic Enzyme. Recall from above, when a Citrate molecule is e xported, the other product of ATP-Citrate Lyase, shown in reaction 9, is oxaloacetate. While this is used as a TCA intermediate in the mitochondria, in the cytoplasm OAA can be converted into pyruvate via Malic Dehydrogenase and Malic Enzyme. This reactions are:

<sup>23</sup> Wherin one glucose molecule yields one NADPH molecule.

$$OAA + NADH \rightarrow Malate + NAD^{+}$$
 (12)

$$Malate + NADP+ \rightarrow NADPH + CO_2 + 2Pyruvate$$
 (13)

In sum this reaction uses a cytosolic NADH and the released oxaloacetate to generate NADPH and two pyruvate molecules, which can then be re-oxidized in the mitochondria. This means that for each Citrate released and Acetyl-CoA generated by ATP-Citrate Lyase, one NADPH is regenerated (again, at a cost of an NADH). This

helps to maintain the NADPH pool. Therefore we can suspect that the 8 Acetyl-CoA's needed in reaction 11 will come with 8 NADPH molecules, so on average we need 6 more from the pentose phosphate shunt. This means we need the equivalent of 6 molecules of glucose to go through the pentose phosphate pathway, and still need to generate 7 ATP, 8 NADH (equivalent to 20 ATP) and 8 Acetyl-CoA molecules (This requires 4 Glucoses, but generates 12 NADH and 4 ATP during partial oxidation) to power the formation of a single palmitic acid. This means a net requirement of 16 glucose molecules to make a single palmitic acid, at a net gain of only 11 ATP equivalents of energy<sup>24</sup>. Recall that if those same glucose molecules were to undergo complete oxidation it would yield 512 ATP<sup>25</sup>. This is a huge diversion of resources for making a single fatty acid.

# Desaturation of Fatty Acids

While palmitate is the initial fatty acid made by mammals, this fatty acid (C16:0) can be further modified by both elongases and desaturases. Elongases can extend dietary or de novo fatty acids by increments of two by adding another Malonyl-CoA moiety and using two more NADPH molecules. This results in the synthesis of very long chain fatty acids. There are several elongases in the human genome, but mutations in *ELOVL4* have been linked to macular dystrophy, highlighting the importance of fatty acid elongation in the conversion of ALA to DHA [Zhang et al., 2001]. Adipocytes and liver cells express high levels of ELOVL6, which elongates fatty acids to a maximum of 18 carbons. This is one reason<sup>26</sup> why the primary de novo synthesized fatty acids end up as C16:0, C16:1 $\Delta^9$ , C18:0, and C18:1 $\Delta^9$ in these tissues.

Humans have three deaturases that work on the  $\Delta_5$ ,  $\Delta_6$  and  $\Delta$ 9 positions of a fatty acid. The enzymes that catalyze these reactions are FADS1 and FADS2, which we discussed in terms of PUFA metabolism and the Stearoyl-CoA desaturase<sup>27</sup> enzymes. The desaturation is a key part of the assembly of triglycerides and phosphpolipids, as a desaturated fatty acid is preferentially attached to the sn2 position. Both desaturases and elongases can alter the structure of both dietary and endogenously produced fatty acids.

#### Regulation of Fatty Acid Synthesis

Two steps of fatty acid biosynthesis are under relatively acute control. ATP-Citrate Lyase is phosphorylated and activated by insulindependent signaling through Akt [Berwick et al., 2002], wheras Acetyl-CoA Carboxylase is dephosphorylated and activated in an

<sup>&</sup>lt;sup>24</sup> The exact math here is unimportant, since this is simplified scenario, but the point is that you have to use up a lot of glucose without gaining a lot of energy to make a single fatty acid

<sup>&</sup>lt;sup>25</sup> 16 Glucose x 32 ATP/glucose.

<sup>&</sup>lt;sup>26</sup> The other is the activity of SCD1, the  $\Delta^9$  desaturase.

<sup>&</sup>lt;sup>27</sup> Abbreviated SCD.

Akt-dependent manner [Witters and Kemp, 1992]. The inactivation of ACC is due to phosphorylation by both PKA<sup>28</sup> or AMPK<sup>29</sup>, so the role of insulin is to reverse that inhibition. The precise process by which insulin and Akt mediate this dephosphorylation is not clear at this stage.

AT A CHRONIC LEVEL, FATTY ACID BIOSYNTHESIS IS REGULATED BY TRANSCRIPTIONAL CONTROL OF SEVERAL ENZYMES. SREBP1c, discussed earlier, promotes the synthesis of Fatty Acid Synthase, ACC, ATP-Citrate Lyase, SCD1 and several of the elongases [Horton et al., 2002, Moon et al., 2012]. Another transcriptional regulator is the more recently described ChREBP<sup>30</sup>. This transcription factor is activated by excessive carbohydrates, and especially in the liver, results in the activation of many of the same lipid synthetic proteins<sup>31</sup>. ChREBP is also thought to play a key role in fructose's strong lipogenic effects in the liver [Kim et al., 2016].

## Lipodistrophy Results from Impaired Triglyceride Storage

We may generally consider fatty acid formation and storage as negative, as increased adiposity is causal of obesity, which now affects over a third of Americans [Flegal et al., 2016]. However fat synthesis and storage is also essential for normal function. Disorders or impairments of either adipocyte generation, or lipid storage result in a disease termed *lipodistrophy*. In this condition, fatty acids are unable to be stored effectively as triglycerides in adipose depots and therefore are ectopically deposited in tissues such as liver and muscle, promoting their impairment and inducing insulin resistance. A recent genetic association study showed that among normal weight seeming individuals with insulin resistance and hyperlipidemia, variants in lipogenic and adipogenic genes were associated with this phenotype [Lotta et al., 2016]. This may also be part of the reason why some ethic groups, such as those of Chinese descent, have increased diabetes and liver disease risk, at adiposity levels that pose less risk for those of European descent [Chiu et al., 2011].

- <sup>28</sup> In response to glucagon or adrenaline signaling
- <sup>29</sup> The AMP-Activated Protein Kinase, In response to energy needs.

- 30 Carbohydrate Response Element **Binding Protein**
- 31 More information about ChREBP can be found in Baraille et al. [2015].

# Cholesterol Synthesis

HMG-CoA Reductase is the Rate Limiting Step for Cholesterol Synthesis

Sensing and Regulation of Sterol Biosynthesis by SREBP2

The Relationship Between Dietary Fat and Cholesterol Synthesis

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