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## Omega-3 Polyunsaturated Fatty Acids as a Treatment Strategy for Nonalcoholic Fatty Liver Disease

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### Abstract

Obese and type 2 diabetic (**T2DM**) patients have a high prevalence of nonalcoholic fatty liver disease (**NAFLD**). **NAFLD** is a continuum of chronic liver diseases ranging from benign hepatosteatosis to nonalcoholic steatohepatitis (**NASH**), cirrhosis and primary hepatocellular cancer (**HCC**). Because of its strong association with the obesity epidemic, **NAFLD** is rapidly becoming a major public health concern worldwide. Surprisingly, there are no FDA approved **NAFLD** therapies; and current therapies focus on the co-morbidities associated with **NAFLD**, namely, obesity, hyperglycemia, dyslipidemia, and hypertension. The goal of this review is to provide background on the disease process, discuss human studies and preclinical models that have examined treatment options. We also provide an in-depth rationale for the use of dietary  $\omega$ 3 polyunsaturated fatty acid ( **$\omega$ 3 PUFA**) supplements as a treatment option for **NAFLD**. This focus is based on recent studies indicating that **NASH** patients and preclinical mouse models of **NASH** have low levels of hepatic C<sub>20-22</sub>  $\omega$ 3 PUFA. This decline in hepatic PUFA may account for the major phenotypic features associated with **NASH**, including steatosis, inflammation and fibrosis. Finally, our discussion will address the strengths and limitations of  $\omega$ 3 PUFA supplements use in **NAFLD** therapy.

### 1. Introduction

Nonalcoholic fatty liver disease (**NAFLD**) is a continuum of chronic liver diseases ranging from benign hepatosteatosis to nonalcoholic steatohepatitis (**NASH**), cirrhosis and primary hepatocellular cancer (**HCC**) (Lauby-Secretan, 2016; Vernon et al., 2011) (Fig 1). **NAFLD** is the most common chronic fatty liver disease in developed countries (Bellentani et al., 2010); and is strongly associated with obesity (Cohen et al., 2011; Farrell, 2006; Lauby-Secretan, 2016). The Centers for Disease Control and Prevention estimates that ~80 million adults (2015a) and ~13 million children (2015b) in the US are obese (BMI  $\geq$  30); and all

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#### 10. Conflict of interest statement.

The authors declare that there are no conflicts of interest.

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have elevated risk for developing NAFLD. Considering its strong association with obesity, NAFLD is rapidly becoming a major public health problem worldwide (Chalasani et al., 2012a; Kleiner et al., 2005; Loomba, 2013).

NASH, the progressive form of NAFLD was first described in 1980 (Ludwig, 1980). Multiple factors contribute to the onset of NAFLD and its progression to NASH, including lifestyle, gender, ethnicity, genetic polymorphisms (Browning, 2004; Lian et al., 2016; Park, 2008; Smagris et al., 2015; Smagris et al., 2016), health status and systemic factors linked to inflammation and metabolic control (Browning et al., 2004; Cohen et al., 2011; Kanth, 2016; Romeo et al., 2008; Smagris et al., 2015) (Fig 2). The top four risk factors for NAFLD are obesity, dyslipidemia, type 2 diabetes (**T2DM**) and metabolic syndrome (**MetS**) (Alberti et al., 2005; Chalasani et al., 2012b). The prevalence of steatosis and NASH in the obese and T2DM populations is high (60%) (Prashanth et al., 2009). T2DM and chronic liver disease are also risks factors for other chronic diseases, such as cardiovascular disease (Adams et al., 2005; Ekstedt et al., 2006; Sessa, 2017; Soderberg et al., 2010). NASH patients often have elevated blood lipids (dyslipidemia) and metabolic disturbances in cholesterol metabolism similar to patients with coronary artery disease (Kerr, 2012; Lonardo, 2015; Min et al., 2012; Min, 2012; Misra, 2009; Wouters et al., 2008).

Ten to 30% of patients with benign steatosis develop NASH; and NASH patients have higher mortality rates than individuals with benign steatosis; and both are higher than the general population. Twenty to 30% of NASH patients progress to cirrhosis; and over a 10 year period, cirrhosis and liver related deaths occur in 20% and 12% of NASH patients, respectively (Heimbach, 2014; McCullough, 2004). Cirrhosis is a risk factor for HCC (Cohen et al., 2011; McCullough, 2006), the 5<sup>th</sup> and 7<sup>th</sup> most common cancer in men and women, respectively (Bosetti, 2014; Forner et al., 2012; Sanyal et al., 2010). Cirrhosis resulting from NASH is anticipated to be the leading cause of liver transplantation in the United States by 2020. Health care costs linked to NASH are predicted to reach \$35 billion by 2025 (McCullough, 2011).

Government, academia and the pharmaceutical industry have made significant investments in the development of safe and efficacious treatments for NAFLD. Current therapies, however, rely on lifestyle modifications (diet & exercise) and treating the co-morbidities associated with NAFLD (Banini, 2017). The goal of this review is to provide background on the disease process and briefly discuss human studies and preclinical models that have examined the onset and progression of NAFLD as well as specific treatment options. Clinical and preclinical studies have established that NAFLD is associated with significant changes in the type and abundance of hepatic lipids. While hepatic saturated fatty acids (**SFA**) and monounsaturated fatty acids (**MUFA**) increase,  $\omega$ 3 and  $\omega$ 6 PUFA decrease (Arendt, 2015, 2009; Depner, 2013a, b; Petit et al., 2012; Puri et al., 2007; Zheng et al., 2012). Preclinical studies have established that the NAFLD-associated decline in hepatic  $\omega$ 3 PUFA is more severe than the decline in  $\omega$ 6 PUFA (Depner, 2013a; Lytle, 2016, 2017). These changes in lipids are also associated with major changes in urinary and hepatic oxidized lipids (**oxylipins**) that are generated by enzymatic and non-enzymatic mechanisms (Depner, 2012, 2013b). Such changes in the type and abundance of hepatic lipids may set the stage for the adverse events associated with NAFLD. Our discussion will include an in-

depth analysis of how changes in hepatic PUFA affect NAFLD markers, i.e., steatosis, inflammation and fibrosis. We will also provide a brief review of clinical and preclinical studies assessing the impact of dietary  $\omega$ 3 PUFA on NAFLD. Finally, we will address the strengths and limitations of using  $\omega$ 3 PUFA supplementation in NAFLD therapy.

## 2. Why livers store excess fat

Livers store more fat when there is a breakdown of the homeostatic mechanisms controlling lipid metabolism. NAFLD is characterized by the chronic and excessive accumulation of neutral lipids [triglycerides (**TAG**), cholesterol esters (**CE**)] in liver cells (Angulo and Lindor, 2002; Neuschwander-Tetri and Caldwell, 2003) (Fig 1). The key metabolic pathways controlling hepatic lipid metabolism include *de novo* lipogenesis (**DNL**), fatty acid oxidation (**FAO**), formation, assembly & secretion of very low density lipoproteins (**VLDL**), TAG catabolism and the uptake of non-esterified fatty acids (**NEFA**) mobilized from adipose stores.

Parks and colleagues used isotopic methods to determine the source of hepatic lipids in NAFLD patients (Lambert, 2014). Of the TAG in the liver, 59% was derived from NEFA mobilized from adipose tissue, 26% was derived from DNL and 15% was derived from the diet (Donnelly et al., 2005). Mobilization of adipose lipid stores represent the major source of NEFA entering hepatic TAG, particularly in obese patients (Zhang, 2014).

### 2.a. Lipid Mobilized from Adipose Tissue

Mobilization of adipose tissue lipid involves the catabolism of TAG stored in lipid droplets and the export of NEFA to the circulation. This process is mediated by several proteins, including adipocyte triglyceride lipase (**ATGL**), comparative gene identification-58 (**CGI58**), hormone sensitive lipase, monoacylglycerol lipase and perilipin (Brasaemle, 2007; Lass et al., 2011). NEFA released from lipid droplets bind fatty acid binding proteins (**FABP**) and are exported to the systemic circulation as NEFA where they are transported in association with albumin and other proteins. NEFA enter cells via transporters (e.g., fatty acid transport protein-1, **FATP1**; cluster of differentiation-36, **CD36**) or diffusion. Adipocyte TAG catabolism increases in adrenergic stimulation, fasting, exercise and insulin resistance; and decreases with feeding and increases in plasma insulin in healthy individuals.

After entry into cells, NEFA are rapidly converted to fatty acyl-CoAs and assimilated into complex lipids, i.e., neutral (TAG & CE) and membrane lipids. Excess neutral lipid is stored in lipid droplets and contributes to the micro- and macro-steatosis seen in histology of biopsied liver (Fig 2). Obese individuals have a large pool of adipose tissue contributing to blood NEFA. As such, obesity, coupled with T2DM and insulin resistance exacerbates lipid mobilization and NEFA flux from adipose tissue to various tissues, including the liver.

### 2.b. De Novo Lipogenesis

The core enzymes involved in DNL and very long chain fatty acid synthesis include ATP citrate lyase (**Acyl**), acetyl CoA carboxylase (**Acc1**), fatty acid synthase (**Fasn**) and fatty acid elongases (**Elovl 1, 3, 5, 6 & 7**), while the key enzyme involved in MUFA synthesis is

stearoyl CoA desaturase (e.g., **Scd1**) (Fig. 3). Scd1 is a highly regulated enzyme that desaturates saturated fatty acids (**SFA**) to form monounsaturated fatty acids (**MUFA**, e.g., 16:1, $\omega$ 7; 18:1, $\omega$ 7 and 18:1, $\omega$ 9) (Fig 3). Palmitate (16:0), stearate (18:0) and oleate (18:1, $\omega$ 9) are prominent hepatic SFA and MUFA. Humans with NAFLD have increased DNL and palmitoleic acid (16:1, $\omega$ 7) reflecting increased MUFA synthesis (Lambert, 2014; Lee, 2015). SFA and MUFA are substrates for complex lipid synthesis, including triglycerides, diacylglycerols, cholesterol esters, phosphoglycerolipids, plasmalogens and sphingolipids. TAG and CE are stored in lipid droplets, while phosphoglycerolipids, plasmalogens and sphingolipids are membrane lipids.

High sugar consumption from sugary beverages, for example, has been implicated as a major contributor to the obesity epidemic (Chung, 2014; Stanhope and Havel, 2008). Glucose induces DNL through insulin-dependent and insulin-independent mechanisms. Increased blood glucose resulting from sugar (glucose-fructose) or starch (glucose) consumption stimulate insulin secretion from pancreatic  $\beta$ -cells. Insulin, acting through its membrane receptor, stimulates the conversion of glucose to neutral lipids which are stored in adipose tissue and liver (Fig 3) (Lim et al., 2010). Feeding rodents diets high in simple sugar (sucrose) and low in PUFA elevates expression of enzymes involved in hepatic DNL and MUFA synthesis (Jump et al., 1994; Landschulz et al., 1994; Liimatta et al., 1994). Diets supplemented with PUFA ( $\omega$ 3 or  $\omega$ 6) suppress pathways involved in DNL (e.g., fatty acid synthase, **Fasn**) and glycolysis (e.g., L-pyruvate kinase, **L-PK**) (Filhoulaud, 2013; Jump et al., 1994; Jump et al., 2013; Landschulz et al., 1994; Towle, 2005).

### 2.c. Insulin and Glucose Control of DNL and MUFA Synthesis

The mechanisms for hormonal and nutrient control of DNL and MUFA synthesis are complex; and involve the regulation of transcription factors that alter the expression of proteins involved in DNL, fatty acid elongation and MUFA synthesis. Key transcription factors controlling DNL and MUFA synthesis include: sterol regulatory element binding protein-1c (**SREBP1c**) (Hua et al., 1993; Kim and Spiegelman, 1996; Vasandani et al., 2002; Yokoyama et al., 1993); and carbohydrate regulatory element binding protein (**ChREBP**) & its heterodimer partner, Max-like factor-X (**MLX**) (Stoekman et al., 2004; Towle, 2005; Uyeda et al., 2002).

Dietary glucose stimulates insulin secretion from  $\beta$ -cells. Increases in blood insulin activate hepatic cellular signaling pathways through the insulin receptor and downstream signaling components, including insulin receptor substrate (**IRS**) and several kinases: phosphoinositol-3-kinase (**PI3K**), rac- $\beta$ -serine-threonine kinase (**Akt2**), extracellular signal regulated kinase (**Erk1/2**) and target of rapamycin-complex 1 (**mTorc1**). The molecular basis for insulin control of SREBP1 and its regulation of transcription of the Acyl, Acc1, Fasn and Scd1 genes involves increases in transcription of the SREBP1c gene, SREBP1c mRNA stability, proteolytic processing of the precursor of SREBP1c to the mature (nuclear) form of SREBP1c, and inhibition of proteasomal degradation of the nuclear SREBP1 (Botolin et al., 2006; Jump et al., 2013; Li et al., 2012; Owen et al., 2012). As such, insulin increases hepatic nuclear content of SREBP1c, which stimulates transcription of genes

involved in fatty acid synthesis. The net result is increased production of long chain SFA and MUFA (Jump et al., 2013).

The second key transcription factor complex controlling DNL and MUFA synthesis is the ChREBP/MLX heterodimer. Glucose functions as a feed-forward regulator of DNL through an insulin-independent pathway. Increased hepatic glucose metabolism alters the phosphorylation status of ChREBP leading to increased nuclear abundance of the CHREBP/MLX heterodimer (Baraille, 2015; Davies et al., 2008; Ma et al., 2006; Sakiyama et al., 2008; Uyeda and Repa, 2006). Like SREBP1c, the CHREBP/MLX heterodimer binds promoters of *Acc1*, *Fasn*, *Scd1*, L-pyruvate kinase (**L-PK**) leading to the increased gene transcription and increased flow of glucose-derived metabolites into SFA and MUFA synthesis (Jump et al., 2013).

Dietary PUFA, in contrast, inhibit hepatic DNL (Jump et al., 2013), an effect first described by Allman and Gibson (Allmann, 1969). Dietary PUFA also inhibit MUFA synthesis (Landschulz et al., 1994) and the elongation and desaturation of PUFA (Wang et al., 2005; Wang et al., 2006). C<sub>20-22</sub> ω<sub>3</sub> PUFA suppress the expression of the enzymes involved in DNL and MUFA synthesis by targeting mechanisms controlling the nuclear content of SREBP1c and the ChREBP/MLX heterodimer (Botolin and Jump, 2003; Jump, 1993; Jump et al., 2013; Landschulz et al., 1994; Liimatta et al., 1994; Mater et al., 1999; Xu et al., 2006). Interestingly, very long chain dietary ω<sub>3</sub> PUFA (C<sub>20-22</sub>) do not interfere with insulin action or the capacity of insulin to decrease blood glucose. Docosahexaenoic acid (**DHA**, 22:6,ω<sub>3</sub>), the major ω<sub>3</sub> PUFA accumulating in cells inhibits the phosphorylation of Akt2-S<sup>473</sup> site, but not the Akt2-T<sup>308</sup> site. The Akt-T<sup>308</sup> site is phosphorylated by PDK1, a downstream target of insulin signaling. Akt2-S<sup>473</sup>, in contrast, is phosphorylated by multiple kinases (Jump et al., 2013). The specific kinase and/or phosphatase involved in DHA control of Akt2-S<sup>473</sup> phosphorylation status have not been identified to date. The overall effect of DHA is to lower SREBP1c nuclear content through proteasome and Erk1/2-dependent mechanisms (Botolin et al., 2006). PUFA lower the ChREBP/MLX heterodimer by suppressing the nuclear content of MLX, an obligate heterodimer partner for ChREBP binding to promoters of target genes (Xu et al., 2006). Lowering hepatic nuclear content of SREBP1 and ChREBP/MLX suppresses the transcription of the *Acc1*, *Fasn*, *Scd1* and L-PK genes leading to the reduced capacity of glucose conversion to SFA and MUFA.

Dietary PUFA also stimulate FAO by binding directly to the peroxisome proliferator activated receptor-α, (**PPARα**), i.e., a fatty acid-regulated nuclear receptor (Pawar and Jump, 2003; Pawar et al., 2002; Ren et al., 1997; Xu et al., 1999). PPARα regulates gene expression by binding promoters of target genes in association with the retinoid X receptor, i.e., PPARα/RXR. Binding of the PPARα/RXR heterodimeric complex to promoters stimulate the recruitment of co-regulators that are involved in controlling the transcription of the target genes. PUFA bind to and activate PPARα result in increased expression of key enzymes involved in fatty acid oxidation in mitochondria, peroxisomes and microsomes. Representative PPARα/RXR target genes in these 3 compartments include carnitine palmitoyl transferase 1, acyl CoA oxidase and cytochrome P450 4A, respectively (Jump et al., 2013; Ren et al., 1997).

More recent studies have established that dietary PUFA and endogenous production of PUFA regulate TAG catabolism (Tripathy, 2014). Both  $\omega$ 3 or  $\omega$ 6 C<sub>20-22</sub> PUFA increase TAG catabolism by increasing hepatic abundance of ATGL (mRNA and protein) and CGI58 (protein). ATGL and CGI58 act together with perilipin, hormone sensitive lipase, monoacylglycerolipase to promote the hydrolysis of TAG to form NEFA and glycerol. While PUFA induction of FAO requires PPAR $\alpha$  (Ren et al., 1997), PUFA regulation of TAG catabolism involves PPAR $\beta$  (Tripathy, 2014). As such, the type and abundance of hepatic fatty acids play critical roles in controlling hepatic DNL & MUFA synthesis as well as FAO & TAG catabolism.

## 2.d. Diet

NAFLD is a disease of excess calorie consumption (Chung, 2014; Cohen et al., 2011; Leslie, 2015); particularly calories from carbohydrates and fats (SFA and MUFA). Calorie dense diets like the “cafeteria” or “western” diet are linked to the obesity & NAFLD epidemic (Cordain, 2005; Ferolla, 2015; Malhotra, 2015; Pagliassotti et al., 1996; Pickens et al., 2009). The western diet, for example, is moderately high in fat (saturated and trans-fat), simple sugar (sucrose/fructose) and cholesterol. Excessive consumption of carbohydrate, particularly simple sugars (fructose, glucose and sucrose), leads to hexose conversion to glycerol 3-phosphate and acetyl CoA, substrates for SFA, MUFA and TAG synthesis (Fig 3).

While diets like the western diet may be sufficient in essential fatty acids [EFA: linoleic acid (LA, 18:2, $\omega$ 6) and  $\alpha$ -linolenic acid (ALA, 18:3, $\omega$ 3)] dietary EFAs typically represent a small fraction of total dietary fat consumed. Moreover, LA is more abundant than ALA in the western diet (Simopoulos, 2002). These EFAs are converted to C<sub>20-22</sub> PUFA, a process that requires desaturation (Fads1, Fads2), elongation (Elovl2, Elovl5) and peroxisomal  $\beta$ -oxidation (Fig 4). Expression of these desaturases and elongases is regulated by SREBP1c and PPAR $\alpha$ , but not ChREBP/MLX. Peroxisomal  $\beta$ -oxidation is regulated by PPAR $\alpha$  (Jump et al., 2013; Wang et al., 2006).

ALA is the major  $\omega$ 3 PUFA in the human diet. Its conversion to C<sub>20-22</sub>  $\omega$ 3 PUFA in humans, however, is inefficient, ~1–4% of dietary ALA is converted to C<sub>20-22</sub>  $\omega$ 3 PUFA. This inefficiency is due, at least in part, to the high level of dietary LA, relative to dietary ALA, and substrate competition at the enzyme (Fads2) level. Interestingly, when DHA is in excess it is retroconverted to eicosapentaenoic acid (EPA, 20:5, $\omega$ 3) and docosapentaenoic acid ( $\omega$ 3 DPA, 22:5, $\omega$ 3) through a peroxisomal process called retroconversion (Fig 4) (Sprecher, 2000). A recent report indicates that DHA retroconversion to 20:5, $\omega$ 3 predominates over Elovl2- and Fads2-mediated synthesis of EPA to tetracosahexaenoic acid (24:6, $\omega$ 3) (Fig 4) (Park, 2016). While the retroconversion pathway is also inefficient, i.e., ~2% of DHA is converted to EPA in human, this pathway provides a mechanism to increase cellular EPA from dietary DHA (Plourde et al., 2011).

The retroconversion pathway is important physiologically because C<sub>20-22</sub>  $\omega$ 3 PUFA are feedback inhibitors of PUFA synthesis; PUFA suppress the expression of an elongase (Elovl5) and desaturases (Fads1, Fads2) involved in PUFA synthesis (Wang et al., 2005). Clinical studies have established that supplementing the human diet with EPA increases blood levels of EPA and  $\omega$ 3 DPA, but not DHA (Allaire, 2016; Itakura et al., 2011).

Supplementing the human diet with DHA, however, increases blood levels of DHA,  $\omega$ 3 DPA and EPA (Park, 2016). In fact, the only effective way to significantly increase blood and tissue levels of DHA is to supplement the diet with DHA. However, some (Asztalos, 2016; Wei, 2011), but not all investigators (Allaire, 2016) have reported that supplemental DHA increases blood levels of LDL-cholesterol, a risk factor for cardiovascular disease.

### 3. Progression of benign steatosis to NASH

While factors promoting benign steatosis to NASH are not fully established (McCullough, 2006; Vernon et al., 2011), aberrant hepatic metabolism associated with the excessive accumulation of lipid in liver is likely a major driver of this process. Ingestion of excess fat and simple sugar promotes lipotoxicity, oxidative stress, formation oxylipins and advanced glycation end products (AGEP). Together these products contribute to liver cell damage resulting in hepatic inflammation and fibrosis (Angulo, 2002; Ito, 2007; Jialal, 2014; Neuschwander-Tetri, 2010; Sunny, 2017). Hepatic injury resulting inflammation and oxidative stress is monitored clinically by quantifying blood levels of enzymes (alanine aminotransferase, **ALT**; aspartate aminotransferase, **AST**;  $\gamma$ -glutamyl transferase,  **$\gamma$ GT**) released from injured hepatic cells (Cohen et al., 2011; Farrell, 2006; Hashimoto et al., 2011).

In addition to these metabolic events, the progression of benign steatosis to NASH is a multi-cellular process involving hepatocytes, Kupffer cells (resident hepatic macrophage), cholangiocytes (bile duct cells), stellate cells [vitamin A storage and a source of extracellular matrix (**ECM**)] and infiltrating leukocytes (monocytes, macrophage and T-cells) (Day, 1998; LaBrecque et al., 2014; Schuppan, 2013; Tilg and Moschen, 2010).

#### 3.a. Aberrant Macronutrient Metabolism

**3.a.i. Lipotoxicity and Oxidative Stress**—Excessive hepatic content of neutral lipid is associated with lipotoxicity (Sunny, 2017). The predominant fatty acids in neutral lipids are SFA and MUFA; these fatty acids are substrates for sphingolipid (ceramide) and diacylglycerol synthesis. Ceramides promote apoptosis, while diacylglycerols activate protein kinase C.

Turnover of hepatic lipid stores resulting from TAG catabolism or increased NEFA mobilized from adipose depots provides substrates for  $\beta$ -oxidation. However, NEFA can overload the capacity of mitochondria to  $\beta$ -oxidize fatty acids. The outcome of mitochondrial overload is an increase in acyl-carnitines, incomplete fatty acid oxidation, impaired ketogenesis and uncoupled mitochondrial respiration. Uncoupling of mitochondrial respiration generates reactive oxygen species (**ROS**) (Alkhoury, 2009; Sunny, 2017). ROS include highly reactive compounds, such as superoxides, hydrogen peroxide, hydroxy radicals and singlet oxygen (Koves, 2008). These compounds can form adducts with lipids, proteins and nucleic acids and alter cell function.

PUFA are prone to non-enzymatic oxidation. Recent metabolomic studies have established that NASH is associated with increased hepatic levels non-enzymatically derived oxylipins, such as 9,10-hydroxyoctadecenoic acid (9,10-DiHOME), 9- and 13-hydroxyoctadecadienoic

acid (9- & 13-HODE). These fatty acids are generated from C<sub>18</sub> PUFA. NASH is also associated with a decline in urinary oxylipins derived from C<sub>20-22</sub> PUFA, including F2- and F3-isoprostanes and N4-neuroprostanes. The decline in the F2-, F3-isoprostanes & N4-neuroprostanes likely reflects the decline in hepatic (and possibly whole body) C<sub>20-22</sub> PUFA. The appearance of non-enzymatically derived oxylipins in cells, blood or urine serve as markers of tissue and whole body oxidative stress. While some reports suggests iso- and neuro-prostanes are bio-active lipids (Milne and Morrow, 2006; Milne et al., 2011; Signorini et al., 2013) their role of in the onset and progression of NAFLD remains to be defined.

**3.a.ii. Enzymatically-derived oxylipins**—PUFA are substrates for enzymatic oxidation leading to the formation of a large array of bio-active oxylipins including hydroxy-, epoxy-, di- and tri- hydroxy lipids. PUFA assimilated into the sn1 and sn2-positions of membrane phosphoglycerolipids affect membrane fluidity and membrane receptor function (Jump, 2002). Excision of PUFA from membrane phosphoglycerolipids by phospholipases (PLC or PLA2) serve as substrates for oxylipin synthesis (Fig 4) (Calder, 2015; Jump, 2002; Jump et al., 2013). These oxylipins are synthesized by highly regulated enzyme-dependent pathways. The cyclooxygenases (Cox1 & Cox2) are involved in generating prostaglandins, while the lipoxygenases (Lox-5, Lox-12, Lox-15) generate hydroxy-fatty acids. Epoxygenases (Cyp2C, Cyp2J) generate epoxy-fatty acids, while epoxide hydroxylases (EPHX-1 & 2) generate di-hydroxy fatty acids.

Western diet induced NASH is associated with a significant induction of Cox2, Lox5 and Lox15 (Depner, 2013b). Metabolomic analyses established that changes in expression of these enzymes was associated with increased hepatic content of Cox and Lox products, including 6-keto-PGF1 $\alpha$  (a non-enzymatic degradation product of prostacyclin I<sub>2</sub>) and 12-hydroxyeicosatetraenoic acid (12-HETE). In addition, NASH was associated with a decline in hepatic 17, 18-dihydroxyeicosatetraenoic acid (17, 18-DiHETE) and 18-hydroxyeicosapentaenoic acid (18-HEPE). 18-HEPE is a resolvin precursor; resolvins are involved in the resolution of inflammation (Serhan, 2015). 17,18-DiHETE is a product of epoxide hydrolase catalysis of 17,18-epoxyeicosatetraenoic acid (17,18-ETE). 17,18-ETE is the epoxy product of EPA catalyzed by Cyp2C. Changes in epoxy- and dihydroxy PUFA were associated with decreased expression of Cyp2C29, Cyp2C37, but no change in the epoxy-hydrolases (EPHX1 or EPHX2).

Enzymatically derived oxylipins regulate a vast number of pathways that, in many cases, are initiated by ligand (oxylipin) binding to membrane or nuclear receptors that regulate signaling pathways, including the control of inflammation, blood pressure, metabolism and gene expression (Calder, 2015; Farooqui, 2012; Funk, 2001; Jump et al., 2013; Milne et al., 2008; Schuck, 2014; Serhan et al., 2008; Serhan, 2015). Thus, diet-induced changes in hepatic PUFA content, coupled with changes in Cox, Lox, epoxygenase and epoxy hydrolyase activity likely play a major role in the onset and progression of NASH.

In this regard, C<sub>18-20</sub>  $\omega$ 6 PUFA are pro-inflammatory, while C<sub>20-22</sub>  $\omega$ 3 PUFA are anti-inflammatory fatty acids (Fig 4) (Calder, 2015). These characteristics of  $\omega$ 3 and  $\omega$ 6 PUFA are link to the capacity of PUFA to regulate Cox and Lox expression and the bioactivity of the  $\omega$ 6 PUFA versus  $\omega$ 3 PUFA products of Cox and Lox. When compared to  $\omega$ 6 PUFA,  $\omega$ 3



PUFA are poor substrates for Cox and Lox. Moreover, Cox and Lox products of  $\omega$ 3 PUFA typically have weak activity when compared to  $\omega$ 6 PUFA-derived Cox and Lox products. As such, the relative abundance of  $\omega$ 6 to  $\omega$ 3 PUFA in cells impacts the cellular inflammatory status (Calder, 2015; Simopoulos, 2002; Spadaro, 2008). Studies in humans (Lou, 2014) and mice (Depner, 2013a; Lytle, 2017) indicate that a high blood and hepatic  $\omega$ 6/ $\omega$ 3 PUFA ratio is associated with inflammation and NAFLD progression.

**3.a.iii. Advanced glycation end products (AGEP)**—AGEP arising from foods or from aberrant carbohydrate metabolism affect the onset and progression of NAFLD (Gugliucci, 2017; Leung, 2016). For example, the formation of AGEP arising from fructose has been implicated in fructose-induced NASH (Thornalley, 1996; Wei et al., 2013). A metabolomic analysis of a preclinical model of western diet-induced NASH established that hepatic S-lactoylglutathione was decreased (Depner, 2013b), while hepatic glutathione levels were unaffected. S-lactoylglutathione is a degradation product of methylglyoxal and methylglyoxyl is involved in AGEP formation. AGEP are formed from excessive cellular hexose, triose-phosphate and ketone bodies. AGEP form adducts with proteins and lipids impairing cellular function (Gugliucci, 2017).

This brief overview reveals the diverse pathways for generating metabolites that have adverse effects on liver health status. A key goal of therapy is to minimize the substrate availability and/or the activation of these pathways to attenuate the formation of these harmful metabolites.

### 3.b. Inflammation

A major driver of hepatic inflammation is the combined action of circulating factors that promote inflammation (Cani et al., 2007; Depner, 2013b; Harte et al., 2010) as well as necroinflammation resulting from liver injury and cell death. Cell death can result from many of the adverse events described above, including lipotoxicity, oxidative stress (ROS and reactive nitrogen species), AGEP, ER-stress, unfolded protein response, ceramide and PKC induced apoptosis.

Circulating factors promoting inflammation can arise from the gut. For example, endotoxemia is often observed in NASH patients (Higgins et al., 2011). Endotoxin (lipopolysaccharide) is a lipophilic cell wall component of gram negative bacteria. The appearance of microbial components in the blood can significantly affect liver health (Boursier, 2015; Goel, 2014). Multiple factors affect the appearances of gut-derived microbial components in the circulation including bacterial overgrowth and disruption of intestinal tight junctions (Boursier, 2015). Dietary fat also affects the appearance of gut derived microbial components in the circulation. In this instance, bacteria-derived lipophilic components are incorporated into chylomicron and delivered to the lymph and blood (Grunfeld, 2009). The appearance of these factors in the blood activates innate immune responses and elevated levels of circulating chemokines and cytokines (Erridge et al., 2007; Harte et al., 2010; Laugerette et al., 2011).

Circulating levels of chemokines and cytokines are mediated, at least in part, by the activation of toll-like receptors (**TLR**). Endotoxin is a toll-like receptor-4 (TLR4) agonist,

while lipoteichoic acid, a cell wall component of gram positive bacteria, is a TLR2 agonists. These agonists bind CD14, a circulating and membrane associated protein produced in multiple cell types. Ligand-bound CD14 binds to and activates TLR2 and TLR4 signaling pathways. TLR2 heterodimerizes with TLR6 in cell membranes; these heterodimers activate caspase 8 and promote apoptosis. TLR2 also heterodimerizes with TLR1; this heterodimer activates PI3K-, Akt- and IKK/NF $\kappa$ B-linked mechanisms leading to increased nuclear abundance of NF $\kappa$ B. NF $\kappa$ B (p50 and p65) is a key transcription factor controlling the expression of multiple proteins involved in inflammation (Vallabhapurapu and Karin, 2009). CD14-endotoxin-activated TLR4 forms homodimers in the plasma membrane and activate pathways that increase nuclear NF $\kappa$ B and AP1 complexes (Fos & Jun). These events induce the expression of multiple chemokines and cytokines involved in innate immunity (Cengiz, 2015; Lee, 2011; Miura, 2014; Wada, 2016). As such, the appearance of lipophilic microbial components in the blood may play a role in the progression of benign steatosis to NASH.

TLRs are expressed in multiple cell types, including resident hepatic macrophage (Kupffer cells) and cells infiltrating the liver including leukocytes [monocytes, polarized macrophage M<sub>1</sub> (inflammatory); M<sub>2</sub> (anti-inflammatory), & T-cells]. The presence of these cells in the liver and their activation by systemic factors leads to increased local production of cytokines affecting the disease process. Pro-inflammatory agents derived from M<sub>1</sub> macrophage include IL6, IL1 $\beta$ , TNF $\alpha$ , while anti-inflammatory agents derived from M<sub>2</sub> macrophage include IL10. M<sub>1</sub> and M<sub>2</sub> macrophage may also play a role in ECM remodeling since these cells are a source of matrix metalloproteases (**Mmp**).

### 3.c. Fibrosis

Fibrosis (scarring) is a tissue repair and remodeling process that is induced in response to cellular injury (Chalasanani et al., 2012c). This process involves hepatic stellate cell activation and recruitment of myofibrillar cells that produce ECM, including multiple collagen subtypes [e.g., collagen 1A1, 1A2, 2A; Col1A1 is the primary collagen subtype produced in human NASH], elastin, smooth muscle  $\alpha$ 2 actin (smActin), and remodeling enzymes, i.e., crosslinking enzymes [lysyl oxidases, (Lox); lysyl oxidase-like subtypes (LoxL)], protease inhibitors (Timp 1 & 2; plasminogen activator inhibitor 1 [PAI1]) (Friedman, 2008; Koyama, 2015; Rosenbloom, 2013) and macrophage-derived Mmps. Hepatic fibrosis is a risk factor for cirrhosis; and cirrhosis is a risk factor for HCC. In fact, hepatic fibrosis severity correlates with overall mortality and liver-related events, such as cirrhosis and liver transplantation (Angulo, 2015; Dulai, 2017). As such, there is considerable interest in identifying therapies that will prevent the onset, stop the progression and reverse fibrosis.

Transforming growth factor  $\beta$  (**TGF $\beta$** ) is a key regulator of fibrosis (Schuppan, 2013). TGF $\beta$  regulates fibrosis through a canonical and non-canonical pathway (Massague, 2012). In the canonical pathway, TGF $\beta$  regulates the function of the **Smad** [homolog of the drosophila protein, mothers against decapentaplegic and the C. elegans protein for small body size, SMA] family of transcription factors. Activated TGF $\beta$ -receptors (TGF $\beta$ R1 and TGF $\beta$ R2) form a membrane-associated heterodimer with serine/threonine kinase activity. TGF $\beta$  binding to receptors recruits Smads to the membrane leading to Smad2 and Smad3 phosphorylation. Phospho-Smads are translocated to nuclei in association with Smad4.

Phospho-Smads bind promoters and induce expression of target genes, like collagen1A1 (Col1A1), plasminogen activated inhibitor-1 (PAI1), cMyc and Smad 7, an inhibitor of TGF $\beta$  signaling (Akhurst, 2012; Hayashi, 1998; Massague, 2012; Nyati, 2015; Yoshida, 2012).

In the non-canonical pathway, activated TGF $\beta$  receptors regulate EGF receptor signaling that activates RAS/MapK, TGF $\beta$ -activated kinase (Tak1), PI3K and Akt pathways leading to increased expression of proteins involved in fibrosis (Massague, 2012; Nyati, 2015). TGF $\beta$  also interacts with other growth factor regulated pathways, including platelet-derived growth factor (PDGF), connective tissue growth factor (CTGF). Together, these signaling pathways regulate the production of ECM, epithelial-mesenchymal transition (EMT), cell migration and cellular invasion (Yoshida, 2012). Clearly, the onset of NAFLD and its progression to NASH is a very complex process involving multiple cell types and regulatory factors.

#### 4. Current treatment strategies for NAFLD and NASH

NAFLD and its progressive form, NASH, are major chronic liver diseases. While these liver diseases have attracted considerable attention from the academic, medical and pharmaceutical communities, the best strategies for managing NAFLD and NASH remain to be established (Chalasani et al., 2012c; Chan et al., 2007; Leslie, 2015; Vilar-Gomez, 2015). Current strategies focus on lifestyle management (exercise and diet) (Glass, 2015; Sofi, 2014) or pharmaceutical approaches targeting mechanisms regulating weight loss [pentoxifylline, phosphodiesterase inhibitor], blood glucose [metformin, SGLT2 inhibitor, GLP1 agonist], hypertension (telmisartan), carbohydrate and lipid metabolism [ligands for nuclear receptors PPAR( $\alpha$ ,  $\beta$ ,  $\gamma$ ) & FXR] and fibrosis (pirfenidone) (Musso, 2012). Some new, but unapproved pharmaceutical agents, target fatty acid synthesis (ACC inhibitor), oxidative stress (Nox4 antagonist), inflammation (TLRs and the Mcp1 receptor)/CCR2 antagonist and collagen cross-linking (LoxL2 inhibitors) (Kappler, 2015; Koyama, 2015).

Recent meta-analyses of clinical trials have provided evidence in support of the use of weight-loss, exercise, low carbohydrate diets, PPAR $\gamma$  agonist (pioglitazone), FXR agonist (obeticholic acid), vitamin E and  $\omega$ 3-PUFA in the management of NAFLD and NASH (He, 2016a; Musso, 2017, 2012, 2010; Sato, 2015; Sawangjit, 2016; Singh, 2015a, b; Wang, 2015).

Of the pharmaceutical agents tested, all but  $\omega$ 3-PUFA are defined, single biochemicals. Omega-3 fatty acids used in clinical trials range from fish or seal oils to highly refined pharmaceutical grade  $\omega$ 3 PUFA. Pharmaceutical grade  $\omega$ 3 PUFA include a mix of EPA + DHA ethyl esters (Lovaza<sup>TM</sup>, GlaxoSmithKline) or EPA, as ethyl esters [Vascepa<sup>TM</sup>, Amarin Corporation] or a free fatty acid [Epanova<sup>TM</sup>, Astrazenaca]. We are unaware of a pharmaceutical grade DHA used in NAFLD/NASH clinical trials. Instead, DHA-enriched triglycerides have been used in children with NAFLD (Nobili, 2014).

The choice of which  $\omega$ 3 PUFA to use in NAFLD therapy is an important one. Preclinical and clinical studies reveal that EPA and DHA do not have equivalent effects on NAFLD (Depner, 2013a), inflammation (Allaire, 2016; Depner, 2013a) or fibrosis (Depner, 2013a;

Lytle, 2015). As discussed above, EPA is not efficiently converted to DHA *in vivo* (Depner, 2013a; Itakura et al., 2011; Lytle, 2015). Accordingly, we will examine the evidence for the use of  $\omega$ 3 PUFA in the NAFLD therapy. Included in this discussion is the rationale for using  $\omega$ 3 PUFA in children (Boyraz, 2015; Janczyk, 2015; Lassandro, 2015; Nobili, 2013, 2011, 2014; Pacifico, 2015) and adults (Capanni, 2006; Cussons, 2009; Dasarathy, 2015; Li, 2015; Musso, 2010; Nogueira, 2016; Qin, 2015; Sanyal, 2014; Scorletti, 2014; Spardaro, 2008; Zhu, 2008). We also include meta-analyses of these studies (He, 2016b; Lu, 2016; Musso, 2012; Parker, 2012).

## 5. Rationale for using $\omega$ 3 PUFA in NAFLD and NASH therapy

Lipidomic analyses of human liver have established that NAFLD is associated with an increase in all major classes of lipids, including neutral (diacylglycerol, triacylglycerol & cholesterol esters) and membrane lipids (phosphoglycero- and sphingo-lipids) (Puri et al., 2007). The fatty acyl composition of these complex lipids, however, is enriched in SFA and MUFA and depleted of PUFA ( $\omega$ 3 and  $\omega$ 6) (Allard, 2007; Arendt, 2015; Di Minno et al., 2012; Parker et al., 2012; Petit et al., 2012; Zheng et al., 2012). Lipidomic analysis of a preclinical model have replicated these findings. In mice,  $\omega$ 3 PUFA (EPA,  $\omega$ 3 DPA and DHA) appear more vulnerable to diet-induced suppression than C<sub>20</sub>  $\omega$ 6 PUFA, e.g., arachidonic acid (ARA, 20:4, $\omega$ 6) (Depner, 2013a, b; Lytle, 2017). While the low PUFA content in livers of NASH patients may be due to low PUFA consumption (Allard, 2007), the depletion of hepatic PUFA in mouse models of NASH is due to the suppression of hepatic PUFA synthesis by high fat diets (Depner, 2013a, 2012, 2013b; Tripathy et al., 2010; Wang et al., 2006).

Low hepatic C<sub>20-22</sub>  $\omega$ 3 PUFA sets the stage for dysregulated hepatic function affecting lipid synthesis and turnover (Jump et al., 2013) (Wang et al., 2005) (Tripathy, 2014), as well as inflammation and fibrosis (Depner, 2013a; Lytle, 2015, 2017). These changes in hepatic function are linked to changes in hepatic nuclear content (SREBP1, ChREBP/MLX, NF $\kappa$ B and Smad3) or function (PPAR $\alpha$ , PPAR $\beta$ ) of major transcription factors controlling these pathways.

The suppression of inflammation by  $\omega$ 3 PUFA is more complex than just controlling NF $\kappa$ B content. EPA and DHA suppress inflammation by: a) competing with C<sub>20</sub>  $\omega$ 6 PUFA for cyclooxygenase and lipoxygenase-mediated inflammatory eicosanoid production; b) forming Cox generated  $\omega$ 3 eicosanoids that bind weakly to eicosanoid receptors; and 3) forming anti-inflammatory/pro-resolving  $\omega$ 3-PUFA derived oxylipins, e.g., resolvins, protectins, maresins (Fig 4) (Calder, 2015; Depner, 2013a, b; Jump et al., 2013; Lytle, 2017).

As noted previously, DHA is more effective than EPA at lowering systemic inflammatory markers in humans (Allaire, 2016). In this regard, DHA, but not EPA, attenuates western diet-induced hepatic fibrosis by targeting the TGF $\beta$ -Smad3 pathway (Lytle, 2015). DHA suppresses western diet-induced phospho-Smad3 accumulation in hepatic nuclei, a key mediator of TGF $\beta$  induction of ECM production. DHA also attenuates the expression of multiple transcripts linked to fibrosis, including collagen subtypes, ECM remodeling, tissue inhibitors of metalloproteases (TIMPs), matrix metalloproteases (MMPs) and lysyl oxidases.

In contrast to targeted pharmaceutical approaches, the foregoing discussion establishes DHA as a pleiotropic regulator of cell function. DHA has broad effects on multiple signaling pathways controlling liver lipid metabolism, inflammation and fibrosis making it an ideal nutraceutical for NAFLD therapy. Equally important, C<sub>20-22</sub> ω<sub>3</sub> PUFA have proven to be safe and efficacious nutraceuticals. Omega-3 PUFA are used to treat severe hypertriglyceridemia (Barter and Ginsberg, 2008; Pratt et al., 2009) and DHA is included in pre- and post-natal supplements.

## 6. Clinical assessment of ω<sub>3</sub> PUFA in NAFLD and NASH therapy

### 6.a. Children

The Centers for Disease Control and Prevention estimates 13 million children in the US are obese (2015b). This population is vulnerable to developing chronic fatty liver disease. Four clinical trials have examined the impact of ω<sub>3</sub> PUFA supplementation on children with NAFLD. The mean age of the patient population ranged from >5 to <19 years old and included male and females. All patients were diagnosed with chronic fatty liver disease as measured by plasma ALT (> 40 U/L) (Table 1)(Boyraz, 2015; Janczyk, 2015; Nobili, 2013, 2011, 2014; Pacifico, 2015). The ω<sub>3</sub> PUFA type and dose used to treat NAFLD varied: DHA (250 and 500 mg/d), EPA + DHA (450 – 1300 mg/d, matched to body weight) or a mix of ω<sub>3</sub> PUFA-ethyl esters (1000 mg/d). The duration of treatment ranged from 3 to 24 months.

When compared to the placebo group, plasma markers of liver injury (ALT & AST) and dyslipidemia (TAG) were reduced; blood DHA levels increased and hepatosteatosis declined in 3 of the 4 clinical studies in the patients receiving ω<sub>3</sub> PUFA therapy (Boyraz, 2015; Nobili, 2013, 2011, 2014; Pacifico, 2015). Janczyk et al (Janczyk, 2015) did not report blood ω<sub>3</sub> PUFA levels prior to or following treatment. Moreover, they reported no change in plasma triglycerides or hepatic NAFLD markers in response to ω<sub>3</sub> PUFA therapy. The lack of an effect on plasma triglycerides, a well-established marker of C<sub>20-22</sub> ω<sub>3</sub> PUFA action may be due to: a) no dyslipidemia in the patient population; b) high blood levels of C<sub>20-22</sub> ω<sub>3</sub> PUFA in the placebo and treatment groups prior to intervention or; c) failure of the placebo and treatment groups to adhere to the intervention. As such, interventions require an assessment of established markers of ω<sub>3</sub> PUFA action, like blood TAG and DHA content, to establish efficacy of treatment as well as treatment compliance.

There are several limitations in these studies. First, most studies evaluated a low number of participants (participants ranged from 20–56 per group). Three studies used ultrasound and 2 studies used histology to assess steatosis and NASH activity score (NAS). Two studies reporting on liver histology (Nobili, 2014; Pacifico, 2015) indicated that supplemental DHA (250–500 mg/d) decreased hepatosteatosis and/or the NAS. Since no study used EPA alone, it is unclear if children would have benefited from EPA treatment. Nevertheless, the results of 3 of the 4 clinic trials indicate that DHA or the combination of EPA + DHA attenuates liver injury (plasma ALT) and reduces hepatosteatosis. These clinical studies provide reasonable evidence in support of future intervention studies to evaluate the efficacy of C<sub>20-22</sub> ω<sub>3</sub> PUFA therapy in children at risk for the progression of benign steatosis to NASH.

## 6.b. Adults

Three to five percent of the US population have some level of chronic fatty liver disease with liver injury, i.e., NASH (Fig. 1). We examined 11 clinical trials assessing the capacity of  $\omega$ 3 PUFA to reduce NASH markers in adults (Table 2) (Argo, 2015; Capanni, 2006; Cussons, 2009; Dasarathy, 2015; Li, 2015; Nogueira, 2016; Qin, 2015; Sanyal, 2014; Scorletti, 2014; Spardaro, 2008; Zhu, 2008). The studies included male and female patients in both placebo and treatment groups. The mean age of the participants ranged from 33 to 58 years of age. Prior to intervention, all participants had evidence of hepatosteatosis as recorded by ultrasound, proton magnetic resonance imaging, magnetic resonance imaging or histology. Most patients had evidence of liver injury with ALT  $\geq$  40 or NAS  $\geq$  4. The treatment groups received supplemental  $\omega$ 3 PUFA ranging in type and dose: 150 mg/d EPA + 200 mg/d DHA to 50 ml of  $\omega$ 3 PUFA [1:1 EPA to DHA]. One group received seal oil at 2 g/d (Zhu, 2008). The duration of treatment ranged from 2 to 18 months.

All but 2 studies (Dasarathy, 2015; Sanyal, 2014) reported that the  $\omega$ 3 PUFA intervention improved plasma and liver parameters linked to NASH at the conclusion of the treatment. These improvements included a decline in liver injury (ALT) and hepatosteatosis. Hepatic inflammation was reported in 3 studies (Li, 2015; Nogueira, 2016; Qin, 2015); and markers of inflammation were reduced in patients receiving EPA + DHA. Only one group reported improvement in fibrosis scores (Li, 2015). One study reported no improvement in whole body, plasma or hepatic parameters of NASH following  $\omega$ 3 PUFA therapy. This study used a controlled diabetic patient population. The authors reported that patients on vitamin E and/or thiazolidinediones, treatment options for diabetes and NAFLD, were excluded from the study. However, the authors did not report other treatment options used to control diabetes in these patients.

Sanyal et al (Sanyal, 2014) reported no benefit of supplemental EPA (at 1.8 or 2.7 g/d) in any blood or hepatic marker of NASH. The lack of an effect on plasma TAG is disturbing since EPA is used to treat hypertriglyceridemia (Bays et al., 2011). The Japan EPA Lipid Intervention Study (JELIS) trial (Itakura et al., 2011), a multi-center, placebo-controlled, randomized, double blind 12-week study in patients with hypertriglyceridemia reported that EPA (as an ethyl ester) decreased TAG blood levels in hypertriglyceridemic patients. This group also reported that patients taking the EPA supplement had increased plasma EPA and DPA, but no increase in plasma DHA. As described previously, supplemental C<sub>20-22</sub>  $\omega$ 3 PUFA inhibits expression of enzymes (Elovl5, Fads2) involved in converting EPA to DHA (Fig 4) (Wang et al., 2005; Wang et al., 2006).

## 6.c. Meta-Analyses

Four meta-analyses of randomized control trials of  $\omega$ 3 PUFA treatment of NAFLD were examined (Table 3) (He, 2016b; Lu, 2016; Musso, 2010; Parker, 2012). These meta-analyses examined 3–9 clinical trials each, with up to 561 total patients. There is considerable overlap of these meta-analyses with the clinical trials discussed in the previous section. The  $\omega$ 3 PUFA dose ranged from 0.83 to 6.4 g/d and the treatment period was 2 to 18 months in duration. The majority of parameters examined were plasma markers of liver damage (ALT, AST and  $\gamma$ GT) and dyslipidemia (plasma TAG, cholesterol, LDL-C and HDL-C). Only one

study reported on liver fat, i.e., hepatosteatosis (Lu, 2016). The overall outcome indicates that  $\omega$ 3 PUFA treatment improves plasma markers of hepatic damage and TAG, but yielded inconsistent effects on plasma markers for cholesterol (cholesterol, LDL-C and HDL-C). Supplemental  $\omega$ 3 PUFA lowered plasma TAG in patients with hypertriglyceridemia. While limited in the NASH markers assessed, the meta-analyses support the use of  $\omega$ 3 PUFA as an intervention to attenuate dyslipidemia and liver injury associated with NASH severity.

#### 6.d. Summary of Clinical Studies

Investigators using C<sub>20-22</sub>  $\omega$ 3 PUFA as fish oil, a mix of EPA + DHA ethyl esters or DHA (as TAG), but not EPA ethyl esters have shown that supplemental C<sub>20-22</sub>  $\omega$ 3 PUFA attenuates plasma and hepatic markers of NASH, including plasma TAG, hepatic injury (ALT and AST), hepatosteatosis and inflammation. Attenuation of hepatic fibrosis, however, was reported in only 1 study [Table 2, (Li, 2015)]. Lovaza™ (~50:50 mix of EPA- & DHA-ethyl esters) treatment succeeded in reducing hepatic steatosis, but did not improve fibrosis scores. EPA-ethyl esters failed to reduce steatosis or fibrosis (Sanyal, 2014).

These clinical trials reveal a range of effects of  $\omega$ 3 PUFA on NASH which may depend on the type of  $\omega$ 3 PUFA used in the study. Several explanations can account for these outcomes. First, EPA and DHA are not equally effective in regulating the multiple pathways linked to NASH. Clinical and preclinical studies have established that DHA is superior to EPA in controlling steatosis, inflammation & fibrosis (Allaire, 2016; Depner, 2013a; Jump et al., 2013; Lytle, 2015). A recent clinical study established that DHA was a more robust anti-inflammatory nutraceutical than EPA (Allaire, 2016). Second, the failure of EPA to lower liver fat and fibrosis in humans (Sanyal, 2014) is likely due to its poor conversion to DHA *in vivo* (Itakura et al., 2011). Both EPA and DHA suppress the expression of enzymes (*Elovl5*, *Fads1*, *Fads2*) required for PUFA synthesis (Wang et al., 2005). In preclinical studies, DHA, but not EPA, suppressed diet-induced hepatic fibrosis by targeting the TGF $\beta$  pathway (Depner, 2013a; Lytle, 2015). Third, some clinical trials did not establish the efficacy of the  $\omega$ 3 PUFA treatment to increase blood levels of EPA and/or DHA or lower blood TAG (Sanyal, 2014). Fourth, fibrosis remission follows a slower timeline of remission than steatosis. Western diet-fed obese mice develop NASH. When these obese-NASH mice are placed on a low fat-low sucrose diet, obesity and hepatosteatosis resolves before fibrosis (Lytle, 2016). Finally, DHA is retroconverted to EPA in rodents and humans (Depner, 2013a; Park, 2016; Plourde, 2011; Sprecher, 2000). As such supplemental DHA increases both EPA and DHA in tissues, while EPA only increases cellular and blood levels of EPA and DPA (Depner, 2013a; Itakura et al., 2011). EPA and DPA fail to attenuate NASH markers in preclinical studies (Depner, 2013a).

### 7. Development of preclinical models for NAFLD and NASH

Preclinical models of human disease provide an opportunity to study the onset, progression and remission of disease in a well-controlled laboratory environment. Moreover, these models are subject to rigorous interrogation at the physiological, biochemical and molecular level. Such models can be used to evaluate clinical approaches for disease management and identify novel biomarkers to track human disease.

The goal of preclinical studies on NASH is to use an animal model that recapitulates human disease. The ideal preclinical animal model of NASH in the MetS patient should include the following characteristics, obesity, diabetes, hepatosteatosis, inflammation and fibrosis. Several models of NAFLD have been developed where liver disease is induced using specific diets or genetic manipulation (Table 4). Animal models of hepatic fibrosis induced by carbon tetrachloride (CCL4) injection or bile duct ligation are not included in this discussion because these models are not representative of NASH in the MetS patient.

Wild type C57BL/6J male mice fed a high fat diet (HFD, 60% calories as fat [91% lard; 9% soybean oil, Research Diets]) or a high fat-high cholesterol diet [HFHC, 54% calories as fat + cholesterol (0.5 % w/w)] become obese and develop glucose intolerance (diabetes), hepatosteatosis and mild hepatic inflammation (Jump, 2016). Mild hepatic fibrosis develops in the HFHC-diet fed mice, but not the HFD-fed mice, implicating a role for dietary cholesterol in the onset of hepatic fibrosis (Depner, 2012). The atherogenic + cholate (AD + C) and methionine-cholate-deficient (MCD) diets promote hepatosteatosis and inflammation without obesity or evidence of diabetes, i.e., fasting hyperglycemia (Kajikawa, 2010; Suzuki-Kemuriyama, 2016).

*Ob/Ob* mice are hyperphagic resulting from deficient leptin expression; leptin is a satiety hormone produced in adipose tissue. These mice become obese when fed a low fat chow (control or maintenance) diet. Livers of *Ob/Ob* mice become fatty (steatotic) with evidence of mild inflammation (Ibrahim, 2016). Little fibrosis develops in *Ob/Ob* mice because leptin plays a role in the development of fibrosis (Takahashi, 2012). Melanocortin receptor-4 null mice (*Mcr4*<sup>-/-</sup>) mice, like the *Ob/Ob* mice, are hyperphagic and develop diabetes and NASH with fibrosis when fed a HFD (Itoh, 2011) or western diet (Konuma, 2015).

LDL-receptor null mice (*Ldlr*<sup>-/-</sup>) (Bieggs et al., 2012; Depner, 2013a) and the cross of C57BL/6J x 129AS1Svlmj mice [also called Diamond mice (diet-induced animal model of non-alcoholic fatty liver disease)] (Asgharpour, 2016) are lean with no evidence of hyperphagia, diabetes or liver disease when fed a low fat or chow maintenance diet. When fed a western diet, however, both genotypes develop robust obesity, diabetes, hepatosteatosis with significant inflammation and fibrosis. The western diet from Research Diets is moderately high in fat (41% total calories; 95% milk fat, 5% corn oil), hexoses (23% calories) and cholesterol (0.15% w/w). The obesity epidemic in western societies is linked, at least in part, to the consumption of high calorie diets, like the western or cafeteria diets (Arendt, 2015; Cordain, 2005; Depner, 2013a, b; Ferolla, 2015; Lytle, 2016; Malhotra, 2015; Pagliassotti et al., 1996; Pickens et al., 2009).

Of the models described above, *Ldlr*<sup>-/-</sup>, *Mcr4*<sup>-/-</sup> and Diamond mice become obese and develop diabetes and NASH when fed a western diet. These preclinical models appear to recapitulate NASH in obese humans. Although not reported in other preclinical models, *Ldlr*<sup>-/-</sup> mice fed a western diet also develop endotoxemia (Depner, 2013b). These mice also have increased blood levels of agonist activating toll-like receptors (TLR2 and TLR4) (Lytle, 2015). As such, this is another feature that recapitulates human NASH (Harte et al., 2010).



## 8. Preclinical studies of $\omega$ 3 PUFA effects on NAFLD and NASH

C57BL/6J, *Ldlr*<sup>-/-</sup>, *Mcr4*<sup>-/-</sup> and liver-specific *Pten*<sup>-/-</sup> null mice (*L-Pten*<sup>-/-</sup>) have been used to assess the impact of EPA, DHA or the combination of EPA & DHA on liver health (Table 5). While *Ldlr*<sup>-/-</sup> and *Mcr4*<sup>-/-</sup> mice were fed a western diet without or with  $\omega$ 3 PUFA; the C57BL/6J mice were fed the AD + C diet without and with  $\omega$ 3 PUFA. The *L-Pten*<sup>-/-</sup> mice were maintained on a chow maintenance diet without or with  $\omega$ 3 PUFA.

Supplemental  $\omega$ 3 PUFA varied from 2% to 10% of total calories. The western diet is 4.7 kcal/g. Mice consume ~3 grams of food/day or 14.1 kcals. At the 2% level mice consume ~0.28 kcal of  $\omega$ 3 PUFA/day. This level of consumption is similar to the level of  $\omega$ 3 PUFA consumed by humans ( $\omega$ 3 PUFA at 1.8% energy per day) ingesting 2000 kcal/day and taking 4 grams of  $\omega$ 3 PUFA/day to treat hypertriglyceridemia (Barter and Ginsberg, 2008; Pratt et al., 2009). The duration of feeding the test diets ranged from 4 to 72 weeks.

All preclinical models developed steatosis and hepatic inflammation. The level of hepatic fibrosis, however, varied from mild to moderate. No mice developed robust fibrosis resembling human cirrhosis. *L-Pten*<sup>-/-</sup> mice fed a chow diet, however, developed a HCC phenotype that was evident after 40 weeks on the chow diet.

In all 4 preclinical studies,  $\omega$ 3 PUFA supplementation lowered steatosis. This result is expected based on the well-established capacity of  $\omega$ 3 PUFA to target transcription factors controlling the expression of enzymes involved in hepatic lipid synthesis, oxidation and storage. Dietary DHA was better than EPA at lowering hepatic markers of inflammation (Depner, 2013a; Lytle, 2017), including osteopontin, IL2rn, IL1 $\alpha$ , IL1 $\beta$ , Mcp1, TNF $\alpha$ , CD68. Interestingly, the endotoxemia induced by the western diet was not attenuated by DHA (Depner, 2013b). Instead, DHA suppressed hepatic expression of CD14 (mRNA and protein), as well as the expression of TLR2 and TLR4. DHA also suppressed the nuclear abundance of NF $\kappa$ B-p50 as well as its precursor mRNA (Depner, 2013a). These studies establish that DHA attenuates the capacity of cells to respond to inflammation, rather than lowering plasma levels of TLR4 agonist (Depner, 2013b; Lytle, 2015).

At the 2% calorie dose, DHA, but not EPA, attenuated western diet-induced fibrosis as examined by gene expression, Col1A1 protein abundance and histology (Depner, 2013a; Lytle, 2015). Moreover, this effect was linked to DHA attenuation of hepatic TGF $\beta$  signaling. DHA attenuated western diet-induced phospho-Smad3 accumulation in hepatic nuclei. DHA attenuated expression of multiple genes associated with fibrosis, i.e., collagens, Timp 1 & 2, lysyl oxidase, and Lox-like oxidases, subtypes 1–3 (Lytle, 2015). At 10% energy, EPA decreased the severity of HCC in *L-Pten*<sup>-/-</sup> mice. Taken together, these findings established that  $\omega$ 3 PUFA have the capacity to decrease the severity of multiple markers of NASH including steatosis, inflammation and fibrosis.

All of the studies presented in Table 5 use  $\omega$ 3 PUFA in the prevention of diet induced liver disease. A recent study tested whether DHA could reverse NASH in mice with pre-existing disease (Lytle, 2017). *Ldlr*<sup>-/-</sup> mice were fed the western diet for 22 weeks to induce NASH. Mice were either maintained on a western diet supplemented with olive oil or switched to a western diet containing DHA at 2% energy; the two diets were isocaloric. After 8 weeks on

these diet, mice had similar body weights, but blood lipids (triglycerides and cholesterol) in the western diet + DHA group were significantly lower when compared to the western diet + olive oil group. Moreover, hepatosteatosis, inflammation and fibrosis was significantly attenuated in the western diet + DHA group versus the western diet + olive oil group. In fact, the level of hepatic inflammation and fibrosis was less than that seen in mice fed the western diet for only 22 wks. As such, DHA not only blocked western diet-induced NASH progression, it also promoted disease remission.

Lytle et al also carried out a comparative study where mice fed the western diet for 22 weeks were switched to a chow diet (Lytle, 2017). In both a previous study (Lytle, 2016), and the more recent study (Lytle, 2017), switching obese mice with NASH from a western diet to a chow diet reversed nearly all adverse events associated with diet-induced NASH, including obesity, hyperglycemia, dyslipidemia, hepatic lipid content, inflammation, oxidative stress and fibrosis. While this approach can effectively treat NASH in humans, compliance to weight loss therapy may be a complicating factor (Burgess, 2017).

## 9. Future usage of $\omega$ 3 PUFA in NASH therapy

NAFLD and NASH represent major public health concerns worldwide. While changes in diet and lifestyle are effective therapies, such therapies are often difficult to sustain over the long term (Burgess, 2017). As such, safe and efficacious pharmaceutical approaches are needed to prevent the onset of NAFLD and its progression to NASH, cirrhosis and HCC.

We have presented evidence in support of  $\omega$ 3 PUFA supplements containing DHA as a treatment strategy for NAFLD. This strategy is supported by data from the clinical trials in which children and adults were treated with DHA alone or EPA + DHA (Tables 1–3). In nearly every study, dietary  $\omega$ 3 PUFA supplementation lowered blood triglycerides and reduced liver steatosis and injury. Unfortunately, measures of hepatic inflammation and fibrosis were not included in all clinical studies. The preclinical studies of NASH (Tables 4 & 5), however, provided more compelling evidence for the use of DHA in the suppression of diet-induced steatosis, inflammation and fibrosis.

A feature of both the clinical and preclinical studies is that EPA and DHA are not equivalent in their effects on systemic inflammation (Allaire, 2016; Cottin et al., 2011) or NASH parameters (Boyras, 2015; Janczyk, 2015; Nobili, 2013, 2011, 2014; Pacifico, 2015). The preclinical studies in mice (Depner, 2013a, 2013b; Lytle, 2017) support the findings that DHA is superior to EPA in the management of NASH markers, i.e., steatosis, inflammation and fibrosis.

While encouraging, there are limitations to  $\omega$ 3 PUFA usage in the obese diabetic patient. These supplements are not effective therapies for weight loss or the control of blood glucose. As is clear from the recent study by Lytle, et al, supplemental DHA has no effect on body weight or blood glucose (Lytle, 2017). Only after returning the obese mice to a low fat-chow diet was body weight and blood glucose returned to levels seen in the lean control mice (Lytle, 2017). As such,  $\omega$ 3 PUFA supplementation should be combined with other therapeutic approaches to combat obesity and hyperglycemia in the obese and T2DM

patient. Despite this limitation, DHA addition to the western diet blocks NAFLD progression (Lytle, 2017). This is good news for obese patients who struggle with weight loss therapy.

Finally, we have a good understanding of how  $\omega$ 3 PUFA lower liver lipids and attenuate inflammation (Calder, 2015; Jump et al., 2013). In contrast, we have an incomplete understanding of how  $\omega$ 3 PUFA attenuate fibrosis, particularly when DHA attenuates expression of a broad array of proteins involved in fibrosis (Lytle, 2015, 2017). One study suggest that DHA acts directly on human stellate cells to suppress expression of Col1A1 (Lytle, 2015). This effect, however, is not robust. DHA suppresses hepatic TGF $\beta$  signaling in a preclinical model, but this effect may be indirect and involve DHA control of other factors, such as circulating cytokines affecting TGF $\beta$  signaling. As such, future studies are needed to clarify the *in vivo* mechanisms for DHA suppression of TGF $\beta$  signaling and hepatic fibrosis.

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## Abbreviations

<b>ACC</b>	acetyl CoA carboxylase
<b>ACYL</b>	ATP citrate lyase
<b>Akt2</b>	rac- $\beta$ serine-threonine kinase
<b>ALA</b>	$\alpha$ -linolenic acid
<b>ALT</b>	alanine aminotransferase
<b>ARA</b>	arachidonic acid, 20:4, $\omega$ 6
<b>AST</b>	aspartate aminotransferase
<b>ATGL</b>	adipocyte triglyceride lipase
<b>CD36</b>	cluster of differentiation-36
<b>CE</b>	cholesterol esters
<b>CGI58</b>	comparative gene identification-58
<b>ChREBP</b>	carbohydrate regulatory element binding protein
<b>Col1A1</b>	collagen 1A1
<b>COX</b>	cyclooxygenase
<b>DHA</b>	docosahexaenoic acid
<b>DNL</b>	<i>de novo</i> lipogenesis

<b>ECM</b>	extracellular matrix
<b>Elovl</b>	elongase
<b>EGF</b>	epidermal growth factor
<b>EPA</b>	eicosapentaenoic acid
<b>Erk1/2</b>	extracellular signal regulated kinase
<b>FABP</b>	fatty acid binding protein
<b>FASN</b>	fatty acid synthase
<b>FADS</b>	fatty acid desaturase
<b>FAO</b>	fatty acid oxidation
<b>FXR</b>	farnesyl X receptor
<b>γGT</b>	γ-glutamyl transferase
<b>HCC</b>	hepatocellular carcinoma
<b>LA</b>	linoleic acid
<b>LDLR</b>	low density lipoprotein receptor
<b>MCD</b>	methionine choline deficient diet
<b>MetS</b>	metabolic syndrome
<b>MLX</b>	Max-like factor-X
<b>MMP</b>	membrane metalloprotease
<b>mTorc</b>	target of rapamycin-complex
<b>MUFA</b>	monounsaturated fatty acids
<b>NAFLD</b>	nonalcoholic fatty liver disease
<b>NASH</b>	nonalcoholic steatohepatitis
<b>NAS</b>	NASH activity score
<b>NEFA</b>	non-esterified fatty acids
<b>PI3K</b>	phosphoinositol-3-kinase
<b>PPAR</b>	peroxisome proliferator activate receptor
<b>PUFA</b>	polyunsaturated fatty acids
<b>ROS</b>	reactive oxygen species
<b>SCD1</b>	stearoyl CoA desaturase-1

<b>SFA</b>	saturated fatty acids
<b>Smad</b>	homolog of the drosophila protein, mothers against decapentaplegic and the <i>C. elegans</i> protein for small body size, SMA
<b>SREBP</b>	sterol regulatory element binding protein
<b>T2DM</b>	type 2 diabetes mellitus
<b>TAG</b>	triglyceride
<b>TGF</b>	transforming growth factor
<b>TLR</b>	toll-like receptors
<b>Timp</b>	tissue inhibitor of metalloprotease
<b>VLDL</b>	very low density lipoprotein

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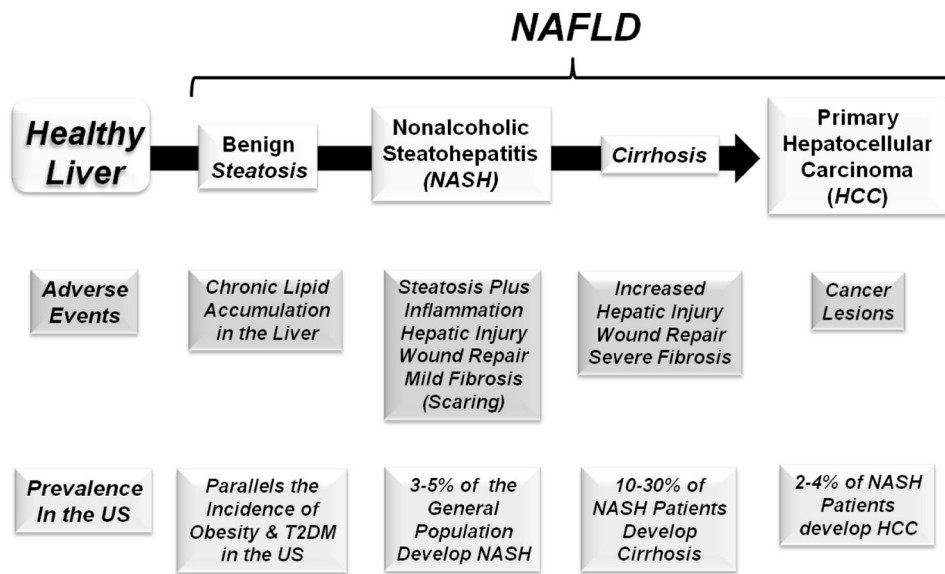
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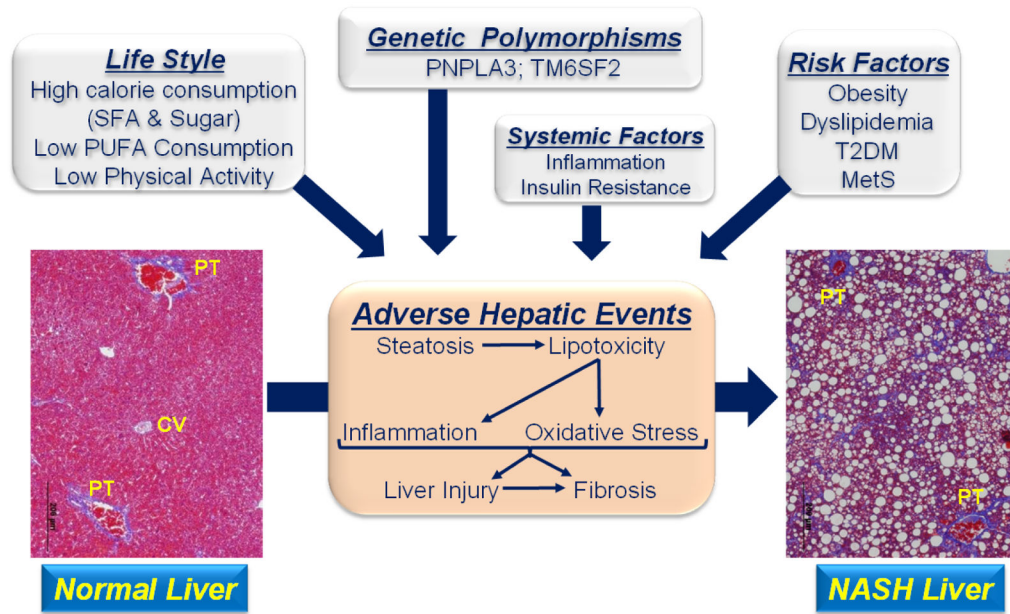
**Figure 1. NAFLD is a continuum of hepatic fatty liver diseases**  
See text for description.

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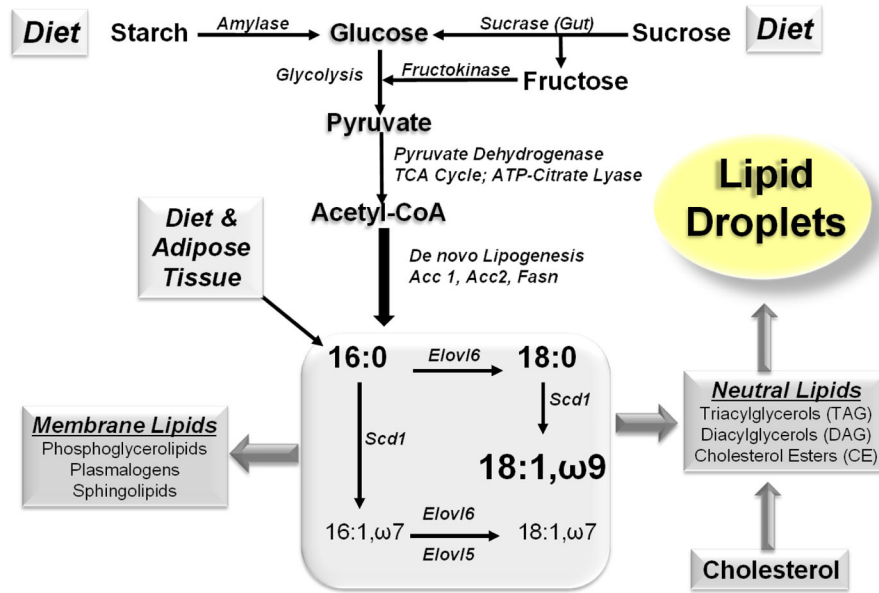
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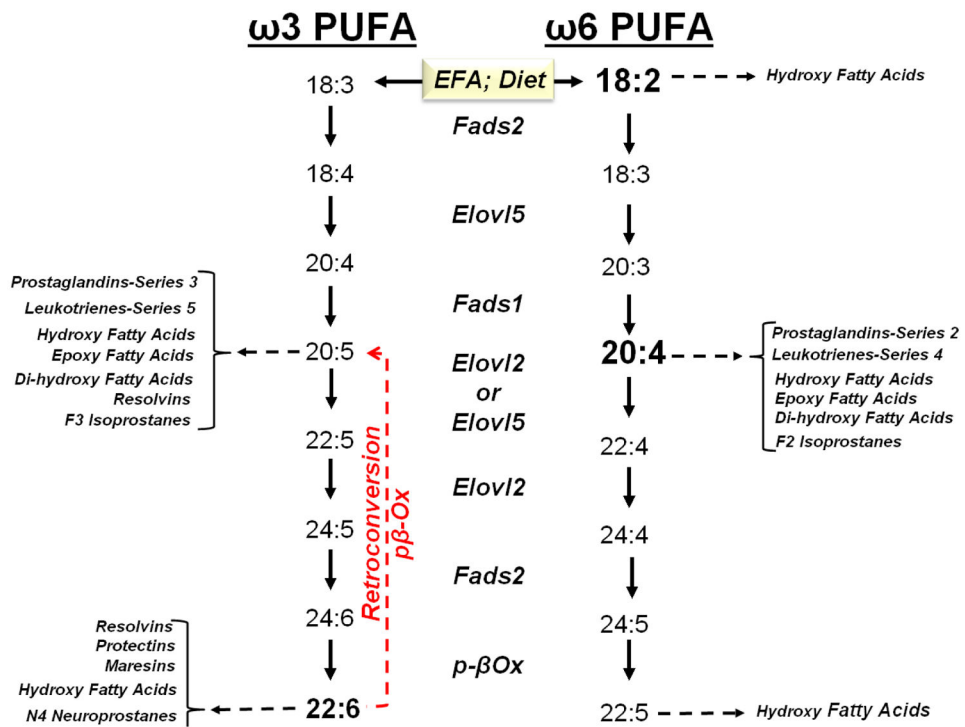
**Figure 2. Factors contributing to the onset and progression of NASH**

Histology from livers of *Ldlr*<sup>-/-</sup> mice fed a western diet for 30 wks (Lytle, 2017). Livers were fixed in formalin, imbedded sliced and stained with trichrome. Steatosis in the NASH liver appears as white circles, while fibrosis appears as blue branching strands of extracellular matrix. CV, central vein; PT, portal track; magnification = 4 x.



**Figure 3. Dietary factors and pathways leading to fat storage**

The diagram illustrates how dietary carbohydrate and fat is assimilated into complex lipids. Lipids are assimilated into membrane lipids or stored in lipid droplets. The diagram does not illustrate the impact of fatty acid oxidation or TAG export as VLDL.



**Figure 4. Conversion of essential fatty acids to C<sub>20-22</sub> ω3 and ω6 PUFA**  
 Essential fatty acids include linoleic acid (LA, 18:2,ω6) and α-linolenic acid (ALA, 18:3,ω3). Long chain PUFA are substrates for oxylipin formation. Oxylipins are generated by enzymatic (cyclooxygenase, lipoxygenase & CYP2C/F) and non-enzymatic processes.

**Table 1**

Clinical trials assessing the impact of  $\omega$ 3 PUFA on Children with NAFLD<sup>1</sup>.

Study (Ref)[Year]	ClinicalTrials.gov	Population	Treatment	Treatment	Key Outcomes
	NCT#	Baseline Values Patients/group	$\omega$ 3 PUFA (Dose, mg/d);	Duration (Months)	$\omega$ 3 PUFA vs Placebo
Nobili et al [2011]	885313	Age: <18 yrs 20 per group ALT > 40 U/L Biopsy Proven NAFLD	Placebo vs DHA 250 & 500 mg/d	6	Plasma ALT ↓ Insulin Sensitivity ↑ Blood DHA ↑ Hepatosteatosis [US] ↓
Nobili et al [2013]	885313	Age: <18 yrs 8M/12F per group ALT 57 U/L	Placebo vs DHA 250 & 500 mg/d	6, 12, 18, 24	Plasma ALT ↓ Blood DHA ↑ Hepatosteatosis [US] ↓
Nobili et al [2014]	885313	Age: <18 yrs Mean Age, 12 8M/12F per group ALT > 57 U/L	Placebo vs DHA 250 & 500 mg/d	6, 12, 18, 24	Serum Cytokines ↓ Blood DHA ↑ NAS ↓ Hepatic GPR120 mRNA ↑ NFκB-pSer <sup>311</sup> ↓
Janczyk, et al [2015]	1547910	Age: >5, <19 yrs Placebo group = 30 $\omega$ 3 PUFA group = 34 ALT 54 U/L	Placebo vs EPA + DHA 450–1300 mg/d Weight Adjusted	3, 6	Blood $\omega$ 3 PUFA [NR] Plasma Triglycerides, [NC] Plasma Adiponectin <sup>†</sup> ; AST ↓ Hepatosteatosis [US]; [NC]
Pacifico, et al [2015]		Placebo: Mean Age: 11 16 Males $\omega$ 3 PUFA: Mean Age: 11 14 males	Placebo vs 250 mg oil/d 39% DHA	6	Body weight ↓; Waist circum. ↓ Plasma ALT [NC]; Insulin ↓ Plasma Triglycerides, ↓ Blood DHA, ↑ Hepatosteatosis [H], ↓
Boyratz, et al [2015]		Placebo: Mean age = 13.8 yrs 27M; 29F ALT, > 56U/L $\omega$ 3 PUFA Mean Age, 13.3 yrs 28 M; 24F ALT, >58 U/L	Placebo vs $\omega$ 3 PUFA Marincap, 1 g/d [FA-EE 18% EPA 12% DHA]	12	Plasma ALT ↓; AST ↓ Fasting Insulin ↓; HOMA-IR ↓ Plasma Triglycerides, ↓ Blood $\omega$ 3 PUFA [NR] Hepatosteatosis [US] ↓

Abbreviations: NAS, NASH activity score; H, Histology; US, hepatic ultrasound; NR, not reported; NC, no change; FA-EE, fatty acid ethyl esters.

**Table 2**

Clinical trials assessing the impact of  $\omega$ 3 fatty acids on Adults with NAFLD<sup>1</sup>.

Study (Ref)[Year]	ClinicalTrials.gov NCT#	Population Baseline Values	Treatment $\omega$ 3 PUFA (Dose, mg/d)	Treatment Duration (Months)	Key Outcomes
Capanni, et al [2006]		Placebo: 9M 5F Mean Age, 58 yrs ALT, 39 U/L $\omega$ 3 PUFA: 23M 19F Mean Age, 57 yrs Mean ALT, 36 U/L	Placebo vs $\omega$ 3 PUFA 375 mg EPA 625 mg/DHA/d	12	$\omega$ 3 PUFA vs Placebo Blood EPA & DHA $\uparrow$ Plasma Triglycerides $\downarrow$ Plasma ALT $\downarrow$ ; AST $\downarrow$ Plasma Glucose $\downarrow$ Steatosis [US] $\downarrow$
Zhu, et al [2008]		Placebo: 50M 18F Mean Age, 51 yrs ALT, 59 U/L $\omega$ 3 PUFA: 11M 7F Mean Age, 50 yrs Mean ALT, 56 U/L	Placebo vs $\omega$ 3 PUFA Seal Oil 2 g/d	2, 3, 4, 6	Blood EPA & DHA [NR] Plasma Triglycerides $\downarrow$ Plasma ALT $\downarrow$ ; AST $\downarrow$ Steatosis [US] $\downarrow$
Spadaro, et al [2008]		Placebo: 8M 10F Mean Age, 58 yrs ALT, 39 U/L $\omega$ 3 PUFA: 23M 19F Mean Age, 57 yrs Mean ALT, 36 U/L	Placebo vs $\omega$ 3 PUFA 375 mg EPA 625 mg/DHA/d	12	Blood EPA & DHA NR Plasma Triglycerides $\downarrow$ Plasma ALT $\downarrow$ ; $\gamma$ GT $\downarrow$ Plasma TNF $\alpha$ $\downarrow$ HOMA-IR $\downarrow$ Steatosis [US] $\downarrow$
Cussons, et al [2009]	620526	PCOS Patients Placebo: 6F Mean Age, 33 yrs ALT, 59 U/L $\omega$ 3 PUFA: 6F Mean Age, 33 yrs Mean ALT, 32 U/L	Placebo vs $\omega$ 3 PUFA 4g/d 27% EPA, 56% DHA 7% other fatty acids	6	Blood EPA & DHA [NR] Plasma Triglycerides $\downarrow$ Steatosis [HMRS] $\downarrow$
Scorletti, et al [2014]		Placebo: 35M 17F Mean Age, 54 yrs ALT 56 U/L $\omega$ 3 PUFA: 25M 26F Mean Age, 49 yrs ALT, 54 U/L	Placebo vs Omacor <sup>TM</sup> [46.5% EPA-EE 37.5% DHA-EE 16% other fatty acids]	15-18	Blood EPA & DHA $\uparrow$ Plasma Triglycerides $\downarrow$ HDL-C $\downarrow$ Steatosis [US] $\downarrow$ Fibrosis [H] NC
Sanyal, et al [2014]	681408	Placebo: 24M 51F Mean Age, 51 yrs	Placebo vs EPA-EE at 1.8 g/d or 2.7 g/d	12	No significant effect on any blood or hepatic marker of NASH

Study (Ref)[Year]	Clinical/Trials-gov NCT#	Population Baseline Values	Treatment ω3 PUFA (Dose, mg/d)	Treatment Duration (Months)	Key Outcomes ω3 PUFA vs Placebo
Argo, et al [2015]	681408	<p>Mean ALT 54 U/L 1.8 g/d EPA-EE 33M 54F</p> <p>Mean Age, 48 yrs Mean ALT, 51 U/L 2.7 g/d EPA-EE 25M 61F</p> <p>Mean Age, 46 yrs Mean ALT, 52 U/L</p> <p>Placebo: 7M 10F</p> <p>Mean Age, 47 yrs ALT, 60.8 U/L</p> <p>Liver Fat (MRI) 11.3% NAS: 4.9</p> <p>ω3 PUFA: 6M 11F</p> <p>Mean Age, 46 yrs ALT, 72.8 U/L</p> <p>Liver Fat (MRI) 18.3% NAS: 45.4</p> <p>Fibrosis Score: 1.6</p> <p>Fibrosis Score: 2.1</p>	<p>Placebo vs ω3 PUFA 3g/d 70% ω3 PUFA; 35% EPA, 25% DHA</p>	12	<p>Plasma ω6:ω3 ratio ↓ Steatosis [MRI] ↓ Fibrosis [NC]</p>
Li, et al [2015]		<p>Placebo: 14M 25F</p> <p>Mean Age, 50 yrs ALT, 91 U/L</p> <p>ω3 PUFA: 13M 26F</p> <p>Mean Age, 54 yrs Mean ALT, 89 U/L</p>	<p>Placebo vs 50 ml ω3 PUFA 50% EPA 50% DHA per day</p>	6	<p>Blood EPA &amp; DHA NR Plasma ALT ↓; AST ↓ Plasma Triglycerides ↓ Plasma Cholesterol ↓ Hepatic Steatosis [H] ↓ Hepatic Inflammation [H] ↓ Hepatic Fibrosis [H] ↓</p>
Dasarathy, et al [2015]	323414	<p>Controlled Diabetics with NASH</p> <p>Placebo: 2M 17F</p> <p>Mean Age, 50 yrs ALT, 49 U/L NAS 4</p> <p>ω3 PUFA: 6M 12F</p> <p>Mean Age, 52 yrs Mean ALT, 48 U/L NAS &gt; 4</p>	<p>Placebo vs ω3 PUFA 2.2 g/d EPA 1.4 g/d DHA</p>	12	<p>ω3 PUFA provided no significant benefit over placebo in patients with diabetes</p>
Qin, et al [2015]	ChiCTR-TRC 12002380	<p>Placebo-Corn Oil 25M 9F</p> <p>Mean Age, 44 Alt 33 U/L</p> <p>ω3 PUFA 26M 10F</p> <p>Mean Age, 46 Alt 32 U/L</p>	<p>Placebo Corn oil in 4 g oil capsules ω3 PUFA: 728 mg EPA + 516 mg DHA in 4g oil capsules</p>	3	<p>Blood/Plasma Parameters: EPA &amp; DHA ↑ Total Cholesterol ↓ Triglycerides ↓ ApoB ↓ Plasma Alt ↓ Plasma γGT ↓ Adiponectin ↓ LT-B4 ↓ FGF21 ↓</p>

Study (Ref)[Year]	Clinical/Trials.gov NCT#	Population Baseline Values	Treatment ω3 PUFA (Dose, mg/d)	Treatment Duration (Months)	Key Outcomes
Nogueira, et al [2016]	1992809	Placebo: 5M 18F Mean Age, 54 yrs ALT, 47 U/L NAS 5.4 ω3 PUFA: 5M 25F Mean Age, 53 yrs Mean ALT, 45 U/L NAS 5.0	Placebo vs ω3 PUFA 0.6 g/d ALA 0.15 g/d EPA 0.2 g/d DHA	12	ω3 PUFA vs Placebo CK18 ↓ PGE2 ↓ Blood EPA & DHA ↑ Plasma Triglycerides ↓ Plasma ALT ↓; AST ↓ Hepatic lobular Inflammation ↓

/ Abbreviations: γGT, γ-glutamyl transferase; EE, ethyl esters; US, ultrasound, HMRS, hepatic proton magnetic resonance spectroscopy; MRI, magnetic resonance imaging, H, histology; NAS, NASH activity score; NR, not reported; LT-B4, leukotriene-B4; FGF21, fibroblast growth factor 21; CK18, cytokeratin 18 fragments M30; PGE2, prostaglandin E2.



**Table 3**

Summary of meta-analyses of RCTs for  $\omega$ 3 PUFA treatment of Adults with NAFLD<sup>1</sup>.

	Clinical Trials	Patients	$\omega$ 3 PUFA Dose, g/d	Duration (Months)	Meta-Analysis		Outcomes	
					IV, Random	95% CI <sup>L</sup>		
Musso, et al (2010)								
Plasma ALT	3	182	2 to 4	2 to 6	-32.24	-39.99	-24.48	$\omega$ 3-PUFA
Parker, et al, (2012)					Hedge's G			
Liver Fat	7	331	0.83 to 5	2 to 12	-0.965	-1.348	-0.582	$\omega$ 3-PUFA
Plasma ALT					-0.563	-1.153	-0.031	$\omega$ 3-PUFA
Plasma AST					-0.971	-1.815	-0.127	$\omega$ 3-PUFA
Lu, et al (2016)					M-H Random			
Liver Fat	5	268	0.83 to 5	12	3.6	1.31	9.98	$\omega$ 3-PUFA
Plasma ALT	8	530	0.83 to 4	2 to 18	-4.97	-11.14	1.2	NE
Plasma AST	7	496	0.83 to 4	2 to 18	-2.01	-8.72	4.7	NE
HDL-C	7	472	0.83 to 9	2 to 18	5.11	-0.03	11	$\omega$ 3-PUFA
LDL-C	6	433	0.83 to 9	2 to 18	1.28	-4.06	6.63	NE
Total Cholesterol	7	474	0.83 to 4	2 to 18	-3.65	-10.4	3.09	NE
Triglycerides	9	561	0.83 to 4	2 to 18	-35.55	-53.9	-17.19	$\omega$ 3-PUFA
He, et al (2016)					IV Fixed			
Plasma ALT	6	396	0.83 to 6.4	6 to 18	-7.61	-12.38	-2.39	$\omega$ 3-PUFA
Plasma $\gamma$ GT	3	181	0.83 to 2.0	6 to 12	-8.28	-18.38	1.83	$\omega$ 3-PUFA
Triglycerides	7	442	0.83 to 6.4	6 to 18	-43.96	-51.21	-36.71	$\omega$ 3-PUFA
Plasma AST	5	362	0.83 to 6.4	6 to 18	-6.89	-17.71	3.92	$\omega$ 3-PUFA
Plasma Cholesterol	6	396	0.83 to 6.4	6 to 18	-13.41	-21.44	-5.38	$\omega$ 3-PUFA
LDL-C	6	360	0.83 to 6.4	6 to 18	-7.31	-14.26	0	$\omega$ 3-PUFA
HDL-C	6	398	0.83 to 6.4	6 to 18	6.97	2.05	11.9	$\omega$ 3-PUFA

<sup>1</sup> Abbreviations: CI<sup>L</sup>, confidence interval-low value; CI<sup>H</sup>, confidence interval-high value; NE, no effect; ALT, alanine aminotransferase; AST, aspartate, aminotransferase,  $\gamma$ GT,  $\gamma$ -glutamyl aminotransferase.

Table 4

Comparison of preclinical models of NAFLD<sup>1</sup>

	<u>Obesity</u>	<u>Diabetes</u>	<u>Hepatosteatosis</u>	<u>Inflammation</u>	<u>Fibrosis</u>	<u>Reference</u>
<b>Dietary Models</b>						
<u>Genetic Background</u>		<u>Diet</u>				
Mice, C57BL/6J	+	+	+	-/+	-	Jump, et al., 2016
Mice, C57BL/6J	+	-/+	+	+	+	Jump, et al, 2016
Mice, C57BL/6J	-	-	+	+	-	Suzuki-Kemuriyama, et al., 2016
Rats, Wistar	-	-	+	+	+	Ibrahim, et al., 2016
<b>Genetic Models</b>						
<u>Genetic Background</u>		<u>Diet</u>				
<i>Ob/Ob</i>	+	+	+	+	-	Ibrahim, et al., 2016
<i>Ldlr</i> <sup>-/-</sup>	-	-	-	-	-	Depner, et al 2013
<i>C57BL/6J x 129AS1/Sv1mJ</i>	-	-	-	-	-	Asgharpour, et al., 2016
<i>Ldlr</i> <sup>-/-</sup>	+	+	+	+	+	Depner, et al., 2013
<i>C57BL/6J x 129AS1/Sv1mJ</i>	+	+	+	+	+	Asgharpour, et al, 2016
<i>Mcr4</i> <sup>-/-</sup>	+	+	+	+	+	Konuma, et al., 2015 Itoh, et al., 2011

<sup>1</sup> Abbreviations: HFD, high fat diet, 60% energy as fat; HFHC, high fat-high cholesterol diet; LFD-Chow ( 10% calories as fat) AD + C, atherogenic diet + cholate; MCD, methionine-choline deficient diet; *Ob/Ob*, leptin deficient mouse; *Mcr4*<sup>-/-</sup>, melanocortin receptor-4 null mouse; *Pten*<sup>-/-</sup>, phosphatase and tensin homolog null mouse; *Ldlr*<sup>-/-</sup>, low density lipoprotein receptor null mouse.

Table 5

Effect of dietary  $\omega$ 3 PUFA on the onset and progression of NASH and primary hepatocellular carcinoma (HCC)<sup>1</sup>.

<u>Mouse Model</u>	<u>Diet</u>	<u><math>\omega</math>3 PUFA Dose<sup>2</sup></u>	<u>Treatment Duration wks</u>	<u>Hepatic Phenotype</u>	<u><math>\omega</math>3 PUFA Effect</u>	<u>Reference</u>
<i>Ldlr</i> <sup>-/-</sup>	Western	EPA or DHA 2% energy	16	NASH [S, I, OS, F]	DHA>EPA Decreased S, I, OS, F	Depner, et al [2013]
<i>Mcr4</i> <sup>-/-</sup>	Western	EPA 9.6% energy	24	NASH [S, I, F]	EPA Decreased S, F, NAS	Nonuma, et al [2015]
<b>C57BL/6J Wild type</b>	AD + C	EPA or DHA 9% energy	4	Mild NAFLD [S, I, OS, F]	EPA = DHA Decreased S, I, OS, F	Suzuki-Kemuriyama, et al [2016]
<i>L-Pten</i> <sup>-/-</sup>	Chow	EPA 10% energy	40 & 72	HCC	EPA Decreased S, I, OS, CP-A	Ishii, et al. [2009]

<sup>1</sup> Abbreviations: AD + C, atherogenic diet + cholate; S, steatosis; I, inflammation; F, fibrosis; OS, oxidative stress; NAS, NASH activity score; CP-A, cell proliferation-apoptosis; HCC, primary hepatocellular carcinoma.

<sup>2</sup> The dose of  $\omega$ 3 PUFA in some reports was reported as weight % of diet. To provide consistency amongst studies, the  $\omega$ 3 PUFA dose was recalculated and represented as % energy in the diet.