Dysregulated lipid metabolism links NAFLD to cardiovascular disease

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ABSTRACT

Background: Non-alcoholic fatty liver disease (NAFLD) is rapidly becoming a global health problem. Cardiovascular diseases (CVD) are the most common cause of mortality in NAFLD patients. NAFLD and CVD share several common risk factors including obesity, insulin resistance, and type 2 diabetes (T2D). Atherogenic dyslipidemia, characterized by plasma hypertriglyceridemia, increased small dense low-density lipoprotein (LDL) particles, and decreased high-density lipoprotein cholesterol (HDL-C) levels, is often observed in NAFLD patients.

Scope of review: In this review, we highlight recent epidemiological studies evaluating the link between NAFLD and CVD risk. We further focus on recent mechanistic insights into the links between NAFLD and altered lipoprotein metabolism. We also discuss current therapeutic strategies for NAFLD and their potential impact on NAFLD-associated CVD risk.

Major conclusions: Alterations in hepatic lipid and lipoprotein metabolism are major contributing factors to the increased CVD risk in NAFLD patients, and many promising NASH therapies in development also improve dyslipidemia in clinical trials.

Keywords Dyslipidemia; NAFLD; Cardiovascular disease; Metabolic syndrome

1. INTRODUCTION
Non-alcoholic fatty liver disease (NAFLD) prevalence is estimated at nearly 25% worldwide due to its close association with metabolic disorders such as obesity and type 2 diabetes (T2D) [1] and will soon nearly 25% worldwide due to its close association with metabolic disorders. NAFLD encompasses a wide spectrum of liver pathologies ranging from isolated hepatic steatosis (non-alcoholic fatty liver, NAFL) to non-alcoholic steatohepatitis (NASH), which combines steatosis with hepatocyte ballooning and inflammation and favors development of fibrosis [2]. NAFLD can be divided into three stages with increasing severity: hepatic steatosis, NASH without fibrosis, and NASH with fibrosis. Disease progression from steatosis to NASH and fibrosis is heterogeneous and occurs over years or even decades. Consequently, the mechanisms involved in the evolution of NAFLD remain poorly understood. However, numerous lines of evidence point to alterations in hepatic and extra-hepatic lipid metabolism as central drivers [3]. For example, mutations in several genes involved in the control of lipid metabolism (for example, patatin-like phospholipase domain containing 3, PNPLA3 [4], transmembrane 6 superfamily member 2, TM6SF2 [5], farnesyl-diphosphate farnesyltransferase 1, FDFT1 [6], and membrane bound O-acyltransferase domain containing 7, MBOAT7 [7]) are associated with increased risk of NAFL or NASH. Importantly, NASH also increases the risk of extra-hepatic complications, especially cardiovascular diseases (CVD), which are among the most common causes of death in NASH patients [8]. Indeed, the alterations in hepatic lipid metabolism that lead to NAFLD also drive the development of atherogenic dyslipidemia, especially elevated plasma triglycerides (TG) and remnant lipoprotein cholesterol levels, and small dense LDL particles that infiltrate the arterial wall and promote the development of atherosclerotic plaques. Altered glucose metabolism and insulin resistance, also hallmarks of NAFLD, can further exacerbate CVD risk in these patients. In this review, we summarize and analyze current data on the links between NAFLD and CVD with a specific focus on the mechanisms of dyslipidemia and their effect on outcomes for patients with NASH.

2. EPIDEMIOLOGY
Although NAFLD shares many common risk factors with CVD (that is, obesity, insulin resistance, T2D, and atherogenic dyslipidemia), there is emerging evidence that NAFL and NASH directly impact CVD risk. This section highlights recent studies assessing the relative contributions of insulin resistance, dyslipidemia, and inflammation to NAFLD-associated CVD.
2.1. NAFLD drives CVD risk independent of obesity, insulin resistance, T2D, and atherogenic dyslipidemia

Several recent studies have investigated the risk of fatal and non-fatal cardiovascular (CV) events in cohorts of NAFLD patients, and there is general agreement that all stages of NAFLD (isolated steatosis and NASH, among others) can increase the risk of CV events such as myocardial infarction, stroke, revascularization, or death. For example, compared to individuals without NAFLD, patients with fatty liver already show an elevated risk of CV events independent of metabolic syndrome risk factors, and this risk further increases when fibrosis is present [9,10]. Surprisingly, even in patients with body mass index (BMI) < 25 kg/m², ultrasound defined-NAFLD is associated with a higher incidence of CV events, indicating that NAFLD may act independent of obesity [11]. Importantly, these observations have been partly confirmed in recent meta-analyses. A first meta-analysis of 16 studies showed an increased risk of non-fatal CV events in NAFLD patients compared to those without NAFLD [12]. However, the risk of CVD mortality primarily increased in patients with severe NAFLD, as defined by the concomitant presence of imaging-diagnosed steatosis and elevated gamma-glutamyl transpeptidase (GGT), high NAFLD fibrosis scores (NFS), high hepatic fluorodeoxyglucose (FDG) uptake, and/or high histological fibrosis stage. A second smaller meta-analysis [13] generally confirmed these findings, despite methodological weaknesses such as inclusion of low numbers of NASH patients (discussed in detail in Liu et al. [14]). Surprisingly, a third meta-analysis found increased liver-related but not CVD mortality in patients with NAFLD [14].

One major limitation of these studies lies in the diagnostic criteria of NASH. The previously described meta-analyses combined studies using a variety of non-invasive diagnostic measures and relatively few patients with histological evaluation of NASH (the current gold standard). Further long-term assessment of larger numbers of histologically diagnosed patients is essential to improve our understanding of the causes of mortality in NASH. NASH diagnosis may result in more aggressive management, potentially leading to reduced CVD-related events and mortality [14].

In conclusion, these studies suggest that NAFLD increases the risk of CV events independent of other major CVD risk factors (such as dyslipidemia, hypertension, obesity, T2D, and insulin resistance) and suggest that other factors link the liver to arterial wall pathologies. CV events such as myocardial infarction or stroke are the result of dysfunction of the vascular system leading to atherosclerosis. This raises the question of whether NAFLD increases CV risk by favoring the development of atherosclerosis.

2.2. Association of NAFLD and atherosclerosis

Atherosclerosis, characterized by the development of neo-intimal plaques in large arteries, drives CV events such as myocardial infarction and stroke. Atherosclerotic plaques develop progressively over several decades, during which the plaque composition changes through distinct processes such as lipid-deposition, inflammation, fibrosis and calcification (reviewed in detail by Libby et al. [15]). Evaluation of subclinical atherosclerosis commonly relies on non-invasive detection of plaque features such as lipid deposition, calcification, and inflammatory cells. Common methods include measuring the coronary artery calcification (CAC) score via computed tomography (CT) and estimating vascular inflammation by FDG position emission tomography (PET). Other methods evaluate plaque size by measuring carotid intima-media thickness (CIMT) or arterial stiffness via brachial-ankle pulse wave velocity (ba-PWV). Some invasive procedures also exist (for example, intravascular ultrasonography, optical coherence tomography, and invasive angiography), although these are generally not used for the initial investigation of atherosclerosis [15]. All these methods have been validated as measuring markers of CV risk.

In line with epidemiological evidence of increased risk of CV events, several studies demonstrated increased subclinical atherosclerosis in NAFLD patients by measuring calcification and aortic stiffness. Cross-sectional studies have shown that ultrasound or magnetic resonance spectroscopy (MRS)-diagnosed NAFLD patients display higher CAC scores than those without NAFLD [16,17], even among patients with BMI < 25 kg/m² [16]. This was further confirmed in a meta-analysis including 12 studies [18]. Interestingly, the annual rate of CAC progression was significantly higher in participants with ultrasound defined-NAFLD compared to those without NAFLD at baseline and was further increased in subjects with elevated NFS independent of obesity, hypertension, dyslipidemia, and diabetes [19]. CIMT is also increased in patients with NAFLD (diagnosed by ultrasound or CT) independent of dyslipidemia and hypertension [17,20]. Interestingly, a prospective study showed that increased CIMT and ba-PWV at baseline were associated with a higher risk of developing NAFLD (assessed by ultrasound) and ba-PWV was associated with a higher likelihood of fibrosis as assessed by the NFS, fibrosis-4 (FIB4), and aspartate transaminase (AST) to platelet ratio index (APRI) scores [21]. This suggests that vascular dysfunction could also drive NAFLD development. However, more studies with bona fide histological assessment of NASH and fibrosis are needed to further explore this hypothesis. Interestingly, some studies failed to find an association between ultrasound-defined NAFLD and subclinical atherosclerosis by measuring ba-PWV [20], carotid to femoral (cf-) PWV [22], or CAC-defined calcification (higher CAC cutoff >10) [23] after adjustment for other risk factors.

While most studies focused on coronary artery plaque calcification, an investigation by CT of several locations (carotid artery, coronary artery, thoracic aorta, iliac artery, renal artery, celiac trunk, and superior mesenteric artery) found a positive association of NAFLD (diagnosed by CT) with calcification in the thoracic aorta and celiac trunk but not in the coronary artery [24]. Moreover, this study demonstrated that NAFLD patients are more susceptible to developing multi-arterial calcification. Surprisingly, two studies indicated that NAFLD is associated with non-calcified plaques measured by cardiac CT rather than with calcified plaques, the former possibly being more vulnerable to rupture and subsequent CV events [23,25]. In line, hepatic steatosis (diagnosed by CT or ultrasound) is associated with increased plaque FDG-PET/CT, an indicator of plaque vulnerability [26,27].

In conclusion, the majority of studies indicate that NAFLD increases the risk of atherosclerosis and seems to favor the development of unstable plaques, adding to traditional CV risk factors such as obesity, diabetes, hypertension, and dyslipidemia. However, adjustments for systemic inflammation (for example, by plasma C-reactive protein (CRP) or interleukin 6 (IL-6) levels) have not always been correctly performed. Considering the role of inflammation as a driver of atherosclerosis, future studies should focus on the contribution of inflammation vs other factors in defining the NAFLD-associated increase in CVD risk. Still, genetic evidence points to NAFLD-associated dyslipidemia as an important contributor to elevated CVD risk. Intriguingly, certain genetic polymorphisms favoring the development of NAFLD unexpectedly decrease CV risk (discussed as follows).

2.3. Genetic variation dissociates NAFLD and CVD incidence

A number of genetic studies have identified variants associated with an increased risk of NAFLD development. Among the most frequent are PNPLA3 rs738409 I148M, which affects remodeling of fatty acid chains in liver TG [26], TM6SF2 rs58542926 E167K, which reduces
hepatic VLDL secretion [5], and MBOAT7 rs641738 [7], an enzyme involved in phospholipid acyl-chain remodeling, with respective minor (risk) allele frequencies of 24%, 7%, and 43% (based on CARDioGRAMplusC4D consortium data [29]). While these variants were identified based on MRS-screened hepatic steatosis, the impact of the PNPLA3 and TM6SF2 variants has been confirmed in histologically diagnosed NAFLD [30,31]. However, although steatosis in NAFLD is thought to be the result of increased lipogenesis and increased flux of adipose-derived FA [32,33], there are currently no data linking these variants to these pathways.

Given the impact of these variants on hepatic lipid metabolism, their modulatory role in CVD risk has also been investigated. The PNPLA3 [34–36] and TM6SF2 [34,36–38] NAFLD risk variants are associated with a lower-risk lipoprotein profile characterized by lower plasma TG and/or LDL-C levels. Interestingly, the effect size of the TM6SF2 E167K variant on plasma TG levels is comparable to that of a lipoprotein lipase (LPL) activating variant [34]. This translates into a modest protection of E167K carriers (~5–15% depending on the study) against CVD development [34,39–41]. The effect on CVD risk is less clear for PNPLA3 I148M carriers. Association studies have shown either no effect [29,42–44] or modest protection [34] from coronary artery disease (CAD) when comparing I148M carriers to non-carriers. Two studies that specifically matched for NAFLD presence (by histology or ultrasound) found no protection from CAD in I148M carriers [43,45].

Unexpectedly, genetic polymorphisms associated with NAFLD tend to decrease CVD risk, indicating that high levels of liver fat, by any mechanism, are not sufficient to increase CVD risk. Nevertheless, these findings indicate a role for NAFLD-driven dyslipidemia as an important mediator of elevated CVD risk.

### 3. Dyslipidemia: Linking Hepatic Lipid Metabolism and the Heart

There are likely several mechanisms behind the close association between CVD and NAFLD. Insulin resistance is an important driver of NAFLD and especially NASH. It affects several processes, such as dyslipidemia, hyperglycemia, activation of oxidative stress and inflammation, endothelial dysfunction, and ectopic lipid accumulation, together creating a pro-atherogenic environment favoring CVD development (reviewed in detail Ormazabal et al. [46]). Additionally, hepatic and circulating immune cell populations are highly correlated in patients with NASH [47] and display alterations comparable to those observed in CVD, suggesting a common immune-inflammatory landscape (reviewed in detail by Gehrke et al. [48]).

Atherogenic dyslipidemia, characterized by plasma hypertriglyceridemia, increased small dense LDL particles, and decreased HDL-C levels, is variably present in patients with NAFLD [49–52]. The liver plays a central role in lipoprotein metabolism as it participates in the production and/or clearance of all classes of lipoprotein particles. In

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**Figure 1:** A summary of hepatic lipid metabolism pathways altered in NAFLD and driving dyslipidemia. Increased hepatic TG in NAFLD is a result of several processes. Elevated plasma insulin and glucose levels respectively activate the LXR and ChREBP pathways, which increase de novo lipogenesis (DNL). Through the action of ACC, DNL increases the concentration of malonyl-coA, leading to inhibition of CPT1 and consequently reducing fatty acid oxidation (FAO) and mitochondrial function. In parallel, LXR increases the expression of ANGPTL8 and 3, two inhibitors of LPL. Moreover, ANGPTL8 contributes to increase hepatic TG by decreasing intracellular TG hydrolysis via inhibiting ATGL. Increased hepatic TG content leads to higher TG secretion and, as a consequence, increased plasma TG levels. Increased intracellular cholesterol in the liver inhibits the SREBP2 pathway. SREBP2 increases LDLR and PCSK9 mRNA expression. These changes combined with additional post-transcriptional regulation of PCSK9 lead to reduced membrane-bound LDLR, which leads to decreased LDL uptake by the liver. The ensemble of these changes contribute to increased large VLDL1 and the formation of small dense LDL, which favors foam cell formation and ultimately atherosclerosis. A number of potential NASH therapies directly target metabolic pathways of lipid metabolism. For example, ficosostat inhibits ACC, reducing DNL and hepatic TG accumulation. Statins inhibit HMG-CoA reductase, the rate-limiting enzyme in cholesterol biosynthesis. Other strategies focus on activating different nuclear receptors that more broadly control lipid and glucose metabolism. These include nuclear receptors from the PPAR family, FXR and TR-β. Among these treatments, fibrates are PPARα agonists, pegbelfermin is a FGF21 analog (a PPARα target gene), thiazolidinediones (pioglitazone and rosiglitazone) are PPARγ agonists, elafibranor is a dual-PPARα/δ agonist, lanifibranor is a pan-PPAR agonist, obeticholic acid (OCA) is an FXR agonist, NGM282 is an FGF19 analog (FXR target gene in the intestine), and resmetirom is a TR-β agonist.

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De novo lipogenesis (DNL) is considered an important driver of NAFLD despite relatively low absolute levels of DNL in humans compared to rodents. Hyperinsulinemia in metabolic syndrome leads to excessive DNL via activation of the liver X receptor-alpha-sterol regulatory element-binding protein 1c (LXRα-SREBP1c) cascade [54]. In addition, nutrient signaling through mechanistic targeting of rapamycin complex 1 (mTORC1) by amino acids and carbohydrate response element-binding protein (ChREBP) by glucose and fructose can further stimulate this pathway. In physiological states, mTORC1-mediated signaling can compensate for the loss of insulin action [55]. Conversely, when challenged with a high-fructose diet, receptor-level ablation of hepatic insulin signaling protects from hepatic steatosis in murine models. Importantly, stable isotope studies have indicated that hepatic DNL is elevated in non-diabetic patients with high intrahepatic TG (measured by MRS) [32,33,56] and is closely associated with altered glucose homeostasis in these patients [56]. These results suggest that hyperinsulinemia (even without overt T2D) likely drives steatosis by impaired fatty acid catabolism but also drives VLDL-TG production, leading to dyslipidemia.

### 3.1. Hepatic TG metabolism and hypertriglyceridemia

#### 3.1.1. Hepatic de novo lipogenesis drives liver injury in NAFLD

De novo lipogenesis (DNL) is considered an important driver of NAFLD as well as fatty acid desaturation [57]. DNL is also closely linked with cellular saturated fatty acid levels [58] and ceramide synthesis [54], both of which can lead to hepatocyte apoptosis and metabolic dysfunction. Conversely, mice lacking hepatic ChREBP, a glucose-induced activator of glycolytic and lipogenic gene expression, showed more pronounced liver damage when challenged with a high-fat high-fructose diet [59]. Moreover, hepatic overexpression of Phf2 finger protein 2 (Phf2), a histone demethylase that mediates some of ChREBP’s actions on gene transcription, induced marked steatosis in chow-fed mice. However, Phf2-overexpressing mice were protected from steatosis-associated endothelial reticulum-stress induced inflammation and hepatic metabolic dysfunction likely due to a fatty acid profile enriched in mono-unsaturated fatty acid [60]. When challenged with a high-fat high-sucrose diet, Phf2 overexpression protected against hepatic inflammation and fibrosis. Thus, ChREBP-Phf2-coordinated regulation of lipogenesis as well as fatty acid elongation/desaturation plays an important role in maintaining the balance of saturated to unsaturated fatty acid.

Similar findings resulted from human trials with ACC inhibitors. A proof-of-concept study in obese patients with elevated intrahepatic TG as assessed by non-invasive imaging showed that ACC1 and ACC2 inhibition potently and rapidly decreased liver TG content (−36% in 4 weeks) [61]. However, this was associated with elevated plasma TG due to compensatory activation of mitochondrial glycerol-3-phosphate acyl-CoA acyltransferase 1 (GPAT1), which is induced by intracellular poly-unsaturated fatty acid deficiency [61]. In contrast, elevated DNL observed in patients with high hepatic TG content (~18% measured by MRS) was also associated with increased hepatic VLDL-TG secretion [33]. Thus, increased DNL in NAFLD favors liver fat accumulation by the combination of increased TG synthesis and decreased fatty acid catabolism but also drives VLDL-TG production, leading to dyslipidemia.

#### 3.1.2. VLDL-TG secretion correlates positively with intrahepatic TG content

VLDL are liver-derived, TG-carrying particles that contribute the majority (by mass) of fasting plasma TG levels. In humans, each VLDL particle contains 1 molecule of full-length apolipoprotein (Apo) B (ApoB100), whereas intestinally produced chylomicrons contain the truncated ApoB isoform (ApoB48). Ultracentrifugation-isolated VLDL separates in two subclasses (VLDL1 and VLDL2) differing in size and metabolic function. VLDL1 are larger with higher TG content, while VLDL2 are smaller and denser. Importantly, VLDL1 are reportedly more pro-atherogenic, since they give rise to small dense LDL [62]. Moreover, increased VLDL1 but not VLDL2 particle numbers are observed in overweight diabetic patients, identifying VLDL1 production as a driver of dyslipidemia [62].

NAFLD patients often display hypertriglyceridemia [49–52] and elevated remnant lipoprotein particle concentrations, which are indicative of delayed intravascular TG metabolism [63]. Both features increase cardiovascular risk. Indeed, the risk of major adverse CV events (MACE) inversely correlates with remnant cholesterol levels, with a 0.8 mmol/L reduction in remnant cholesterol levels lowering the risk of MACE by 20% [64]. Intrahepatic TG content directly correlates with hepatic VLDL production in obese non-diabetic patients with normal liver fat content [65,66]. Surprisingly, this direct correlation was lost in patients with intrahepatic TG content higher than 10% (as assessed by MRS) [65,66]. These results suggest that hepatic VLDL-TG secretion can only compensate for increased liver fat content to a certain extent. Beyond this limit of 10%, it is possible that the correct balance of TG and phospholipids, which coat the VLDL particle (discussed to follow), can no longer be maintained, and VLDL production lags behind further increases in hepatic TG content. In parallel, insulin resistance in more severe steatosis and NASH leads to reduced plasma VLDL clearance (discussed to follow) that can further exacerbate plasma hypertriglyceridemia. Together, these factors may explain the co-existence of hepatic steatosis and hypertriglyceridemia in NAFLD patients (Figure 2). Importantly, the molar ratio of VLDL-TG (particle content) to VLDL-ApoB100 (the number of particles) is elevated in non-diabetic obese individuals with high intrahepatic TG (measured by MRS), suggesting that the secreted VLDL particles are larger and resemble the pro-atherogenic VLDL1 [65]. Since other studies indicated that plasma ApoB concentration is a stronger predictor of CVD risk than LDL-C or plasma TG concentrations [67], further investigation is required to dissect the contribution of ApoB particle concentration vs particle content (TG or cholesterol) in determining CVD risk. It is important to bear in mind that the majority of the aforementioned studies focused on differences in intrahepatic TG and not necessarily NASH per se. Moreover, the relative impact of insulin resistance vs elevated intrahepatic TG on hepatic VLDL-TG secretion remains an open question. In insulin-sensitive individuals, insulin inhibits hepatic VLDL production and favors LPL activity, decreasing plasma VLDL [46]. However, in NAFLD patients (defined by MRS) hyperinsulinemia...
Increases VLDL-TG secretion [68], indicative of a loss of insulin-dependent inhibition of VLDL production. Alterations in phospholipid homeostasis may also contribute to altered VLDL regulation in NAFLD. Phosphatidylethanolamine (PE) and phosphatidylcholine (PC) are the major phospholipid components that form the surface monolayer of VLDL (and other) particles. Several studies have shown that PC is crucial for VLDL assembly and secretion [69]. Lowering PE levels can also reduce VLDL secretion as ~30% of hepatic PC production comes from the conversion of PE to PC by phosphatidylethanolamine N-methyltransferase (PEMT) [70]. Moreover, a decreased PC/PE ratio [71] and lower levels of PEMT [72] were observed in patients with NASH. Likewise, diets that reduce PC availability, such as methionine- and choline-deficient diets, reduce hepatic VLDL-TG secretion and are commonly used to induce features of NAFLD in mice. Studies in mice indicate that in addition to steatosis, reduced PC availability may also contribute to hepatic injury in NASH. For example, high-fat diet (HFD)-fed PEMT-deficient mice develop more severe hepatic inflammation and fibrosis, likely due to cholestasis [73]. Similarly, patients with histological NASH and methionine- and choline-deficient-fed mice [74] displayed increased hepatic glutaminase 1 (GLS1), which is associated with reduced PC synthesis. Inhibition of GLS1 in the methionine and choline-deficient diet mouse model reduced oxidative stress by limiting anaplerotic reactions for the tricarboxylic acid (TCA) cycle. By reducing oxidative stress, serine was used preferentially in the 1 carbon cycle to generate methyl donors for use in the conversion of PE to PC and VLDL-TG output was consequently restored [74]. This study provides evidence that oxidative stress induced by hepatic lipid overload could indirectly impact the PC/PE balance, impacting hepatic VLDL-TG output (Figure 2). Together, these results highlight important links between phospholipid metabolism, amino acid metabolism, and oxidative stress in the development of NAFLD-associated dyslipidemia.

3.1.3. Reduced plasma VLDL clearance results in the accumulation of remnant particles in NAFLD

While hepatic VLDL output is increased in NAFLD, triglyceride-rich lipoprotein (TRL, that is, VLDL and chylomicron remnants) clearance is reduced, further exacerbating hypertriglyceridemia in these patients. Circulating TRLs are metabolized by LPL to deliver fatty acids to peripheral tissues. Importantly, several exchangeable apolipoproteins modulate LPL activity, including ApoC2 and ApoA5, which stimulate LPL activity, and ApoC1, ApoC3, and ApoE, which inhibit LPL activity. Additionally, angiopoietin-like proteins ANGPTL3 and ANGPTL4 also inhibit LPL individually, particularly when complexed with ANGPTL8 [75]. Elevated non-fasting plasma TG and remnant cholesterol levels (defined as total cholesterol, LDL-C, and HDL-C) due to genetic ApoA5 polymorphisms causally increased the risk of myocardial infarction (independent of hypertension and T2D) [76]. Moreover, loss of function mutations in ApoC3 [77,78] and ANGPTL4 [79] decreased plasma TG and CVD risk. In line, decreased plasma levels of ANGPTL8 [80] and ANGPTL3 [81] are both associated with decreased CVD risk (defined by the presence of atherosclerotic plaques or myocardial infarction). Conversely, decreased plasma ANGPTL4, which is transcriptionally activated by peroxisome proliferator-activated receptor (PPAR) α, is also associated with carotid artery stenosis (vessel wall thickness measured by magnetic resonance imaging, MRI) [82], suggesting increased CVD risk. In ApoE*3-Leiden mice, ANGPTL4 overexpression decreased foam cell formation and protected against atherosclerosis formation via lipid-independent anti-inflammatory effects [82]. ApoC3 variants clearly impact plasma TG levels but they do not seem to have a consistent effect on hepatic steatosis [83–87]. In line, overexpression of ApoC3 in HFD-fed mice did not change steatosis despite markedly elevating plasma triglycerides due to impaired TG clearance [88]. Thus, ApoC3 more likely plays a role in NAFLD-
associated dyslipidemia. Mechanistically, ApoC3 is transcriptionally activated by glucose [89] and repressed by PPARx, itself reduced in patients with NASH [90]. Since ApoC3-mediated clearance explains ~75% of plasma TG variations in obese non-diabetic patients [91], these findings support the idea that NAFLD favors increased ApoC3 transcription, suppressing TRL clearance. Several studies have observed elevated ANGPTL protein levels in NAFLD patients. Plasma ANGPTL3 levels are higher in biopsy-proven NASH patients [92,93], and plasma [94,95] and hepatic [96] ANGPTL8 are elevated in patients with fatty liver independent of insulin resistance [94]. In addition to inhibiting LPL activity, ANGPTL8 seems to favor TG accumulation in hepatocytes. In vitro, ANGPTL8 knockdown decreased TG accumulation in hepatocytes after fatty acid treatment [96], whereas incubation of hepatocytes with recombinant ANGPTL8 increased TG accumulation likely due to decreased activity of adipose triglyceride lipase (ATGL) [97]. Overall, several regulators of LPL activity are perturbed in patients with NAFLD, which may contribute to increasing the risk of CVD. After TG lipolysis, TG-poor VLDL become remnant particles enriched in cholesterol [98]. Ultrasound-diagnosed NAFLD patients displayed higher fasting remnant cholesterol concentrations, indicating impaired clearance of remnant particles. Moreover, NAFLD patients with above-median concentrations of remnant cholesterol display higher CV event rates than NAFLD patients with below-median concentrations of remnant particles [99]. Indeed, similar to LDL, remnant particles can cross the arterial wall [100]. NAFLD patients also display elevated levels of small dense LDL particles, which are more atherogenic than large buoyant LDL [51,101]. Moreover, NAFL patients (defined by ultrasound or biopsy) display higher activity of cholesteryl ester transfer protein (CETP), an enzyme that exchanges cholesterol between HDL and ApoB particles, thereby favoring the formation of small dense LDL [102,103]. Together, these changes lead to increased abundance and altered metabolism of TRL, which are major contributors to CVD risk in NAFLD.

3.2. Intracellular cholesterol metabolism and elevated plasma LDL-C

3.2.1. Lipid-lowering therapy

Statins have been investigated as a potential therapy for NASH due to their actions in correcting total plasma and LDL-C levels and subsequent CVD risk reduction. Initial studies using indirect measures of NAFLD showed promising results, including decreased alanine transaminase (ALT) and AST levels or reduced steatosis on ultrasound [104]. However, randomized trials evaluating the impact of statin therapy on biopsy-proven NASH are lacking [105]. A post hoc analysis of a randomized, placebo-controlled trial assessing pioglitazone in NASH revealed that patients on statin therapy either at enrollment or starting therapy during the trial displayed lower plasma ALT and liver fat content on MRS [106]. However, histological analysis did not show statistically significant improvements in steatosis, inflammation, ballooning, or fibrosis [106]. Very surprisingly, the use of lipid-lowering agents (statins or other lipid-lowering agents) was reported not to reduce CV mortality in NAFLD patients [107]. However, the groups in this study were not well matched for other co-morbidities increasing CVD risk (T2D or a history of CVD), which could mask the potential beneficial effect of lipid-lowering therapy on CVD risk. A transcriptomic analysis of liver biopsies from obese patients revealed that statin treatment is associated with increased expression of genes involved in DNL [108]. Altogether, while statins are clearly safe drugs to use in NAFLD patients, whether they also improve NAFLD is doubtful.

3.2.2. Receptor-mediated LDL clearance

Unlike mice, humans metabolize the majority of their VLDL particles to LDL. Elevated concentrations of LDL-C, an undisputed CVD risk factor, are often but not always observed in NAFLD patients [50,52]. Several cell surface receptors facilitate the removal of LDL and remnant lipoprotein particles from the circulation by the liver, the best characterized being the LDL receptor (LDLR) and the closely related LDLR-related protein-1 (LRP1). However, LDLR is negatively regulated by proprotein convertase subtilisin/kexin type 9 (PCSK9), a protease whose best-known function is to bind LDLR at the plasma membrane. Upon binding, PCSK9 promotes intracellular degradation of LDLR by blocking its recycling in the endosomal compartment. This leads to reduced LDLR presence at the plasma membrane and less LDL uptake as a consequence. Regulation of LDLR activity by PCSK9 has been shown to impact CVD. Several studies showed dysregulated levels of these proteins in NAFLD. Plasma levels of PCSK9 positively correlated with histological hepatic steatosis [109,110], whereas low levels of hepatic LRP1 expression are associated with poor prognosis in hepatocellular carcinoma, an outcome of NAFLD [111]. Recent preclinical studies suggest that PCSK9 and LRP1 could modulate NAFLD pathogenesis. PCSK9 deficiency exacerbated the development of hepatic steatosis in HFD-fed mice [112]. Moreover, hepatic steatosis decreased cell surface LDLR expression and increased plasma ApoB and total cholesterol in a PCSK9-dependent manner [110]. Indeed, increased intracellular lipid induces endoplasmic reticulum stress, which can activate sterol regulatory element-binding protein 2 (SREBP2), a transcriptional regulator of PCSK9 expression [110]. These results demonstrate the dual role of PCSK9. By decreasing cholesterol uptake through LDLR inhibition, PCSK9 protects the liver from excess intrahepatic cholesterol accumulation. However, this leads to increased plasma LDL-C levels, increasing CVD risk. High-fat high-cholesterol diet-fed liver-specific LRP1-deficient mice showed accelerated NAFLD development characterized by increased steatosis, inflammation, and fibrosis independent of any changes in plasma lipids [113]. Worsening of NAFLD was attributed to a marked increase in hepatic cholesterol in liver-specific LRP1-deficient mice associated with increased cell death and stellate cell activation [113]. The lack of effect on plasma lipoproteins in this model was surprising but is not consistent across all murine models. For example, hepatocyte LRP1 deficiency increased atherosclerosis severity in ApoE/LDLR double-knockout mice despite decreased plasma cholesterol levels [114]. Collectively, these data suggest that dysregulation of receptors mediating hepatic LDL-C uptake induces liver injury by increasing intrahepatic cholesterol. Moreover, insulin signaling strongly induces LRP1 translocation to the membrane, highlighting a further mechanism by which insulin resistance could contribute to remnant accumulation observed in NAFLD patients [115]. Dysregulation of intracellular cholesterol metabolism is also associated with NASH severity in humans, and excess dietary cholesterol drives hepatic inflammation in murine models of NASH [116]. The mechanisms underlying these findings are likely complex. Intracellular cholesterol directly regulates gene transcription by suppressing SREBP2 activity, or indirectly via activating LXRa and farnesoid X receptor (FXR) by cholesterol-derived metabolites (oxysterols and bile acids, respectively). Moreover, the transcriptional regulator tafazzin (TAZ) has been identified as another cholesterol-sensitive pathway implicated in NASH progression [117]. Excess intracellular cholesterol stabilizes TAZ, which in turn activates the expression of pro-fibrotic genes in hepatocytes. In line, in vitro studies showed that increased intracellular cholesterol as a result of LDL loading provoked...
mitochondrial injury and induced several signs of lipotoxicity such as apoptosis, necrosis, and oxidative stress [118]. These studies highlight an important role of alterations in intracellular cholesterol homeostasis as a driver of NASH.

4. THERAPEUTIC STRATEGIES AND THEIR POTENTIAL IMPACT ON CVD RISK IN NAFLD

4.1. Lifestyle interventions and bariatric surgery to treat NAFLD: impact on CVD risk factors

4.1.1. Lifestyle interventions

Lifestyle interventions such as diet modification and increased physical activity are cornerstones for the prevention and treatment of NAFLD. Several dietary intervention strategies have been tested: modification of diet composition (e.g., Mediterranean, low-carbohydrate, low-fat, and ketogenic diets) or feeding behavior (time-restricted feeding or intermittent fasting) with the Mediterranean diet showing the most promising results for reducing hepatic steatosis, dyslipidemia, and other metabolic comorbidities [119]. A recent study of overweight and obese patients showed that consumption of a ketogenic diet reduced intrahepatic fat content (measured by MRS) and markedly increased hepatic mitochondrial activity [120]. This was paralleled by a 25% reduction in plasma TG levels and improved insulin sensitivity despite minimal (~3%) weight loss. Similarly, in NAFLD patients, 2 weeks of a low-carbohydrate diet also strongly decreased liver fat (assessed by MRS) associated with decreased hepatic DNL, increased mitochondrial β-oxidation, ~50% reduced VLDL-TG, improved insulin sensitivity, and decreased plasma IL-6 and tumor necrosis factor α (TNF-α) levels despite only modest weight loss (~2%) [121].

Increasing physical activity has also shown promise for treating NAFLD and associated comorbidities. High-intensity interval training improved several metabolic parameters including liver fat content (measured by MRT), BMI, plasma lipids, and insulin resistance in obese NAFLD patients (diagnosed by ultrasound or CT) with T2D [122]. Similarly, supervised exercise training, including a combination of aerobic and resistance training, effectively decreased body weight, hepatic steatosis, and adipose depot size (measured by MRI) compared to standard lifestyle advice [123]. Moreover, these changes were accompanied by improvements in several CVD risk parameters, including decreased plasma LDL-C, increased clearance of large TG-rich VLDL1 particles, improved insulin sensitivity, and reduced arterial stiffness [123].

Overall, these studies indicate that short-term reductions in liver fat and improvements in CVD risk factors can be achieved independent of weight loss. However, more significant weight loss (>7%) is needed to achieve resolution of biopsy-proven NASH [124,125], and fibrosis regression was observed only after ≥10% weight loss [124]. Future studies should address whether dietary interventions such as ketogenic or Mediterranean diets lead to durable reductions in NASH and CVD events with or without significant weight loss.

4.1.2. Bariatric surgery

Bariatric surgery is highly effective for treating hepatic steatosis and NASH as an alternative to diet and exercise, especially in the context of severe and morbid obesity. Bariatric surgery refers to several types of surgical interventions that mechanically restrict food intake, including gastric bypass (most commonly Roux-en-Y gastric bypass, RYGB), gastric banding, and sleeve gastrectomy. Gastric bypass and gastric banding have been shown to significantly improve histological features of NASH one [126] and five [127] years after the intervention and progressively decrease histological fibrosis in NASH patients [128]. Gastric bypass was the most efficient procedure to improve NASH [128], correlating with the degree of weight loss. Both studies also found durable improvements in dyslipidemia, insulin resistance, and markers of systemic inflammation, indicating that CVD risk may also be reduced. Histological improvements in NASH (steatosis, inflammation, and fibrosis) after RYGB or gastric banding have been confirmed in several smaller studies with median follow up varying from 6 months [129,130] to 40 months [131]. Prior to studies on NASH, bariatric surgery was found to rapidly and markedly improve several CVD risk factors, especially insulin resistance and dyslipidemia (reviewed by Tailleux et al.) [132]. A 12-year follow up study demonstrated that RYGB is associated with persistent weight loss, T2D remission, increased HDL-C, and decreased LDL-C and plasma TG compared to obese patients not undergoing surgical weight loss intervention [133]. Importantly, bariatric surgery also reduced CVD events and mortality [134] among T2D and obese patients. Therefore, it is highly likely that bariatric surgery will also reduce CV events in severely and morbidly obese patients with NASH, although this remains to be directly proven. Moreover, despite these benefits, bariatric surgery is not without risks [135] and is unlikely to become a universal treatment for metabolic diseases, including NAFLD.

4.2. Potential impact of pharmacological interventions targeting NASH on CVD risk

There are currently no approved pharmacological therapies for NASH. However, several clinical phase 2 and 3 trials assessing metabolically oriented strategies are ongoing (Figure 1). Many of these therapies are nuclear receptor agonists or hormone analogs that mainly stimulate metabolic pathways to reduce hepatic fat accumulation and decrease liver injury. Interestingly, the majority of these compounds also showed atheroprotection in preclinical studies. In this chapter, we highlight the current advancement of selected clinical trials for NASH and report specifically on associated effects on dyslipidemia and CVD risk reduction when data are available.

4.2.1. Bile acid metabolism modulators

4.2.1.1. FXR agonists. FXR is a nuclear receptor activated by bile acids that is highly expressed in the enterohepatic system. In addition to suppressing bile acid synthesis, FXR activation protects against lipid accumulation in the liver by decreasing hepatic lipogenesis, favoring fatty acid oxidation as well as decreasing inflammation by repressing nuclear factor kappa B (NF-κB) signaling [136]. Obelithic acid (OCA) is a semi-synthetic steroidal FXR agonist. Phase 2 (FLINT) and phase 3 (REGENERATE) trial data indicate that OCA treatment improves fibrosis without worsening NASH (Clinical-Trials.gov identifier: NCT02548351). However, complete NASH resolution was not achieved. Unfortunately, these benefits are associated with an increase in LDL-C by approximately 20 mg/dL [137,138], increasing the CVD risk score category from low or medium to high risk in 7.6% of patients [139]. Detailed lipoprotein profile analysis in the FLINT trial patients revealed that OCA treatment increased both large buoyant (less atherogenic) and small dense (more atherogenic) LDL particles. Moreover, despite similar total VLDL levels, OCA-treated patients displayed a shift from large VLDL1 toward less atherogenic small VLDL2 particles [140]. Importantly, the OCA-induced LDL-C increase was reversed by concomitant treatment with statins [138,141].

This finding suggests that the OCA-associated increase in LDL-C likely results from FXR-mediated reduction of cytochrome P450 family 7 subfamily A member 1 (CYP7A1) expression in hepatocytes, leading to decreased conversion of intrahepatic cholesterol to bile acids.
Increased intrahepatic cholesterol represses the statin-sensitive SREBP2 pathway, reducing LDLR transcription and decreasing plasma LDL clearance [136]. Several other synthetic FXR agonists with potential selective modulator actions are currently being evaluated in phase 2 clinical trials enrolling patients with NASH. One such compound, cilofexor (ClinicalTrials.gov identifier: NCT02854605), an intestinally targeted, non-steroidal FXR agonist, decreases hepatic steatosis (assessed by MRI) without increasing LDL-C [142]. Further studies should address whether these novel FXR agonists also impact CVD risk.

Agonism of the FXR pathways has the potential to treat fibrotic NASH but the current class of compounds likely needs refinement to improve the metabolic and CV risk profile. Indeed, FXR activation favors reverse cholesterol transport, reduces VLDL secretion, and decreases lipoprotein (Lp) (a) levels but also increases plasma LDL-C by decreasing bile acid synthesis (which leads to decreased LDLR expression) and increasing CETP expression [136].

### 4.2.1.3. Future directions for treatments targeting the bile acid pathways

#### 4.2.1.2. Fibroblast growth factor 19 analogs

The non-tumorigenic fibroblast growth factor (FGF) 19 analog NGM282 is also under evaluation for NASH treatment (ClinicalTrials.gov identifier: NCT02443116). FGF19, a hormone produced by the distal intestine and FXR target gene, participates in a feedback loop to reduce bile acid synthesis. FGF19 also exerts metabolic actions that may be beneficial against NAFLD such as lowering liver TG and reducing plasma TG levels [143]. Interestingly, circulating FGF19 is decreased in NASH patients, providing further therapeutic rationale [144].

In two 12-week trials, NGM282 improved all of the histological parameters of NASH and fibrosis and reduced ALT and AST [145,146]. However, as with OCA, NGM282 treatment increased plasma LDL-C [145] mostly as large, buoyant LDL particles [147]. Plasma TG levels decreased [145] as a result of fewer large VLDL particles [147]. The net effect of these changes on CV risk remains unclear. Preclinical studies of ApoE-deficient mice demonstrated a clear atheroprotective action of NGM282 [148]. In humans, FGF19 levels correlated negatively with CAD (defined by coronary angiography) independent of BMI, hypertension, dyslipidemia, and diabetes [149]. As with OCA, increased plasma LDL-C was blunted by statin treatment [146,147], suggesting that the action mechanism of NGM282 is similar to OCA. Larger phase 3 trials are ongoing.

#### 4.2.1.3. Future directions for treatments targeting the bile acid pathways

Volixibat (an intestinal ASBT inhibitor that prevents reabsorption of bile acids) decreased plasma total cholesterol and LDL-C but did not improve NASH in an interim analysis from a phase 2 trial (ClinicalTrials.gov identifier: NCT02787304) [150]. Bile acid sequestrants, such as colestevam and cholestyramine, improve plasma total cholesterol and LDL-C levels but not CV risk (WHO trial) without improving liver histology (ClinicalTrials.gov identifier: NCT01066364) [151]. Therefore, intra-vascular cholesterol and lipid metabolism appear to be regulated in a dissociated manner from liver histological NAFLD.

#### 4.2.2. PPAR agonists

Several clinical trials are currently assessing agonists of different members of the PPAR family to treat NASH. This family is composed of three isoforms, PPARα, PPARβ (or PPARδ), and PPARγ, which have distinct and partially overlapping functions in the control of lipid and glucose metabolism (for a detailed review, see Dubois et al. [152]). Each PPAR isoform can be specifically activated by appropriately designed synthetic agonists. For example, fibers (PPARα agonists) and thiazolidinediones (TZDs, PPARγ agonists) are used to treat hyperlipidemia and diabetes, respectively.

#### 4.2.2.1. Single PPAR agonists

Since hepatic PPARα expression is downregulated in NASH and restored PPARα activity is observed upon NASH resolution [90], activation of this pathway appears to be a reasonable approach to treat NASH. Fibrates (for example, fenofibrate, gemfibrozil, and pirosamil, among others) mainly target PPARα and thereby improve atherogenic dyslipidemia observed in obesity, T2D, and NAFLD. Pre-statin era and post hoc analysis of the FIELD and ACCORD studies indicated a reduction in CV risk, especially in high TG/low HDL-C subpopulations [153]. Several small studies have found poor efficacy for older fibrates (fenofibrate, gemfibrozil, and clofibrate) on histological improvements for NASH despite correction of dyslipidemia [154]. Pemafibrate, a new PPARα agonist, is currently being evaluated for CV risk in the PROMINENT trial (ClinicalTrials.gov identifier: NCT03071692). Studies of NASH assessed by non-invasive methods (MRI for steatosis, FIB4, NFS, and enhanced liver fibrosis [ELF] tests) are underway.

Thiazolidinediones (TZD) such as pioglitazone and rosiglitazone are PPARγ agonists used in T2D patients to increase peripheral insulin sensitivity. Because of the strong interaction between diabetes and NAFLD, several trials tested their ability to improve NASH. A meta-analysis of 8 randomized clinical trials evaluating pioglitazone and rosiglitazone therapy on biopsy-proven NASH revealed that pioglitazone, but not rosiglitazone, promotes NASH resolution and reduces liver fibrosis [155]. While rosiglitazone treatment reduced steatosis and plasma ALT, it did not improve lobular inflammation, ballooning, or fibrosis [156,157]. Conversely, pioglitazone improved all histological features of NASH and fibrosis, including improving dyslipidemia [158—160]. Pioglitazone is also superior to rosiglitazone in terms of CV risk reduction in T2D patients. Rosiglitazone increases plasma LDL-C and HDL-C levels [161], whereas pioglitazone improves dyslipidemia by increasing plasma HDL-C levels and decreasing plasma TG levels without raising LDL-C [162] or increasing large, buoyant LDL [161,163]. While long-term rosiglitazone treatment of T2D patients may slightly increase the risk of myocardial infarction [164], pioglitazone somewhat decreases the risk of MACE (although the risk of heart failure increases with all glitazones) [165,166]. In biopsy-proven NASH patients, rosiglitazone treatment did not modify plasma TG or HDL-C levels but increased LDL-C [156]. Conversely, pioglitazone treatment resulted in similar plasma lipid improvements in biopsy-proven NASH patients as observed in subjects with T2D [159,167]. In the PIVENS trial, pioglitazone reduced plasma TG levels and VLDL particle size, while plasma HDL-C and LDL particle size increased, especially in patients with NASH resolution [168,169], further underscoring the close association between NASH and atherogenic dyslipidemia. These different effects of rosiglitazone and pioglitazone may be due to the fact that pioglitazone also activates PPARγ [170,171]. Thus, whereas selective PPARγ activators may reduce steatosis by lowering adipose-liver free fatty acid flux as a result of enhanced insulin sensitivity, pioglitazone’s additional beneficial effects on inflammation and fibrosis may be a result of its modest activation of PPARα.

Seladelpar is a PPARδ agonist analyzed in a phase 2 clinical trial (ClinicalTrials.gov identifier: NCT03551522). Importantly, the trial was prematurely stopped after interim analysis due to the presence of abnormal hepatic inflammation during interim analysis, reportedly unrelated to seladelpar treatment, despite improvements in liver function biomarkers. Another trial in patients with mixed dyslipidemia showed that seladelpar decreased plasma LDL-C and TG and increased HDL-C levels. Improvements in insulin sensitivity and...
plasma levels of alkaline phosphatase and GGT, markers of liver injury, were also observed in seladelpar-treated patients [172].

4.2.2.3. FGF21 analog. FGF21, a hepatokine and PPAR agonist is a triple-PPAR agonist that in a phase 2b study demonstrated improvements in histological ballooning and lobular inflammation without worsening of fibrosis, benefits that were most apparent in patients with more severe NASH at baseline (ClinicalTrials.gov identifier: NCT01694849) [173]. Improvements in insulin sensitivity and decreased systemic inflammation markers (measured by CRP, fibrinogen, and haptoglobin) were also observed in elafibranor-treated patients. Decreases in plasma TG and LDL-C were observed with concomitant increases in HDL-C, even among patients treated with statins. However, recently reported interim phase 3 results did not show statistically significant differences in resolving NASH between elafibranor and placebo. Long-term evaluation of CVD risk reduction by elafibranor in NASH patients has not yet been demonstrated.

Saroglitazar is a dual-PPARα/γ agonist currently in phase 3 clinical trials in the US and Europe (ClinicalTrials.gov identifier: NCT03061721) already approved for treating NASH in India. Lanifibranor is a triple-PPARα/γ/δ agonist that was recently reported to reach primary and secondary endpoints by significantly decreasing the Steatosis Activity Fibrosis Score (SAF), inducing NASH resolution and lowering fibrosis in the NATIVE phase 2b trial (ClinicalTrials.gov identifier: NCT03008070). Preclinical results indicated that saroglitazar decreases plasma TG levels, reduces hepatic steatosis, ballooning, and inflammation in the choline-deficient, L-amino acid-defined, high-fat diet NASH mouse model, and protects against carbon tetrachlorethyl (CCL4)-induced fibrosis [174], whereas lanifibranor treatment decreases CCL4-induced hepatic fibrosis, plasma glucose, and TG levels in db/db mice [175].

4.2.2.4. FGF21 analog. FGF21, a hepatokine and PPARα target gene, displays many favorable metabolic actions, including enhancing fatty acid oxidation, reducing lipogenesis, and improving dyslipidemia and insulin sensitivity [176]. Accordingly, a recombinant analog of human FGF21, pegbefelmin (ClinicalTrials.gov identifier: NCT02413372), reduced hepatic fat (assessed by MRI) and plasma transaminases and improved dyslipidemia (reduced plasma TG and LDL-C and increased HDL-C) in a phase 2 study of biopsy-diagnosed NASH patients (no post-treatment biopsy was performed) [177]. In ApoE-deficient mice, FGF21 protects against atherosclerosis progression acting on several pathways by reducing plasma cholesterol levels and cholesterol biosynthesis in the liver and reducing apoptosis in the aortic roots [178,179].

4.2.3. Thyroid hormone receptor β agonists

Thyroid hormones (TH) regulate several pathways of hepatic lipid metabolism mainly through TR receptor β (TR-β), the major isoform in the liver. In NAFLD, TR-β activation induces fatty acid uptake and oxidation and enhances mitochondrial biogenesis and activity. TR-β agonism also favors LDLR-mediated endocytosis and reverse cholesterol transport [180]. At least two TR-β selective agonists are under investigation for NASH, resmetirom (ClinicalTrials.gov identifier: NCT02912260) and VK2809 (ClinicalTrials.gov identifier: NCT04173065), employing distinct pharmacological mechanisms for liver selectivity. Resmetirom treatment decreased liver fat content (assessed by MRI) and NAS on biopsy in patients with biopsy-proven NASH with fibrosis (stages 1–3) [181]. Moreover, resmetirom treatment improved dyslipidemia by decreasing plasma levels of TG, ApoB, Lp(a), ApoCIII, and LDL-C, especially reducing the proportion of small LDL and large VLDL [181]. Thus, TR-β activation reduces both liver fat content and plasma lipid CVD risk factors, which may result in improved CV function.

4.2.4. Other treatments in clinical trials

The glucagon-like peptide-1 (GLP1) analog lixisenatide used for T2D treatment showed promise for NASH treatment with improvements in body weight and glycemic control in 9 biopsy-proven NASH patients (ClinicalTrials.gov identifier: NCT01237119) [182], prompting confirmation in larger studies with histological endpoints. Interestingly, the GLP1 receptor (GLP1-R) agonists liraglutide [183] and semaglutide [184] both reduce CVD risk in T2D patients. Considering the lack of GLP1-R expression in hepatocytes, it is unclear how GLP1 analogs improve NAFLD (either through non-parenchymal actions on the liver or by enhancing systemic insulin sensitivity and weight loss). In Western diet-fed ApoE or LDLR-deficient mice, both liraglutide and semaglutide decrease atherosclerosis through anti-inflammatory effects independent of cholesterol lowering and weight loss [185]. Moreover, GLP1 may prevent macrophage foam cell formation [186]. In high-fat diet-fed ApoE*-3-Leiden CETP-transgenic mice, GLP1-R agonist treatment decreased hepatic steatosis and plasma TG by respectively decreasing DNL and VLDL production [187]. Several drugs in development target the increase in DNL observed in NASH patients [32]. Recent phase 2 studies with the ACC1/ACC2 inhibitor fisposcostat showed reduced hepatic steatosis but increased plasma VLDL particles and TG levels (ClinicalTrials.gov identifier: NCT02856555) [188]. Thus, despite improving hepatic steatosis, fisposcostat may increase CVD risk by increasing the amount of atherogenic particles.

In conclusion, several therapeutic strategies for NAFLD with evidence or potential for CVD protection are emerging. However, a detailed assessment of CV outcomes (MACE and subclinical atherosclerosis) is lacking, and in several short-term studies, changes in plasma lipid profiles were unreported. More importantly, the efficacy of these therapies on NASH remains to be demonstrated. Nevertheless, it remains a major challenge to identify NASH patients at the highest risk of CVD to appropriately target them for management.

5. CONCLUSION

NAFLD is closely associated with CVD. While epidemiological studies do not consistently show increased CVD mortality in NAFLD patients, there is clear evidence for increased (non-fatal) MACE. Alterations in lipid and lipoprotein metabolism are major contributing factors linking NAFLD to CVD. Moreover, many promising NASH therapies in development also improve dyslipidemia in clinical trials. Given the current lack of approved pharmacological therapies for NASH, a clear understanding of the underlying factors that drive elevated CVD risk in NAFLD will be critical for the effective care and management of this growing patient population.

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