Triglyceride-rich lipoproteins as a causal factor for cardiovascular disease

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Abstract: Approximately 25% of US adults are estimated to have hypertriglyceridemia (triglyceride [TG] level ≥150 mg/dL [≥1.7 mmol/L]). Elevated TG levels are associated with increased cardiovascular disease (CVD) risk, and severe hypertriglyceridemia (TG levels ≥500 mg/dL [≥5.6 mmol/L]) is a well-established risk factor for acute pancreatitis. Plasma TG levels correspond to the sum of the TG content in TG-rich lipoproteins (TRLs; i.e., very low-density lipoproteins plus chylomicrons) and their remnants. There remains some uncertainty regarding the direct causal role of TRLs in the progression of atherosclerosis and CVD, with cardiovascular outcome studies of TG-lowering agents, to date, having produced inconsistent results. Although low-density lipoprotein cholesterol (LDL-C) remains the primary treatment target to reduce CVD risk, a number of large-scale epidemiological studies have shown that elevated TG levels are independently associated with increased incidence of cardiovascular events, even in patients treated effectively with statins. Genetic studies have further clarified the causal association between TRLs and CVD. Variants in several key genes involved in TRL metabolism are strongly associated with CVD risk, with the strength of a variant’s effect on TG levels correlating with the magnitude of the variant’s effect on CVD. TRLs are thought to contribute to the progression of atherosclerosis and CVD via a number of direct and indirect mechanisms. They directly contribute to intimal cholesterol deposition and are also involved in the activation and enhancement of several proinflammatory, proapoptotic, and procoagulant pathways. Evidence suggests that non-high-density lipoprotein cholesterol, the sum of the total cholesterol carried by atherogenic lipoproteins (including LDL, TRL, and TRL remnants), provides a better indication of CVD risk than LDL-C, particularly in patients with hypertriglyceridemia. This article aims to provide an overview of the available epidemiological, clinical, and genetic evidence relating to the atherogenicity of TRLs and their role in the progression of CVD.

Keywords: triglycerides, non-high-density lipoprotein cholesterol, hypertriglyceridemia, lipoprotein lipase, chylomicrons, very low-density lipoproteins

Introduction

Robust clinical evidence demonstrates that statin-induced reductions in low-density lipoprotein cholesterol (LDL-C) lead to substantial reductions in cardiovascular disease (CVD) risk in both the primary and secondary prevention settings. Therefore, in patients at high cardiovascular risk due to dyslipidemia, the majority of current guidelines recommend LDL-C as a primary treatment target, with statins as first-line therapy. Despite significant LDL-C lowering with statin therapy, substantial residual cardiovascular risk often remains. This residual risk is thought to be due, in part, to inadequate reduction of LDL-C for a given level of risk, low levels of...
high-density lipoprotein cholesterol (HDL-C), and/or high levels of triglycerides (TG). Additionally, there is increasing evidence that non-high-density lipoprotein cholesterol (non-HDL-C), which is calculated for individuals by subtracting their HDL-C value from total cholesterol (including TG-rich lipoproteins [TRLs]), may be more strongly associated with atherosclerotic risk than LDL-C alone. Approximately 25% of US adults are estimated to have hypertriglyceridemia, defined as a TG level of ≥150 mg/dL (≥1.7 mmol/L). Hypertriglyceridemia often coexists with secondary disorders that are independently associated with increased plasma TG levels, such as type 2 diabetes mellitus, chronic kidney disease, metabolic syndrome, and obesity. Severe hypertriglyceridemia, defined by the 2014 National Lipid Association guidelines as a TG level of ≥500 mg/dL (≥5.6 mmol/L), is a well-established risk factor for acute pancreatitis, and moderately elevated TG levels have been shown to be independently associated with increased CVD risk, even in patients treated effectively with statins to reduce LDL-C.

Plasma TG levels are known to correspond with the levels of TRLs and their remnants. However, hypertriglyceridemia is also often accompanied by further lipoprotein disturbances, including increased very-low-density lipoproteins (VLDLs) and total apolipoprotein (apo) C-III, elevated levels of small, dense LDL-C and total LDL particles, and decreased levels of HDL-C, all of which have been shown to be associated with increased CVD risk. This, along with the fact that cardiovascular outcome studies of TG-lowering agents have produced inconsistent results, means that there remains some uncertainty regarding the direct causal role of TRLs in the progression of atherosclerosis and CVD.

This review aims to provide an overview of the available genetic, epidemiological, and clinical evidence relating to the atherogenicity of TRLs and their role in the progression of CVD.

Search strategy
A search of PubMed was performed using the following search strategy: (“triglyceride-rich lipoproteins” OR “apolipoprotein C-III” OR “remnant lipoproteins” OR “intermediate-density lipoproteins”) AND (“Cardiovascular Diseases”[MeSH]).

The search was limited to English-language publications published between July 2005 and July 2015. The reference lists of articles identified using this search strategy was also searched such that widely referenced, older publications were also screened.

TRL metabolism
Cholesterol esters and TG are the two most important circulating lipids. Owing to their hydrophobic nature, they are combined into lipoprotein particles in association with proteins that allow them to be transported in the plasma. Cholesterol is transported by all lipoproteins and is particularly concentrated in HDL and LDL particles. In general, TGs are transported in the plasma in VLDL, chylomicrons, and their remnants created during metabolism. These TRLs are the largest lipoprotein particles. In addition to size, lipoproteins may also be characterized by the apoproteins they contain, with apoB100 (also known as apoB) being associated with VLDL and LDL and apoA-I associated with HDL. There are several other apoproteins involved in lipoprotein metabolism, many of which are thought to potentially contribute to a number of diseases, such as CVD, multiple sclerosis, and Alzheimer’s disease.

TRLs are highly heterogeneous, differing in size, density, composition, and associated cardiovascular risk. They are composed of a neutral core of TG and cholesterol esters and a surface monolayer comprising phospholipids, free cholesterol, and apoproteins, which participate in the regulation of transport and metabolism of the TRL. The metabolism of TRLs occurs via two principal pathways, an exogenous pathway that originates in the small intestine and an endogenous pathway regulated by the liver (Figure 1). During the exogenous pathway, TGs from dietary fat are absorbed by enterocytes following the ingestion of a meal. Here, they are incorporated into chylomicrons, which are large apoB-containing lipoproteins with a large TG core (80%–95%). Newly synthesized chylomicrons are then exported via perimesenteric lymphatics before entering the circulation, where they acquire apoC-II, apoC-III, and apoE. Once in circulation, chylomicrons are quickly hydrolyzed by lipoprotein lipase (LPL) along the luminal surface of the capillaries. LPL is synthesized by myocytes and adipocytes before being transported to the lumen of the capillaries via GPIHBP1, a small glycosylphosphatidylinositol-anchored protein synthesized by the capillary endothelial cells. LPL requires activation via apoC-II, and its activity is highly regulated by various proteins, including apoC-III, apoA5, and angiopoietin-like proteins 3 and 4. The hydrolysis of chylomicrons via LPL results in the production of free fatty acids and chylomicron remnants. The free fatty acids liberated by lipolysis are oxidized by a variety of cell types, such as skeletal and myocardial myocytes, or stored in adipose tissue. The chylomicron remnants, which are rich in cholesterol esters and apoE, are removed from circulation by the liver via binding to the LDL receptor or the LDL receptor-related protein.
During the endogenous pathway, TGs are synthesized in hepatocytes from free fatty acids and glycerol and then incorporated into the core of apoB-containing VLDL particles. ApoC-I, apoC-II, apoC-III, and apoE are added to the surface of VLDL particles during secretion. Following secretion, VLDLs undergo LPL-mediated hydrolysis in the plasma, generating in succession progressively smaller VLDLs and then intermediate-density lipoproteins (IDLs). Some IDL particles are taken up by the liver, and some undergo further catabolism by LPL and hepatic TG lipase to produce LDL particles.  

Importantly, polymorphisms in genes encoding key components of the TRL biosynthetic pathways are known to be strongly associated with CVD risk.  

**Proposed pathophysiology of TRLs in the progression of atherosclerosis**  
Evidence suggests that TRLs and their remnants and specific markers of TG metabolism, such as LPL and apoC-III, contribute to the progression of atherosclerosis and CVD both directly and indirectly. These proposed pathophysiological mechanisms are summarized in Figure 2.  

It has been suggested that it is primarily the cholesterol content of TRL remnants that directly contributes to the progression of atherosclerosis, rather than the TGs themselves. Like LDL, cholesterol-enriched, TG-depleted TRL remnants are able to penetrate the arterial intima, where they become selectively bound to the connective tissue matrix. Once entrapped in the subendothelial space, TRLs can be scavenged by resident macrophages, thereby contributing to macrophage foam cell formation as well as plaque formation and progression. TRLs are thought to be equally or more atherogenic than LDL. In contrast to LDL, TRL remnants can be taken up directly by arterial macrophages without oxidative modification and, due to their larger size, carry more cholesterol per particle than LDL. TRL remnants have also been shown to promote endothelial dysfunction, which potentiates atherogenesis.
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vascular endothelial permeability, promote leukocyte adhesion

and, at high concentrations, cause cellular injury and death. TRL remnants have also been shown to induce endothelial cell apoptosis via increased secretion of the proapoptotic cytokines, tumor necrosis factor-α, and interleukin-1β, a process that is known to contribute to vascular injury and atherosclerosis. TRLs and their remnants enhance platelet aggregation and clot formation and amplify the coagulation cascade by 1) supporting the assembly of the prothrombinase complex and 2) upregulating the expression of plasminogen activator inhibitor-1 and plasminogen activator inhibitor-1 antigen. They have also been shown to upregulate the endothelial expression of tissue factor, a key initiator of the coagulation cascade. Finally, TRL remnants suppress the atheroprotective and anti-inflammatory effects of HDL and have been shown to significantly correlate with impairment of coronary vasodilation.

Key components of TRL metabolism have also been shown to be associated with CVD, in particular apoC-III, a key contributor to hypertriglyceridemia due to its inhibitory effects on LPL. ApoC-III is thought to contribute to the progression of atherosclerosis via a number of mechanisms: it can impaire VLDL binding to cellular receptors, resulting in small, dense LDL particle formation, and has been shown to induce the expression of proinflammatory mediators and to stimulate monocyte activation and the adhesion of monocytes to endothelial cells. It is also thought to induce apoptosis and may accumulate on atheroprotective HDL particles on secretion, thus rendering them dysfunctional.

**Genetic evidence**

The fact that monogenic disorders of TG metabolism, such as hyperlipoproteinemia type 3, predispose individuals to CVD suggests that raised TG and remnant cholesterol levels contribute to this process. Conversely, a recent Mendelian randomization study based on data from 10,208 individuals included in the Copenhagen City Heart Study found that subjects with genetically confirmed reduction in non-fasting plasma TG levels had reduced all-cause mortality.

**Figure 2** Proposed pathophysiology of triglyceride-rich lipoproteins in the progression of atherosclerosis.


**Abbreviations:** LPL, lipoprotein lipase; TRL, triglyceride-rich lipoproteins; TRL-R, triglyceride-rich lipoprotein remnants.

LPL-mediated TRL hydrolysis results in a high concentration of lipolytic products, such as oxidized free fatty acids, along the vascular endothelium or within the arterial intima. These lipolytic products, along with TRLs themselves, are also known to activate a number of proinflammatory, procoagulant, and proapoptotic signaling pathways that play a fundamental role in the pathogenesis of atherosclerosis. Oxidized free fatty acids are known to increase the expression of inflammatory interleukins and cytokines, leading to endothelial inflammation, while TRL remnants have been shown to upregulate the endothelial expression of intercellular adhesion molecule-1 and vascular cell adhesion molecule-1. These proatherogenic adhesion molecules facilitate the transendothelial migration of leukocytes to sites of inflammation. Consequently, their TRL remnant-mediated activation leads to endothelial monocyte adhesion and an enhanced inflammatory response. TRL remnants have also been shown to induce early monocyte and neutrophil activation.

Additionally, TRL remnants are known to increase the production of reactive oxygen species, which can increase vascular endothelial permeability, promote leukocyte adhesion, and, at high concentrations, cause cellular injury and death. TRL remnants have also been shown to induce endothelial cell apoptosis via increased secretion of the proapoptotic cytokines, tumor necrosis factor-α, and interleukin-1β, a process that is known to contribute to vascular injury and atherosclerosis. TRLs and their remnants enhance platelet aggregation and clot formation and amplify the coagulation cascade by 1) supporting the assembly of the prothrombinase complex and 2) upregulating the expression of plasminogen activator inhibitor-1 and plasminogen activator inhibitor-1 antigen. They have also been shown to upregulate the endothelial expression of tissue factor, a key initiator of the coagulation cascade. Finally, TRL remnants suppress the atheroprotective and anti-inflammatory effects of HDL and have been shown to significantly correlate with impairment of coronary vasodilation.

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An additional meta-analysis involving 188,578 genotyped individuals with 185 different single nucleotide polymorphisms found that the strength of a variant’s effect on TG levels strongly correlated with the magnitude of its effect on coronary artery disease, even after adjustment for effects on LDL-C and HDL-C. These results support the hypothesis that TRLs causally influence cardiovascular risk.\(^\text{16}\)

Sequence variants in several key genes involved in the metabolism of TRLs, such as those encoding LPL and the proteins that regulate it, appear to be strongly associated with CVD risk.\(^\text{10,33}\) For example, apoC-III, an apolipoprotein playing a central role in TG metabolism by inhibiting LPL, is overexpressed in hypertriglyceridemia and is significantly associated with cardiovascular risk.\(^\text{84}\) In one study, which evaluated 18,666 genes in 3,734 participants, four loss-of-function mutations were identified in \(APOC3\), the gene encoding apoC-III. Heterozygous carriers of any of these mutations had 46% lower circulating levels of apoC-III, corresponding to 39% lower plasma TG levels, and a 40% lower risk of coronary heart disease than noncarriers.\(^\text{39}\) Similarly, a second study, which analyzed data from 75,725 participants, found that heterozygosity of loss-of-function mutations in \(APOC3\) were associated with a mean reduction in nonfasting TG levels of 44% and a corresponding 41% decrease in the incidence of ischemic vascular disease compared with wild-type individuals.\(^\text{60}\) In addition, mutations in the gene encoding apoAV (\(APOA5\)), an activator of LPL, have also been shown to be associated with CVD risk. Carriers of nonsynonymous \(APOA5\) mutations have higher plasma TG levels, lower HDL-C levels, but similar overall cholesterol levels, compared with noncarriers. Carriers of these mutations were shown to have a 2.2-fold higher risk of myocardial infarction (MI) and coronary artery disease compared with noncarriers.\(^\text{50}\) Furthermore, polymorphisms in the \(APOA5\) promoter region, which led to decreased \(APOA5\) expression, were strongly related to increased plasma TG levels and a concordant increase in coronary heart disease risk.\(^\text{64}\) Finally, loss-of-function variants in the gene encoding angiopoietin-like protein 4 (\(ANGPTL4\)), an inhibitor of LPL, were found to be associated with substantially decreased TG levels and decreased coronary heart disease risk.\(^\text{62}\)

Genetically elevated levels of TG and remnant lipoprotein cholesterol are also associated with increased low-grade inflammation, marked by elevated C-reactive protein levels. As this association was not observed for genetically elevated LDL-C, this suggests that the inflammatory component of atherosclerosis may be driven by elevated TRLs and remnant cholesterol.\(^\text{63}\) Taken together, these findings are consistent with the hypothesis that elevated TG levels, and consequently TRLs, are causally associated with CVD.

### Epidemiological evidence

In the Pravastatin or Atorvastatin Evaluation and Infection Therapy-Thrombolysis In Myocardial Infarction 22 (PROVE IT-TIMI 22) trial, it was demonstrated that, among patients receiving statin therapy following acute coronary syndrome, an on-treatment fasting TG level <150 mg/dL (<1.7 mmol/L) was associated with a reduction in recurrent coronary heart disease risk versus higher TG levels (Table 1), even after adjustment for HDL-C and LDL-C levels (hazard ratio 0.8; \(P=0.025\)).\(^\text{10}\) A wealth of epidemiological evidence exists, demonstrating that both fasting and nonfasting TG levels are significant predictors of cardiovascular events, even in individuals who have already achieved guideline-recommended LDL-C levels with lipid-lowering therapy.\(^\text{64–68}\)

However, nonfasting TG levels are thought to be a much stronger predictor of cardiovascular events than fasting TG levels. The Women’s Health Study (n=26,509) showed that both fasting and nonfasting TG levels were strongly associated with an increased risk of cardiovascular events, independent of baseline cardiovascular risk factors (age, blood pressure, smoking, and use of hormone therapy). However, after adjustment for TC and HDL-C levels and indicators of insulin resistance, the association between fasting TG levels and the risk of cardiovascular events was no longer significant (\(P=0.90\)). In contrast, the association between nonfasting TG levels and cardiovascular risk remained strong even after adjustment for other lipid parameters and markers of insulin resistance (\(P=0.006\)).\(^\text{67}\) Likewise, two prospective cohort studies using data from the Copenhagen City Heart Study found that the cumulative incidence of cardiovascular events (ischemic stroke, MI, ischemic heart disease) and all-cause mortality were strongly associated with increasing nonfasting TG levels (all \(P<0.001\)) (Table 1). That being said, these associations were not adjusted for other lipid parameters.\(^\text{66,68}\)

A number of studies have found that the association between plasma TG levels (both fasting and nonfasting) and cardiovascular risk is often attenuated once adjusted for other lipid parameters, including HDL-C and non-HDL-C. An analysis conducted by the Emerging Risk Factors Collaboration, which included data from 302,430 individuals from 68 long-term prospective studies, demonstrated that there was a significant and stepwise association between fasting and nonfasting TG levels and CVD risk. However, this association was no longer significant after adjustment for HDL-C and non-HDL-C (Table 1).\(^\text{64}\) Likewise, in a combined analysis...
of the Incremental Decrease in End Points through Aggressive Lipid Lowering (IDEAL) and Treating to New Targets (TNT) trials in patients achieving low LDL-C (<70 mg/dL [1.8 mmol/L]), CVD risk increased incrementally with increasing on-treatment fasting TG level, with patients in the highest quintile experiencing a 63% higher rate of cardiovascular events than those in the lowest quintile ($P<0.001$).

However, this association was also attenuated ($P=0.044$).

Table 1: Key studies investigating the association between triglycerides and cardiovascular disease

<table>
<thead>
<tr>
<th>Study</th>
<th>Population (sample size)</th>
<th>Key findings</th>
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<tbody>
<tr>
<td>ERFC: individual record analysis of 68 long-term prospective studies[^6]</td>
<td>No prior CHD (n=302,430)</td>
<td>After adjustment for nonlipid risk factors, TG levels were significantly associated with the incidence of CHD (HR 1.37, 95% CI 1.31–1.42). The association was no longer significant after adjustment for HDL-C and non-HDL-C (HR 0.99, 95% CI 0.94–1.05).</td>
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<tr>
<td>Post hoc analysis of two statin trials: IDEAL and TNT[^33]</td>
<td>CHD and ACS, on potent statin therapy (n=15,779)</td>
<td>Risk of CVE occurring after the first year of the trials increased as a function of increasing on-treatment TG, with patients in the 5th quintile of TG having a 63% increase in events versus patients in the first quintile after adjusting for age and sex ($P&lt;0.001$). After adjustment for HDL-C and apoB/apoAl, the association was attenuated ($P=0.044$).</td>
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<tr>
<td>PROVE IT-TIMI 22[^30]</td>
<td>Hospitalized for ACS, on potent statin therapy (n=4,162)</td>
<td>On-treatment fasting TG &lt;150 mg/dL was associated with a reduction in CHD risk versus high TG (HR 0.73, 95% CI 0.62–0.87; $P&lt;0.001$). For each on-treatment 10-mg/dL decrement in TG, the incidence of death, MI, and recurrent ACS was lowered by 1.6% or 1.4% after adjustment for LDL-C ($P&lt;0.001$) or non-HDL-C ($P=0.01$), respectively.</td>
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<td>Prospective Copenhagen City Heart Study: ischemic stroke[^46]</td>
<td>General population of Denmark (n=13,956)</td>
<td>The cumulative incidence of ischemic stroke increased with increasing levels of baseline nonfasting TG in both sexes ($P&lt;0.001$). For men, age-adjusted HRs ranged from 1.4 (95% CI 0.9–2.1) in those with baseline TG of 89–176 mg/dL to 3.2 (95% CI 1.7–6.2) in those with TG ≥443 mg/dL versus men with TG &lt;89 mg/dL. For women, HRs ranged from 1.3 (95% CI 1.0–1.8) in those with baseline TG of 89–176 mg/dL to 5.1 (95% CI 1.7–14.8) in those with TG ≥443 mg/dL versus women with TG &lt;89 mg/dL.</td>
</tr>
<tr>
<td>Women’s Health Study[^67]</td>
<td>Healthy US women (n=26,509)</td>
<td>Baseline fasting and nonfasting TG were both strongly associated with CVE. Fasting TG was not significantly associated with CVE after adjustment for TC and HDL-C and measures of insulin resistance. However, nonfasting TG remained significantly associated with CVE after adjustment for TC and HDL-C ($P=0.006$).</td>
</tr>
<tr>
<td>Prospective Copenhagen City Heart Study: MI, IHD, and all-cause mortality[^48]</td>
<td>General population of Copenhagen, Denmark (n=13,981)</td>
<td>Levels of remnant lipoprotein cholesterol increased with increasing non-fasting TG. The cumulative incidence of MI, IHD, and all-cause mortality increased with increasing nonfasting TG ($P&lt;0.001$). For men, HRs for MI increased from 1.6 (95% CI 1.1–2.3) for TG of 88.5–176.1 mg/dL to 4.6 (95% CI 2.7–8.0) for TG ≥443 mg/dL versus those with TG &lt;88.5 mg/dL. For women, HRs for MI increased from 2.2 (95% CI 1.6–3.2) for TG of 88.5–176.1 mg/dL to 16.8 (95% CI 6.8–41.6) for TG ≥443 mg/dL versus those with TG &lt;88.5 mg/dL.</td>
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<tr>
<td>Post hoc analysis of the Framingham Heart Study[^19]</td>
<td>Participants of the Framingham Heart Study without CVD at baseline (n=3,501)</td>
<td>High plasma TG levels (&gt;150 mg/dL), in the absence of high LDL-C (&gt;130 mg/dL) or low HDL-C (&lt;40 mg/dL) levels, were not significantly associated with an increased risk of CVD events ($P=0.13$).</td>
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<td>ARIC study[^20]</td>
<td>Participants of the ARIC Study without CVD at baseline (n=12,339)</td>
<td>In women, TG was significantly associated with CHD risk after adjustment for age, race, LDL-C, apoB, apoA-1, and HDL-C subfractions (RR 1.29, $P&lt;0.01$). However, TG was not significantly associated with CHD risk in men.</td>
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<td>PROCAM study[^21]</td>
<td>Men and women aged 16–65 years (n=19,698)</td>
<td>In a logistic function analysis, log-transformed TG levels showed a significant association with CHD incidence ($P&lt;0.01$). However, after adjustment for HDL-C, this association was no longer significant.</td>
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Abbreviations: ACS, acute coronary syndrome; apo, apolipoprotein; ARIC, The Atherosclerosis Risk in Communities; CHD, coronary heart disease; CI, confidence interval; CVD, cardiovascular disease; CVE, cardiovascular events; ERFC, Emerging Risk Factors Collaboration; HDL-C, high-density lipoprotein cholesterol; HR, hazard ratio; IDEAL, Incremental Decrease in End Points through Aggressive Lipid Lowering; IHD, ischemic heart disease; LDL-C, low-density lipoprotein cholesterol; MI, myocardial infarction; non-HDL-C, non-high-density lipoprotein cholesterol; PROCAM, Prospective Cardiovascular Munster; PROVE IT-TIMI 22, Pravastatin or Atorvastatin Evaluation and Infection Therapy-Thrombolysis In Myocardial Infarction 22; RR, relative risk; TC, total cholesterol; TG, triglyceride; TNT, Treating to New Targets.
adjustment for HDL-C and apoB/apoA1 (Table 1). Elevated TG levels are closely associated with higher levels of non-HDL-C and apoB and low levels of HDL-C, and this may explain why this association is weakened after adjustment for these parameters.

Elevated remnant cholesterol levels, which directly correlate with elevated levels of TRLs, have also been shown to be associated with CVD. Using data from 73,513 subjects, Varbo et al. found that every 88.6 mg/dL (1 mmol/L) increase in remnant cholesterol was associated with a 2.8-fold increase in CVD risk, independent of low HDL-C.

**Available treatment options to reduce TG levels and their potential impact on cardiovascular outcomes**

Although the clinical definition of the severity of hypertriglyceridemia differs among guidelines, the majority of guidelines define severe hypertriglyceridemia as a TG level of ≥500 mg/dL (≥5.6 mmol/L). In such cases, guidelines recommend the initiation of TG-lowering therapy to reduce the risk of pancreatitis. Guidelines also acknowledge that a TG level of <150 mg/dL (<1.7 mmol/L) is desirable, and if elevated TG or non-HDL-C levels remain following lifestyle intervention and statin therapy, a number of guidelines recommend the use of TG-lowering agents, primarily fibrates, niacin, or omega-3 fatty acids (Table 2).

**Fibrates**

Fibrates decrease TG levels by ~36%, non-HDL-C levels by ~6%–16%, and LDL-C levels by ~8% and increase HDLC levels by ~10%. Of note, however, fibrate-induced increases in LDL-C may occur in patients with severe hypertriglyceridemia. To date, cardiovascular outcome studies of fibrates have produced varied results, with some studies suggesting a small benefit, particularly in patients with other factors besides hypertriglyceridemia, such as low HDL-C or metabolic syndrome, and others showing no benefit. A meta-analysis including data from 45,058 participants from 18 clinical trials showed that fibrate therapy was associated with a significant decrease in major cardiovascular events (relative risk [RR] reduction 10%; P=0.048), although this did not translate into a benefit for all-cause mortality (RR reduction 0%; P=0.92). In another meta-analysis of 7,389 patients with high TG levels (>200 mg/dL [2.3 mmol/L]), fibrate therapy was associated with a 25% decrease in vascular events, and in 5,068 patients with both high TG and low HDL-C levels (<40 mg/dL [1 mmol/L]), a 29% decrease in vascular events was observed.

**Table 2 Summary of available triglyceride-lowering therapies**

<table>
<thead>
<tr>
<th>TG-lowering agent</th>
<th>Proposed mechanisms of action</th>
<th>Lipid-modifying effects (%)</th>
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<tr>
<td><strong>Fibrates</strong>&lt;sup&gt;2,72,73&lt;/sup&gt;</td>
<td>Weak agonists of PPAR-&lt;sub&gt;α&lt;/sub&gt; Decrease production of apoC-III Increase LPL expression Increase VLDL, IDL, and LDL apoB100 catabolism Decrease CETP activity Increase HDL&lt;sub&gt;α&lt;/sub&gt; and HDL&lt;sub&gt;β&lt;/sub&gt;</td>
<td>TG: –36&lt;br&gt;Non-HDL-C: –6 to –16&lt;br&gt;HDL-C: 10&lt;br&gt;LDL-C: –8</td>
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<tr>
<td><strong>Niacin</strong>&lt;sup&gt;2,72,122&lt;/sup&gt;</td>
<td>Decrease TG synthesis Inhibit hepatic DGAT-2 Accelerate hepatic apoB degradation Decrease hepatic secretion of VLDL and LDL Decrease hepatic apoAI catabolism Inhibit removal of HDL-apoAI</td>
<td>TG: –20&lt;br&gt;Non-HDL-C: –7 to –39&lt;br&gt;HDL-C: 16&lt;br&gt;LDL-C: –12</td>
</tr>
</tbody>
</table>

Abbreviations: apo, apolipoprotein; CETP, cholesteryl ester transfer protein; DGAT-2, diacylglycerol acyltransferase 2; DHA, docosahexaenoic acid; EPA, eicosapentaenoic acid; HDL, high-density lipoprotein; HDL-C, high-density lipoprotein cholesterol; IDL, intermediate-density lipoprotein; IDL-C, low-density lipoprotein; LDL, low-density lipoprotein; LDL-C, low-density lipoprotein cholesterol; LPL, lipoprotein lipase; Lp-PLA2, lipoprotein-associated phospholipase A2; non-HDL-C, non-high-density lipoprotein cholesterol; PPAR<sub>α</sub>, peroxisome proliferator-activated receptor alpha; TG, triglyceride; VLDL, very low-density lipoprotein.
Fibrate therapy is associated with a number of adverse effects, including increases in creatinine levels, myopathy, and, in rare cases, rhabdomyolysis, especially when used in combination with other lipid-lowering therapies. Gemfibrozil, in particular, has been shown to increase exposure to and reduce the renal clearance of statins by inhibiting their glucuronidation, potentially leading to severe side effects. However, fenofibrate appears to be better tolerated than other fibrates, with no cases of rhabdomyolysis observed with fenofibrate–statin combination therapy in two large-scale clinical trials. Fenofibrate is often recommended for use in combination with statin therapy in patients requiring additional non-HDL-C lowering. Studies have so far failed to demonstrate any significant reduction in cardiovascular risk with fenofibrate–simvastatin combination therapy, compared with simvastatin monotherapy. However, in a subgroup of patients with a TG level in the upper third (≥204 mg/dL [≥2.30 mmol/L]) and an HDL-C level in the lower third (≤34 mg/dL [≤0.88 mmol/L]), there was a nonsignificant trend (P=0.06) for reduction in cardiovascular risk.

Niacin

Prescription strength niacin is indicated to reduce elevated total cholesterol, LDL-C, apoB, and TG levels and to increase HDL-C in patients with primary hyperlipidemia, severe hypertriglyceridemia, and mixed dyslipidemia. Niacin has been shown to decrease TG levels by 20%, LDL-C levels by 12%, and non-HDL-C levels by 7%–39% and to increase HDL-C levels by an average of 16%. Despite improvements in coronary atherosclerosis and carotid intima-media thickness, niacin does not appear to impact risk for cardiovascular events when added to statin therapy. That being said, one post hoc analysis showed that among patients with TG >200 mg/dL (≥2.3 mmol/L) and HDL-C <32 mg/dL (<0.8 mmol/L), niacin may reduce cardiovascular events by 37% (P<0.05).

The use of niacin is often limited due to the high incidence of associated adverse effects. The most common adverse effect, cutaneous vasodilatation or “flushing,” reportedly occurs in up to 70% of patients receiving niacin therapy and often leads to treatment discontinuation. Although attempts have been made to reduce the incidence of niacin-induced flushing using laropiprant, a specific antagonist of the prostaglandin D2 receptor, this combination therapy does not significantly reduce the risk of major cardiovascular events and has been shown to actually increase the incidence of adverse effects. Additional adverse effects, including hyperglycemia, insulin resistance, hyperuricemia, myopathy, pruritus, and elevations in liver enzymes, are also associated with niacin monotherapy or statin–niacin combination therapy.

Omega-3 fatty acids

Three prescription omega-3 fatty acid formulations are currently approved in the US, including formulations comprising omega-3 carboxylic acids, a mixture of long-chain omega-3 fatty acids in free fatty acid form, primarily eicosapentaenoic acid (EPA), docosahexaenoic acid (DHA), and docosapentaenoic acid (Epanova®); omega-3 fatty acid ethyl esters, a mixture of long-chain omega-3 fatty acid ethyl esters, primarily EPA and DHA (Lovaza®, Omtryg®, and some generics); and icosapent ethyl (EPA ethyl esters) (Vascepa®). These prescription omega-3 fatty acids have been shown to reduce plasma TG levels by 25%–45%, VLDL-C levels by 20%–42%, and non-HDL-C levels by 8%–14% in patients with severe hypertriglyceridemia. Increases in HDL-C levels of 5%–9% have also been observed with DHA-containing formulations. Higher doses of each formulation and higher baseline TG levels are associated with greater TG reductions. Of note, DHA-containing formulations are thought to increase LPL expression, leading to increased TG removal from circulating VLDL and chylomicron particles. This results in the increased production of IDL particles, some of which undergo further catabolism by LPL to produce LDL particles. Consequently, DHA-containing formulations have been shown to significantly increase LDL-C levels in patients with severe hypertriglyceridemia by up to 45%. However, these increases are accompanied by reductions in non-HDL-C. This may be of particular importance, as non-HDL-C levels are more strongly associated with the risk of cardiovascular events than LDL-C, particularly in patients with hypertriglyceridemia.

Like fibrates, cardiovascular outcome studies of omega-3 fatty acids have also produced inconsistent results. A meta-analysis including data from 63,030 individuals from 20 clinical trials demonstrated that omega-3 fatty acid therapy did not have an impact on a composite cardiovascular end point or total mortality (P=0.24 and P=0.28, respectively) but was associated with a significantly decreased rate of vascular death (RR 0.86; 95% confidence interval, 0.75–0.99; P=0.03). In subgroup analysis, protection against the composite cardiovascular end point with omega-3 fatty acid use was observed in those trials that enrolled patients with high baseline TG levels (≥150 mg/dL [≥1.7 mmol/L]; RR 0.82; 95% confidence interval 0.74–0.91) versus those with lower baseline TG levels (P=0.006). Notably, many of the clinical
Emerging TG-lowering therapies

As studies further elucidate the proatherogenic mechanisms of TRLs, new treatment targets are beginning to emerge, and a number of novel therapeutic agents are currently in clinical development. These agents include antisense apoC-III inhibitors and LPL gene replacement therapy. Volanesorsen (formerly ISIS-APOCIII Rx) is an antisense apoC-III inhibitor currently undergoing Phase III clinical trials. It inhibits hepatic apoC-III synthesis by binding to apoC-III messenger RNA, thereby promoting its degradation. Phase II clinical trials conducted in patients with LPL deficiency demonstrated that Volanesorsen successfully reduced levels of apoC-III, TG, and non-HDL-C. However, concerns have been raised that the inhibition of hepatic apoC-III synthesis could lead to hepatic lipid accumulation.

Another agent with promise in reducing TG levels is alipogene tiparvovec (AAV1-LPLS447X), a nonreplicating and nonreplacing adeno-associated viral vector that delivers copies of the human LPL gene to muscle tissue. It has been approved in Europe for adult patients diagnosed with familial LPL deficiency, a disorder characterized by severe hypertriglyceridemia and increased risk of pancreatitis. However, it has not yet been approved in the US. Intramuscular administration of alipogene tiparvovec has been shown to be generally well tolerated and is associated with clinical improvement and reduced incidence and severity of acute pancreatitis.

Mipomersen and lomitapide are both approved by the US Food and Drug Administration for the treatment of homozygous familial hypercholesterolemia. However, as both agents interfere with TRL synthesis in the liver, along with substantial reductions in LDL-C, they have also been shown to diminish plasma TG levels. Mipomersen is an antisense oligonucleotide that blocks the translation of the apoB-100 gene, reducing its synthesis and consequently decreasing the circulation of atherogenic apoB-100-containing lipoproteins, including VLDL, IDL, and LDL. Lomitapide is a microsomal TG transfer protein inhibitor that blocks the microsomal TG transfer protein-mediated transfer of lipids to apoB, and therefore, results in significant reductions in VLDL, LDL-C, TG, and non-HDL-C. Currently, the sole indication for both mipomersen and lomitapide is homozygous familial hypercholesterolemia. However, in the future they may be found to have a place in the treatment of hypertriglyceridemia once their effect on TRLs has been fully defined.

Additional treatment considerations

Several guidelines recommend evaluating TG levels in the fasting state. However, nonfasting TG levels significantly correlate with increased levels of remnant lipoprotein cholesterol, are associated with a stepwise increase in the incidence of cardiovascular events, and have been shown to be a superior predictor of cardiovascular risk compared with fasting TG levels. Therefore, the measurement of nonfasting TG levels in the absence of a high-fat meal is now suggested by the American Heart Association and the European Atherosclerosis Society.

Additionally, although a number of guidelines recommend LDL-C as a primary treatment target, there is increasing evidence that non-HDL-C, the sum of the total cholesterol...
carried by all atherogenic lipoproteins (including LDL, IDL, and VLDL and chylomicrons and their remnants) provides a better indication of cardiovascular risk than LDL-C.\textsuperscript{6} Therefore, the use of non-HDL-C as a treatment target is now advocated by several guidelines, particularly in patients with hypertriglyceridemia.\textsuperscript{8,69,70} It is also worth noting that LDL-C levels are ordinarily estimated by the Friedewald equation.\textsuperscript{116} The Friedewald equation is inapplicable for patients with fasting TG > 400 mg/dL; however, it has also been shown to underestimate LDL-C levels in the presence of TG levels \(\geq 150\) mg/dL, particularly in patients with low LDL-C. Therefore, alternative evaluation is warranted in high-risk patients with elevated TG levels.\textsuperscript{117,118}

**Conclusion**

Although some studies have failed to report a link between raised TG levels and CVD when adjusting for other lipid parameters, there is now a large body of evidence indicating that elevated TG levels are independently associated with an increased incidence of cardiovascular events. Genetic studies have further clarified the causal association between TRLs and CVD, with variants in several key genes involved in TRL metabolism, such as LPL and its regulators, shown to be strongly associated with cardiovascular risk. Moreover, it has been shown that the strength of a variant’s effect on TG levels correlates with the magnitude of the variant’s effect on CVD. TRLs are known to contribute to the progression of atherosclerosis and CVD via a number of direct and indirect mechanisms. For example, they directly contribute to intimal cholesterol deposition and are also involved in the activation and enhancement of several proinflammatory, proapoptotic, and procoagulant pathways. Consequently, a number of guidelines now recommend the use of TG-lowering agents, primarily fibrates, niacin, or omega-3 fatty acids, if elevated TG or non-HDL-C levels remain following lifestyle intervention and statin therapy. Although cardiovascular outcome studies of TG-lowering agents have produced inconsistent results, post hoc analyses of large-scale clinical trials have shown a clinical benefit of TG-lowering agents in patients with hypertriglyceridemia, those with atherogenic dyslipidemia, and patients with features of metabolic syndrome. However, further large-scale clinical trials conducted in patients with clinically defined hypertriglyceridemia are required to further clarify this clinical benefit.

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**References**

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