HDL-cholesterol levels and cardiovascular risk: acCETPing the context

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This editorial refers to 'CETP genotype predicts increased mortality in statin-treated men with proven cardiovascular disease: an adverse pharmacogenetic interaction' by J.J. Regieli et al., on page 2792

High levels of plasma high-density lipoprotein cholesterol (HDL-C) are associated with a decreased risk of cardiovascular disease. Cholesterol ester transfer protein (CETP) facilitates the transfer of cholesteryl esters from HDL to triglyceride-rich lipoproteins, and its inhibition raises HDL-C levels. Accordingly, interest has focused on CETP inhibition to augment current lipid-lowering strategies.

In a substudy from the Regression Growth Evaluation Statin Study (REGRESS) angiographic trial cohort, Regieli and colleagues evaluated 812 statin-treated participants and found that the 60% who were carriers of the TaqIB-B2 CETP gene allele had ~20% lower CETP and 15% higher HDL-C serum concentrations than those without the allele, and a markedly increased risk of atherosclerotic disease mortality. The hazard ratio per each B2 copy was 1.60 (P = 0.01). The authors concluded that statin use in the setting of low CETP levels worsens clinical outcome in patients with proven cardiovascular disease. In light of the Investigation of Lipid Level Management to Understand its Impact in Atherosclerotic disease (ILLUMINATE) trial—which showed a 61% increase in combined mortality and cardiovascular events in 15 000 individuals at high risk for coronary events who were treated with atorvastatin plus torcetrapib, a CETP inhibitor, compared with atorvastatin alone— the Regieli study highlights the fact that much remains unknown about the interaction between HDL-C metabolism and cardiovascular disease.

CETP is a plasma glycoprotein that shuttles cholesteryl esters, phospholipids, and triglycerides between HDL-C, very low-density lipoprotein and low-density lipoprotein (LDL). CETP genotypes that result in moderate inhibition of CETP activity are associated with increased serum HDL-C levels and reduced coronary risk. In contrast, pharmacological CETP inhibition has not proven to be clinically beneficial. In statin-treated patients, concomitant CETP inhibition with torcetrapib did not slow the progression of atherosclerosis and was associated with a 25% increase in cardiovascular events and 58% increase in mortality, despite increased HDL-C levels. Although this increase in events has been attributed to the ‘off-target’ effects of torcetrapib (i.e. increased systemic arterial pressure and serum aldosterone), the possibility remains that combination statin and CETP inhibition therapy is harmful. There are several possible mechanisms that may account for the negative outcomes.

While CETP synthesis is generally assigned to liver and adipose tissue, many organs express CETP mRNA, including spleen, bone marrow, adrenal gland, intestine, kidney, lung, prostate, brain, heart, and skeletal muscle. Apart from its role in regulating arterial pressure and serum aldosterone, the possibility remains unknown about the interaction between HDL-C metabolism and cardiovascular disease.

CETP inhibition may render them pro-inflammatory.

While the current study demonstrates that the B2 CETP gene allele is associated with increased mortality in comparison with...
the B1 allele, it suggests but does not prove that ‘the efficacy of statin therapy to reduce cardiovascular risk depends on CETP genotype’ as the authors conclude. Since all patients in the study received statin therapy, the association of CETP genotype and response to statin therapy cannot be determined without no-statin control groups. The suggestion that statin therapy in the setting of low CETP levels is associated with increased mortality, however, is not without merit. The reason for the varying results may relate to differences in (i) study design; (ii) the types of study patients enrolled; (iii) medication compliance; or (iv) duration of follow-up. It does not, however, appear to be related to differences in the percentage of patients in each study treated with a statin, as proposed by Regieli and co-workers; when the statin-treated patients within each study are assessed, the TaqIB genotype was not associated with coronary artery disease risk.

It is important to note that the study by Regieli et al. included no diabetic patients and exclusively evaluated men. Thus, whether their observations extend to diabetic patients or women requires additional investigation. CETP transfers oestradiol esters from HDL-C to LDL, where it may serve an anti-oxidant role. Whether CETP genotypes influence this transfer to modify the cardiovascula risk still needs to be determined.

In summary, the study by Regieli et al. uses a pharmacogenomic approach to suggest an interaction between statin therapy and CETP genotype. A pharmacogenomic approach has proven clinically useful in predicting the anti-coagulation response to warfarin therapy and the risk of statin-induced myopathy. However, before CETP genotyping is utilized to decide which patients should receive statin therapy, the results of the current study require confirmation in a properly performed, appropriately powered, prospective study inclusive of both placebo-treated and statin-treated patients with genotypes reflective of the general population. In addition, future studies are needed to uncover the mechanisms and circumstances whereby CETP inhibition—due to genetic variation or pharmacological manipulation—may increase cardiac risk despite favourable changes in HDL-C levels.

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References