

EDITORIAL COMMENT

Lipoprotein(a) and Cardiovascular Disease in Heterozygous Familial Hypercholesterolemia



Should We Also Blame the LDL Receptor?*

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Heterozygous familial hypercholesterolemia (FH) is associated with an elevated lifetime burden of cardiovascular disease (CVD) (1). However, despite early exposure to elevated plasma low-density lipoprotein cholesterol (LDL-C) levels, there is heterogeneity at the onset of CVD in this population. In addition to the presence of traditional risk factors such as hypertension, smoking, and low plasma high-density lipoprotein cholesterol levels, elevated lipoprotein(a) [Lp(a)] levels have been associated in some (2,3), but not all (4,5), studies with a greater prevalence of clinical and subclinical CVD in patients with FH. However, these studies have been limited by their cross-sectional or retrospective design and also, with exception of 1 study (3), by the small number of included subjects.

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A recent patient-level meta-analysis of prospective studies, as well as a large Mendelian randomization evaluation (6,7), have established an independent association between Lp(a) levels and the risk of CVD in the general population. This might occur because Lp(a) is an LDL-like lipoprotein. Nonetheless, Lp(a) carries an additional apolipoprotein, apo(a) (8), that inhibits fibrinolysis and may be the link between lipid transport and coagulation systems in the atherogenic process (8). In addition to its high cholesterol content and antifibrinolytic action, Lp(a) should also induce atherosclerosis by its pro-inflammatory action secondary to

increased binding and transport of oxidized phospholipids (9). Plasma Lp(a) levels are strongly determined genetically, and production rates predominate over catabolism as their main determinants (10,11).

In this issue of the *Journal*, Alonso et al. (12) elegantly report on the association between Lp(a) levels as a predictor of CVD and the interaction with the severity of *LDLR* mutations in a well-characterized Spanish cohort of patients with FH in SAFEHEART (Spanish Familial Hypercholesterolemia Cohort Study). The investigators initially found that Lp(a) levels were higher in patients with FH than in their nonaffected normolipidemic relatives. The investigators then showed that patients with FH and CVD had higher Lp(a) levels than either unrelated subjects with FH, but without CVD, or normolipidemic relatives without CVD. When subjects carrying the most frequent null and defective *LDLR* mutations were compared, those with null mutations had higher Lp(a) levels and a greater incidence of CVD. Moreover, for those with *LDLR* null mutations, who are in theory those with the highest risk of CVD due to more elevated LDL-C levels from birth (1), Lp(a) levels were higher in those with CVD. Also, subjects with null mutations and Lp(a) levels >50 mg/dl, a value associated with an elevated risk of CVD in the general population (13), had the shortest free survival time from onset of CVD. Finally, Lp(a) levels were independently associated with CVD in the whole FH cohort.

This is a well-performed proof-of-concept study with a large number of study subjects and is carefully analyzed. All subjects with FH had a molecular diagnosis, and the non-affected control subjects were their relatives. Of course, the cross-sectional nature of the study precludes the final conclusion that an elevated Lp(a) level is indeed an independent risk factor for CVD in patients with FH, as it is for the general population (6,7). However, this study advances the knowledge in the field and adds important information regarding a possible interaction between more severe *LDLR* mutations and higher Lp(a) levels with a greater risk of CVD.

The lack of a role of *LDLR* in catabolism of Lp(a) is intriguing, because this lipoprotein encloses apoB-100, the natural binder of *LDLR* within its particle. Studies in both subjects with homozygous FH (10) and transgenic mice (11) failed to show reduced clearance from plasma of Lp(a) in the absence of the *LDLR*. Indeed, statins that increase hepatic *LDLR* expression do not show a significant effect on Lp(a) plasma levels (13). However, the role of a putative influence of the *LDLR* on Lp(a) catabolism cannot be discarded because subjects with homozygous FH present with higher Lp(a) levels than both subjects with heterozygous FH and normolipidemic subjects (14). The finding of the current study (12) of higher Lp(a) levels in those with more frequent *LDLR* null mutations suggests that this association might not be attributable to chance only. However, one cannot discard that those with more severe *LDLR* mutations can also inherit greater apo(a) production capacity and therefore have higher Lp(a) plasma levels.

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Recent evidence showing that antibodies that block PCSK9 (proprotein convertase subtilisin/kexin type 9), a protein that facilitates the degradation of LDLR in the liver, reduce, not only LDL-C and apoB-100, but also Lp(a) (15) plasma levels in patients with FH adds more controversy to the matter. Should we also blame LDLR for higher Lp(a) plasma levels and possibly a greater risk of CVD in subjects with FH?

Notwithstanding biological controversies, the study by Alonso et al. (12) suggests an interaction between the more severe *LDLR* mutations and Lp(a) as a cause of CVD in patients with FH. The prospective phase of SAFE-HEART is needed to confirm the findings of the current evaluation.

Despite the strong evidence between increased Lp(a) plasma levels and a greater risk of CVD in the general population (6,7) and the suggested elevated risk of CVD shown in the study by Alonso et al. (12) and studies from others in subjects with FH (2,3), there is very little evidence that a reduction in Lp(a) level prevents CVD (16). It is important to show that newer therapies that reduce Lp(a) levels in addition to LDL-C levels (15,17) will prevent CVD.

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