So Far, PCSK9 Inhibitors Work for All Heterozygous FH Patients

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In heterozygous FH patients with biallelic null mutations, evolocumab fails to reduce LDL-C.¹ Does something similar occur in heterozygous FH patients with PCSK9 mutations? PCSK9 mutations affect PCSK9’s affinity for the LDL receptor (LDLR). Could they also result in variable affinities for a PCSK9 monoclonal antibody?

Although studies of alirocumab and evolocumab have already been done in heterozygous FH patients, they did not require genetic screening for inclusion.²³ In the current issue of Circulation: Cardiovascular Genetics, Hopkins et al⁴ describe the results of an interventional study of alirocumab (150 mg every 2 weeks) administered to 13 FH patients harboring PCSK9 gain-of-function mutations. The study represents the first such investigation in humans with gain-of-function PCSK9 mutations.

Alirocumab achieved 50% to 70% LDL-C reductions—with 12 of 13 patients getting to an LDL-C <70 mg/dL—similar to prior studies, including those of heterozygous FH patients (Figure).¹⁻³⁻⁶ The high affinity of monoclonal antibodies to PCSK9 and the high therapeutic concentrations of antibody likely overwhelm any difference in affinity that a PCSK9 mutation may cause.

Clearly, as long as interference occurs in the PCSK9:LDLR interaction, a therapeutic response results. In fact, previous in vitro studies showed that mutant PCSK9 proteins can be blocked by a peptide that inhibits the PCSK9:LDLR interaction.⁷ Alirocumab seems safe for PCSK9 FH patients. No serious adverse events directly attributable to the study drug and no reports of myalgia or neurocognitive disorders—both of which have been noted in other studies⁵,⁶—occurred.

A few questions still remain. Will evolocumab have similar efficacy for PCSK9 FH patient? It has a different epitope and, thus, the potential exists for variable affinity to mutant PCSK9. Also, the study by Hopkins et al lacks comparator groups of FH patients with LDLR or apolipoprotein B (APOB) mutations, so no direct comparison of the efficacy of PCSK9 monoclonal antibodies has been reported thus far.

More than 10 variants are associated with FH, with the p.S127R and p.D374Y mutations being most studied functionally. The aspartic acid residue at position 374 lies in the binding interface between the LDLR and PCSK9, and its substitution to tyrosine results in increased affinity between PCSK9 and LDLR at the cell surface. The substitution of serine residue at position 127 to arginine does not affect binding affinity at the cell surface; how it results in downregulation of LDLR from the surface remains unclear.

Because of the rarity of gain-of-function PCSK9 mutations (<2% of genetic hypercholesterolemias’), few descriptions exist of their phenotype, especially in comparison with the more common genetic cause of FH (LDLR and APOB).
Previous studies with a single gain-of-function mutation in \textit{PCSK9} (D374Y) have noted higher levels of LDL-C and earlier onset of coronary heart disease than heterozygous mutations in either \textit{LDLR} or \textit{APOB}.\footnote{8}

Hopkins et al\textsuperscript{9} assembled clinical characteristics of the largest cohort (n=160) to date of patients with FH-causing \textit{PCSK9} mutations. A few issues may limit the interpretation of their data. Controls were not selected from the same population as cases; because coronary heart disease rates vary throughout the world, some bias may have been introduced into their analysis of premature coronary heart disease. Also, some of the mutations had LDL-C levels lower than typical for FH (156 mg/dL in 6 patients with Pro71Leu mutations from the Netherlands, for example) making it unclear whether these are true disease-causing mutations.

Overall, it seems that \textit{PCSK9} gain-of-function mutations cause more severe hypercholesterolemia than \textit{LDLR} mutations. Such information may be useful for ongoing FH genetic screening studies\footnote{10} and future bio-banking efforts, where sequencing of \textit{PCSK9} can be reserved for patients with the highest LDL-C who lack any mutations in \textit{LDLR} or \textit{APOB}.

Although several questions remain unanswered about the use of PCSK9 inhibitors, they seem safe and efficacious at lowering LDL-C levels in most patients, regardless of the underlying genetics. Of course, the results of ongoing cardiovascular outcome studies will ultimately have the largest effect on both prescriber and payer practices.

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