

Putting Into Perspective the Hazards of Untreated Familial Hypercholesterolemia

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In this issue of *JAHA*, Kjaergaard et al¹ describe a 21-year follow-up of 118 heterozygous carriers of low-density lipoprotein (LDL) receptor (*LDLR*) mutations causing heterozygous familial hypercholesterolemia (FH) together with 102 of their noncarrier relatives. These FH patients and their relatives had been identified between 1992 and 1994 through cascade screening, starting from *LDLR* mutation-carrying probands in 32 families. Lipid-lowering treatment with statins was recommended to all mutation-carrying FH participants. Primary outcomes were tracked in the Danish National Patient Registry and included death from any cause, myocardial infarction, coronary revascularization, ischemic stroke, transient ischemic attack, and peripheral artery disease. FH patients and unaffected relatives with occurrence of any of these outcomes prior to the baseline were excluded from the analysis. A set of controls were matched 10:1 by birth year and sex from the Danish Civil Registration System.

Despite the vast majority of the FH patients being treated with statins, probably for most of the follow-up period, risk of the primary outcome remained elevated with a hazard ratio 1.65 (95% CI 1.17–2.33). Not unexpectedly, the risk specifically for coronary events was considerably higher among the FH patients (hazard ratio 5.91, 95% CI 3.83–9.10), while no significant excess risk was seen for stroke or total mortality. Indeed, total mortality was somewhat less in FH patients and considerably less in their unaffected relatives, perhaps because of the increased attention to healthy lifestyle these families often display. The strengths of the study include the

long-term and comprehensive follow-up together with carefully defined FH cases and unaffected relatives. Limitations are the relatively small size of the cohort as well as incomplete information about lipid-lowering therapy (only available after 2004) and standard coronary risk factors.

This study comes on the heels of new recognition of a much higher prevalence of FH than previously appreciated, making it the most common, serious monogenic disorder in humans. New, objective screening projects in large US populations, with genetic testing done without regard to lipid levels, place the prevalence of FH mutation carriers (with mutations in *LDLR*, *APOB*, and *PCSK9*) in 3 different studies at 1 in 204,² 1 in 211,³ and 1 in 222⁴; more than double older estimates of 1 in 500. Similar estimates for prevalence of FH are reported for European populations.^{5–8} Yet, FH remains seriously underdiagnosed and inadequately treated.

Risk of premature coronary disease among *untreated* FH remains of major interest for public health planning and to better appreciate the need for early identification and treatment. Accurate estimates of cumulative risk for coronary artery disease (CAD) by age and sex in untreated FH as compared with non-FH subjects are also of interest for purposes of calculating the likelihood of having FH in a newly developed algorithm for clinical diagnosis of FH.⁹ The estimates of cardiovascular disease and CAD incidence and associated hazard ratios by Kjaergaard et al herein¹ are hampered by incompletely documented effects of prior treatment. The same may be said of the several other recent overall estimates of risk associated with genetically defined FH.^{2–4} None of the overall estimates of risk in FH mutation carriers versus noncarriers in these studies should be taken as risk associated with untreated FH. Perhaps the best estimate of risk of CAD in untreated FH in these studies comes from Khera et al,³ who calculated a hazard ratio of 22.3 ($P < 0.0001$) for CAD among FH mutation carriers with LDL cholesterol (LDL-C) ≥ 190 mg/dL (compared with a reference group of noncarriers with LDL-C < 130 mg/dL). Interestingly, in this study FH mutation carriers had ≈ 2 - to 3-fold higher risk compared with noncarriers at the same current LDL-C levels. Evidence was presented that FH mutation carriers had experienced longer

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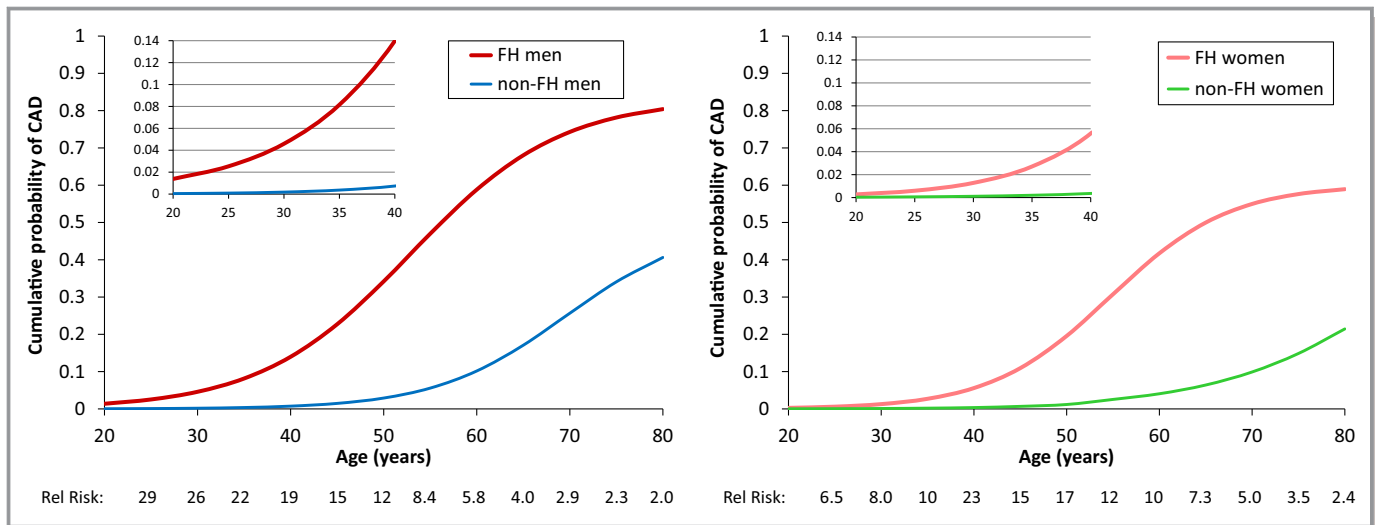


Figure. Cumulative probability of developing coronary artery disease (CAD) in men (left) and women (right) with heterozygous familial hypercholesterolemia (FH) as compared with unaffected relatives or the general population (non-FH). Relative risk (Rel Risk), as the ratio of cumulative CAD risk in FH divided by non-FH, is shown below each curve.

exposure to higher LDL-C levels, suggesting a cause for the increased risk in FH. For this reason, without a genetic or reliable clinical diagnosis of FH, even large data sets from pooled prospective studies would be expected to underestimate the risk of FH when only LDL-C cut points are utilized to determine risk.¹⁰

Perhaps the largest study estimating CAD risk in FH patients and the effects of lipid-lowering was performed among 1950 previously untreated FH patients identified by 1990 by the Dutch Lipid Network group.¹¹ Of these, 413 FH patients (mean age 41.7) were started on statins while 1537 (mean age 38.2) continued off treatment. Remarkably, after 12.5 years of follow-up, $\approx 67\%$ of the untreated group had developed coronary disease. Use of statins led to an adjusted 82% risk reduction ($P < 0.001$).

Older estimates of CAD risk attributable to FH from the pre-statin era provide important insights. The severe consequences of not treating FH sufficiently early with appropriate medication, as happens all too often, should be considered paramount when planning a clinical approach to FH. In addition, as noted above, estimates for age- and sex-specific cumulative risk for CAD in FH and non-FH subjects are used in a new algorithm for clinical diagnosis of FH.⁹ To estimate CAD risks in untreated FH compared with the general population, data were utilized from previously published estimates of cumulative CAD risk for FH patients (excluding angina only) and their unaffected relatives^{12,13} supplemented by lifetime cumulative estimates for CAD in the general population.^{14,15} Adjustment downward for reported coronary disease in very young women in the general population was made because of the frequent

(>50%) finding of nonatherosclerotic disease in this subgroup (because of microvascular disease, arteritis, embolic and thrombotic events such as those associated with birth control and smoking, and fibromuscular dysplasia with spontaneous dissection, all of which are much more common in young women than men).^{16–18}

The resulting smoothed logistic curves are shown in the Figure. Relative risks for CAD in FH as compared to non-FH exceed 25 in young men. Even higher relative risk estimates for CAD death, greater than 40- to 100-fold, were reported for young, untreated FH in the Simon-Broome Registry before the statin era.^{19–21} The diminishing relative risk with age, as general population rates rise, is consistent with other reports, even among partially treated FH as shown in a large Norwegian registry of genetically verified FH.²² Note that the cumulative risk for a CAD event in untreated FH reaches $\approx 20\%$ by age 42 in men and by age 50 in women. Therefore, untreated FH may be considered a “coronary risk equivalent” by age 32 in men and age 40 in women. The much lower apparent risks reported by Kjaergaard et al¹ herein may therefore be considered a qualified success attributable to the extensive use of statins motivated by the early screening efforts among these FH patients. Nevertheless, the 5-fold residual risk for coronary disease should serve as a strong impetus for aggressive finding and treatment of FH patients at an early age.

Disclosures

None.

References

- Kjærgaard KA, Christiansen MK, Schmidt M, Olsen MS, Jensen HK. Long-term cardiovascular risk in heterozygous familial hypercholesterolemia relatives identified by cascade screening. *J Am Heart Assoc*. 2017;6:e005435. DOI: 10.1161/JAHA.116.005435.
- Do R, Stitzel NO, Won HH, Jorgensen AB, Duga S, Angelica Merlini P, Kiezun A, Farrall M, Goel A, Zuk O, Guella I, Asselta R, Lange LA, Peloso GM, Auer PL, Project NES, Girelli D, Martinelli N, Farlow DN, DePristo MA, Roberts R, Stewart AF, Saleheen D, Danesh J, Epstein SE, Sivapalaratnam S, Hovingh GK, Kastelein JJ, Samani NJ, Schunkert H, Erdmann J, Shah SH, Kraus WE, Davies R, Nikpay M, Johansen CT, Wang J, Hegele RA, Hechter E, Marz W, Kleber ME, Huang J, Johnson AD, Li M, Burke GL, Gross M, Liu Y, Assimes TL, Heiss G, Lange EM, Folsom AR, Taylor HA, Olivieri O, Hamsten A, Clarke R, Reilly DF, Yin W, Rivas MA, Donnelly P, Rossouw JE, Psaty BM, Herrington DM, Wilson JG, Rich SS, Bamshad MJ, Tracy RP, Cupples LA, Rader DJ, Reilly MP, Spertus JA, Cresci S, Hartiala J, Tang WH, Hazen SL, Allayee H, Reiner AP, Carlson CS, Kooperberg C, Jackson RD, Boerwinkle E, Lander ES, Schwartz SM, Siscovick DS, McPherson R, Tybjaerg-Hansen A, Abecasis GR, Watkins H, Nickerson DA, Ardisino D, Sunyaev SR, O'Donnell CJ, Altshuler D, Gabriel S, Kathiresan S. Exome sequencing identifies rare LDLR and APOA5 alleles conferring risk for myocardial infarction. *Nature*. 2015;518:102–106.
- Khera AV, Won HH, Peloso GM, Lawson KS, Bartz TM, Deng X, van Leeuwen EM, Natarajan P, Emdin CA, Bick AG, Morrison AC, Brody JA, Gupta N, Nomura A, Kessler T, Duga S, Bis JC, van Duijn CM, Cupples LA, Psaty B, Rader DJ, Danesh J, Schunkert H, McPherson R, Farrall M, Watkins H, Lander E, Wilson JG, Correa A, Boerwinkle E, Merlini PA, Ardisino D, Saleheen D, Gabriel S, Kathiresan S. Diagnostic yield and clinical utility of sequencing familial hypercholesterolemia genes in patients with severe hypercholesterolemia. *J Am Coll Cardiol*. 2016;67:2578–2589.
- Abul-Husn NS, Manickam K, Jones LK, Wright EA, Hartzel DN, Gonzaga-Jauregui C, O'Dushlaine C, Leader JB, Lester Kirchner H, Lindbuchler DM, Barr ML, Giovanni MA, Ritchie MD, Overton JD, Reid JG, Metpally RP, Wardeh AH, Borecki IB, Yancopoulos GD, Baras A, Shuldiner AR, Gottesman O, Ledbetter DH, Carey DJ, Dewey FE, Murray MF. Genetic identification of familial hypercholesterolemia within a single U.S. health care system. *Science*. 2016;354:1550.
- Nordestgaard BG, Chapman MJ, Humphries SE, Ginsberg HN, Masana L, Descamps OS, Wiklund O, Hegele RA, Raal FJ, Defesche JC, Wiegman A, Santos RD, Watts GF, Parhofer KG, Hovingh GK, Kovanen PT, Boileau C, Averno M, Boren J, Bruckert E, Catapano AL, Kuivenhoven JA, Pajukanta P, Ray K, Stalenhoef AF, Stroes E, Taskinen MR, Tybjaerg-Hansen A. Familial hypercholesterolaemia is underdiagnosed and undertreated in the general population: guidance for clinicians to prevent coronary heart disease: consensus statement of the European Atherosclerosis Society. *Eur Heart J*. 2013;34:3478–3490a.
- Sjouke B, Kusters DM, Kindt I, Besseling J, Defesche JC, Sijbrands EJ, Roeters van Lennep JE, Stalenhoef AF, Wiegman A, de Graaf J, Fouchier SW, Kastelein JJ, Hovingh GK. Homozygous autosomal dominant hypercholesterolaemia in the Netherlands: prevalence, genotype-phenotype relationship, and clinical outcome. *Eur Heart J*. 2015;36:560–565.
- Benn M, Watts GF, Tybjaerg-Hansen A, Nordestgaard BG. Mutations causative of familial hypercholesterolaemia: screening of 98 098 individuals from the Copenhagen General Population Study estimated a prevalence of 1 in 217. *Eur Heart J*. 2016;37:1384–1394.
- Wald DS, Bestwick JP, Morris JK, Whyte K, Jenkins L, Wald NJ. Child-parent familial hypercholesterolemia screening in primary care. *N Engl J Med*. 2016;375:1628–1637.
- Hopkins PN. Genotype-guided diagnosis in familial hypercholesterolemia: population burden and cascade screening. *Curr Opin Lipidol*. 2017;28:136–143.
- Perak AM, Ning H, de Ferranti SD, Gooding HC, Wilkins JT, Lloyd-Jones DM. Long-term risk of atherosclerotic cardiovascular disease in US adults with the familial hypercholesterolemia phenotype. *Circulation*. 2016;134:9–19.
- Versmissen J, Oosterveer DM, Yazdanpanah M, Defesche JC, Basart DC, Liem AH, Heeringa J, Witteman JC, Lansberg PJ, Kastelein JJ, Sijbrands EJ. Efficacy of statins in familial hypercholesterolaemia: a long term cohort study. *BMJ*. 2008;337:a2423.
- Stone NJ, Levy RI, Fredrickson DS, Verter J. Coronary artery disease in 116 kindred with familial type II hyperlipoproteinemia. *Circulation*. 1974;49:476–488.
- Hopkins PN. Encouraging appropriate treatment for familial hypercholesterolemia. *Clin Lipidol*. 2010;5:339–354.
- Lloyd-Jones DM, Wilson PW, Larson MG, Beiser A, Leip EP, D'Agostino RB, Levy D. Framingham risk score and prediction of lifetime risk for coronary heart disease. *Am J Cardiol*. 2004;94:20–24.
- JBS3 Board. Joint British Societies' consensus recommendations for the prevention of cardiovascular disease (JBS3). *Heart*. 2014;100:ii1–ii67.
- Sanghani M, Gulati M. Sex differences in the pathophysiology, treatment, and outcomes in IHD. *Curr Atheroscler Rep*. 2015;17:511.
- Gulati M, Shaw LJ, Bairey Merz CN. Myocardial ischemia in women: lessons from the NHLBI WISE study. *Clin Cardiol*. 2012;35:141–148.
- Vanzetto G, Berger-Coz E, Barone-Rochette G, Chavanon O, Bouvaist H, Hacini R, Blin D, Machecourt J. Prevalence, therapeutic management and medium-term prognosis of spontaneous coronary artery dissection: results from a database of 11,605 patients. *Eur J Cardiothorac Surg*. 2009;35:250–254.
- Scientific Steering Committee on behalf of the Simon Broome Register Group. Risk of fatal coronary heart disease in familial hypercholesterolaemia. Scientific Steering Committee on behalf of the Simon Broome Register Group. *BMJ*. 1991;303:893–896.
- Scientific Steering Committee on behalf of the Simon Broome Register Group. Mortality in treated heterozygous familial hypercholesterolaemia: implications for clinical management. *Atherosclerosis*. 1999;142:105–112.
- Neil A, Cooper J, Betteridge J, Capps N, McDowell I, Durrington P, Seed M, Humphries SE. Reductions in all-cause, cancer, and coronary mortality in statin-treated patients with heterozygous familial hypercholesterolaemia: a prospective registry study. *Eur Heart J*. 2008;29:2625–2633.
- Mundal L, Iglund J, Ose L, Holven KB, Veierod MB, Leren TP, Retterstol K. Cardiovascular disease mortality in patients with genetically verified familial hypercholesterolemia in Norway during 1992–2013. *Eur J Prev Cardiol*. 2017;24:137–144.

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