Cardiometabolic Risk

High Intestinal Cholesterol Absorption Is Associated With Cardiovascular Disease and Risk Alleles in *ABCG8* and *ABO*

Evidence From the LURIC and YFS Cohorts and From a Meta-Analysis

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Objectives	This study sought to determine whether high intestinal cholesterol absorption represents a cardiovascular risk factor and to link ABCG8 and ABO variants to cardiovascular disease (CVD).
Background	Plant sterol-enriched functional foods are widely used for cholesterol lowering. Their regular intake yields a 2-fold increase in circulating plant sterol levels that equally represent markers of cholesterol absorption. Variants in <i>ABCG8</i> and <i>ABO</i> have been associated with circulating plant sterol levels and CVD, thereby suggesting atherogenic effects of plant sterols or of cholesterol uptake.
Methods	The cholestanol-to-cholesterol ratio (CR) was used as an estimate of cholesterol absorption because it is independent of plant sterols. First, we investigated the associations of 6 single nucleotide polymorphisms in <i>ABCG8</i> and <i>ABO</i> with CR in the LURIC (LUdwisghafen RIsk and Cardiovascular health study) and the YFS (Young Finns Study) cohorts. Second, we conducted a systematic review and meta-analysis to investigate whether CR might be related to CVD.
Results	In LURIC, the minor alleles of rs4245791 and rs4299376 and the major alleles of rs41360247, rs6576629, and rs4953023 of the ABCG8 gene and the minor allele of rs657152 of the ABO gene were significantly associated with higher CR. Consistent results were obtained for rs4245791, rs4299376, rs6576629, and rs4953023 in YFS. The meta-analysis, including 6 studies and 4,362 individuals, found that CR was significantly increased in individuals with CVD.
Conclusions	High cholesterol absorption is associated with risk alleles in <i>ABCG8</i> and <i>ABO</i> and with CVD. Harm caused by elevated cholesterol absorption rather than by plant sterols may therefore mediate the relationships of <i>ABCG8</i> and <i>ABO</i> variants with CVD. (J Am Coll Cardiol 2013;62:291–9) © 2013 by the American College of Cardiology Foundation

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Abbreviations and Acronyms
CI = confidence interval
CVD = cardiovascular disease
GWA = genome-wide association
LDL = low-density lipoprotein
RR = risk ratio
SMD = standardized mean difference
SNP = single nucleotide polymorphism

The American Heart Association has recommended plant sterolenriched foods (e.g., margarines) for cholesterol lowering (1). Moreover, the European Heart Association and the European Atherosclerosis Society have mentioned plant sterol-enriched functional foods as cholesterol lowering agents in their guidelines for the management of dyslipidemia (2). The regular intake of plant sterols reduces low-density lipoprotein (LDL) cholesterol by about 13 mg/dl (3) but also raises circulating plant sterols from

about 1 mg/dl (4) by approximately 2-fold (5). Patients with sitosterolemia, a rare genetic disorder caused by mutations in the ATP-binding cassette transporters G5 and G8 (ABCG5 and ABCG8) (6), have up to 50-fold increased circulating plant sterols and may develop early onset cardiovascular disease (CVD) (7). Hence, it has been suggested that plant sterols are atherogenic (8-10). These concerns have been reinforced by the detection of plant sterols in carotid atherosclerotic plaques (11). In addition, plant sterol intake was related to increased plant sterol content in aortic valve cusps (12). Two recent studies have confirmed that consumption of plant sterols as part of a dietary portfolio and as an adjunct to treatment with ezetimibe has favorable effects on the lipid profile (13,14). Nevertheless, in continuance of the long-lasting safety discussion, these studies also garnered critical comments (15,16). The debate was further fueled by a genome-wide association (GWA) study showing that

common variants in the ABCG8 (major allele of rs41360247 and minor allele of rs4245791) and ABO (minor allele of rs657152) genes increased both circulating plant sterols and cardiovascular risk (17). Correlations of variants in ABCG8 (minor allele of rs4299376) and ABO with CVD subsequently have been replicated in other large-scale GWA studies (18-20). However, because circulating plant sterols are markers for cholesterol uptake (21), the genetic data may also indicate adverse vascular effects of high cholesterol absorption.

The current study consisted of a genetic analysis and a meta-analysis; its purpose was to investigate whether high intestinal cholesterol absorption represents a cardiovascular risk factor and to link ABCG8 and ABO variants to CVD. We used the ratio of circulating cholestanol-to-cholesterol to estimate levels of intestinal cholesterol absorption independently of plant sterol concentrations (22).

Methods

Genetic analyses. STUDY DESIGN AND PARTICIPANTS. Genetic association studies were performed in the LURIC (LUdwigshafen RIsk and Cardiovascular health study) and the YFS (Young Finns Study) (23,24).

LURIC is a cross-sectional and prospective German cohort study designed to investigate biochemical and genetic cardiovascular risk factors. A total of 3,316 participants referred for coronary angiography were recruited between July 1997 and January 2000 at the Ludwigshafen Heart Center (23). Measurements of lathosterol, cholestanol, campesterol, and sitosterol were completed in 1,257 LURIC participants who did not receive stating and did not have type 1 diabetes (25). Individuals in this subgroup with available data on ABCG8 or ABO single nucleotide polymorphisms (SNPs) were included in the current analyses.

YFS is a Finnish population-based, 27-year follow-up study on the evolution of cardiovascular risk factors from childhood to adulthood (24). The first cross-sectional study was conducted in 1980 at 5 centers and included 3,596 participants in the age groups of 3, 6, 9, 12, 15, and 18 years who were randomly chosen from the national population registry. In 2001, a total of 2,620 individuals, who were then aged 24 to 39 years, were studied. The sterol and lipid determinations used in the current analysis were taken from the year 2001 participants. Sterol and genetic data were available in 434 subjects.

Both studies were approved by the local ethical committees and performed according to the Declaration of Helsinki. Informed written consent was obtained from all participants (23,24). Diabetes mellitus was categorized according to the 2009 criteria of the American Diabetes Association (26).

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LABORATORY ANALYSES. All laboratory measurements were performed on fasting blood samples. In LURIC, cholesterol was measured with enzymatic reagents from WAKO (Neuss, Germany) on a WAKO 30 R or Olympus AU640 analyzer (Tokyo, Japan) (23). Lipoproteins were separated by a combined ultracentrifugation precipitation method (beta-quantification).

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