JACC: ASIA © 2023 THE AUTHOR. PUBLISHED BY ELSEVIER ON BEHALF OF THE AMERICAN COLLEGE OF CARDIOLOGY FOUNDATION. THIS IS AN OPEN ACCESS ARTICLE UNDER THE CC BY-NC-ND LICENSE (http://creativecommons.org/licenses/by-nc-nd/4.0/).

EDITORIAL COMMENT

What Can Be Seen From "Intracranial-Vascular"-Susceptibility Genetic Factor in "Cardiovascular-Susceptible" Familial Hypercholesterolemia*

A New Clue

Atsushi Nohara, MD, PHD

amilial hypercholesterolemia (FH) is 1 of the most common genetic diseases, with a genetic defect in LDL-receptor function.¹ Highly elevated serum LDL-C levels from birth are the cause of premature coronary artery disease (CAD) in these patients. Early diagnosis is essential to prevent their early deaths, but FH has been still underdiagnosed and undertreated in many countries, also in east Asia, despite efforts to raise awareness up to the present.

It has been a kind of mystery why elevated LDL-C in FH clearly accelerates atherosclerosis in coronary arteries but not in intracranial arteries. Without appropriate intensive therapies, patients with homozygous FH, a severe form of FH with 2 pathogenic variants, frequently die young of cardiovascular disease² but rarely of intracranial artery disease,³ with some exceptional cases. Also, in a large cohort of individuals with genetically verified heterozygous FH, the risks of cerebrovascular disease and ischemic stroke were not increased.⁴

Many LDL-lowering clinical trials have shown a clear risk reduction in cardiovascular disease but less reduction in cerebrovascular disease. Acceleration of

ISSN 2772-3747

atherosclerosis with LDL-C seems not the same in coronary arteries as in intracranial arteries, and the underlying mechanisms at the molecular level are not well elucidated.

DISCOVERY OF RNF213 GENE IN MOYAMOYA DISEASE

Moyamoya disease is a rare cause of stroke, and RNF213 is the major susceptibility gene in people from east Asia.⁵ It is characterized by progressive stenosis of the terminal portion of the internal carotid arteries and compensatory capillary collateral arteries. The Arg4810Lys variant of the RNF213 gene is strongly associated with moyamoya disease, but its penetrance rate is <1%. By contrast, the Arg4810Lys variant had been reported to have an important role in the development of intracranial major artery stenosis/occlusion (ICASO).⁶

THE RNF213 GENE VARIANT INCREASED ICASO IN FH IN A DIFFERENT MANNER THAN IN CAD

In this issue of *JACC: Asia*, Noda et al⁷ report that FH patients show an increased prevalence and severity of ICASO associated with the Arg4810Lys variant. FH may be a kind of naturally occurring accelerated atherosclerosis model in which affected individuals have elevated LDL-C from birth. It is intriguing that CAD-associated indices in FH, like tendon xanthomas, pathogenic LDLR gene variants,⁸ or "severe FH" with double mutations in LDLR and PCSK9 genes,⁹ were not associated with ICASO in FH. The Arg4810-Lys variant in RNF213 gene in FH was not associated with CAD in this study. This RNF213 gene variant

^{*}Editorials published in *JACC: Asia* reflect the views of the authors and do not necessarily represent the views of *JACC: Asia* or the American College of Cardiology.

From the Department of Clinical Genetics, Ishikawa Prefectural Central Hospital, Ishikawa, Japan.

The author attests they are in compliance with human studies committees and animal welfare regulations of the author's institution and Food and Drug Administration guidelines, including patient consent where appropriate. For more information, visit the Author Center.

seems basically a cerebrovascular-susceptive genetic factor.

The authors speculate that the Arg4810Lys variant in the RNF213 gene may downregulate PECAM-1, an important shear stress sensor, and may cause inflammatory changes and mitotic abnormalities in endothelial cells. The underlying mechanisms in the differences between coronary arteries and intracranial arteries should be elucidated in future studies.

Only exceptional cases of FH show symptomatic intracranial artery disease in young individuals, but this study could reveal a genetic background for these cases. This study may reveal 1 clinical clue to this mystery and may enable the development of new strategies for intracranial artery disease.

FUNDING SUPPORT AND AUTHOR DISCLOSURES

The author has reported that he has no relationships relevant to the contents of this paper to disclose.

ADDRESS FOR CORRESPONDENCE: Dr Atsushi Nohara, Department of Clinical Genetics, Ishikawa Prefectural Central Hospital, Kuratsuki-higashi 2-1, Kanazawa, Ishikawa 920-8530, Japan. E-mail: atsushinohara@mac.com.

REFERENCES

1. Mabuchi H. Half a century tales of familial hypercholesterolemia (FH) in Japan. *J Atheroscler Thromb.* 2017;24:189-207.

2. Nohara A, Tada H, Ogura M, et al. Homozygous familial hypercholesterolemia. *J Atheroscler Thromb*. 2021;28:665–678.

3. Tromp TR, Hartgers ML, Hovingh GK, et al. Worldwide experience of homzygous familial hypercholesterolaemia: retrospective cohort study. *Lancet.* 2022;399:719-728.

4. Hovland A, Mundal LJ, Igland J, et al. Risk of ischemic stroke and total cerebrovascular disease in familial hypercholesterolemia. *Stroke*. 2019;50: 172-174. https://doi.org/10.1161/STROKEAHA.118. 023456

5. Ihara M, Yamamoto Y, Hattori Y, et al. Moyamoya disease: diagnosis and interventions. *Lancet Neurol.* 2022;21:747-758.

6. Okazaki S, Morimoto T, Kamatani Y, et al. Moyamoya disease susceptibility variant RNF213 p. R4810K increases the risk of ischemic stroke attributable to large-artery atherosclerosis. *Circulation.* 2019;139:295-298.

7. Noda K, Hattori Y, Hori M, et al. Amplified risk of intracranial artery stenosis/occlusion associated with *RNF213* p.R4810K in familial hypercholesterolemia. *JACC: Asia.* 2023;3(4):625-633.

8. Tada H, Kawashiri MA, Nohara A, Inazu A, Mabuchi H, Yamagishi M. Impact of clinical signs and genetic diagnosis of familial hypercholesterolaemia

on the prevalence of coronary artery disease in patients with severe hypercholesterolaemia. *Eur Heart J.* 2017;38(20):1573–1579.

9. Santos RD, Gidding SS, Hegele RA, et al. Defining severe familial hypercholesterolaemia and the implications for clinical management: a consensus statement from the International Atherosclerosis Society Severe Familial Hypercholesterolemia Panel. *Lancet Diabetes Endocrinol*. 2016;4:850–861.

KEY WORDS familial hypercholesterolemia, genetic interaction, genetic mutation, intracranial artery stenosis/occlusion, RNF213 p.R4810K