

Filamin A Variant as a Possible Second-Hit Gene Promoting Moyamoya Disease-like Vascular Formation Associated With RNF213 p.R4810K Variant

Ikeuchi, Yasuhito

From the Department of Neurology, Japanese Red Cross Fukuoka Hospital, ■Department of Human Genetics (N. Miyake, N. Matsumoto), Yokohama City University Graduate School of Medicine, ■Department of Human Genetics, Research Institute, National Center for Global Health and Medicine, ■Department of Medicine and Clinical Science, Graduate School of Medical Sciences, Kyushu University

Kitayama, Jiro

From the Department of Neurology, Japanese Red Cross Fukuoka Hospital, ■Department of Human Genetics (N. Miyake, N. Matsumoto), Yokohama City University Graduate School of Medicine, ■Department of Human Genetics, Research Institute, National Center for Global Health and Medicine, ■Department of Medicine and Clinical Science, Graduate School of Medical Sciences, Kyushu University

Sahara, Noriyuki

From the Department of Neurology, Japanese Red Cross Fukuoka Hospital, ■Department of Human Genetics (N. Miyake, N. Matsumoto), Yokohama City University Graduate School of Medicine, ■Department of Human Genetics, Research Institute, National Center for Global Health and Medicine, ■Department of Medicine and Clinical Science, Graduate School of Medical Sciences, Kyushu University

Okata, Takuya

From the Department of Neurology, Japanese Red Cross Fukuoka Hospital, ■Department of Human Genetics (N. Miyake, N. Matsumoto), Yokohama City University Graduate School of Medicine, ■Department of Human Genetics, Research Institute, National Center for Global Health and Medicine, ■Department of Medicine and Clinical Science, Graduate School of Medical Sciences, Kyushu University

出版情報 : Neurology Genetics. 8 (5), 2022-10. Ovid Technologies (Wolters Kluwer Health)

他ージョン :

権利関係 : Creative Commons Attribution-NonCommercial-NoDerivatives 4.0 International



Filamin A Variant as a Possible Second-Hit Gene Promoting Moyamoya Disease–like Vascular Formation Associated With *RNF213* p.R4810K Variant

Yasuhito Ikeuchi, MD,* Jiro Kitayama, MD, PhD,* Noriyuki Sahara, MD, Takuya Okata, MD, Noriko Miyake, MD, PhD, Naomichi Matsumoto, MD, PhD, Takanari Kitazono, MD, PhD, and Tetsuro Ago, MD, PhD

Correspondence

Dr. Ago
ago.tetsuro.544@m.kyushu-u.ac.jp

Neurol Genet 2022;8:e200017. doi:10.1212/NXG.0000000000200017

Abstract

Background and Objective

The objective of this case report was to identify a second-hit gene that may promote Moyamoya disease (MMD)–like vascular formation in an individual having the *RNF213* p.R4810K variant.

Methods

We performed magnetic resonance imaging and genetic analyses of *RNF213* and *FLNA* in a 21-year-old woman, who showed Ehlers-Danlos–like symptoms and developed a first-ever unprovoked seizure, and of her healthy parents.

Results

We identified bilateral periventricular nodular heterotopia (PNH) as the cause of seizures and MMD-like vascular formation in the patient. The patient had the *RNF213* p.R4810K variant. Exome analysis identified c.4868delG in the X-linked *FLNA* gene encoding filamin A p.G1623Vfs*41, which could explain PNH and Ehlers-Danlos–like symptoms. Her mother had the same *FLNA* variant and had asymptomatic bilateral PNH, whereas her father had the *RNF213* variant and had normal cerebrovascular structure.

Discussion

The family study suggested that the *FLNA* variant promoted MMD-like vascular formation in a patient having the *RNF213* variant, while the *RNF213* variant amplified the phenotypic changes elicited by the *FLNA* abnormality. Collectively, we identified a gene abnormality in filamin A, a target of *RNF213*-mediated proteasomal degradation, that may promote MMD-like vascular formation as a possible second-hit gene in individuals having the *RNF213* p.R4810K variant.

*These authors contributed equally to this work.

From the Department of Neurology (Y.I., J.K., N.S., T.O.), Japanese Red Cross Fukuoka Hospital, Fukuoka, Japan; Department of Human Genetics (N. Miyake, N. Matsumoto), Yokohama City University Graduate School of Medicine, Yokohama, Japan; Department of Human Genetics (N. Miyake), Research Institute, National Center for Global Health and Medicine, Tokyo, Japan; Department of Medicine and Clinical Science (T.K., T.A.), Graduate School of Medical Sciences, Kyushu University, Fukuoka, Japan

Funding information and disclosures are provided at the end of the article. Full disclosure form information provided by the authors is available with the full text of this article at [Neurology.org/NG](https://neurology.org/NG).

The Article Processing Charge was funded by the authors.

This is an open access article distributed under the terms of the Creative Commons Attribution-NonCommercial-NoDerivatives License 4.0 (CC BY-NC-ND), which permits downloading and sharing the work provided it is properly cited. The work cannot be changed in any way or used commercially without permission from the journal.

Moyamoya disease (MMD) is characterized by chronic stenosis or occlusion of the terminal portion of the internal carotid artery accompanied by the formation of abnormal vascular networks. Its familial and regional prevalence in East Asia suggests genetic involvement: A genome-wide association study identified the *RNF213* (NM_001256071.3) c.14429G>A (p.R4810K) variant as a susceptible gene for MMD.¹ This variant is found in 80% of patients with MMD in Japan, whereas 1.8% of healthy controls have this variant.² Because MMD develops in approximately 1 per 10,000 Japanese individuals, it can be estimated that 1%–2% of patients having the variant would develop MMD. However, recent reports have demonstrated that the *RNF213* variant is frequently identified in stroke and nonstroke patients with large-artery atherosclerosis, thereby expanding its impact beyond MMD.² Experiments in mice with genetic deletion of *Rnf213* or carrying human *RNF213* c.14429G>A knock-in did not show any abnormal cerebrovascular structures. Therefore, it is believed that additional genetic or environmental factors are required to develop MMD in individuals having *RNF213* p.R4810K variant, that is, the two-hit theory in MMD.³

Case Description

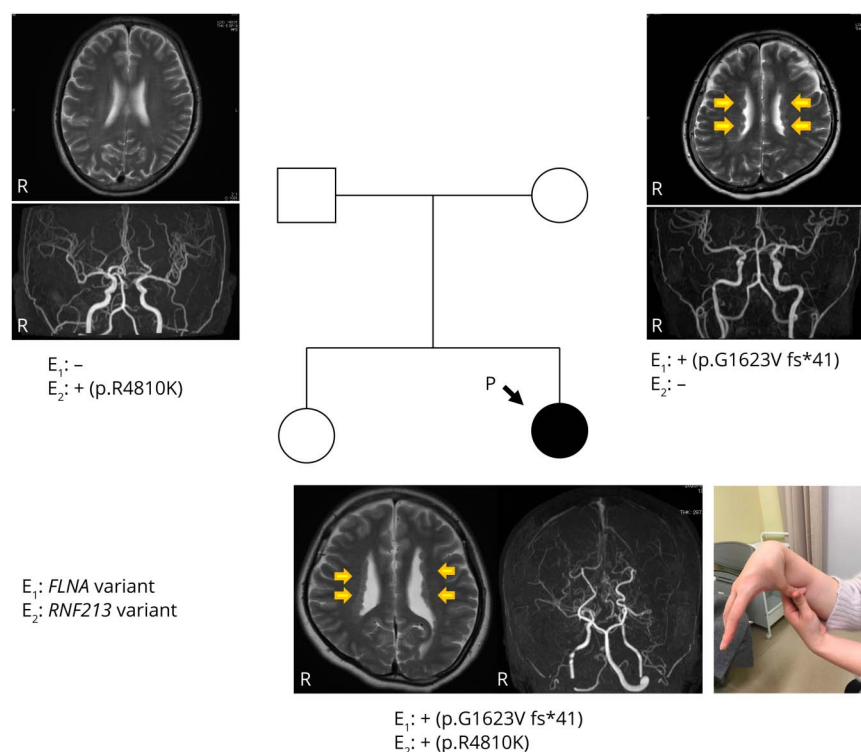
A 21-year-old woman developed generalized tonic-clonic seizures when she visited our hospital due to headaches and transient loss of consciousness. IV administration of diazepam easily controlled seizures. She had no mental retardation or known CNS disorders except depression and migraine. We observed

joint hypermobility (Figure 1) and ecchymoma in her lower limbs, with a medical history of patellar luxation and spontaneous pneumothorax. MRI revealed bilateral periventricular nodular heterotopia (PNH) (Figure 1). Magnetic resonance angiography (MRA) revealed stenosis of the terminal portion of the bilateral internal carotid artery and MMD-like abnormal vascular networks (Figure 1). Transthoracic echocardiography revealed mild aortic regurgitation. Bilateral PNH accompanied by Ehlers-Danlos-like symptoms resembled the variant phenotype of the X-linked gene *FLNA* (NM_001456.4), encoding filamin A.⁴ An exome analysis of the patient identified a previously unknown heterozygous variant in *FLNA* (c.4868delG), creating a protein with Gly1623Val substitution with early termination, omitting the C-terminal 983 amino acids (p.G1623V fs*41) (Figure 2). We also sequenced *RNF213* and found the heterozygous c.14429G>A variant (Figure 2). We further performed brain MRI/MRA and genetic analyses in the patient's parents. Her mother had the same *FLNA* variant, but not the *RNF213* variant, with asymptomatic bilateral PNH and normal cerebrovascular structure (Figures 1 and 2). By contrast, her father did not show any abnormalities on MRI and MRA while having the heterozygous *RNF213* variant (Figures 1 and 2).

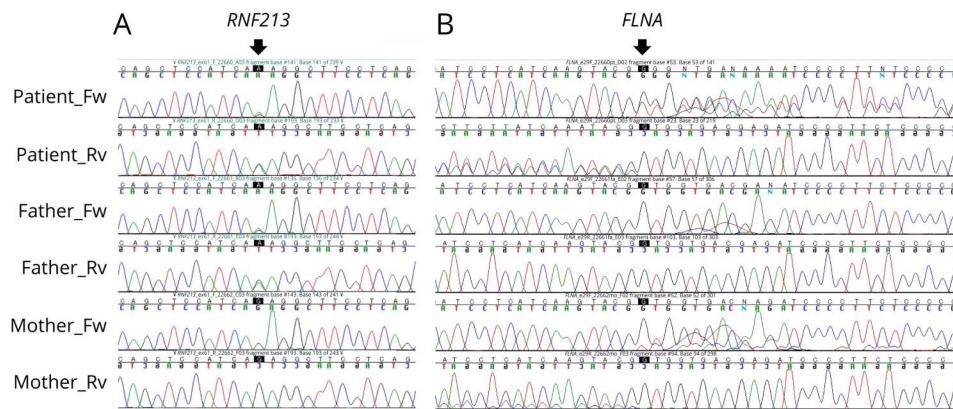
Discussion

In this study, we clearly demonstrated that the patient inherited the *RNF213* variant from her father and the *FLNA* variant from her mother (Figures 1 and 2). Because only the

Figure 1 Pedigree and Neuroimaging



(Patient) Bilateral PNH (yellow arrows) on T2-weighted images and narrowing of the bilateral internal carotid artery terminal portion accompanied by MMD-like vascular formation on MRA. Hypermobility of the thumb is shown. (Mother) Asymptomatic bilateral PNH (yellow arrows) without MMD-like vascular formation and Ehlers-Danlos-like symptoms. (Father) No abnormalities on MRI/MRA. MMD = Moyamoya disease; MRA = magnetic resonance angiography; PNH = periventricular nodular heterotopia.

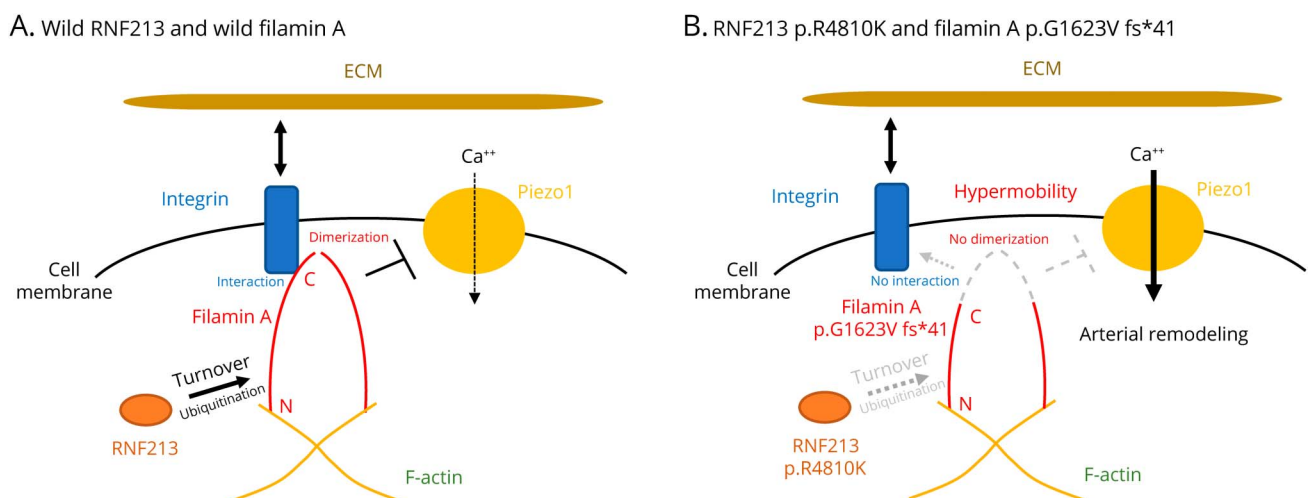
Figure 2 Sequencing of *RNF213* and *FLNA*

(A) A single nucleotide substitution of G to A at 14429 in the *RNF213* gene (c.14429G>A, p.Arg4810Lys) is identified in the patient and her father. (B) A single nucleotide deletion at 4868 in the *FLNA* gene (c.4868delG), leading to p.Gly1623Val fs*41 is identified in the patient and her mother. Fw = forward; Rv = reverse.

patient having the 2 variants manifested symptomatic PNH accompanied by Ehlers-Danlos-like symptoms and MMD-like vascular formation, each gene abnormality alone may cause only subtle phenotypic changes while their coexistence may mutually amplify them. It is important that the direct interaction between the 2 molecules is known: Filamin A undergoes RNF213-mediated ubiquitination and proteasomal degradation for its turnover.² Because ligase activity of *RNF213* p.R4810K is decreased, the abnormal filamin A may accumulate in cells expressing *RNF213* and filamin A.² Filamin A regulates cellular mobility through its dimerization and interaction with various molecules, including integrin, via its C-terminal portion, which further interacts with extracellular

matrix proteins, while the N-terminal portion participates in actin-binding and ubiquitination (Figure 3).⁴ The C-terminally truncated filamin A may impair the migration of newborn neurons from the periventricular zone to the cortex during development, resulting in PNH. In adults, the filamin A variant may cause flow-mediated or pressure-mediated hypermobility of vascular smooth muscle cells, resulting in increased calcium influx through the mechanosensor channel, thereby inducing arterial remodeling and stenosis of cerebral arteries (Figure 3).⁵

We cannot completely exclude that the *FLNA* abnormality coexists independently of MMD in the patient. Even if it is related to MMD, it may be uncommon in MMD. There has

Figure 3 Putative Mechanism Underlying Hypermobility of Vascular Smooth Muscle Cells Leading to Arterial Remodeling in the Patient Having *RNF213* p.R4810K and *FLNA* p.G1623V fs*41

(A) Filamin A forms a homodimer and interacts directly with integrin via its C-terminal portion, while its N-terminal portion interacts with filamentous actin (F-actin) and undergoes RNF213-mediated ubiquitination for its turnover. Integrin interacts with ECM, thereby preventing hypermobilization of vascular smooth muscle cells. (B) The filamin A variant with p.G1623V fs*41, which lacks its C-terminal portion, can neither form a homodimer nor interact with integrin, thereby causing hypermobilization of vascular smooth muscle cells. The hypermobilization of the cells increases the calcium influx through the mechanosensor channel Piezo1 that leads to arterial remodeling. Because the *RNF213* p.R4810K variant has a decreased ubiquitin ligase activity that is required for the degradation and turnover of filamin A, the filamin A variant would accumulate and amplify its phenotypic changes as a dominant-negative form in the cells expressing the *RNF213* p.R4810K variant. ECM = extracellular matrix proteins.

been 1 case series that briefly described a 3-year-old male patient having an *FLNA* abnormality among 54 patients with MMD.⁶ Although *FLNA* variants presenting with PNH often show X-linked dominant inheritance, we should note that some *FLNA* variants are transmitted in X-linked recessive mode and their phenotypic changes may be affected by X-inactivation.⁷

Collectively, we identify *FLNA* as a possible second-hit gene that may affect MMD-like vascular formation in an individual having the *RNF213* p.R4810K variant. *RNF213* also functions as an E3-ubiquitin ligase for itself and other molecules, including NFAT.² Moreover, *RNF213* contributes to lipid metabolism.² Thus, besides *FLNA*, there may be other *RNF213*-related genes whose abnormality can elicit MMD-like vascular formation associated with the *RNF213* p.R4810K variant.

Acknowledgment

The authors thank the patient and her parents who provided consent for the publication of the case report. The authors also thank Editage (editage.com) for the English language editing.

Study Funding

This work was supported in part by the Japan Agency for Medical Research and Development (AMED) under Grant numbers JP20ek0109486, JP21ek0109549, JP21cm0106503, and JP21ek0109493 (to N. Matsumoto); the Japan Society for the Promotion of Science (JSPS) KAKENHI (Grant number JP19H03621 to N. Miyake); and the Takeda Science Foundation (to N. Matsumoto and N. Miyake).

Disclosure

The authors report no disclosures relevant to the manuscript. Full disclosure form information provided by the authors is available with the full text of this article at Neurology.org/NG.

Publication History

Received by *Neurology: Genetics* April 18, 2022. Accepted in final form June 30, 2022. Submitted and externally peer reviewed. The handling editor was Alexandra Durr, MD, PhD.

Appendix Authors

Name	Location	Contribution
Yasuhito Ikeuchi, MD	Department of Neurology, Japanese Red Cross Fukuoka Hospital, Fukuoka, Japan	Drafting/revision of the manuscript for content, including medical writing for content; major role in the acquisition of data; Study concept or design; analysis or interpretation of data

Appendix (continued)

Name	Location	Contribution
Jiro Kitayama, MD, PhD	Department of Neurology, Japanese Red Cross Fukuoka Hospital, Fukuoka, Japan	Drafting/revision of the manuscript for content, including medical writing for content; major role in the acquisition of data; Study concept or design; analysis or interpretation of data
Noriyuki Sahara, MD	Department of Neurology, Japanese Red Cross Fukuoka Hospital, Fukuoka, Japan	Drafting/revision of the manuscript for content, including medical writing for content; major role in the acquisition of data; analysis or interpretation of data
Takuya Okata, MD	Department of Neurology, Japanese Red Cross Fukuoka Hospital, Fukuoka, Japan	Drafting/revision of the manuscript for content, including medical writing for content; major role in the acquisition of data; analysis or interpretation of data
Noriko Miyake, MD, PhD	Department of Human Genetics, Yokohama City University Graduate School of Medicine, Yokohama, Japan; Department of Human Genetics, Research Institute, National Center for Global Health and Medicine, Tokyo, Japan	Major role in the acquisition of data; analysis or interpretation of data
Naomichi Matsumoto, MD, PhD	Department of Human Genetics, Yokohama City University Graduate School of Medicine, Yokohama, Japan	Drafting/revision of the manuscript for content, including medical writing for content; analysis or interpretation of data
Takanari Kitazono, MD, PhD	Department of Medicine and Clinical Science, Graduate School of Medical Sciences, Kyushu University, Fukuoka, Japan	Drafting/revision of the manuscript for content, including medical writing for content
Tetsuro Ago, MD, PhD	Department of Medicine and Clinical Science, Graduate School of Medical Sciences, Kyushu University, Fukuoka, Japan	Drafting/revision of the manuscript for content, including medical writing for content; Study concept or design; analysis or interpretation of data

References

- Kamada F, Aoki Y, Narisawa A, et al. A genome-wide association study identifies *RNF213* as the first Moyamoya disease gene. *J Hum Genet*. 2011;56(1):34-40.
- Mineharu Y, Miyamoto S. *RNF213* and *GUCY1A3* in Moyamoya disease: key regulators of metabolism, inflammation, and vascular stability. *Front Neurol*. 2021;12:687088.
- Fujimura M, Sonobe S, Nishijima Y, et al. Genetics and biomarkers of Moyamoya disease: significance of *RNF213* as a susceptibility gene. *J Stroke*. 2014;16(2):65-72.
- Feng Y, Walsh CA. The many faces of filamin: a versatile molecular scaffold for cell motility and signalling. *Nat Cell Biol*. 2004;6(11):1034-1038.
- Retailleau K, Duprat F, Arhatte M, et al. *Piezo1* in smooth muscle cells is involved in hypertension-dependent arterial remodeling. *Cell Rep*. 2015;13(6):1161-1171.
- Amlie-Lefond C, Ellenbogen RG. Factors associated with the presentation of moyamoya in childhood. *J Stroke Cerebrovasc Dis*. 2015;24(6):1204-1210.
- Kyndt F, Gueffet JP, Probst V, et al. Mutations in the gene encoding filamin A as a cause for familial cardiac valvular dystrophy. *Circulation*. 2007;115(1):40-49.