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Serum α_1 -antitrypsin level and phenotype associated with familial moyamoya disease

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T. Amano () S. Inoha · C.-M. Wu T. Matsushima · K. Ikezaki Department of Neurosurgery, Graduate School of Medical Sciences, Kyushu University, 3–1-1 Maidashi, 812–8582 Higashi-ku, Fukuoka, Japan e-mail: amano@ns.med.kyushu-u.ac.jp Tel.: +81-92-6425524 Fax: +81-92-6425526 Abstract Background: Obstructive vascular lesions at the terminal portion of the internal carotid arteries are thought to be the primary and essential lesions in moyamoya disease. The etiology remains unknown. To detect possible mediators of the thickened intima of moyamoya disease, we measured serum alpha-1antitrypsin (α 1-AT) levels and characterized the phenotype of patients with familial moyamoya disease. Patients and methods Fifty-six individuals were examined, including 29 patients with moyamoya disease from 14 families. Serum α 1-AT levels were analyzed by electroimmunoassay and genomic phenotype by isoelectric focusing. *Results:* All individuals had a normal α 1-AT phenotype. The average serum α 1-AT level in moyamoya disease patients was significantly higher than that of normal individuals, although both were within the normal range. *Conclusions:* These findings suggest that serum α 1-AT level may be a marker, rather than an etiologic factor, indicating the progression of moyamoya disease.

Keywords Moyamoya disease · Alpha-1-antitrypsin

Introduction

The primary lesion in moyamoya disease is a progressive stenotic or occlusive change involving the circle of Willis and its major branches, including the distal segments of the internal carotid arteries [16]. The etiology of moyamoya disease remains unknown, although many clinical, pathological, pathophysiological, and genetic studies have been carried out.

Congenital factors have been considered to be important due to the high incidence among Asians and occasional familial occurrence. There is, however, no simple and useful marker to distinguish those who will develop moyamoya disease in familial cases, although several genetic factors seem to be significantly associated with moyamoya disease [2, 14, 17, 18, 32].

Intimal thickening due to smooth muscle cell proliferation and migration [16] is a primary structural abnormality in the major stenotic cerebral arteries in moyamoya disease. Moreover, similar lesions have been observed in vessels of the heart, kidney, and other organs [12] suggesting that systemic etiological factors are associated with moyamoya disease. Although the origin of moyamoya disease and the reason why it is limited to the major cerebral vessels remain unclear, several factors associated with the vascular wall, such as nitric oxide (NO), platelet-derived growth factor (PDGF) [1, 3], vascular endothelial growth factor (VEGF), or elastase [30] may be biochemically related to the etiology of moyamoya disease. Yamamoto et al. [29] have reported that elastin levels in smooth muscle cells are higher in patients with moyamoya disease. Elastin and collagen are the main components of the arterial wall, and an imbalance between a serum proteinase and its inhibitor may give rise to vascular matrix degeneration [6, 19]. Alpha-1-antitrypsin (α 1-AT) is a serine protease inhibitor that primarily functions to protect tissue by inhibiting elastase or collagenase [21]. Quantitatively high levels of α 1-AT can result in structural abnormalities of the vascular wall by excessive proliferation of smooth muscle cells and by synthesis of vascular matrix components such as elastin and collagen. Low levels of α 1-AT, on the other hand, can result in the destruction of elastin, collagen, and other connective tissue by increasing relative protease activity [21]. Ikari et al. [11] showed that α 1-AT is a critical antiapoptotic factor for vascular smooth muscle cells, while a deficiency of α 1-AT has been reported to increase the risk of developing intracranial aneurysms [24, 25, 28] and fibromuscular dysplasia [26].

We hypothesized that serum levels and a genetic abnormality of α 1-AT may affect the development of moyamoya disease.

Patients and methods

We studied 56 Japanese individuals (26 males and 30 females) from 14 families, each with 3–5 subjects. Each subject or the parents of the 14 families received informed consent and agreed to participate in this study. Twenty-nine (10 males and 19 females) had a definitive diagnosis of moyamoya disease (based on guidelines for the diagnosis of moyamoya disease established by the Research Committee on Spontaneous Occlusion of the Circle of Willis of the Ministry of Health and Welfare of Japan) [15].

Serum α 1-AT levels were determined by electroimmunoassay [20, 26]. α 1-AT phenotypes were determined by isoelectric focusing in polyacrylamide gels at pH values between 3.5 and 5.0, using genomic DNA extracted from peripheral blood [20, 26].

Statistical analysis was performed using the Student's *t*-test. Differences with a p value <0.05 were considered statistically significant.

Results

The numbers of subjects with each α 1-AT phenotype are listed in Table 1. The 'others' group in Table 1 consists of PiMM2, PiM1M1, PiM1M2, or PiGM alleles. All individuals had the normal M phenotypes except for one with a rare normal PiGM allele. None of the subjects had abnormal phenotypes such as PiZ or PiS, causing quantitative deficiency of α 1-AT levels. Thus, no specific α 1-AT phenotype appears to be associated with moyamoya disease.

The average serum α 1-AT levels of normal individuals and of individuals with moyamoya disease were 131.1±23.6 (mean ± SD) and 152.0±39.1 mg/dl respectively (Table 1). Patients with moyamoya disease showed significantly higher values than normal individuals (p<0.05). The average serum α 1-AT levels in both groups, however, were within the normal range (from 100 to 190 mg/dl). Only 4 individuals had values above 190 mg/dl (3 moyamoya patients and 1 normal individual, data not shown). None of the subjects had decreased levels of serum α 1-AT (below 100 mg/dl).

Discussion

Moyamoya disease occurs predominantly in people of Asian origin and the incidence of familial occurrence has been estimated to be 10% [9, 17, 34]. A massive case control study by the Research Committee on Moyamoya Disease in Japan failed to show any positive associations with environmental agents, including previous infections [10, 13, 33]. Therefore, certain genetic factors are likely to be related to the etiology of moyamoya disease, but have thus far remained unknown.

We investigated serum α 1-AT levels and phenotypes in familial moyamoya disease. More than 75 different α 1-AT alleles are known, which qualitatively and quantitatively affect a1-AT protein [8, 21]. Most normal individuals have the M phenotype (M, M1, or M2), expressing normal levels of α 1-AT. The most common alleles associated with a quantitative deficiency of α 1-AT are PiZ and PiS, resulting in pulmonary emphysema or liver cirrhosis [5, 7, 26]. In contrast, particular alleles associated with an excess of α 1-AT have not been identified. A deficiency of serum α 1-AT associated with genetic factors may cause degradation of the arterial wall, through inadequate protection against protease activity [5, 22, 26]. In the present study, however, all individuals had the normal a1-AT phenotype including a rare normal GM phenotype, indicating that there was no specific genetic abnormality of α 1-AT in association with familial moyamoya disease. Thus, it is not possible to use α 1-AT phenotype to predict those susceptible family members who may develop moyamoya disease.

On the other hand, the average serum α 1-AT level of moyamoya disease patients was comparatively higher, although it was not above the normal range. α 1-AT is also recognized as an acute phase reactant similar to C reactive protein (CRP), increasing in response to the

Table 1 The number of alpha-
1-antitrypsin phenotype and the
average serum alpha-1-anti-
trypsin level (αI -AT alpha-1-
antitrypsin, ns not significant)

α 1-AT phenotype	Moyamoya disease		Normal individuals	
	n	Serum α1-AT level (mg/dl) ^a	n	Serum α1-AT level (mg/dl) ^a
М	7	112.9±6.2 (ns)	5	108.8±7.5
MM	11	155.5±35.9*	7	131.9±11.5
MM1	8	151.4±47.7*	11	131.0±29.4
Others	3	136.0±11.8	4	147.5±47.9
Total	29	152.0±39.1**	27	131.1±23.6

^a Mean±SD **p*<0.01; ***p*<0.05 general stimulus of inflammation [4]. Previous reports have suggested that inflammatory stimuli and the subsequent response of inflammatory cells may produce a proliferative response in smooth muscle cells, leading to the thickened intima in patients with moyamoya disease [23, 27, 31]. In our study, serum samples were obtained from moyamoya disease subjects during the development of ischemic changes. The quantitative changes in serum α 1-AT might reflect tissue damage or inflammation in the progressive state of moyamoya disease, rather than being a primary cause of moyamoya disease. However, continuously high levels of α 1-AT may also result in structural abnormalities of the vascular wall due to excessive proliferation of smooth muscle cells and synthesis of vascular matrix components such as elastin and collagen. Thus, further study is necessary to clarify:

- 1. The relationship between serum α 1-AT level and severity of ischemia in moyamoya disease
- 2. The chronological changes in serum α 1-AT levels
- 3. The involvement of other serine protease inhibitors

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