

Association of Intracranial Aneurysms With Aortic Aneurysms in 125 Patients With Fusiform and 4253 Patients With Saccular Intracranial Aneurysms and Their Family Members and Population Controls

Arttu Kurtelius, BM;* Nelli Väntti, BM;* Behnam Rezai Jahromi, MD; Olli Tähtinen, MD, PhD; Hannu Manninen, MD, PhD; Juha Koskenvuo, MD, PhD; Riikka Tulamo, MD, PhD; Satu Kotikoski, BM; Heidi Nurmonen, BM; Olli-Pekka Kämäräinen, MD, PhD; Terhi Huttunen, MD, PhD; Jukka Huttunen, MD, PhD; Mikael von und zu Fraunberg, MD, PhD; Timo Koivisto, MD, PhD; Juha E. Jääskeläinen, MD, PhD;[†] Antti E. Lindgren, MD, PhD[†]

Background—Varying degrees of co-occurrence of intracranial aneurysms (IA) and aortic aneurysms (AA) have been reported. We sought to compare the risk for AA in fusiform intracranial aneurysms (fIA) and saccular intracranial aneurysms (sIA) disease and evaluate possible genetic connection between the fIA disease and AAs. Additionally, the characteristics and aneurysms of the fIA and sIA patients were compared.

Methods and Results—The Kuopio Intracranial Aneurysm Database includes all 4253 sIA and 125 fIA patients from its Eastern Finnish catchment population, and 13 009 matched population controls and 18 455 first-degree relatives to the IA patients were identified, and the Finnish national registers were used to identify the individuals with AA. A total of 33 fIA patients were studied using an exomic gene panel of 37 genes associated with AAs. Seventeen (14.4%) fIA patients and 48 (1.2%) sIA patients had a diagnosis of AA. Both fIA and sIA patients had AAs significantly more often than their controls (1.2% and 0.5%) or relatives (0.9% and 0.3%). In a competing risks Cox regression model, the presence of fIA was the strongest risk factor for AA (subdistribution hazard ratio 7.6, 95% CI 3.9–14.9, P<0.0005). One likely pathogenic variant in COL5A2 and 3 variants of unknown significance were identified in MYH11, COL11A1, and FBN1 in 4 fIA patients.

Conclusions—The prevalence of AAs is increased slightly in sIA patients and significantly in fIA patients. fIA patients are older and have more comorbid diseases than sIA patients but this alone does not explain their clinically significant AA risk. (*J Am Heart Assoc.* 2019;8:e013277. DOI: 10.1161/JAHA.119.013277.)

Key Words: abdominal aortic aneurysm • aortic aneurysm • fusiform intracranial aneurysm • genetics • intracranial aneurysm • saccular intracranial aneurysm • thoracic aortic aneurysm

neurysmal subarachnoid haemorrhage at an incidence of 7.9 per 100 000 and in an average age of 50 to 60 years, 1 is a devastating form of stroke: 20% of aneurysmal subarachnoid haemorrhage patients die instantly 2 and, when admitted alive, up to 30% within 12 months. 3 In most cases, the cause is rupture of a saccular intracranial aneurysm (sIA), formed during life in some 3% of general population. 4 Risk

factors for sIA include age, female sex, smoking, hypertension, sIA family history, and polycystic kidney disease.⁵ Fusiform IAs (fIAs), like aortic aneurysms (AA), are dilatations of arterial segments^{6,7}: fIAs are uncommon, and remain poorly characterized and difficult to treat.⁷

Aortic aneurysms (AA) are increasingly prevalent amongst aging population especially in aging, smoking, hypertensive

From the NeuroSurgery, NeuroCenter (A.K., N.V., S.K., H.N., O.-P.K., T.H., J.H., M.v.u.z.F., T.K., J.E.J., A.E.L.) and Department of Clinical Radiology (O.T., H.M.), Kuopio University Hospital, Kuopio, Finland; Institute of Clinical Medicine, Faculty of Health Sciences, University of Eastern Finland, Kuopio, Finland (A.K., N.V., O.T., H.M., S.K., H.N., O.-P.K., T.H., J.H., M.v.u.z.F., T.K., J.E.J., A.E.L.); Departments of Neurosurgery (B.R.J.), and Vascular Surgery (R.T.) Helsinki University Hospital, University of Helsinki, Finland; Blueprint Genetics, Helsinki, Finland (J.K.); Department of Clinical Physiology and Nuclear Medicine, Turku University Hospital, Turku, Finland (J.K.).

Correspondence to: Arttu Kurtelius, BM, Neurosurgery of NeuroCenter, Kuopio University Hospital, P.O. Box 1777, 70211 Kuopio, Finland. E-mail: arttu.kurtelius@gmail.com

Received May 15, 2019; accepted July 12, 2019.

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^{*}Dr Kurtelius and Dr Väntti are co-first authors.

[†]Dr Jääskeläinen and Dr Lindgren are co-last authors.

Clinical Perspective

What Is New?

- Prevalence of diagnosed aortic aneurysms in patients with intracranial aneurysms, assessed through the Finnish nationwide diagnosis register, is increased in comparison to the matched population controls and first-degree relatives of the patients with intracranial aneurysms.
- Patients with fusiform intracranial aneurysms, constituting 3% of all patients with intracranial aneurysms, are older and have a greater burden of comorbid diseases and cardiovascular risk factors in comparison with patients with saccular intracranial aneurysms.
- Presence of a fusiform intracranial aneurysm is strongly associated with aortic aneurysms, independent of other investigated cardiovascular risk factors.

What Are the Clinical Implications?

- Fusiform intracranial aneurysms are associated with a clinically significant co-occurrence rate of aortic aneurysms, and we recommend screening patients with fusiform intracranial aneurysms for aortic aneurysms.
- Saccular intracranial aneurysm patients share risk factors with aortic aneurysm patients, but routine screening does not seem to be indicated because of low co-occurrence rate.

men. 8-10 The incidence varies from 6 to 10.4 per 100 000 for thoracic aortic aneurysms (TAA) to 0.89 to 176.08 per 100 000 for abdominal aortic aneurysms (AAA), the prevalence increasing steeply towards the most aged groups. Familial background precedes 20% of thoracic aortic aneurysms (TAA) and positive family history is a known risk factor for abdominal aortic aneurysm (AAA). 8-10

Increased prevalence of IA in AA patients, as well as AA in IA patients, has been suggested in several studies. $^{12-19}$ However, these cohorts did not contain patients and controls from defined catchment populations and, in most cases, flAs were not distinguished from slAs.

We analyzed, in retrospect, the characteristics of the 125 flA patients with 134 flAs as compared with the 4253 slA patients with 6097 slAs from a defined population in the Kuopio Intracranial Aneurysm Patient and Family Database, which comprises all the IA patients of Eastern Finland diagnosed or treated in Kuopio University Hospital (KUH) since 1980. We studied the prevalence of AA in these flA patients and slA patients and in their first-degree relatives, and in the patients' matched population controls, using the Finnish national clinical registries. Hypothesizing that flAs and AAs, both fusiform in shape, share genomic risk factors, we systematically reviewed the literature for family trees containing IA and AA patients and for patient cohorts with patients with both IA and AA. Finally, we

sequenced 33 flA patients with an aortic gene panel of 37 AA-related genes.

Methods

Data that support the findings of this study are available to qualified researchers who meet the criteria provided by the board of the Kuopio Intracranial Aneurysm Patient and Family Database. The Finnish National Institute for Health and Welfare provides data for all researchers who meet the criteria set by the institute.

KUH Catchment Population

KUH Neurosurgery has been a sole provider of full-time acute and elective neurosurgical services for the KUH catchment population in Eastern Finland since 1977. Between 1980 and 2015, the geographic area of the KUH hospital districts did not change, the population decreased from 882 671 to 815 021, the median age increased from 37 to 42 years in men and from 40 to 45 years in women whereas the proportion of men has remained at 49%. ^{21,22}

Kuopio IA Patient and Family Database

The database includes all cases of unruptured and ruptured IA patients admitted to KUH since 1980 for angiography and treatment, unless moribund or very aged. The database has been prospective since 1990. The database is administrated by a full-time nurse coordinator who interviews all new patients and codes this information, including family history, and the clinical data from hospital periods and follow-up visits into an extensive list of variables. The criterion for familial IA disease was at least 2 affected first-degree relatives. The genealogy of the verified IA patients has been extensively mapped through parish records up to the 17th century. The phenotype, concomitant diseases, genetics, and outcome of Eastern Finnish sIA patients have been analyzed in several studies.

Study Population

The Kuopio IA Database includes only patients with intracranial aneurysm(s) in 4-vessel angiography (digital subtraction angiography, computed tomography angiography, or magnetic resonance angiography), initially verified by both a neurovascular neurosurgeon and a neuroradiologist. The 125 flA patients were agreed to carry a fusiform or dolichoectatic aneurysm in a re-review by a neurosurgeon (B.R.J.) and an interventional neuroradiologist (O.T.) (Figure 1). Intracranial artery dissections were excluded. The size of the flAs was

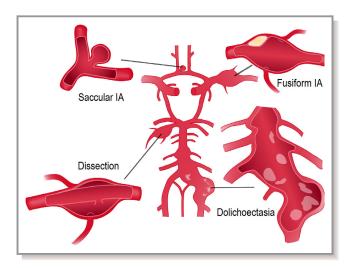


Figure 1. The major intracranial extracerebral arteries with the circle of Willis (middle). The saccular intracranial aneurysm (left upper) forms at the branching sites of the arteries, whereas the uncommon fusiform intracranial aneurysm (right upper) mainly involves the arterial trunks (see Figure 5). The dolichoectasia (right lower) usually involves the vertebrobasilar trunk. The acute dissection with false lumen (left lower), more frequent in the cervical arteries, is also shown. Note the atheromatous plaque of the fusiform intracranial aneurysm wall and the calcification of the dolichoectatic artery. IA indicates intracranial aneurysm.

defined as the greatest length or largest perpendicular diameter, whichever larger.

A total of 450 and 17 825 first-degree relatives were identified to the 125 flA patients and the 4253 slA patients, respectively, using the personal identity codes and the Finnish Population Register Centre (Figure 2). 22,25 A total of 340 and 12 669 matched population controls were assigned to the 125 flA patients and the 4253 slA patients, matched by age, sex, year, and municipality at the IA diagnosis. The index date for the matching was the date of the admission for IA or aneurysmal subarachnoid haemorrhage, with all controls alive at that point. Although we aimed at a 3:1 ratio (controls:patients), this ratio was not achieved in 35 flA and 90 slA patients.

Data Fusion From National Registries With Personal Identification Codes

The drugs prescribed for the patients, the family members and the controls, purchased from apothecaries, as well as the hospital diagnoses and the causes of death were imported from the Finnish national registries. The Finnish electronic hospital diagnosis registry (Care Register for Health Care HILMO, managed by the Finnish Institution for Health and Welfare), which was cross-linked with the Kuopio IA Database, covers all secondary and tertiary centers in Finland and includes all medical specialties. The hospital diagnoses from 1969 to 2014 and medication reimbursement statistics from 1995 to 2014 were available.

The prevalence of hypertension and atherosclerotic disease were investigated with both the hospital discharge register and the national medicine reimbursement statistics. Hypertension was defined as either: (1) diagnosis of hypertension or hypertensive disease in the diagnosis registry; or (2) use of prescribed antihypertensive medication (ATC codes C02, C03A, C04, C08 and C09). The general limit for diagnosis of hypertension has been $\geq 140/\geq 90$ mm Hg during the time period covered by the study. Atherosclerotic disease and hyperlipidemia were defined as either: (1) diagnosis of atherosclerotic disease (peripheral atherosclerosis, atherosclerotic heart disease, atherosclerotic disease of cerebral arteries); or (2) use of prescribed lipid-lowering drugs (ATC code C10).

Diagnosed Aortic Aneurysms in the Study Population

The diagnosed aortic aneurysms in the 4378 IA patients and their 17 278 first-degree relatives and 13 009 population controls were obtained from the Finnish national hospital discharge register (Care Register for Health Care). The *International Classification of Diseases, Ninth Revision (ICD-9)* and *ICD-10* codes for aortic dissections and aneurysm (441.0–441.9 and I71.00–I71.9.) were used. Furthermore, the chest radiographs, as well as thoracic and abdominal computed tomography (CT) or ultrasound imaging available for 80 (64%) flA patients were re-reviewed by a neurosurgeon (B.R.J.) and an interventional radiologist (H.M.) for presence of aortic aneurysms. The patient records of the IA patients with AA or dissection were obtained whenever possible to ascertain the diagnosis and the type of AA.

Genotyping With Aortic Aneurysm Gene Panel

Of the 41 flA patients alive, 33 were available for genotyping with informed and signed consent. Four of these 33 patients were diagnosed with AA. First-degree relatives of the patients were not available for screening. Targeted sequencing was performed using OS-seq technology described earlier^{26,27} using a commercial Aorta Panel consisting of 37 AA genes: ABCC6, ACTA2, ADAMTS2, CBS, COL1A1, COL1A2, COL2A1, COL3A1, COL5A1, COL5A2, COL9A1, COL9A2, COL11A1, COL18A1, EFEMP2, ELN, ENPP1, FBLN5, FBN1, FBN2, FKBP14, FLNA, GATA5, MFAP5, MYH11, NOTCH1, PLOD1, SKI, SLC2A10, SLC39A13, SMAD3, TGFB2, TGFB3, TGFBR1, TGFBR2, TNXB, ZNF469 (www.blueprintgenetics.com). All protein coding exons and exon-intron boundaries (± 15 bps) are covered in each targeted gene and the panel has high sensitivity to detect single nucleotide variants, small insertions and deletions up to 50 bps as well as exon level copynumber variations. Variant classification followed the 2015 ACMG (American College of Medical Genetics and Genomics) guideline.²⁸

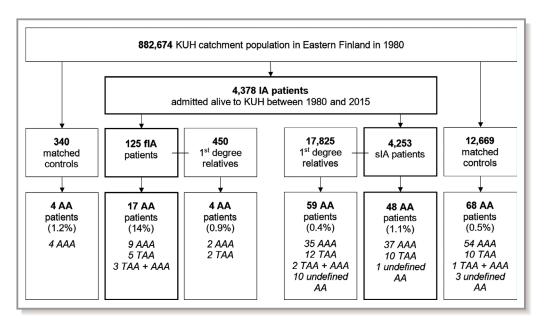


Figure 2. Flowchart of the study population: 4378 patients with intracranial aneurysm (IA) disease admitted to the Kuopio University Hospital with an unruptured IA or first aneurysmal subarachnoid hemorrhage from the Eastern Finnish catchment population from 1980 to 2015. A total of 450 and 17 825 first-degree relatives were identified to the 125 patients with fusiform IA and the 4253 patients with saccular IA, respectively, using the personal identity codes and the Finnish Population Register Centre. A total of 340 and 12 669 matched population controls were assigned to the 125 fusiform IA patients and the 4253 saccular IA patients, matched by age, sex, year, and municipality at the IA diagnosis. In 35 fusiform IA patients and 90 saccular IA patients, the 3:1 ratio was not achieved. AA indicates aortic aneurysm; AAA, abdominal aortic aneurysm; IA, intracranial aneurysm; KUH, Kuopio University Hospital; TAA, thoracic aortic aneurysm; undefined AA, AA of undefined site.

Shortly, pathogenicity of the identified variants were assessed by considering the predicted consequence, the biochemical properties of the codon change, the degree of evolutionary conservation as well as a number of reference population databases and mutation databases such as, but not limited, to the 1000 Genomes Project, gnomAD, ClinVar and the Human Gene Mutation Database. For missense variants, in silico variant prediction tools such as SIFT, PolyPhen, MutationTaster were used to assist with variant classification.

Statistical Analysis

Categorical variables were compared between groups with χ^2 and Fisher exact tests as appropriate. Continuous variables were tested with the Student t-test or Mann-Whitney U test as appropriate. As aortic aneurysms and death might share risks factors, competing risks multivariate Cox regression was performed to determine risk factors for the diagnosis of aortic aneurysms among patients with IA. The variables used in the model were sex, hypertension, any diagnosed atherosclerotic disease or prescribed lipid-lowering drug, type 2 diabetes mellitus, IA rupture status, and IA morphology. Schoenfeld residuals were used to confirm the model's assumption of proportionality. The

results are expressed as subdistribution hazard ratios (SHRs) with 95% Cls. Two-tailed P<0.05 was considered significant.

Analyses were performed using IBM SPSS Statistics, version 22.0 (IBM SPSS, Armonk, NY) and the R environment for statistical computing, including cmprsk library.

Literature Review

Two literature searches were performed in September 2018 for English articles published between 1995 and 2016, a period of non-invasive IA imaging by CT and magnetic resonance (MR). Firstly, all cohorts with concurrent IA and AA were searched with the terms: (aneurysm OR dolichoecta*) AND (aorta OR aortic OR abdominal OR thoracic) AND (intracranial OR cerebral OR cns OR brain OR subarachnoid*) AND (prevalen* OR inciden* OR epidemiology OR population). This gave 3969 hits). All abstracts were reviewed, and the case reports and duplicates were excluded. The remaining articles were searched for cohorts that reported the angiographically identified IAs in the patients with diagnosed AA and cohorts that reported diagnosed AAs in the patients with IA. The final 8 cohorts presented the proportion of IAs among the AA patients or AAs among the IA patients (Figure 3, Table 1). 12-19

Secondly, all illustrated IA family trees with co-occurrence of AA and AA family trees with co-occurrence of IA were searched with the terms: (intracranial OR cerebral OR cns OR brain OR subarachnoid*) AND (aortic OR aorta) AND aneurysm* AND (gene OR genetic* OR genome OR family OR famili* OR kindred OR generation OR hereditary OR heritable OR inherited OR phenotyp* OR mutation* OR variant* OR polymorphism OR linkage OR chromosomal). This gave 314 articles. Some family trees seemed to be duplicates but were not separable by the available data, and may consequently be counted twice, and some did not differentiate saccular IAs from other types of IAs. The final 8 cohorts presented 27 family trees (Figure 4, Table 2).^{29–37}

Ethical Approvals

The Kuopio IA Patient and Family Database has been approved by the Research Ethics Committee the KUH, the Finnish Ministry of Social Affairs and Health, and the National Institute

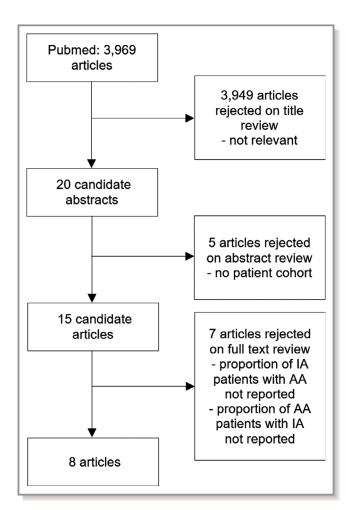


Figure 3. Literature search for patient cohorts with concurrent intracranial aneurysms and aortic aneurysms. AA indicates aortic aneurysm; IA, intracranial aneurysm.

for Health and Welfare. Written informed consent was obtained from all the IA patients registered to the database.

Results

Our Study Population

There were 125 patients with 134 flAs and 4253 patients with 6097 slAs (Figures 2 and 5, Tables 3 and 4). Of the 125 flA patients, 22 (18%) carried also slAs. For the 125 flA patients, there were 450 first-degree relatives and 340 matched population controls, and for the 4253 slA patients 17 825 first-degree relatives and 12 669 matched controls (Figure 2, Table 3).

Comparison of fIA Patients and sIA Patients

Aneurysmal subarachnoid haemorrhage had occurred in 26% of the 125 flA patients and 69% of the 4253 slA patients (P<0.0005) (Table 3). Other clinically notable differences between the fIA and sIA patients were: age at diagnosis of IA (median 64 and 54 years, P<0.0005); male predominance (52% versus 45%, P=0.11); multiple flAs versus multiple slAs (6% versus 28%, P<0.0005); hypertension (91% versus 75%, P=0.001); and type 2 diabetes mellitus (22% versus 13%, P=0.012). However, autosomal dominant polycystic kidney disease occurred equally often (2% versus 1%, P=0.663). None of the fIA patients with autosomal dominant polycystic kidney disease had concurrent sIA. There were both fIA and sIA patients who belonged to an sIA family (6% versus 14%, P=0.007). No patients with a diagnosis of Marfan syndrome, Loeys-Dietz syndrome, or Ehlers-Danlos syndrome were identified in either patient group.

Comparison of flAs and slAs

In a comparison of the 134 flAs and 6097 slAs, including the sIAs of the 22 fIA patients with both fIA and sIA, the largest sizes of the 102 unruptured flAs were significantly larger than those of the 3145 unruptured sIAs (17 mm versus 4 mm, P<0.0005). The 32 ruptured flAs were significantly larger than the ruptured 2952 slAs as well (10 mm versus 7 mm, P=0.01). Among the unruptured IAs, the 102 flAs preferred the arterial trunks of vertebrobasilar tree (59%) and the trunk of the internal carotid artery (ICA) (23%) whereas the 3145 slAs were most frequent on the middle cerebral artery bifurcation (45%) and the intracranial ICA (25%) (Figure 5, Table 4). Among the ruptured IAs, the 32 fIAs preferred the vertebrobasilar artery trunks (50%) and the ICA trunk (25%) while the 2925 slAs were most frequent on the middle cerebral artery bifurcation (33%) and the anterior communicating artery (31%) (Figure 5, Table 4).

Table 1. Published Relevant Cohorts of Patients With Intracranial Aneurysms and Aortic Aneurysms

Reference	Number of Patients	Imaging Modality	Proportion of Patients With Both IA and AA		
IA patients with AAs					
Flemming et al ¹² 2005	159 patients with flA or dolichoectasia*	Not reported	AAA 29/159 (18%) sIA 16/159 (10%)		
Miyazawa et al ¹³ 2007	181 IA patients [†]	Abdominal ultrasound	AAA 13/181 (7%)		
Goyal et al ¹⁵ 2015	317 IA patients [†]	TTE or TEE	TAA 15/317 (5%)		
Brinjikji et al ¹⁸ 2016	139 patients with vertebrobasilar dolichoectasia [†] 25 with diffuse intracranial dolichoectasia [†]	Not reported	AAA 12/139 (14%) AAA 10/25 (63%)		
AA patients with IAs		•			
Kuzmik et al ¹⁴ 2010	212 TAA patients 52 retrospective and 160 prospective	MRA or CTA	IA [†] 9/52 (17%) and IA [†] 10/160 (6%)		
Lee et al ¹⁹ 2017	133 AA patients 25 aortic dissection patients	MRA or CTA	sIA 25 and fIA 2/133 (20%) sIA 8/25 (30%)		
Shin et al ¹⁶ 2015	660 AA patients	MRA or CTA	IA 71/660 (12%) flAs excluded		
Rouchaud et al ¹⁷ 2016	1081 AA patients	CTA or MRA or DSA	sIA 128/1081 (12%) <i>20</i> dissections or fIAs excluded		

CTA indicates computed tomography angiography; flA, fusiform intracranial aneurysm; DSA, digital subtraction angiography; MRA, magnetic resonance angiography; slA, saccular intracranial aneurysm; TEE, transoesophageal echocardiography; TTE, transthoracic echocardiography.

*Partially same patient cohort.

In the 125 flA patients, 29 flAs (23%, median size 32 mm) caused symptoms of brain or cranial nerve compression in contrast to only 54 (1.3%, median size 28 mm) slAs in the 4253 slA patients (Table 3). Of all 67 basilar trunk flAs, 18 (27%,

Pubmed: 314 articles 296 articles rejected on preliminary review - no family tree 18 candidate articles 9 articles rejected on full text review - no concurrent IA and AA patients - identification of IA patients not possible 9 articles

Figure 4. Literature search for family trees with both intracranial aneurysm and aortic aneurysm patients. AA indicates aortic aneurysm; IA, intracranial aneurysm.

median size 38 mm) caused brainstem compression and 3 obstructive hydrocephalus. In the anterior circulation, 2 symptomatic fIAs were on the middle cerebral artery and 6 on the ICA.

AA in 125 flA Patients

There were 17 (14%) flA patients with a diagnosed AA (4 TAA, 12 AAA) or suspected AA (4 widened thoracic aortas in chest radiographs [2 patients] or thoracic echocardiographs [2 patients] indicative of TAAs) and 1 patient with an aortic dissection (Figure 2, Table 3). Of the 17 AAs, 3 (17%) had ruptured. Of the 17 patients, 12 (71%) were men, 16 (94%) had diagnosed hypertension, and 9 (53%) had diagnosed atherosclerotic disease. The median ages at the diagnosis of flA and AA were 64 and 63 years (Figure 6A); in 8 patients, the AA diagnosis preceded the flA diagnosis. The 125 flA patients had significantly more often AA than their 450 first-degree relatives (14% versus 0.9%, *P*<0.0005) and their 340 matched population controls (14% versus 1.2%, *P*<0.0005) (Table 3).

AA in 4253 sIA Patients

There were 48 (1.1%) sIA patients with diagnosed AA (37 AAA, 10 TAA, 1 undefined AA) and 2 patients with diagnosed aortic dissection (Figure 2, Table 3). Of the 48 AAs, 8 (17%) had ruptured. Of the 48 patients, 37 (79%) were men, 46 (96%) had diagnosed hypertension, and 22 (48%) had diagnosed atherosclerotic disease. The median ages at the diagnosis of

[†]Fusiform and saccular intracranial aneurysm were not distinguished.

Table 2. Published Family Trees Including Patients With Both Intracranial and Aortic Aneurysms

	Family Members	IA Patients With No AA	IA Patients With AA	AA Patients With No IA	Implicated Genes
Schievink et al ²⁹ 1997	19	2 IA*	1 IA* + TAA	0	
Cannon-Albright et al ³⁰ 2003	69	7 IA*	0	4 AA*	
Nahed et al ³¹ 2005	19	2 IA*	1 IA* + TAA	0	
Kim et al ³² 2005	28	4 IA*	0	4 AA*	
	16	3 IA*	0	3 AA*	
	24	4 IA*	0	2 AA*	
	22	3 IA*	1 IA* + AA*	1 AA*	
	25	5 IA*	0	1 AA*	
	16	1 IA*	0	10 AA*	
Regalado et al ^{33, 34}	19	1 IA*	1 IA* + TAA	6 TAA	
	32	1 IA*	0	7 TAA	
	8	2 sIA	0	2 TAA, 3 AAA	
	17	1 IA*	1 IA* + TAA	4 TAA, 1 AAA	
	14	4 IA*	0	3 TAA	
	10	1 IA*	1 IA* + TAA	1 TAA, 1 AAA	
	6	1IA*	0	2 TAA	
	6	1 IA*	1 IA* + TAA	2 TAA	
	19	2 sIA	0	6 TAA	
	17	2 IA*	0	2 TAA	
	13	2 sIA	0	2 TAA	
	7	1 IA*	0	2 TAA, 1 AAA	
	11	1 IA*	0	4 TAA	
	24	3 sIA	0	6 TAA	
	17	1 IA*	1 IA* + TAA	4 TAA, 1 AAA	SMAD3
Luukkonen et al ³⁵ 2012	20	1 IA	0	3 AA*	NTM
Bertoli-Avella et al ³⁶ 2015	23	0 IA	1 IA* + AA*	2 AA*	TGFB3
Mazzella et al ³⁷ 2017	22	1 flA	0	5 TAA	TGFB2
Total	523	57 IA	9 IA + AA	95 AA	

AA indicates aortic aneurysm; AAA, abdominal aortic aneurysm; flA, fusiform intracranial aneursym; slA, saccular intracranial aneurysm; TAA, thoracic aortic aneurysm *The distinction between flA vs slA or TAA vs AAA not reported.

sIA and AA were 57 years and 66 years (P=0.006); in 9 patients, the AA diagnosis preceded the sIA diagnosis (Figure 6B). The 4253 sIA patients had significantly more often AA than their 17 825 first-degree relatives (1.1% versus 0.3% P<0.0005) and their 12 669 matched population controls (1.1% versus 0.5%, P<0.0005). Diagnosed or suspected AAs were significantly more prevalent in the 125 fIA patients than in the 4253 sIA patients (14% versus 1.1%, P<0.0005).

Multivariate Analysis of Independent Risk Factors for AA in fIA and sIA Patients

In the multivariable regression analysis, fusiform IA morphology (SHR 7.6, 95% CI 3.9-14.9, P<0.0005), diagnosed

atherosclerotic disease or use of lipid-lowering medication (SHR 4.4, 95% CI 2.3–8.2, P<0.0005), and male sex (SHR 3.1, 95% CI 1.5–6.4, P=0.002) were independent risk factors for AA. However, hypertension (SHR 6.4, 95% CI 0.9–47.2, P=0.07), type 2 diabetes mellitus (SHR 1.4, 95% CI 0.7–2.7, P=0.36), and history of IA rupture (SHR 0.4, 95% CI 0.2–1.8, P=0.26) were not.

Sequencing of fIA Patients

Thirty-three fIA patients were sequenced using an AA gene panel of 37 genes. Six fIA patients each carried a different rare variant (minor allele frequency <0.01): splice-donor variant *COL5A2* c.322+1G>C (NM_000393.3), four missense variants each in one patient: *SKI* c.1582G>C, p.(Ala528Pro)

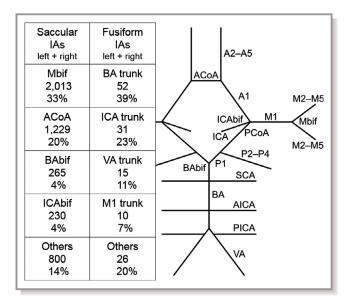


Figure 5. The most common locations of the 134 fusiform intracranial aneurysms and the 6097 saccular intracranial aneurysms in descending order of frequency (the left side and the right side combined). The schematic illustration of the major intracranial arteries with the circle of Willis shows the major arterial bifurcations and segments (compare with Figure 1.). Anterior circulation: the internal carotid artery up to the internal carotid artery bifurcation; A1 the proximal segment of the anterior cerebral artery; the anterior communicating artery; A2 to A5, the distal segments of the anterior cerebral artery; M1, the proximal segment of the middle cerebral artery; the bifurcation of the middle cerebral artery; M2 to M5, the distal segments of the middle cerebral artery. Posterior circulation: the vertebral artery; the posterior inferior cerebellar artery; the basilar artery; the anterior inferior cerebellar artery; the superior cerebellar artery; the basilar tip bifurcation; P1, the proximal segment of the posterior cerebral artery; P2-P4, the distal segments of the posterior cerebral artery; the posterior communicating artery. A1 indicates the proximal segment of the anterior cerebral artery; A2 to A5, the distal segments of the anterior cerebral artery; ACoA, the anterior communicating artery; AICA, the anterior inferior cerebellar artery; BA, the basilar artery; BAbif, the basilar tip bifurcation; IA, intracranial aneurysm; M1, the proximal segment of the middle cerebral artery; Mbif, the bifurcation of the middle cerebral artery; M2 to M5, the distal segments of the middle cerebral artery; P1, the proximal segment of the posterior cerebral artery; P2-P4, the distal segments of the posterior cerebral artery; PCoA, the posterior communicating artery; PICA, the posterior inferior cerebellar artery; SCA, the superior cerebellar artery; VA, the vertebral artery.

(NM_003036.3); FBN1 c.3571G>A, p.(Asp1191Asn) (NM_000138.4); MYH11 c.4903T>A, p.Ser1635Thr) (NM_022844. 2); COL11A1 c.278G>T, p.Gly93Val) (NM_080629.2) and a synonymous variant COL3A1 c.1038C>T, p.(Ser346=) (NM_000090.3) (Table 5). The splice-donor variant in COL5A2 was classified as likely pathogenic. There are 21 individuals heterozygous for the COL5A2 c.322+1G>C variant in the Genome Aggregation Database (gnomAD, n>120 000 exomes

and >15 000 genomes). This variant affects consensus splice site, and all 5 components of Alamut splicing software (SSF, MaxEnt, NNSPLICE, GeneSplicer, HSF) predict that this variant breaks the wild-type donor site but does not generate a near-by in-frame cryptic splice site. It has been detected in clinical testing in one affected individual by GeneDx (ClinVar 213136), and 2 additional affected patients with either ascending aortic dissection or multiple aneurysms including IA have been identified by Blueprint Genetics (Juha Koskenvuo, MD PhD, unpublished data, 2019). One of the tested healthy family members did not carry the variant, which further supports pathogenicity of the variant. However, variant's allele frequency is relatively high, at least suggesting that it may not be fully penetrant. MYH11, COL11A1, and FBN1 variants were classified as of uncertain significance whereas SKI and COL3A1 were deemed to be likely benign.

Literature Review

We identified 27 published family trees with co-occurrence of IAs and AAs (Table 1). In 21 family trees the morphology of the IAs, whether fusiform or saccular, was not reported. There were 9 IA patients with AA and 57 IA patients with no AA. There were 95 AA (58 TAA, 7 AAA, 30 unspecified AA) patients without IA.

Discussion

Comparative Characteristics of fIA and sIA Diseases

To our knowledge, this is the first population-based study comparing the characteristics of the fIA disease (125 patients with 134 fIAs) and the sIA disease (4253 with 6097 sIAs). Fusiform IAs are rare, comprising 2.2% of all IAs in our study, and sizeable published series are few. Like aortic aneurysms, fIAs are dilatations of arterial segments, while sIAs are saccular pouches usually formed at the artery forks (Figure 1). The site distribution of the sIAs and the fIAs differs significantly 38,39: in our study, 56.7% of the fIAs but only 7.6% of the sIAs were located on the vertebrobasilar artery complex (Figure 5, Table 4).

The fIA patients were significantly older than the sIA patients, less likely to present with IA rupture, and more likely diagnosed with hypertension, type 2 diabetes mellitus, hyperlipidemia, or atherosclerotic disease (Table 3). In the fIA patients of this cohort, hypertension and type 2 diabetes mellitus were even more frequent (91% and 22%) than in the previously reported fIA cohorts. 12,40,41 Autosomal dominant polycystic kidney disease, an unusual monogenic risk factor for the sIA disease, 24 previously reported to occur in some 2% of patients with vertebrobasilar fIAs, 12 occurred equally in the fIA and the sIA patients (Table 3). Multiple fIAs on arterial trunks were less frequent than multiple sIAs on bifurcations (6% versus 28%) (Table 3). In previous cohorts of patients with

Table 3. Characteristics of the 4378 Patients With Intracranial Aneurysms

Total																						
P Value—Total flA Vs Total sIA*	<0.0005	<0.0005		<0.0005			0.106	0.007				<0.0005		0.273						0.08	<0.0005	
12 669 Matched Controls to the sIA Patients	0 fIA 58 sIA (33 aSAH)	56 (52 to 73) at sIA		:			7029 (55%)	:	:	:	:	54 (0.4%)	12 (0.1%)	10 (0.1%)	1 (0.0%)	:	1 (0.0%)	:	3 (0.0%)	18 (0.1%)	86 (0.7%)	13 (0.1%)
17 825 First Degree Relatives to the SIA Patients	0 flA 224 slA (76 aSAH)	47 (42 to 54) at sIA		:			8706 (49%)	:	:		::	35 (0.2%)	11 (0.1%)	12 (0.1%)	4 (0.0%)		2 (0.0%)		10 (0.1%)	5 (0.0%)	64 (0.4%)	15 (0.1%)
flAs	Total sIA n=4253	54 (45 to 63)		:			2352 (55%)	573 (14%)	:	÷	1191 (28%)	37 (1%)	8 (0.2%)	10 (0.2%)	:	:	÷		1 (0.0%)	2 (0.0%)	20 (1%)	8 (0.0%)
4253 Saccular IA Patients With No flAs	Ruptured sIA n=2923 (69%)	52 (43 to 62)		:			1598 (55%)	314 (11%)	:		811 (28%)	13 (0.4%)	6 (0.2%)	2 (0.1%)		:	:		:	2 (0.1%)	16 (0.1%)	5 (0.0%)
4253 Saccular IA	Unruptured sIA n=1330 (31%)	57 (48 to 66)		1164 (88%)	54 (4%)	112 (8%)	754 (57%)	259 (20%)	:	:	380 (29%)	24 (2%)	2 (0.2%)	8 (0.6%)	:	:	:		:	:	27 (2%)	2 (0.0%)
340 Matched Controls to the flA Patients	0 flA 1 slA	75 at sIA		:			160 (47%)	:	:	:	:	4 (0.6%)	:	:		:	:		:	:	4 (1%)	:
450 First Degree Relatives to the fIA Patients	0 flA 1 slA	47 at sIA					222 (49%)	:	:	:	:	2 (0.4%)	:	2 (0.4%)		:	:		.:	2 (0.4%)	6 (1%)	
s 22 flA Patients	Total flA n=125	64 (55 to 72)		:			60 (48%)	7 (5.7%)	22 sIA (18%)	8 (6%)	(%2) 6	6 (2%)	2 (2%)	1 (1%)	:	4 (4%)	3 (2%)	1 (1%)	:	1 (1%)	18 (14%)	3 (2%)
125 Fusiform IA Patients Including 22 flA Patients With Concomitant slAs	Ruptured flA n=32 (26%)	59 (48 to 73)		:			21 (66%)	2 (6%)	:	1 (3%)	(%0) 0	:		÷		1 (3%)	:		:	:	1 (3%)	:
125 Fusiform IA Patien With Concomitant slAs	Unnuptured flA n=93 (64%)	66 (57 to 72)		64 (69%)	29 (31%)	:	39 (42%)	5 (5%)	:	7 (8%)	9 (10%)	9 (10%)	2 (2%)	1 (1%)	:	3 (4%)	3 (3%)	1 (1%)	:	1 (1%)	17 (18%)	3 (3%)
	Variables for 4378 IA Patients From 1980 to 2015	Median age at IA diagnosis, y (25 and 75 percentiles)	Presentation	Incidental	Symptomatic	sIA family screening	Women, n (%)	slA family, n (%)	Concomitant sIA, n (%)	Multiple flAs (≥2), n (%)	Multiple slAs (≥2), n (%)	AAA total	Ruptured	Confirmed TAA total	Ruptured	Suspected TAA	AAA + TAA total	Ruptured	Unspecified AA	Dissection	AA or dissection	Ruptured

Continued

Table 3. Continued

Total										
P Value—Total fIA Vs Total sIA*	<0.0005	0.61		0.663				0.001	0.012	<0.0005
12 669 Matched Controls to the sIA Patients	0 fIA 58 sIA (33 aSAH)	70 (63 to 76)	n=10 391	7 (0.1%)	1 (0.0%)	0	0	7989 (77%)	1742 (17%)	5132 (49%)
17 825 First Degree Relatives to the SIA Patients	0 flA 224 slA (76 aSAH)	66 (58 to 72)	n=15 652	33 (0.2%)	3 (0.0%)	0	0	8780 (56%)	2270 (14%)	5764 (37%)
flAs	Total sIA n=4253	65 (59 to 73)	n=2722	51 (1%)	0	0	0	2044 (75%)	361 (13%)	345 (13%)
4253 Saccular IA Patients With No fIAs	Ruptured sIA n=2923 (69%)	÷	n=1580	32 (1%)	0	0	0	1123 (71%)	187 (12%)	158 (10%)
4253 Saccular IA	Unruptured sIA n=1330 (31%)	÷	n=1142	19 (1%)	0	0	0	921 (81%)	174 (15%)	187 (16%)
340 Matched Controls to the flA Patients	0 flA 1 slA	76 (73 to 81)	n=251	0	0	0	0	245 (97%)	61 (24%)	172 (68%)
450 First Degree Relatives to the fIA Patients	0 flA 1 slA	50 (46 to 54)	n=405	0	0	0	0	146 (45%)	34 (8%)	98 (24%)
g 22 flA Patients	Total fIA n=125	63 (58 to 73)	П=89	2 (2%)	0	0	0	81 (91%)	20 (22%)	37 (30%)
125 Fusiform IA Patients Including 22 flA Patients Mith Concomitant slAs	Ruptured flA n=32 (26%)	:	п=20	1 (3%)	0	0	0	18 (90%)	5 (25%)	3 (9%)
125 Fusiform IA Patien With Concomitant slAs	Unruptured flA n=93 (64%)	:	69=u	1 (1%)	0	0	0	63 (91%)	15 (22%)	23 (25%)
	Variables for 4378 IA Patients From 1980 to 2015	Median age at AA diagnosis, y (25 and 75 percentiles)	Comorbid diseases in IA patients from 1995 to 2014	ADPKD, n (%)	Marfan, n (%)	Loeys-Dietz, n (%)	Ehlers-Danlos, n (%)	Hypertension, n (%)	Type 2 diabetes mellitus, n (%)	Atherosclerotic disease or hyperlipidemia

AA indicates aortic aneurysm; AAA, abdominal aortic aneurysm; ADPKD, autosomal dominant polycystic kidney disease; aSAH, aneurysmal subarachnoid haemorrhage; flA, fusiform intracranial aneurysm; IAA, thoracic aortic aneurysm.

*Statistical significance calculated for the difference of the total flA and sIA groups.

Table 4. Site and Size Distribution of the 134 Fusiform and the 6097 Saccular Intracranial Aneurysms

	134 flAs			6097 sIAs	6097 sIAs					
Variables	Unruptured n (median size)*	Ruptured n (median size)*	Total n (median size)*	Unruptured n (median size)*	Ruptured n (median size)*	Total n (median size)*				
Total	102 (17 mm)	32 (10 mm)	134 (15 mm)	3145 (4 mm)	2952 (7 mm)	6097 (6 mm)				
25 and 75 percentiles	9 to 35 mm	6 to 19 mm	8 to 28 mm	3 to 7 mm	5 to 11 mm	4 to 9 mm				
A1	0	2 (15 mm)	2 (15 mm)	21 (3 mm)	14 (5 mm)	35 (3 mm)				
ACoA	1 (15 mm)	1 (7 mm)	2 (11 mm)	390 (4 mm)	909 (7 mm)	1299 (6 mm)				
A2 to A5	0	0	0	158 (3 mm)	153 (6 mm)	311 (4 mm)				
ICA to ICAbif	23 (10 mm)	8 (9 mm)	31 (10 mm)	790 (4 mm)	626 (7 mm)	1416 (6 mm)				
M1 to Mbif	9 (9 mm)	1 (30 mm)	10 (9 mm)	1424 (4 mm)	963 (9 mm)	2387 (6 mm)				
M2 to M5	4 (7 mm)	3 (23 mm)	7 (8 mm)	113 (3 mm)	25 (6 mm)	138 (3 mm)				
PCoA	0	0	0	3 (11 mm)	2 (8 mm)	5 (8 mm)				
P1	1 (9 mm)	0	1 (9 mm)	14 (3 mm)	11 (6 mm)	25 (3 mm)				
P2 to P4	4 (8 mm)	1 (7 mm)	5 (8 mm)	2 (13 mm)	8 (5 mm)	10 (5 mm)				
BA	46 (25 mm)	6 (8 mm)	52 (25 mm)	131 (7 mm)	134 (9 mm)	265 (8 mm)				
SCA	1 (8 mm)	0	1 (8 mm)	54 (3 mm)	25 (7 mm)	79 (4 mm)				
AICA	0	0	0	0	1 (8 mm)	1 (8 mm)				
PICA	6 (10 mm)	2 (9 mm)	8 (10 mm)	33 (4 mm)	67 (5 mm)	100 (5 mm)				
VA	7 (28 mm)	8 (10 mm)	15 (19 mm)	9 (12 mm)	10 (9 mm)	19 (10 mm)				
Others	0	0	0	3 (2 mm)	4 (8 mm)	7 (3 mm)				

A1 indicates the proximal segment of the anterior cerebral artery; A2-A5, the distal segments of the anterior cerebral artery; ACoA, the anterior communicating artery; AICA, the anterior inferior cerebral artery; BA, the basilar artery; ICA, the internal carotid artery; ICAbif, the ICA bifurcation; M1, the proximal segment of the middle cerebral artery; M2-M5, the distal segments of the middle cerebral artery; Mbif, the bifurcation of the middle cerebral artery; others, not classified; P1, the proximal segment of the posterior cerebral artery; P2-P4, the distal segments of the posterior cerebral artery; PCoA, the posterior communicating artery; PICA, the posterior inferior cerebellar artery; SCA, the superior cerebellar artery; VA, the vertebral artery.

vertebrobasilar dolichoectasia, the reported frequency of diffuse dolichoectatic lesions has varied considerably with a range of 16% to 45%. ^{18,40} Different morphologies make size comparison of the flAs and slAs impractical; however, unruptured and ruptured flAs were larger than unruptured and ruptured slAs (Table 4), and the pronounced size difference underlines and explains the different rates of aneurysm symptomaticity. Because of their larger size and preference for the posterior fossa, the flAs often caused symptoms by compression: brainstem compression, cranial nerve symptoms, and obstructive hydrocephalus.

Few studies on the histopathology and molecular biology of the fIA wall, resected during surgery or at autopsy, are available. In 1 study, walls of 8 fIAs displayed fragmentation of internal elastic lamina, neoangiogenesis within thickened intima, intramural haemorrhage and thrombus formation. Emiliar changes are observed not only in sIAs but also in AAAs and TAAs. Fusiform IAs have been linked to atherosclerotic risk factors, but the association of atherosclerosis with fIAs or its role as causal factor is complex and unclear.

The distinctions between and the terminology of the spectrum of conditions that manifest as different forms of non-saccular dilatations of intracranial arteries are often unclear. Dolichoectasia, elongation and dilatation of intracranial arteries (Figure 1), is most common in the vertebrobasilar complex. In neuroimaging (CT, CT angiography, MR imaging, MR angiography, digital subtraction angiography), dolichoectasia can be difficult to distinguish from fusiform IAs, and the two may represent the same disease spectrum and pathogenesis. Acute dissection is considerably less frequent in the intracranial arteries than in the cervical arteries, and is a rare cause of subarachnoid haemorrhage or brain infarction. There are data indicating that intracranial dissection may develop into a chronic fusiform aneurysm.

AA in flA and slA Patients

To our knowledge, this is also the first population-based study to investigate the occurrence of aortic aneurysms in both fIA and sIA patients (Figures 2 and 6), and in the first-degree relatives and the matched population controls for the both

^{*}Median of largest sIA diameter or median of largest fIA length or diameter.

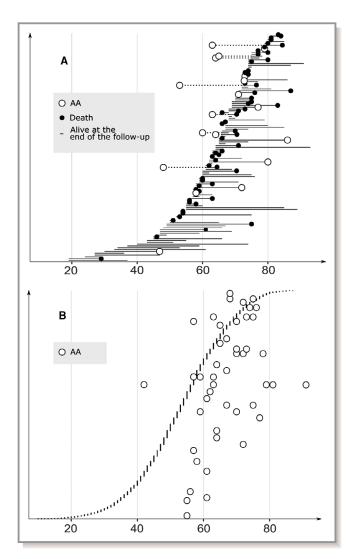


Figure 6. A, Distribution of the ages, in ascending order, of the 125 fusiform intracranial aneurysm patients (32 ruptured, 93 unruptured) at the diagnosis of the fusiform intracranial aneurysm. The ages at the diagnosis their 17 aortic aneurysms are shown (white circles). The lifelines of the fusiform intracranial aneurysm patients from either diagnosis (fusiform intracranial aneurysm or AA) until the AA diagnosis, death (without AA diagnosis, black circles) or the last follow-up (without AA diagnosis, end of lifeline) are shown. B, Distribution of the ages, in ascending order, of the 4253 saccular intracranial aneurysms patients (2923 ruptured, 1330 unruptured) at the diagnosis of their saccular intracranial aneurysms. The ages at the diagnosis of their 48 aortic aneurysms are shown (white circles). The lifelines of the saccular intracranial aneurysm patients with diagnosed AA are omitted for the sake of clarity. The median ages at the saccular intracranial aneurysm diagnosis (dashed vertical line) and AA diagnosis (continuous vertical line) are shown. AA indicates aortic aneurysm.

patient groups (Figure 2, Table 3). The 125 flA patients had significantly more often AA than their 450 first-degree relatives (14% versus 0.9%) and their 340 matched population controls (14% versus 1.2%). In a recent retrospective study of

139 patients with fIA or dolichoectasia, similar proportion of the patients (12% and 14%) had concomitant AAA (Table 1), equal to 14% in this study. In other studies, fIAs may have been excluded or not distinguished from sIAs (Table 1). The risk of AA seems to warrant the screening of aortas of all fIA patients with MR angiography or CT angiography.

The 4253 sIA patients had clearly fewer diagnosed AAs during the follow-up than the fIA patients (1.1% versus 14%) (Figures 2 and 6), but, nevertheless, more than their 17 825 first-degree relatives (1.1% versus 0.3%) and their 12 669 matched population controls (1.1% versus 0.5%) (Figure 2, Table 3). The 48 sIA patients with AA were, on average, men, diagnosed with hypertension, with a substantial portion carrying atherosclerotic diseases. Despite the older age and higher burden of comorbid diseases in the fIA patients, as compared with the sIA patients, the fusiform IA morphology was the most important risk factor for AA in a multivariate model alongside the recognized risk factors, indicating that the increased AA risk in the flA patients is not contributable to the traditional risk factors alone. No robust data on smoking were available for the multivariate model. The identified risk factors for AA in IA patients and the equivocal or protective role of diabetes mellitus against AA are well in line with previous research.43

Genetics of fIA Disease in Relation to AA Disease

We hypothesized that the rare flAs share, in addition to their fusiform shape, genomic and acquired risk factors with AAs. In our review of the literature, we identified 27 families containing both IA patients and AA patients (Table 2). Among the 523 members of these families, 57 IA patients and 95 AA (58 TAA, 7 AAA, 30 unspecified AA) patients but only 9 patients with both IA and AA were identified. In most cases, the morphology of the IAs, whether fusiform or saccular, was not reported. A previous meta-analysis of genome-wide association studies (GWAS) of IA, TAA and AAA cohorts did not find evidence for shared genetic risk of IAs, TAAs and AAAs; however, in that study, flA patients were not distinguished from the sIA patients.⁴⁸ The genetics of the sIA disease remain elusive.⁵

In the present study, none of the 125 flA patients had been diagnosed with inherited traits predisposing to AAs, including Marfan, Loeys-Dietz, or Ehlers-Danlos syndromes. We were unable to identify any flA families, but 6% of the flA patients belonged to an slA family. Interestingly, concomitant flA and slA were seen in 18% of the flA patients, suggesting susceptibility of these patients to aneurysms in general. Considerable co-occurrence of flAs and slAs has been reported previously in some studies. We sequenced 33 flA patients with a panel of 37 genes associated to AA, and found only one likely pathogenic variant (*COL5A2* c.322+1G>C),

Table 5. Rare Variants Found Among 33 Patients With Fusiform Intracranial Aneurysm in the Genetic Screening of 37 Genes Associated With Aortic Aneurysms

Gene	Effect on Genomic DNA	Effect on Protein Transcript	Allele Frequency in gnomAD	Predicted Functional Consequence	Clinical Significance
SKI	c.1582G>C	p.(Ala528Pro)	7.68e-05 Missense variant		Likely benign
FBN1	c.3571G>A	p.(Asp1191Asn)	1.08e-05	Missense variant	VUS
COL3A1	c.1038C>T	p.(Ser346=)	3.97e-05	Synonymous variant	Likely benign
MYH11	c.4903T>A	p.(Ser1635Thr)	0	Missense variant	VUS
COL11A1	c.278G>T	p.(Gly93Val)	0	Missense variant	VUS
COL5A2	c.322+1G>C	N/A	7.58e-05	Splice donor variant	Likely pathogenic

gnomAD indicates a reference database of exomic sequences; VUS, variant of unknown significance.

which would likely predispose to classic Ehlers-Danlos syndrome or TAA or thoracic aortic dissection.

The molecular pathologies of TAA and AAA are divergent. ⁴⁹ A substantial proportion of TAAs occur in association with a defined genetic syndrome or an identifiable single-gene mutation. A focused or genome-wide testing of the proband and, in the case of a positive finding, the first-degree relatives, is recommended in patients presenting with syndromic features, positive family history or absence of traditional cardiovascular risk factors. Even though hereditary risk to AAA is well-established, known causative single-gene mutations are few. ⁵⁰ Consequently, family history affects the decision threshold of screening for the first degree relatives of AAA patients. ⁹ However, genetic counselling is recommended if the AAA disease cannot be solely explained by a non-genetic cause. ¹⁰

Our Strengths

Our study has several strengths. The publicly-funded Finnish healthcare system is universal, which reduces bias to minimum in population-based disease databases. Neurosurgery of the Kuopio University Hospital (KUH) is the sole provider of acute and elective neurosurgical services for Eastern Finland, including unruptured and aneurysmal SAH patients. The Kuopio IA Patient and Family Database has incorporated clinical data for both the IA patients and their relatives and matched population controls from the national registries, using the Finnish personal identity codes. The data quality of the Finnish national health registers has been shown to be good.⁵¹

Our Limitations

However, the study has some limitations. Aortic aneurysms have not been screened in Finland either on a population level, in patients with intracranial aneurysms or in patients with stroke. Unruptured IAs, with an estimated lifetime prevalence of 2% to 3%, and AAs, with an estimated lifetime prevalence of 1% to 8%, are increasingly diagnosed as incidental findings in the imaging of the head or the body for other reasons in the elderly

population. As both sIA and fIA patients may have had shortened survival, their controls may have accrued longer follow-up with age-related increasing prevalence of AA, possibly leading to some attenuation in our finding of increased AA prevalence in IA patients. Our data contained 68 AAs in IA patients but only 67 could be accurately verified. Consequently, our available data did not allow exact anatomical classification of all the AAs. Furthermore, the specificity of chest radiograph for TAA, used in 2 flA patients as the basis diagnosis, is not high enough for definite diagnosis of TAA. The determination of the disease status of the comorbid diseases was in many cases based on medication use without knowledge of the indication of the prescription. Among the 125 flA patients, only a small subgroup was available for the genotyping. The genotyping included only genes the mutations of which are known to associate with AA.

Suggested Further Research

Fusiform IAs that involve perforators, branches and bifurcations of the cerebral arteries are difficult to treat with the present microsurgical, endovascular and bypass techniques. Since flAs and AAs share the fusiform shape and, possibly, pathogenetic mechanisms, and as flAs and AAs partially cooccur, cellular and molecular biology data from AA walls may help to develop pharmacological treatment that would stabilize the flA wall and reduce its growth. In slA patients, the risk of AA diagnosis was low in our study, and may not justify screening for AA of slA patients, but it remains an underestimate in the absence of prospective screening studies of slA patients for incidental AA.

Conclusions

In our study, without any screening for AA, $\approx 1\%$ of the sIA patients had diagnosis for AA, in contrast to 0.3% of their first-degree relatives and 0.5% of their matched population controls. These data suggest that screening for AA in sIA

patients is not generally indicated. In fIA patients the risk of AA was 14%. This warrants screening of aortas of all fIA patients with MR angiography or CT angiography. The AA gene panel with 37 previously known AA-associated genes in 33 fIA patients identified likely disease-causing variant only in 1/33 (3%) of the patients. This observation suggests that fIA and AA may share genetic background in some cases, but larger cohorts are needed to validate or dispute this association.

Sources of Funding

This study was funded by The Finnish Cultural Foundation, The North Savo Regional Fund of Finnish Cultural Foundation, The Petri Honkanen Foundation, The Päivikki and Sakari Sohlberg Foundation, The Maire Taponen Foundation, The Olvi Foundation, The Pro Humanitate Foundation, the Academy of Finland, and the Kuopio University Hospital.

Disclosures

Dr Koskenvuo is a cofounder and director of Blueprint Genetics. All other authors declare no disclosure.

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