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Cerebral Amyloid Angiopathy in Down Syndrome and Sporadic and Autosomal-Dominant Alzheimer's Disease

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Cerebral amyloid angiopathy in Down syndrome and sporadic and autosomal dominant Alzheimer's disease

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Abstract

Introduction—We aimed to investigate if cerebral amyloid angiopathy (CAA) is more frequent in genetically determined than in sporadic early onset forms of Alzheimer disease (EOAD).

Methods—Neuroimaging features of CAA, *APOE*, and cerebrospinal fluid-A β 40 levels were studied in subjects with Down syndrome (DS, n=117), autosomal dominant AD (ADAD, n=29), sporadic EOAD (n=42), and healthy controls (n=68).

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Results—CAA was present in 31%, 38% and 12% of cognitively impaired DS, symptomatic ADAD, and sporadic EOAD subjects, and in 13% and 4% of cognitively unimpaired DS individuals and healthy controls, respectively. *APOE-ε4* genotype was borderline significantly associated with CAA in sporadic EOAD ($p=0.06$), but not with DS or ADAD. There were no differences in Aβ40 levels between groups or between subjects with and without CAA.

Discussion—CAA is more frequently found in genetically determined AD than in sporadic EOAD. Cerebrospinal fluid-Aβ40 levels are not a useful biomarker for CAA in AD.

Keywords

Cerebral amyloid angiopathy; sporadic early onset Alzheimer disease; autosomal dominant Alzheimer disease; Down syndrome; neuroimaging; cerebrospinal fluid biomarkers

Introduction

Most cases of Alzheimer's disease are sporadic, and caused by complex interactions between genetic and environmental factors. In approximately 5% of cases, Alzheimer's disease can present clinically before the age of 65 (early-onset Alzheimer's disease-EOAD) [1]. These patients frequently present with non-amnesic phenotypes and faster clinical decline than older sporadic Alzheimer's disease cases [1]. In 0.1-0.5% of cases, Alzheimer's disease is transmitted with an autosomal dominant pattern of inheritance (ADAD) due to the presence of mutations in presenilin 1 (*PSEN1*), presenilin 2 (*PSEN2*), or amyloid precursor protein (*APP*) genes [2]. Down syndrome (DS) is also recognized as a form of genetically determined AD, mainly caused by the *APP* gene triplication [2]. Despite the different genetic background, the Alzheimer's disease neuropathological findings in sporadic EOAD, ADAD and DS are very similar [3,4].

Cerebral Amyloid Angiopathy (CAA) is a major cause of lobar intracerebral hemorrhage (ICH) in the elderly and is present in up to 90% of Alzheimer's disease brains at autopsy [3]. Previous neuropathological studies have suggested a more severe CAA in ADAD than in sporadic Alzheimer's disease [4]. CAA in some *APP* mutations or duplication carriers drives the clinical presentation [4], and is also consistently observed in subjects with DS [5]. The modified Boston criteria for CAA-related hemorrhage (mBCAA) have been validated to attribute *in vivo* an ICH to CAA based on several neuroimaging features, and are frequently used in clinical practice [6]. There are no previous studies systematically assessing the CAA neuroimaging features in DS and ADAD.

Aβ40 is the major form of Aβ deposited in the vessel walls in individuals with CAA. Low levels of Aβ40 and Aβ42 have been found in the CSF of subjects with sporadic CAA [7]. However, scarce and contradictory data are available about the CAA CSF biomarker profile in sporadic Alzheimer's disease patients [8–10], and no previous studies have assessed this profile in DS or ADAD. Moreover, the *APOE-ε4* genotype is a risk factor for both sporadic Alzheimer's disease and sporadic CAA [11], as it increases Aβ deposition in both the parenchyma and blood vessels [12]. However, the effect of the *APOE* genotype in Alzheimer's disease dementia within DS and ADAD is controversial, and there are no studies assessing the influence of the *APOE* genotype on CAA in ADAD or DS [13].

The differences in the CAA neuroimaging features and CSF biomarkers profile in the different forms of Alzheimer's disease are thus, not established. Our primary objective was to determine the CAA presence assessing the fulfillment of the mBCAA and the CSF A β 40 levels in three different Alzheimer's disease populations: DS, ADAD and EOAD. We hypothesized that patients with genetically determined Alzheimer's disease would have more CAA neuroimaging and biochemical features than EOAD.

Materials and methods

Study design and participants

A total of 256 subjects were recruited from 5 centers: Hospital of Sant Pau, Hospital Clínic de Barcelona, and Barcelona Down Medical Center in Barcelona, Spain; and the Sanders-Brown Center on Aging in Kentucky and the Down Syndrome Biomarker Initiative (DSBI) project in San Diego, United States of America. Four study groups were evaluated: EOAD, ADAD, DS, and healthy controls (HC).

EOAD (N=42)—Patients were recruited at the Memory Unit of Hospital de Sant Pau from the Sant Pau Initiative on Neurodegeneration (Barcelona SPIN cohort) [14]. We used the International Working Group-2 diagnostic criteria for AD with *in vivo* evidence of AD based on CSF biomarkers [2]. This group included 19 individuals with prodromal AD (p-EOAD) and 23 subjects with probable AD dementia (d-EOAD).

ADAD (N=29)—Participants were recruited from the Genetic counseling program for familial dementias (PICOGEN) at the Hospital Clínic de Barcelona [15]. Fifteen symptomatic carriers (CDR = 0.5) carrying 9 different *PSEN1* mutations (M139T, S169P, L173F, G209E, L235R, K239N, L282R, L286P, I439S) and one symptomatic carrier of the *APP*I716T mutation were included. The symptomatic carriers were further classified as prodromal ADAD (pAD-ADAD, *n*=5) and ADAD dementia (dADAD, *n*=11). Twelve pre-symptomatic mutation carriers (CDR= 0) carrying 7 different *PSEN1* mutations (M139T, S169P, L173F, R220G, K239N, L282R, I439S) and one pre-symptomatic carrier of *APP* I716T were labeled as asymptomatic ADAD. We used the IWG2 diagnostic criteria for AD [2].

DS (N=117)—Adults with DS were recruited from three centers, the Down Alzheimer Barcelona Neuroimaging Initiative (DABNI) in the Barcelona Down Medical Center [16]; the Sanders-Brown Center on Aging; and the DSBI pilot project [17]. Adapted neuropsychological batteries (detailed in the Appendix section), covering all cognitive domains classified DS subjects into “without cognitive decline” (asymptomatic DS, N=91), prodromal AD (pAD-DS, N=13), and AD dementia (dAD-DS, N=13). pAD-DS and dAD-DS were also labeled as symptomatic DS.

Healthy controls (N=68)—Participants were recruited at Hospital de Sant Pau (*n*=60) and Hospital Clínic de Barcelona (*n*=8) enrolled among patients' caregivers. They did not have cognitive complaints, scored 0 on CDR, had normal neuropsychological evaluation, and normal core Alzheimer's disease CSF biomarkers [18,19].

Procedures

Medical records were reviewed for potential confounders and effect modifiers: age, sex, and presence of arterial hypertension, neuropsychological information on disease severity (MMSE for EOAD and ADAD and the Cambridge Examination for mental Disorders of Older People with DS and Others with Intellectual Disabilities-Cognitive Scale - CAMCOG-DS scale for DS) were recorded and sent to the coordinating center (Hospital de Sant Pau) with the CSF biomarkers (A β 42, A β 40) and neuroimaging data.

The study was approved by the local Ethics Committees following the ethical standards recommended by the Helsinki Declaration. All participants and/or their caregivers gave their written informed consent.

Neuroimaging assessments

The inclusion criteria for all participants included a 1.5 or 3T MRI scan including T2*-GRE or SWI, axial fluid attenuated inversion recovery (FLAIR) and coronal T1-weighted sequences in the five centers involved GRE or SWI sequences were assessed for the presence of the main CAA neuroimaging features: localization and number of lobar microbleeds, presence of cortical superficial siderosis (cSS), and lobar ICH. We evaluated the fulfillment of mBCAA in all participants regardless of the age criterion included in the criteria set (>55 years) [6]. White matter hyperintensities (WMH) were semi-quantitatively assessed in FLAIR sequences according to the Fazekas score [20]. MTA was evaluated in coronal T1-weighted images through the Scheltens scale [21]. MTA was scored bilaterally and the highest score was considered for the analyses.

The radiological evaluation was performed by two raters (either MCI or MB; neurologists with expertise in cognitive disorders and SG; neuroradiologist) blinded to the clinical data. Inter-rater reliability was above 90% and discrepancies within ratings were solved by consensus.

CSF biomarkers and APOE genotype

The inclusion criteria for EOAD and healthy controls included CSF data, but not for ADAD and DS. Details of analysis are described elsewhere [14,18]. In short, commercially available ELISA kits were used to determine CSF-A β 40 and CSF-A β 42 levels (Millipore and Fujirebio-Europe, respectively), following the manufacturers' recommendations.

APOE genotyping was performed by PCR amplification of the exon four fragment containing the two polymorphisms (rs429358 and rs7412) that encode the three common *APOE* isoforms. The following oligonucleotides: APOE-F: 5'-ACTGGAGGAACAACACTGACCC-3' and APOE-R: 5'-CTGCCCATCTCCTCCATC-3', were used and final PCR products were purified and Sanger sequenced using BigDye terminator chemistry (Applied Biosystems). Sequences were run on an Applied Biosystems® 3130 Genetic Analyzer and resulting electropherograms were visually inspected using Sequencher (version 4.1, Gene Codes Corp.).

Statistical analysis

Statistical analyses were performed with the Statistical Package for the Social Sciences v19 software (IBM corp. <http://www-01.ibm.com/software/es/analytics/spss/>). The primary objectives of this study were to compare across groups the frequency of the mBCAA and the CSF-A β 40 levels and were analyzed with the exact Fisher and Mann-Whitney test respectively.

The secondary objectives were to assess the white matter lesions measured by the Fazekas scale and the hippocampal atrophy measured by the Scheltens scale and were analyzed with the exact Fisher test. Spearman correlation coefficients were calculated between age, clinical stage, hippocampal atrophy and white matter lesions and the different study groups. With the purpose of improving the statistical power, prodromal and demented groups in EOAD, ADAD and DS were merged when analyzed. All significance tests were two-sided with the statistical significance set at 5%.

Results

Table 1 displays the demographics, clinical features, CSF data, and *APOE* genotype of the participants. CSF data were available in 71% (N=182) of the subjects (including all healthy controls and patients with EOAD). Symptomatic ADAD and symptomatic DS subjects were younger than patients with EOAD (48.4, 54.4, 61.1 years of age respectively; $p<0.001$).

The *APOE-ε4* allele frequency was higher in EOAD than in any other group. However, no differences were observed between symptomatic or asymptomatic subjects within the ADAD or DS groups.

Table 2 shows the neuroimaging results across the different groups. The fulfillment of the mBCAA criteria was more frequent in symptomatic DS, 31% (N=8), and in symptomatic ADAD, 38% (N=6), than in EOAD 24% (N=19) ($p=0.055$ and 0.026 respectively). When present, the most frequent CAA neuroimaging features were lobar microbleeds in 91.2% (N=31), followed by cSS in 29.4% (N=10) and ICH in 8.8% (N=3). All three features were more frequent in the symptomatic than in asymptomatic subjects within all groups (Table 2).

The symptomatic ADAD and DS groups had a higher proportion of lobar microbleeds than the EOAD group ($p=0.02$, $p=0.046$ respectively). cSS and ICH were statistically associated ($p=0.004$). In those who had cSS, 20% (N=2) had also an ICH, and cSS was present in 67% (N=2) of those with ICH. Symptomatic DS had a higher proportion of subjects with cSS and lobar ICH than the EOAD group, but this difference did not reach statistical significance ($p=0.056$). The position of the mutation (pre or post codon 200) did not significantly impact the presence of lobar microbleeds in *PSEN1* carriers (37 vs 25%, $p=0.4$). One of the asymptomatic ADAD subjects included in our study had a massive lobar ICH after recruitment into this study that led to its death in a stage of moderately severe dementia.

The mean time lag between MRI and CSF sampling was 5.3 months. Symptomatic participants had lower CSF-A β 42 levels than asymptomatic subjects within all groups (Figure 1A). No significant differences were detected in CSF-A β 40 levels between the

different groups or between symptomatic and asymptomatic subjects within each group (Figure 1B).

All EOAD patients fulfilling the mBCAA were *APOE-ε4* carriers and the *APOE-ε4* allele was significantly more frequent in EOAD subjects than in the symptomatic ADAD ($p=0.015$) or symptomatic DS ($p=0.071$) subjects fulfilling the mBCAA criteria. In sporadic EOAD, there was a trend for the association between mBCAA fulfillment and *APOE-ε4* genotype ($p=0.06$).

There were no differences in CSF-Aβ40 levels between those subjects who fulfilled the mBCAA criteria and those who did not (or the presence of lobar microbleeds, cSS, or ICH) in any group.

Symptomatic subjects presented higher Fazekas scores than asymptomatic subjects in all groups: EOAD patients had higher Fazekas scores than healthy controls ($p<0.001$); symptomatic ADAD higher than asymptomatic ADAD ($p=0.022$) or HC ($p=0.05$); and symptomatic DS higher than asymptomatic DS ($p=0.011$) and healthy controls ($p<0.001$) (Figure 2 and table 2).

There was a significant positive correlation between age and Fazekas score in the whole sample ($r=0.337$, $p<0.001$), in ADAD ($r=0.407$, $p=0.031$) and in DS ($r=0.506$, $p=0.000$) groups. This correlation was also found in asymptomatic DS ($r=0.495$, $p<0.001$) and in healthy controls ($r=0.323$, $p=0.007$) (Figure 3).

Age positively correlated with Scheltens scores in the whole sample ($r=0.229$, $p<0.001$), healthy controls ($r=0.276$, $p=0.023$), ADAD ($r=0.412$, $p=0.027$), asymptomatic DS ($r=0.341$, $p=0.001$), and symptomatic DS ($r=0.431$, $p=0.023$). The Scheltens scores increased from asymptomatic to symptomatic subjects within each group. Symptomatic DS patients presented higher MTA scores than EOAD and symptomatic ADAD patients ($p<0.001$ in each comparison) and asymptomatic DS subjects higher than HC ($p<0.001$). There were no differences in Scheltens scores between EOAD and symptomatic ADAD.

Discussion

We found that DS and ADAD have a more severe CAA than EOAD as measured by the mBCAA criteria, but CAA did not impact the CSF-Aβ40 levels. The *APOE-ε4* allele might be associated with CAA in EOAD, but does not seem to have an effect in DS or ADAD.

There are previous studies assessing the prevalence of lobar microbleeds in ADAD and EOAD [22] [23,24], but, to our knowledge, none has specifically assessed and compared the mBCAA between the different AD populations. The mBCAA were more frequent in DS and ADAD, suggesting a more severe CAA, as shown in pathological studies [5,25]. The most frequent CAA neuroimaging feature was the presence of lobar microbleeds, as previously described [26]. The frequency of lobar microbleeds in ADAD (and healthy controls) was in agreement with the literature (ranging from 25% to 66%) [23,24][26], but we found a lower frequency of lobar microbleeds in EOAD (9.5%) than that reported in late onset AD (20-30%) [27]. Age might be responsible for this difference. There are no previous studies

assessing the mBCAA criteria or any of its component neuroimaging features in DS. CAA is also consistently observed in DS pathological studies [5,28], but it had been proposed that other genetic factors in DS might protect these subjects from the ICH [29]. We found 38.5% of frequency for lobar microbleeds in symptomatic DS and, more importantly, a frequency of 15.4% for ICH. While this is lower than the reported 30% prevalence for symptomatic ICH in non-trisomic APP duplication carriers, it is well above the 3%-3.8% figure for symptomatic ICH in DS reported in the same study [29]. This discrepancy might be explained because many non-fatal ICH might be unnoticed in DS with AD. Of note, both subjects with DS and ICH on the MRI did not present with ICH-related clinical symptoms.

A significant percentage of symptomatic subjects, nonetheless, did not meet the mBCAA and were free of the CAA associated neuroimaging features. This is in contrast with pathological studies, where CAA is found in up to 90% of AD brains, suggesting that the available MRI sequences only identify a subset of AD-CAA subjects [23]. The CAA neuroimaging features might thus detect only the most severe cases or, alternatively, they might select different subgroups of patients [23]. In this respect, cSS, although less frequent than lobar microbleeds, was strongly associated with lobar ICH. cSS might be a particular important marker for severe CAA in AD, as it has been suggested in sporadic CAA [7]. More work in longitudinal studies is needed to confirm the higher risk conferred by cSS for future ICH and cognitive decline.

We also assessed other neuroimaging features associated with CAA, but not included in the mBCAA criteria. Both WMH and MTA are increasingly recognized as core Alzheimer's disease features and as a manifestation of CAA [30]. We found a gradient in WMH extension in all groups [31], but we also found more extended WMH in those subjects fulfilling the mBCAA criteria. WMH also increased with age and in relation with vascular risk factors. We found this correlation also in healthy controls, even though they all had normal core Alzheimer's disease CSF biomarkers and low prevalence of HBP. However, we found a strong correlation between age and WMH in asymptomatic DS despite their younger mean age. This correlation supports the relationship between amyloid deposition and WMH. Not surprisingly, the Scheltens scores increased with symptom severity in all AD populations. Hippocampal atrophy, however, was more severe in DS, even in asymptomatic DS individuals. These results are in agreement with the notion that individuals with DS have smaller hippocampal size from birth, but also show atrophy when Alzheimer's disease develops [17].

Decreased CSF-A β 40 levels might differentiate sporadic CAA from healthy controls and Alzheimer's disease cases [7]. In our study, nevertheless, CSF-A β 40 levels did not discriminate CAA neuroimaging features in any group. This finding could be influenced by the fact that amyloid vascular burden in CAA in ADAD and DS contains not just A β 40, but also A β 42. It is difficult to sort out the contribution of vascular A β 42 deposition from parenchymal plaque deposition except with neuropathological analysis of the post-mortem brain, which was not available in this study [32,33].

To our knowledge, there are only two studies that determine CSF-A β 40 in subjects with Alzheimer's disease with and without lobar microbleeds and show conflicting findings

[9,10]. In any case, our results suggest that CSF-A β 40 levels are not a sensitive biomarker to detect CAA in the context of an Alzheimer's disease process.

The *APOE- ϵ 4* allele confers a higher risk for CAA in the general population and in Alzheimer's disease [11,12]. We also found a trend for an association between *APOE- ϵ 4* genotype and CAA in sporadic early onset Alzheimer's disease. Furthermore, all EOAD subjects with CAA neuroimaging features were *APOE- ϵ 4* carriers. We did not find this relationship in ADAD or DS. The *APOE- ϵ 4* genotype might be thus associated with CAA in EOAD, but not in DS or ADAD. In ADAD and DS, other genetic factors, such as the type or position of the causing mutation in ADAD, might be more important in predicting CAA [4].

Our findings have potential clinical implications. The mBCAA criteria have not been validated in patients <55 years of age. We consider that, at least in ADAD and DS, age should not be an essential requirement for CAA diagnosis. Our results also have substantial implications in Alzheimer's disease clinical trials given the relationship between CAA and amyloid-related neuroimaging abnormalities (ARIA). Vascular amyloid may be a pathophysiological mechanism for ARIA [34,35] and recent studies have shown that after A β immunotherapy there is an increase in CAA severity and an increase in lobar microbleeds associated with removal of plaques [36]. Trials targeting A β commonly use lobar microbleeds and *APOE* genotype to stratify subjects [37]. However, there are no available data on the relationship of CAA neuroimaging abnormalities (and *APOE*) and ARIA in the setting of amyloid-lowering therapy in ADAD and DS. Our data emphasize heterogeneity in prevalence and possibly etiology for CAA, therefore, the recommendations on the exclusions for presence of baseline ARIA-H (microbleeds or hemosiderosis) from sporadic Alzheimer's disease should be taken with caution. On the other hand, the *APOE- ϵ 4* genotype is also commonly used to stratify participants given its influence on ARIA [37], however this strategy might not be as important in ADAD or DS. Finally, our data also suggest that the CSF-A β 40 levels will not be a useful biomarker in these trials.

The higher prevalence of CAA in ADAD and DS might play a role in the conversion to clinical dementia. In sporadic Alzheimer's disease, CAA is an independent contributor to cognitive impairment and can worsen the severity of cognitive dysfunction [38]. Future longitudinal studies are needed to assess the CAA contribution to cognitive impairment in ADAD and DS.

The main strengths of the present study are the relatively large sample size of different rare populations, such as ADAD and DS, as well as the confirmation of the clinical diagnosis with genetics or CSF biomarkers. The study has some limitations. The use of two different imaging techniques is an important limitation when estimating the real prevalence of lobar microbleeds. SWI has a higher sensitivity for hemosiderin, detecting up to 50% more lobar microbleeds than conventional T2*GRE [39]. However, in our study the ADAD group was exclusively investigated using T2*GRE leading to a possible underestimation of the CAA neuroimaging features in these subjects. Finally, most of the EOAD patients were at a stage of mild dementia and we lack neuropathological data.

In conclusion, the CAA-associated neuroimaging features are more frequent in adults with DS and in patients with ADAD than in those with EOAD suggesting a more severe CAA pathology. Our study also shows that the CSF-A β 40 levels are not a reliable biomarker for CAA and that the risk factors for CAA (such as the *APOE- ϵ 4* genotype) might be different in EOAD and genetically determined Alzheimer's disease. These findings should be taken into account in the design of clinical trials with anti-amyloid therapies in people with ADAD or DS.

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Appendix

Neuropsychological batteries assessed in the Down syndrome group.

Each participant received an annual physical and neurological examination. Diagnostic was established after a clinical consensus performed by the clinical neurologist and the neuropsychologist.

- Neuropsychological battery from the Down Alzheimer Barcelona Neuroimaging Initiative (DABNI), Spain.
 - Kauffman Brief Intelligence Test
 - CAMDEX-DS (informant interview) and CAMCOG-DS (subject cognitive assessment)
 - Clued recall test (Free and cued immediate and delayed recall)
 - Digit span (forward/ backward)
 - Picture cancellation task (Devenny)
 - Barcelona Test (assessment of limb apràxia-pantomime of intransitive gestures)
 - Cats and dogs tests
 - Verbal fluency (animals)
 - Abstract thinking (Barcelona test)

- Mental State examination- DS (MEC-DS), screening instrument, not validated
- Neuropsychiatric inventory (12 item NPI)
- Fundació Catalana Síndrome de Down Functional Scale, not validated
- Dementia questionnaire for people with intellectual disabilities (DMR)
- Neuropsychological battery from the Sanders-Brown Center on Aging in Kentucky, USA.
 - Kauffman Brief Intelligence Test
 - Severe Impairment Battery
 - Brief Praxis Test
 - Fuld Object Memory Evaluation
 - Peabody Picture Vocabulary Test (4th ed.)
 - Children's Memory Scale: Dot Locations
 - Category verbal fluency
 - Beery Visual Motor Integration
 - Neuropsychiatric inventory (NPI)
 - WISC-R Block Design
 - Vineland Adaptive Behavior Scale
 - Behavioral Rating of Executive Functions (BRIEF)
 - Adaptive Behavior Assessment System (ABAS)
 - Dementia Questionnaire for people with intellectual disabilities (DMR)
- Neuropsychological battery from the Down Syndrome Biomarker Initiative (DSBI) project in San Diego, USA is detailed in Rafii et al. *Front Behav Neurosci.* 2015, 14:9:239.

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Abbreviations

ADAD Autosomal dominant Alzheimer's disease

APOE	Apolipoprotein-E
APP	Amyloid precursor protein
ARIA	Amyloid-related imaging abnormalities
CAA	Cerebral amyloid angiopathy
CAMCOG-DS	Cambridge Examination for mental Disorders of Older People with DS and Others with Intellectual Disabilities-Cognitive Scale
cSS	Cortical superficial siderosis
d-EOAD	Dementia in early onset Alzheimer's disease
DABNI	Down Alzheimer Barcelona Neuroimaging Initiative
dAD-DS	Alzheimer's disease dementia in Down syndrome
dADAD	Dementia in autosomal dominant Alzheimer's disease
DS	Down syndrome
DSBI	Down syndrome biomarker initiative
EOAD	Early onset Alzheimer's disease
FLAIR	Fluid attenuation inversion recovery
GRE	Gradient echo
ICH	Intracerebral hemorrhage
mBCAA	Modified Boston criteria for cerebral amyloid angiopathy
MMSE	Mini-mental state examination
MTA	Medial temporal atrophy
p-EOAD	Prodromal Alzheimer's disease in early onset Alzheimer's disease
pAD-ADAD	Prodromal Alzheimer's disease in autosomal dominant Alzheimer's disease
pAD-DS	Prodromal Alzheimer's disease in Down syndrome
PSEN1	Presenilin 1
PSEN2	Presenilin 1
SWI	Susceptibility weighted imaging
WMH	White matter hyperintensities

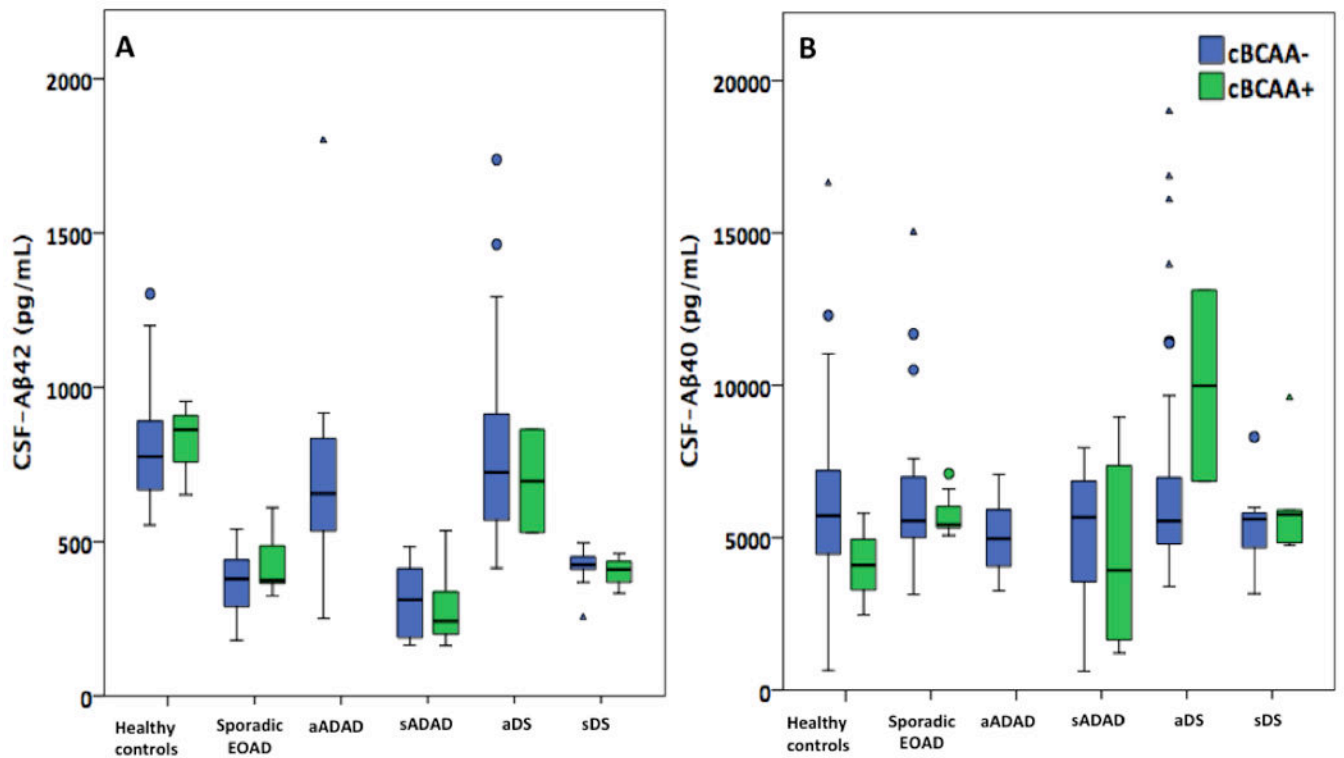


Figure 1. CSF biomarker levels in the clinical groups, according to modified Boston criteria for CAA (cBCAA)

Box plot displaying the distribution of CSF-Aβ42 (A) and CSF-Aβ40 (B) from healthy controls, sporadic EOAD, asymptomatic ADAD (aADAD), symptomatic ADAD (sADAD), asymptomatic DS (aDS), and symptomatic DS (sDS). Subgroups in blue represent those subjects who do not fulfill cBCAA, subgroups coloured in green do fulfill cBCAA for possible or probable CAA.

No differences in levels of CSF-Aβ42 or in CSF-Aβ40 were detected between subjects that fulfilled cBCAA and those who did not within each clinical group.

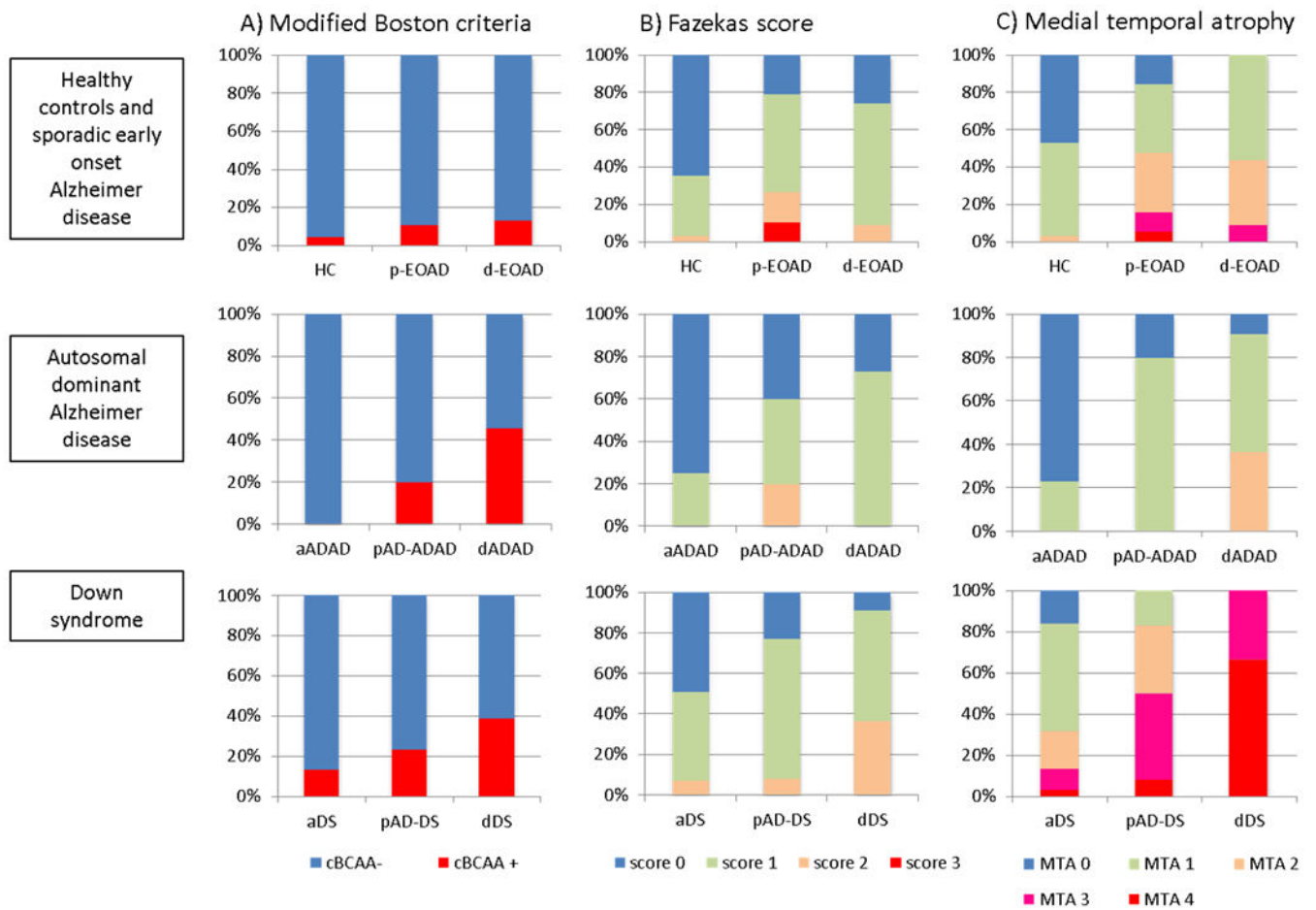


Figure 2. Proportion of subjects in each group according modified Boston criteria, Fazekas score, and medial temporal lobe atrophy

Frequency bar graph showing: A) modified Boston criteria fulfillment, B) Fazekas score and, C) MTA score of each clinical group. Healthy controls and sporadic AD in the first row, ADAD in the second row, DS in the third row.

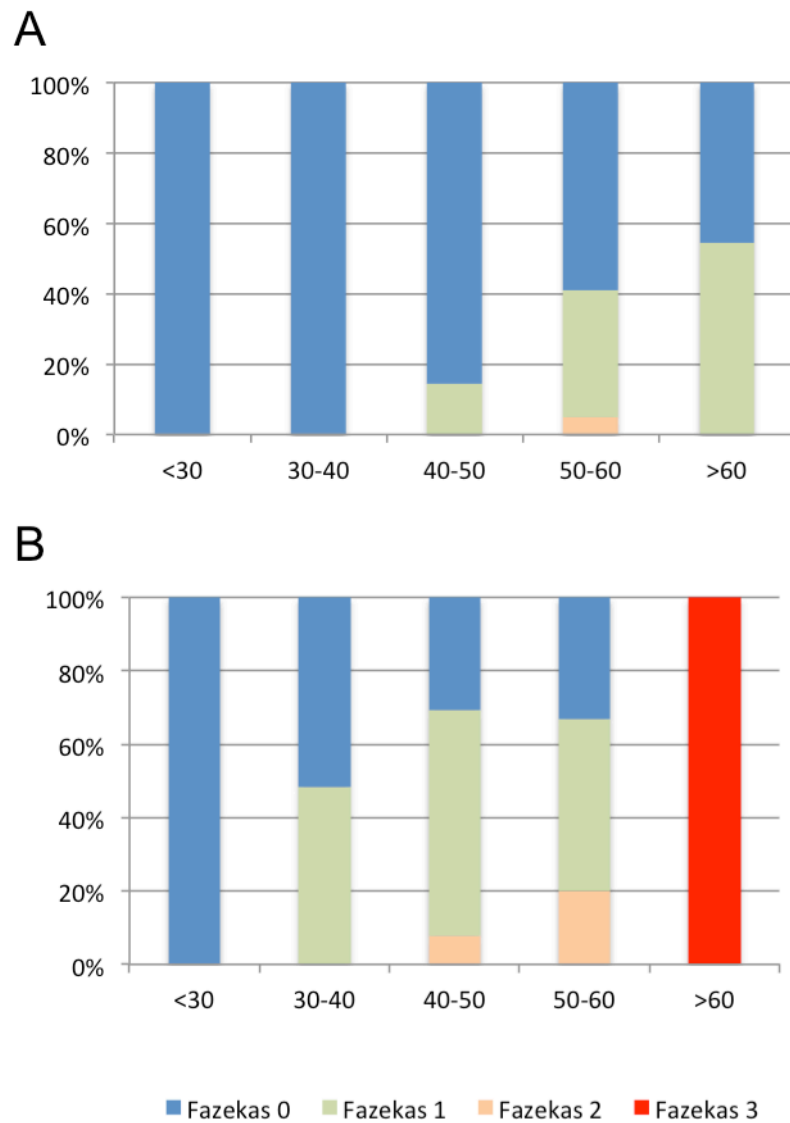


Figure 3. Proportion of subjects according to Fazekas score by range of age

The frequency bar graphs showing the percentage of subjects with each Fazekas categories by age in: A) Healthy controls and, B) asymptomatic Down syndrome subjects.

Table 1
Demographics and CSF biomarker characteristics of the participants

	Healthy controls		Sporadic EOAD		Asymptomatic ADAD		Symptomatic ADAD		Asymptomatic DS		Symptomatic DS	
			d-EOAD				pAD-ADAD				pAD-DS	
	p-EOAD	d-EOAD	pAD-ADAD	dADAD	pAD-DS	dAD-DS						
N	68	23	13	5	11	91	13	13	56.2 (3)	54	54	15.4
Age, years old	54.3 (9)	61.8 (7)	33.6 (11)	54.8 (17)	47.3 (10)	39.6 (16)	51.5 (5)	56.2 (3)	54	54	15.4	33.3
Gender, % men	35.3	39.1	23.1	60	45.5	56	0	0	0	0	0	8.3
High blood pressure	13.2	13	0	0	0	3.1	0	0	0	0	0	8.3
APOE-ε4 carriers, % ^Ω	32.4	43.5	7.7	0	18.2	34.6	20	20	20	20	20	8.3
APOE-ε2 carriers, % ^Ω	4.4	4.3	7.7	0	27.3	19.2	10	10	10	10	10	8.3
MMSE/ Total CAMCOG [‡]	29 (2)	20 (7)	30 (1)	24 (4)	20 (7)	80 (16)	74 (22)	74 (22)	42 (34)	42 (34)	42 (34)	8.3
CSF-Aβ42, pg/mL	779.5 (237)	360.5 (118)	655.5 (412)	217 (242)	279 (201)	724.8 (294)	432.5 (33)	432.5 (33)	400.8 (56)	400.8 (56)	400.8 (56)	8.3
CSF-Aβ40, pg/mL	5617.9 (2738)	6224.5 (1923)	4966.8 (2225)	4837 (3602)	5783.7 (6739)	5559.5 (2088)	5605 (1143)	5605 (1143)	5763.3 (780)	5763.3 (780)	5763.3 (780)	8.3
CSF ratio-Aβ42/40	0.143 (0.07)	0.060 (0.02)	0.130 (0.14)	0.070 (0.01)	0.060 (0.18)	0.106 (0.08)	0.078 (0.02)	0.078 (0.02)	0.069 (0.03)	0.069 (0.03)	0.069 (0.03)	8.3

Unless otherwise specified, values are presented as medians (interquartile range, IQR).

^ΩMedian relative age (IQR) was -11.9 (14.4), 3.15 (6.5) and 3.5 (4.8) in asymptomatic ADAD, pAD-ADAD and dADAD respectively.

^ΩAPOE was available from 68/68 HC, 19/19 p-EOAD, 23/23 d-EOAD, 13/13 asymptomatic ADAD, 5/5 pAD-ADAD, 11/11 dADAD, 78/91 asymptomatic DS, 10/13 pAD-DS, and 12/13 dAD-DS.

[‡]CAMCOG score was available in 55/91 asymptomatic DS subjects, 7/14 pAD-DS subjects, and 3/12 dAD-DS subjects.

[‡]CSF was available from 68/68 HC, 19/19 p-EOAD, 23/23 d-EOAD, 7/13 asymptomatic ADAD, 5/5 pAD-ADAD, 9/11 dADAD, 36/91 asymptomatic DS, 9/13 pAD-DS, and 6/13 dAD-DS.

Key: Aβ42, amyloid-β 1-42; Aβ40, amyloid-β40; APOE, Apolipoprotein E; CAMCOG-DS, Cambridge Examination for mental Disorders of Older People with Down Syndrome and Others with Intellectual Disabilities-Cognitive Scale; CSF, cerebrospinal fluid; dADAD, dementia in autosomal dominant Alzheimer disease; dAD-DS, Alzheimer dementia in Down syndrome; d-EOAD, dementia early onset Alzheimer disease; EOAD, early onset sporadic Alzheimer disease; MMSE, minimal state examination; pAD-ADAD, prodromal Alzheimer disease in autosomal dominant Alzheimer disease; pAD-DS, prodromal Alzheimer disease in Down syndrome; p-EOAD, prodromal sporadic early onset Alzheimer disease.

Table 2

Neuroimaging findings

	Healthy controls		Sporadic EOAD		Asymptomatic ADAD		Symptomatic ADAD		Asymptomatic DS		Symptomatic DS	
			p-EOAD	d-EOAD			pAD-ADAD	dADAD			pAD-DS	dAD-DS
N	68		19	23	13		5	11	91		13	13
MRI (%GRE%/SWI)	38.2/ 61.8		24.1/57.9	39.1/60.9	100/0		100/0	100/0	46.2/53.8		23.1/76.9	30.8/69.2
% Lobar microbleeds	2.9		10.5	8.7	0		20	45.5	12.1		23.1	38.5
% cSS	1.5		0	8.7	0		0	0	3.3		7.7	23.1
Lobar ICH	0		0	0	0		0	0	1.1		0	15.4
Boston criteria	4.4		10.5	13	0		20	45.5	13.2		23.1	38.5
Fazekas												
score 0	64.7		21.1	26.1	75		40	27.3	49.4		23.1	9.1
score 1	32.4		52.6	65.2	25		40	72.7	43.7		69.2	54.5
score 2	2.9		15.8	8.7	0		20	0	5.7		7.7	36.4
score 3	0		10.5	0	0		0	0	1.1		0	0
MTA												
score 0-1	97.1		52.6	56.5	100		100	63.6	68.2		16.7	0
score 2-4	2.9		47.4	43.5	0		0	36.4	31.8		83.3	100

Unless otherwise specified, values are presented as proportions.

* Fazekas score was not available in 4 asymptomatic DS subjects and in 2 dAD-DS subjects.

** MTA score was not available in 1 AD subject, in 3 asymptomatic DS subjects, in 1 pAD-DS subject, and in 1 dAD-DS subject.

Key: Aβ42, amyloid-β 1-42; Aβ40, amyloid-β40, APOE, Apolipoprotein E; CAMCOG-DS, Cambridge Examination for mental Disorders of Older People with Down Syndrome and Others with Intellectual Disabilities-Cognitive Scale; CSF, cerebrospinal fluid; cSS, cortical superficial siderosis; dADAD, dementia in autosomal dominant Alzheimer disease; dAD-DS, Alzheimer dementia in Down syndrome; d-EOAD, dementia in sporadic early onset Alzheimer disease; EOAD, early onset sporadic Alzheimer disease; GRE, gradient echo sequences; HBP, high blood pressure; ICH, intracerebral hemorrhage; MMSE, minimal state examination; MRI, magnetic resonance imaging; MTA, medial temporal atrophy; pAD-ADAD, prodromal Alzheimer disease in autosomal dominant Alzheimer disease; pAD-DS, prodromal Alzheimer disease in Down syndrome; p-EOAD, prodromal sporadic early onset Alzheimer disease; SWI, susceptibility weighted imaging.