

Comorbidities in Early-Onset Sporadic versus Presenilin-1 Mutation-Associated Alzheimer's Disease Dementia: Evidence for Dependency on Alzheimer's Disease Neuropathological Changes

Diego Sepulveda-Falla¹, Carlos Andrés Villegas Lanau², Charles White III³, Geidy E. Serrano⁴, Juliana Acosta-Uribe^{2,5}, Barbara Mejía-Cupajita^{2,5}, Nelson David Villalba-Moreno¹, Pinzhang Lu¹, Markus Glatzel¹, Julia K. Kofler⁶, Bernardino Ghetti⁷, Matthew P. Frosch⁸, Francisco Lopera Restrepo², Kenneth S. Kosik⁵ and Thomas G. Beach⁴

1. Institute of Neuropathology, University Medical Center Hamburg-Eppendorf, Martinistraße 52 20246 Hamburg, Gebäude Nord 27 / Raum 02.005 Telefon: +49 (0)40 7410- 51920
Fax: +49 (0)40 7410-55696

2. Faculty of Medicine, Neuroscience Group of Antioquia, University of Antioquia, Medellin, Colombia

3. Neuropathology Section, Department of Pathology, University of Texas Southwestern Medical Center, Dallas, TX

4. Civin Laboratory for Neuropathology, Banner Sun Health Research Institute, 10515 W Santa Fe Drive, Sun City, AZ 85351

5. Neuroscience Research Institute and Department of Molecular Cellular and Developmental Biology, University of California Santa Barbara

6. Department of Pathology, University of Pittsburgh, Pittsburgh, PA

7. Department of Pathology and Laboratory Medicine, Indiana University School of Medicine, Indianapolis, IN

8. Department of Pathology, Massachusetts General Hospital and Harvard Medical School, Boston, MA

Abstract

Autopsy studies have demonstrated that comorbid neurodegenerative and cerebrovascular disease occur in the great majority of subjects with Alzheimer disease dementia (ADD), and are likely to additively alter the rate of decline or severity of cognitive impairment. The most important of these are Lewy body disease (LBD), TDP-43 proteinopathy and cerebrovascular disease, including white matter rarefaction (WMR) and cerebral infarcts. Comorbidities may interfere with ADD therapeutic trials evaluation of ADD clinical trials as they may not respond to AD-specific molecular therapeutics. It is possible, however, that at least some comorbidities may be, to some degree, secondary consequences of AD pathology, and if this were true then effective AD-specific therapeutics might also reduce the extent or severity of comorbid pathology. Comorbidities in ADD caused by autosomal dominant mutations such as those in the presenilin-1 (*PSEN1*) gene may provide an advantageous perspective on their pathogenesis, and deserve attention because these subjects are increasingly being entered into clinical trials. As ADD associated with *PSEN1* mutations has a presumed single-cause etiology, and the average age at death is under 60, any comorbidities in this setting may be considered as at least partially secondary to the causative AD mechanisms rather than aging, and thus indicate whether effective ADD therapeutics may also be effective for comorbidities. In this study, we sought to compare the rates and types of ADD comorbidities between subjects with early-onset sporadic ADD (EOSADD; subjects dying under age 60) versus ADD associated with different types of *PSEN1* mutations, the most common cause of early-onset autosomal dominant ADD. In particular, we were able to ascertain, for the first time, the prevalences of a fairly complete set of ADD comorbidities in United States (US) *PSEN1* cases as well as the Colombian E280A *PSEN1* kindred. Data for EOSADD and US *PSEN1* subjects (with multiple different mutation types) was obtained from the National Alzheimer Coordinating Center (NACC). Colombian cases all had the E280A mutation and had a set of neuropathological observations classified, like the US cases according to the NACC NP10 definitions. Confirmatory of earlier reports, NACC-defined Alzheimer Disease Neuropathological Changes (ADNC) were consistently very severe in early-onset cases, whether sporadic or in *PSEN1* cases, but were slightly less severe in EOSADD. Amyloid angiopathy was the only AD-associated pathology type with widely-differing severity scores between the 3 groups, with median scores of 3, 2 and 1 in the *PSEN1* Colombia, *PSEN1* US and EOSADD cases, respectively. Apolipoprotein E genotype did not show significant proportional group differences for the possession of an E-4 or E-2 allele. Of ADD

comorbidities, LBD was most common, being present in more than half of all cases in all 3 groups. For TDP-43 co-pathology, the Colombian *PSEN1* group was the most affected, at about 27%, vs 16% and 11% for the US *PSEN1* and sporadic US cases, respectively. Notably, hippocampal sclerosis and non-AD tau pathological conditions were not present in any of the US or Colombian *PSEN1* cases, and was seen in only 3% of the EOSADD cases. Significant large-vessel atherosclerosis was present in a much larger percentage of Colombian *PSEN1* cases, at almost 20% as compared to 0% and 3% of the US *PSEN1* and EOSADD cases, respectively. Small-vessel disease, or arteriolosclerosis, was much more common than large vessel disease, being present in all groups between 18% and 37%. Gross and microscopic infarcts, however, as well as gross or microscopic hemorrhages, were generally absent or present at very low percentages in all groups. White matter rarefaction (WMR) was remarkably common, at almost 60%, in the US *PSEN1* group, as compared to about 18% in the EOSADD cases, a significant difference. White matter rarefaction was not assessed in the Colombian *PSEN1* cases. The results presented here, as well as other evidence, indicates that LBD, TDP-43 pathology and WMR, as common comorbidities with autosomal dominant and early-onset sporadic ADD, should be considered when planning clinical trials with such subjects as they may increase variability in response rates. However, they may be at least partially dependent on ADNC and thus potentially addressable by anti-amyloid or and/anti-tau therapies.

Key Words: Lewy body; alpha-synuclein; TDP-43; white matter rarefaction; cerebral infarcts; clinical trial; aging; autosomal dominant mutation; E280A; Colombia; autopsy; neuropathology

Introduction

Autopsy studies have demonstrated that comorbid neurodegenerative and cerebrovascular disease occur in the great majority of subjects with Alzheimer's disease dementia (ADD), and are likely to additively alter the rate of decline or severity of cognitive impairment [1-5]. Some of these ADD comorbidities including Lewy body disease (LBD), hippocampal sclerosis, TDP-43 proteinopathy and cerebral infarcts, are capable of causing dementia even when ADD neuropathological changes (ADNC) are low or absent. Comorbidities may interfere with the evaluation of ADD clinical trials as they may not respond to ADNC-specific molecular therapeutics. At present, it is not possible to stratify ADD trials for comorbidity presence as there are no proven sensitive and specific diagnostics nor disease-modifying therapies. It is possible, however, that at least some comorbidities may be, to some degree, secondary consequences of ADNC, and if this were true then effective ADNC-specific therapeutics might also reduce the extent or severity of comorbid pathology.

As most reports have focused on the very old, because comorbidities increase with age, it has often been assumed that these would be largely absent in younger, clinical trial-aged sporadic ADD subjects. However, it has recently been shown that comorbidities, particularly LBD, white matter rarefaction (WMR) and TDP-43 proteinopathy, are present even in many cases of early-onset sporadic ADD (EOSADD)[6,7].

Much less is known about the rate of comorbidities in early-onset ADD caused by autosomal dominant mutations in genes such as presenilin-1 (*PSEN1*). This is a critical knowledge gap as ADD in this setting may provide an advantageous perspective on the pathogenesis of ADD comorbidities, and because these subjects are increasingly being entered into clinical trials. It is important, therefore, to understand the comorbidity rates for both early-onset sporadic and inherited ADD so that subject stratification might allow the recognition of significant trial outcomes for subject subsets, even when overall results are negative. As ADD associated with *PSEN1* mutations has a presumed single-cause etiology, and the average age at death is under 60, any comorbidities in this setting may be considered as at least partially secondary to the causative AD mechanisms rather than aging, and thus indicate whether effective ADD therapeutics may also be effective for comorbidities.

In this study, we sought to compare the rates and types of ADD comorbidities between subjects with early-onset sporadic ADD (EOSADD) versus ADD associated with different types of *PSEN1* mutations, the most common cause of early-onset autosomal dominant ADD. In particular, we were able to ascertain, for the first

time, the prevalences of a fairly complete set of ADD comorbidities in United States (US) *PSEN1* cases as well as the Colombian E280A *PSEN1* kindred [8-9]. A previous study used NACC data to profile *PSEN1* mutation ADD [10] but less comorbidity data had been reported at that time.

Methods

The approach used in this study is the same as used in a previous [6] survey of neuropathological comorbidities in sporadic ADD using data from the National Alzheimer Coordinating Center (NACC)[11,12]. Included subject data for sporadic ADD in this study was restricted to those who died under the age of 60 who had dementia and met NIA-AA intermediate or high AD NDNC [13,14], considered to be a sufficient cause of cognitive impairment or dementia; subjects known to have autosomal dominant AD mutations were excluded. Only data from the most recent version of the Neuropathology form (NP10) was used (<https://files.alz.washington.edu/documentation/rdd-np.pdf>), as this is most inclusive of comorbidity data. Also included were comparative data for *PSEN1* ADD subjects from the United States, derived from the same initial NACC dataset, and from Colombia. The US cases included subjects with several different *PSEN1* mutations while the Colombian cases all had the E280A mutation [8,9]; 36 of the latter have been reported in earlier studies [15,16].

Pathology categories investigated included the major AD-specific lesions: senile or amyloid plaques (NACC variable NPTHAL; all plaque types classified by Thal amyloid phase [17], neurofibrillary tangles (NACC variable NPBRAAK; classified by Braak stage [18,19]), neuritic plaques (NACC variable NPNEUR; classified according to CERAD) [20], diffuse plaques (NACC variable NPDIFF; classified analogously to CERAD neuritic plaques), amyloid angiopathy (NPAMY; classified as none, mild, moderate or severe) and NIA-AA AD Neuropathological Change Level (NACC variable NPADNC [13,14].

Comorbid neurodegenerative conditions investigated included LBD (NACC variable NPLBOD; for this study only presence or absence in any brain region was recorded), TDP-43 pathology (NACC variable NPTDPB; TDP-43 pathology present at least in amygdala, hippocampus or entorhinal area), non-AD tauopathy (NACC variable NPFTDTAU; any non-AD tauopathy including progressive supranuclear palsy, corticobasal degeneration, Pick's disease, argyrophilic grains, chronic traumatic encephalopathy, or "other"). For each comorbid pathology and ADD group, the proportion of cases possessing that pathology was determined.

Comorbid cerebrovascular conditions investigated included circle of Willis arteriosclerosis (NACC variable NACCAVAS; for this study only “moderate” and “severe” qualified for presence of the condition), old gross cerebral infarcts (NACC variable NPINF; for this study 1 or more large or lacunar infarcts qualified for the condition), old microscopic infarcts (NACC variable NPOLD; for this study 1 or more old microinfarcts qualified for the condition), old microscopic hemorrhages (NACC variable NPOLDD; for this study 1 or more old microhemorrhages qualified for the condition), arteriolosclerosis (NACC variable NACCARTE; for this study only “moderate” and “severe” qualified for presence of the condition) and white matter rarefaction (NACC variable NPWMR; for this study only “moderate” and “severe” qualified for presence of the condition). For each comorbid pathology, the proportion of cases possessing that pathology, relative to those for whom the pathology’s presence or relative absence was specifically recorded, was determined.

Statistical methods used for continuous data included analysis of variance (ANOVA) with post-hoc paired comparisons by the Tukey test. Chi-square and Fisher exact tests were used for comparisons of proportional data. For all tests, the significance level was set at $p < 0.05$.

Results

Demographic and mutation data for all cases are shown in Table 1. As a result of our selection criteria, all EOSADD cases were less than 60 years old at death, with a mean age of 56.3 years. For USA *PSEN1* cases the mean age at death was 53.4 years while for Colombian cases it was 57.7 years. For US *PSEN1* subjects a single subject died at age 75; all other US *PSEN1* subjects were 65 or less at death. For Colombian subjects, 4 died at 70 or older. All USA *PSEN1* subjects were Caucasian/white; 13 were non-Hispanic while 6 were Hispanic. There were 12 different *PSEN1* mutations in the US cases while all Colombian cases were multiracial-Hispanic and had the E280A mutation.

Table 1. Demographics and mutation genotypes for *PSEN1* and sporadic EOSADD cases.

Group	Age at Death (mean, SD, range)	Sex	<i>PSEN1</i> Mutation Genotypes	
			Mutation	N (Cases)
<i>PSEN1</i> - USA N = 19	53.4; 8.6; 41-75	6 M, 13F	A431E	5
			G206A	3
			A426P	2
			P88L	2
			H163R	2
			P267A	1
			G206E	1
			M233L	1
			N135S	1
			S130L	1
<i>PSEN1</i> - Colombia N = 50	57.7; 7.25; 42-79	15M, 35F	E280A12 (All)	
Sporadic - USA N = 33	56.3; 2.98; 47-59	25M; 8 F	N/A	

Confirmatory of earlier reports, [6-8,10) ADNC is consistently very severe in early-onset cases, whether they are sporadic or autosomal dominant (Table 2). All 3 groups had median scores that were the highest possible for Thal amyloid phase, Braak neurofibrillary stage, diffuse plaque density and NIA-AA ADNC Level. When the mean scores were compared, however, ANOVA showed significant group differences for Thal amyloid phase, diffuse plaque density and ADNC. On pairwise comparisons for most ADD pathologies, *PSEN1* groups had significantly greater scores than the EOSADD group. Analysis of variance and pairwise comparisons for CERAD neuritic plaque density and Braak neurofibrillary stage were not significant.

Amyloid angiopathy, which is very consistently present in ADD, was the only AD-associated pathology type with widely-differing severity scores between the 3 groups, with median scores of 3, 2 and 1 in the *PSEN1* Colombia, *PSEN1* US and EOSADD cases, respectively (Table 2). Analysis of variance showed very significant group differences as well as significant differences between both *PSEN1* groups and the EOSADD group.

Apolipoprotein E genotype, which strongly influences not only the prevalence and age of onset of ADD but also the prevalence of amyloid angiopathy, did not show significant proportional group differences for the possession of an ϵ -4 or ϵ -2 allele, although the US *PSEN1* and EOSADD groups had 1.6-fold and 1.8-fold

greater ϵ -4 possession as compared to the Colombian *PSEN1* group (Table 2). Possession of the ϵ -2 genotype was relatively rare among all groups, between ~ 3% and 14%.

Table 2. Autopsy data for AD-related neuropathological variables in sporadic EOSADD and *PSEN1* ADD cases. *ApoE*- ϵ 4 and *apoE*- ϵ 2 = number and percentage of cases with one or more *ApoE* ϵ -4 or ϵ -2 alleles. Means and medians are shown for the following: NPTHAL = Thal amyloid phase; NPBRAAK = Braak neurofibrillary stage; NPNEUR = CERAD neuritic plaque density; NPDIFF = diffuse plaque density; NPAMY = cerebral amyloid angiopathy density; NPADNC = NIA-AA AD Neuropathological Change level.

Group	ApoE-4	ApoE-2	NPTHAL ¹	NPBRAAK	NPNEUR	NPDIFF ²	NPADNC ³	NPAMY ⁴
<i>PSEN1</i>-US	6/14 42.9%	2/14 14.3%	5; 5	6; 6	3; 3	3; 3	3; 3	2.10; 2
<i>PSEN1</i>-COL	9/34 26.5%	1/34 2.94%	5; 5	6; 6	2.9; 3	3; 3 (n = 36)	3; 3	2.58; 3
Sporadic-US	16/33 48.5%	2/33 6.1%	4.72; 5	5.79; 6	2.85; 3	2.70; 3	2.88; 3	1.42; 1

1. ANOVA $p = 0.002$; Tukey $p < 0.05$ for EOSADD vs both *PSEN1* groups.
2. ANOVA $p = 0.022$; Tukey $p < 0.05$ for EOSADD vs *PSEN1*-Col group.
3. ANOVA $p = 0.012$; Tukey $p < 0.05$ for EOSADD vs *PSEN1* Col group.
4. ANOVA $p = 0.0001$; Tukey $p < 0.05$ for EOSADD vs both *PSEN1* groups.

The proportion of cases lacking any typical ADD comorbidities, classified here as “AD-Only” cases (Table 3), was low for both of the *PSEN1* groups, less than 30% and only a little higher, 44%, for the EOSADD group. The difference in proportions between groups was not significant.

Of ADD comorbidities, LBD was most common (Table 3), being present in more than half of all cases in this study, and was particularly common in the Colombian *PSEN1* and EOSADD cases, at about 70% for each group; the differences in group proportions were not significant, however. Of those cases with Lewy body disease, the brain distribution in the majority of all groups was limbic predominant and amygdala-only, especially in the *PSEN1* groups (Table 4). The brainstem-predominant stage was uncommon in all groups. The olfactory bulb-only stage was not seen in the sporadic group but was significantly more common, at 30%, in the US *PSEN1* group, as compared to 8% for the Colombian *PSEN1* group. The neocortical stage was not seen in the US *PSEN1* group and was only about 10-12% of the EOSADD and Colombian groups; these differences were not significant.

For TDP-43 co-pathology (Table 3), the Colombian *PSEN1* group was the most affected, at about 27%, vs 16% and 11% for the US *PSEN1* and sporadic US cases, respectively. The group differences were significant and the paired comparison between EOSADD and Colombian *PSEN1* cases was significant. For all US

PSEN1 and EOSADD cases, TDP-43 pathology was confined to the amygdala, hippocampus and entorhinal area without any pathology in the neocortex or spinal cord. Colombian *PSEN1* cases were positive in the amygdala with incomplete assessment elsewhere. Notably, hippocampal sclerosis and non-AD tau pathological conditions were not present in any of the US or Colombian *PSEN1* cases, and was seen in only 3% of the EOSADD cases.

Table 3. Autopsy data for EOSADD and *PSEN1* cases, showing proportions of common ADD comorbidities. Numerator is the number of cases meeting criteria for the comorbidity while denominator is the number of AD cases evaluated for the condition; also given is percentage. AD Only = AD without any of the other conditions in this table and without NPINF, NPOLD and NPOLDD from Table 4; NPLBOD = Lewy body disease; NPHIPSCL = hippocampal sclerosis; TDPNOS = TDP-43 pathology in amygdala, hippocampus or entorhinal area; NPFTDTAU = non-AD tau pathology (PSP, CBD, Pick's, argyrophilic grains, other).

Group	AD Only	NPLBOD	NPHIPSCL	TDPNOS ¹	NPFTDTAU
<i>PSEN1</i>-US	2/19; 10.5%	10/19; 52.6%	0/19; 0%	3/19; 15.8%	0/19; 0%
<i>PSEN1</i>-COL	5/17; 29.4%	12/17; 70.6%	0/50; 0%	4/15; 26.7%	0/36; 0%
Sporadic-US	8/18; 44.4%	19/33; 57.6%	1/32; 3.1%	2/18; 11.1%	1/33; 3.0%

1. Chi-square $p = 0.04$; Fisher exact test $p = 0.03$ for sporadic US vs *PSEN1* Col group.

Table 4. Autopsy data for EOSADD and *PSEN1* cases, showing brain stages of cases with Lewy body disease. Numerator is the number of cases meeting criteria for the stage while denominator is the number of AD cases that were positive in any region for Lewy body disease; also given is percentage. LB-OB = olfactory bulb only; LB-BS = brainstem predominant; LB-Amyg = amygdala predominant; LB-Limb = limbic (transitional); LB-Neo = Neocortical (diffuse).

Group	LB-OB ¹	LB-BS	LB-Amyg	LB-Limb	LB-Neo
<i>PSEN1</i>-US	3/10; 30%	0/10; 0%	3/10; 30%	4/10; 40%	0/10; 0%
<i>PSEN1</i>-COL	1/12; 8.3%	1/12; 8.3%	8/12; 66.7%	4/12; 26.7%	2/12; 11.8%
Sporadic-US	0/33; 0%	1/33; 3.0%	10/33; 30.3%	4/33; 12.1%	3/33; 10.0%

1. Fisher exact test $p = 0.002$ for US *PSEN1* vs *PSEN1* Col group.

Cerebrovascular disease categories varied between the EOSADD and *PSEN1* groups (Table 5).

Significant large-vessel atherosclerosis, assessed by NACC as circle of Willis atherosclerosis (NACCAVAS), was present in a much larger percentage of Colombian *PSEN1* cases, at almost 20% as compared to 0% and 3% of the USA *PSEN1* and EOSADD cases, respectively. The group differences were significant as was the comparison between EOSADD cases vs Colombian *PSEN1* cases. Small-vessel disease, or arteriolosclerosis

(NACCARTE) was much more common than large vessel disease, being present in all groups between 18% and 37%; the group differences were not significant on a Chi-square test. The consequences of vascular disease include gross and microscopic infarcts, as well as microscopic hemorrhages, but all were generally absent or present at very low percentages in all groups.

White matter rarefaction, conventionally regarded as a consequence of arteriolosclerosis, was remarkably common, at almost 60%, in the USA *PSEN1* group, as compared to about 18% in the EOSADD cases, a significant difference. White matter rarefaction was not assessed in the Colombian *PSEN1* cases.

Table 5. Autopsy data for sporadic EOSADD and *PSEN1* cases, listing major cerebrovascular comorbidities. The numerator is the number of cases meeting criteria for the condition, while the denominator is the number of cases evaluated for the condition. NACCAVAS = severity of atherosclerosis of circle of Willis > mild; NPINF = gross infarcts including lacunes; NPOLD = old microinfarcts; NPOLDD = old cerebral microhemorrhages; NACCARTE = arteriolosclerosis > mild; NPWMR = white matter rarefaction > mild.

Group	NACCAVAS ¹	NPINF	NPOLD	NPOLDD	NACCARTE	NPWMR ²
<i>PSEN1</i> -US	0/19; 0%	2/19; 10.5%	0/19; 0%	0/19; 0%	7/19; 36.8%	11/19; 57.9%
<i>PSEN1</i> -COL	9/48; 18.75%	0/25; 0%	2/48; 4.2%	1/48; 2.1%	9/36; 25%	N/A
Sporadic-US	1/33; 3.0%	0/33; 0%	2/33; 6.1%	0/29; 0%	6/32; 18.7%	5/28; 17.9%

1. Chi-square test $p = 0.05$ between groups; Fisher exact test $p = 0.04$ for sporadic vs *PSEN1*-COL.
2. Fisher exact test $p = 0.01$ between EOADD and US *PSEN1* groups.

Discussion

This report updates an earlier comparison by Ringman et al [10] of NACC neuropathology scores for *PSEN1* and sporadic ADD cases (of all ages), based on the prior NP9 NACC neuropathology data elements. We used accumulated data from the NP10 version, which is more extensive in its inclusion of ADD comorbidities. Additionally, we included a comparison with *PSEN1* E280A cases from the Colombian kindred, which have not previously been systematically evaluated for ADD comorbidities.

Confirmatory of earlier NACC-based reports [10,21] we find that the mean severity scores for amyloid plaques, tau-tangles, and especially CAA, are uniformly greater in *PSEN1* ADD cases as compared with those for sporadic ADD, but in this study we have also found that sporadic ADD cases dying before age 60 have amyloid and tau pathology that are almost equivalent in severity to that found in *PSEN1* mutation cases. This concurs with previous reports of sporadic ADD histopathology [6,7] documenting the greatest severity in the cases with earliest onset, and progressively decreased severity with advancing age afterwards. Cerebral

amyloid angiopathy (CAA) was the only AD-associated pathology type with markedly differing severity scores between the 3 studied groups, with median scores of 3, 2 and 1 in the *PSEN1* Colombia, *PSEN1* US and EOSADD cases, respectively. The greater CAA severity in *PSEN1* mutation-associated ADD has been previously documented [10] and is of particular relevance because of the greater risk with CAA for amyloid-related imaging abnormalities (ARIA), vasogenic edema and hemorrhage, which are limiting factors for anti-amyloid monoclonal antibody therapy. Other predisposing factors to CAA include the apolipoprotein ϵ -4 allele, hypercholesterolemia and Hispanic ethnicity [22,23], and, within *PSEN1* cases, mutations beyond codon 200 [10]. The greater prevalence of significant CAA in the E280A Colombian cases might therefore be at least partially due to the higher-codon mutation location, greater proportion with Hispanic ethnicity and greater likelihood of hypercholesterolemia in Colombia as compared to the US [24], although the current set of Colombian subjects had a lower ϵ -4 carriage rate. Despite the severe CAA severity, only 1/67 combined *PSEN1* cases, and no EOSADD cases, had gross or microscopic brain hemorrhages.

The distribution of apolipoprotein E (*APOE*) ϵ -4 and ϵ -2 alleles differed between the studied groups. The ϵ -4 allele was most common in the *PSEN1* US group, followed by the EOSADD group. Carriage of an *APOE* ϵ -4 or ϵ -2 allele has been associated with earlier or later clinical onset, respectively, in both sporadic ADD and *PSEN1* ADD subjects [25,26]. The USA *PSEN1* group had a higher proportion of ϵ -4 carriers than the Colombian *PSEN1* group, perhaps contributing to their earlier mean onset age in the current study subjects. There was an extremely low proportion (1/34) of Colombian *PSEN1* cases with the ϵ -2 allele but with this small sample this was not significantly different from the other groups. A single case of the Christchurch *APOE* ϵ -3 mutation, which in some respects has biochemical similarities with the ϵ -2 allele, has recently been reported in the Colombian *PSEN1* kindred [27]; this case had a remarkable 30 year delay of clinical onset as well as reduced levels of ADNC [28]. Other disease-modifying genetic loci have been described for *PSEN1* ADD [10,25,29-31] and these offer important clues for the design of disease-modifying strategies.

ADD comorbidities are not restricted to the oldest-old [6,7] as some comorbidity types are common even in early-onset ADD, whether sporadic or inherited. The percentage of cases with ADNC as the sole pathology, defined as the absence of LBD, TDP-43, non-AD tauopathy, infarcts and microhemorrhages, was low in all

groups. The NACC *PSEN1* cases had the lowest rate of AD as a sole major pathology, at ~ 10% while this rate was ~ 29% for the Colombian *PSEN1* cases and 48% for the EOSADD group.

Of typical ADD comorbidities, LBD is the most common in subjects under age 80 [6], and reaches very high prevalences in this study's *PSEN1* and EOSADD subjects. Colombian *PSEN1* cases had the highest LBD comorbidity rates, at ~ 70%, as compared to the US EOSADD and *PSEN1* group at 57% and 53%, respectively. The rates we report here for the *PSEN1* and US sporadic EOSADD cases are considerably higher than the approximately 30% prevalences reported by Ringman et al [10]; this may be due to interim improvements in the immunohistochemical detection of pathological α -synuclein, and to the addition, in the NACC NP10 dataset, of the olfactory bulb-only LBD stage. In another, more recent study of 12 *PSEN1* USA cases [12], 6 (50%) had comorbid α -synuclein pathology.

As the decline in LBD prevalence with age in sporadic ADD closely parallels the decline in AD pathology severity with age [6], and in fact reaches very high prevalences in these *PSEN1* and EOSADD cases, this might suggest that LBD in this setting is largely dependent on AD pathology. The high reported rates of LBD in not only *PSEN1* ADD but also in other forms of autosomal dominant ADD [32-38], Down's syndrome and even in other, non-A β cerebral amyloidoses [39-42], supports this conclusion.

Comorbid TDP-43 proteinopathy has previously been investigated in human *PSEN1* ADD but not in *PSEN1* with the E280A mutation. Two separate USA studies, of a combined 42 cases, both reported limbic-restricted TDP-43 pathology in 17% of subjects with varying *PSEN1* genotypes [42, 43]. We found that TDP-43 pathology is common in *PSEN1* E280A Colombian cases, affecting 27% of subjects, with lesser but still definite rates of 16% and 11% in US *PSEN1* and EOSADD cases. Concurrent TDP-43 proteinopathy is also found in 14% of Down's syndrome cases [42], further directly implicating ADD pathology. Hippocampal sclerosis, however, which in older people is often associated with hippocampal TDP-43 pathology [44], was present in only 1 EOSADD case and none of the US or Colombian *PSEN1* cases and would therefore seem to require aging as a co-factor. It is likely that non-AD tauopathies, which were rare in these younger subjects, are not common or direct consequences of ADNC and probably are more age-related, as has been shown for sporadic ADD cases with a wide age range [6,7].

Significant large-vessel atherosclerosis, which is usually regarded as age-dependent, was present in a surprisingly large percentage of Colombian *PSEN1* cases, at almost 20% as compared to 0% and 3% of the

US *PSEN1* and sporadic US cases, respectively. The Colombian accentuation may be less related to the E280A mutation or ADD pathology than to the generally higher cardiovascular risk factors in Colombia [22] as compared to the USA. Small-vessel disease, or arteriolosclerosis, was more common than large vessel disease in all 3 of the studied groups, with ascending rates between 18% and 37% going from sporadic to Colombian to US *PSEN1* groups. Arteriolosclerosis has been reported to be unrelated to CAA in the *PSEN1* Colombian kindred [16]. Intriguingly, the most severe forms of arteriolosclerosis are found in CADASIL, an autosomal dominant condition [45] caused by Notch-3 mutations, while *PSEN1* acts enzymatically on all Notch genes [46].

The consequences of cerebrovascular disease include gross and microscopic infarcts, as well as gross and microscopic hemorrhages, but all were generally absent or rare in all groups. As mentioned, it might have been expected that microhemorrhages would be much more common in *PSEN1* ADD due to the high rates of CAA. However, these may be underestimated at autopsy as compared to MRI, or when special efforts are made in postmortem examination [47]. In the Alzheimer's Disease Neuroimaging Initiative, 25% of a mixed group of subjects, with normal cognition, mild cognitive impairment (MCI) and ADD, had microhemorrhages, with a tendency for proportionately more in ADD [48].

White matter rarefaction (WMR), conventionally regarded as a consequence of arteriolosclerosis, was remarkably common, at almost 60%, in the US *PSEN1* group, as compared to 18% in the EOSADD cases. We did not have postmortem estimates of WMR for the Colombian E280A group but previous MR reports [49, 50] found increased mean diffusivity and white matter hyperintensity volumes compared to healthy controls. White matter rarefaction (WMR) has long been noticed to be roughly twice as common in ADD as compared to normal elderly [51-54], and is reportedly pronounced relative to age-matched controls in subjects with *PSEN1* and APP mutations and Down's syndrome [55-58]. The greater WMR in the US *PSEN1* group studied here, relative to the EOSADD group, may be in part due to the higher proportion of significant arteriolosclerosis, 37% vs 18%, in the *PSEN1* group as compared to the EOSADD group but could also be due to the more severe AD histopathological scores as WMR probably has a dual vascular and AD-related pathogenesis [59-61]. Like LBD, WMR is highly prevalent in autosomal dominant ADD and thus this adds to other evidence that it is at least partially dependent on ADNC. Supporting a vascular contribution in this study both arteriolosclerosis and WMR rarefaction were most common in the NACC *PSEN1* cases.

Effective clinical trials depend on accurate estimates of required subject numbers, which largely depend on accurate assumptions for effect size and variability. Variability of clinical rates of decline will require larger subject numbers to overcome. Effect sizes depend on the efficacy of the therapeutic agent, which may be dependent on a matching of molecular mechanisms of agent and pathogenesis. Neuropathological comorbidities in ADD affect cognition but may have a completely different molecular pathogenesis and therefore might not respond to ADNC-specific therapeutics, leading to loss of efficacy, effect size and probability of clinical trial success. Stratifying ADD subjects by presence and types of accompanying comorbidities might result in increased observed effect sizes in some groups as compared to others, potentially “rescuing” failed clinical trials.

An important limitation of this study is the relatively small subject numbers, especially for the US *PSEN1* group. Some statistical associations may therefore be spurious or undetected.

This study is the first to compare rates of ADD comorbidities in the *PSEN1* Colombian E280A kindred with those in other *PSEN1* mutation types as well as with similarly-aged sporadic ADD cases. The results presented here, as well as other evidence, indicates that LBD, TDP-43 pathology and WMR, as common comorbidities with autosomal dominant and early-onset sporadic ADD, should be considered when planning clinical trials with such subjects as they may increase variability in response rates. However, they may be to some extent at least partially dependent on ADNC and thus potentially addressable by anti-amyloid or anti-tau therapies.

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