

# Laminar distribution of $\beta$ -amyloid ( $A\beta$ ) peptide deposits in the frontal lobe in familial and sporadic Alzheimer's disease

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## Abstract

To determine whether genetic factors influence frontal lobe degeneration in Alzheimer's disease (AD), the laminar distributions of diffuse, primitive, and classic  $\beta$ -amyloid ( $A\beta$ ) peptide deposits were compared in early-onset familial AD (EO-FAD) linked to mutations of the amyloid precursor protein (APP) or presenilin 1 (PSEN1) gene, late-onset familial AD (LO-FAD), and sporadic AD (SAD). The influence of apolipoprotein E (Apo E) genotype on laminar distribution was also studied. In the majority of FAD and SAD cases, maximum density of the diffuse and primitive  $A\beta$  deposits occurred in the upper cortical layers, whereas the distribution of the classic  $A\beta$  deposits was more variable, either occurring in the lower layers, or a double-peaked (bimodal) distribution was present, density peaks occurring in upper and lower layers. The cortical layer at which maximum density of  $A\beta$  deposits occurred and maximum density were similar in EO-FAD, LO-FAD and SAD. In addition, there were no significant differences in distributions in cases expressing Apo E  $\epsilon$ 4 alleles compared with cases expressing the  $\epsilon$ 2 or  $\epsilon$ 3 alleles. These results suggest that gene expression had relatively little effect on the laminar distribution of  $A\beta$  deposits in the frontal lobe of the AD cases studied. Hence, the pattern of frontal lobe degeneration in AD is similar regardless of whether it is associated with APP and PSEN1, mutation, allelic variation in Apo E, or with SAD.

**Key words:** laminar distribution,  $\beta$ -amyloid ( $A\beta$ ) peptide deposits, gene mutation.

## Introduction

The neuropathology of Alzheimer's disease (AD) is characterised by the formation of extracellular senile plaques (SP) and intracellular neurofibrillary tangles (NFT) [5,34]. The most important molecular constituent of the SP is  $\beta$ -amyloid ( $A\beta$ ) [28], an approximately 4 kDa peptide arising by constitutive cleavage of a trans-membrane amyloid precursor protein (APP). A variety of  $A\beta$  peptides are formed as a result of secretase cleavage of APP [40]. The most common of these peptides is  $A\beta$ 42, found largely in

discrete  $A\beta$  deposits, whereas the more soluble  $A\beta$ 40 is also found in association with blood vessels [37] and may develop later in the disease [21]. The discovery of  $A\beta$  led to the formulation of the 'amyloid cascade hypothesis' (ACH), one of the most important models of the molecular pathology of AD developed over the last 25 years [28]. Essentially, the ACH proposes that the deposition of  $A\beta$  is the initial pathological event in AD leading to the formation of NFT, cell death, and cortical degeneration [28].

At least four genetic loci are associated with AD: the APP gene on chromosome 21 [17,26], the prese-

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nilin *PSEN* genes on chromosome 14 (*PSEN1*) [47] and chromosome 1 (*PSEN2*) [32], and the apolipoprotein E (*Apo E*) gene on chromosome 19 [44]. *APP* and *PSEN* mutation may alter *APP* metabolism, resulting in increased deposition of A $\beta$  peptide, while allelic polymorphism of *Apo E*, and especially the expression of allele  $\epsilon$ 4, may influence the proportion of the more fibrillogenic A $\beta$ 42 formed in the tissue [32,47]. These genetic factors, however, may not explain the majority of AD cases [27]. Hence, early-onset cases linked to *APP* and *PSEN* mutations may account for less than 5% of total AD [29]. Additional susceptibility genes and environmental factors are therefore likely to be involved, especially in sporadic AD (SAD) [38,39,41]. In isolated Amish communities, for example, 24 markers have been linked to dementia [33] and several other linkage studies have shown the presence of AD susceptibility genes on chromosomes 9, 10, and 12 [48]. Hence, a small number of AD cases have been linked recently to the chromosome 9 open reading frame 72 (*C9ORF72*) gene [55].

Three morphological subtypes of A $\beta$  deposit are commonly observed in AD: 1) diffuse ('pre-amyloid') deposits, in which the A $\beta$  is not in a fibrillar form with a  $\beta$ -pleated conformation, dystrophic neurites (DN) and paired helical filaments (PHF) being largely absent, 2) primitive ('neuritic') deposits, in which the A $\beta$  is in a fibrillar form and is associated with DN and PHF, and 3) classic ('dense-cored') deposits, in which A $\beta$  is highly aggregated to form a central amyloid plaque 'core' surrounded by a 'ring' of DN [3,8,10-12,20]. In the cerebral cortex in AD, A $\beta$  deposits [2] and NFT [54] often exhibit significant variation in density across the cortex from pia mater to white matter, maximum density occurring within different layers [2,54]. The laminar distribution of A $\beta$  deposits may be a consequence of degeneration of neural pathways that have their neurons of origin or axon terminals located within particular layers [22]. The main objective of this study was to determine whether genetic factors influence the laminar distribution of A $\beta$  deposits and therefore result in a specific type of cortical degeneration in the frontal cortex in AD. Hence, laminar distributions of diffuse, primitive, and classic A $\beta$  deposits were studied in three groups of cases: 1) early-onset familial Alzheimer's disease (EO-FAD) linked to mutations of either amyloid precursor protein (*APP717*) or presenilin 1 (*PSEN1: G209V, E280A*) genes, 2) late-onset familial AD (LO-FAD), and 3) sporadic AD (SAD). In addition,

the influence of *Apo E* genotype on the distribution of A $\beta$  deposits was studied.

## Material and methods

### Cases

Alzheimer's disease cases ( $N = 20$ ; details in Table I) were obtained from the Brain Bank, Department of Neuropathology, Institute of Psychiatry, King's College, London, UK. Informed consent was given for the removal of all brain tissue according to the 1996 Declaration of Helsinki (as modified Edinburgh, 2000). Patients were clinically assessed and all fulfilled the 'National Institute of Neurological and Communicative Disorders and Stroke and the Alzheimer Disease and Related Disorders Association' (NINCDS/ADRDA) criteria for probable AD [52]. The histological diagnosis of AD was established by the presence of widespread neocortical SP consistent with the 'Consortium to Establish a Registry of Alzheimer Disease' (CERAD) criteria [34] and 'National Institute on Aging (NIA)-Reagan Institute' criteria [30,34]. The cases were divided into three groups: 1) EO-FAD (onset  $\leq 65$  years) ( $n = 4$ ), 2) LO-FAD ( $\geq 65$  years) ( $n = 6$ ), and 3) SAD ( $n = 10$ ) with no evidence of familial involvement.

### Tissue preparation

A block of the frontal cortex was taken at the level of the genu of the corpus callosum to study the superior frontal gyrus (SFG). Tissue was fixed in 10% phosphate-buffered formal saline and embedded in paraffin wax. Both immunostaining and thioflavin S have been used to stain SP in AD [14,45]. Thioflavin S staining indicates that amyloid in these plaques contains fibrillar material with a  $\beta$ -pleated sheet conformation [14]. By contrast, immunohistochemistry generally reveals more plaques including diffuse A $\beta$  deposits which are mainly thioflavin S-negative [14]. Hence, to label all types of plaque 7  $\mu$ m coronal sections were immunolabelled with a rabbit polyclonal antibody (Gift of Prof. B.H. Ader-ton, Institute of Psychiatry, King's College London) raised to the 12-28 amino acid sequence of the A $\beta$  protein and first used to identify A $\beta$  deposit subtypes in Down's syndrome (DS) [50] but which also effectively distinguishes the major types of A $\beta$  deposit in AD [4,6,7,10,50]. The antibody was used at a dilution of 1 in 1200 and the sections incubated at 4°C overnight. Sections were pretreated with

**Table I.** Demographic and genetic data of the Alzheimer's disease (AD) cases studied

Case	Sex	Age	Onset of disease	Group	Genetics	<i>Apo E</i>
1	M	65	–	EO-FAD	<i>APP717</i>	3/3
2	F	59	–	EO-FAD	<i>APP717</i>	3/3
3	F	61	–	EO-FAD	<i>PSEN1</i>	3/3
4	F	45	40	EO-FAD	<i>PSEN1</i>	2/3
5	F	72	66	LO-FAD	ND	2/3
6	F	86	80	LO-FAD	ND	3/4
7	F	77	72	LO-FAD	ND	3/3
8	F	79	68	LO-FAD	ND	3/4
9	F	85	76	LO-FAD	ND	3/3
10	F	89	–	LO-FAD	ND	3/3
11	M	80	77	SAD	–	3/3
12	F	87	82	SAD	–	3/4
13	F	64	59	SAD	–	4/4
14	F	91	83	SAD	–	3/4
15	M	73	66	SAD	–	2/3
16	F	82	75	SAD	–	ND
17	F	91	85	SAD	–	3/4
18	F	86	83	SAD	–	3/4
19	F	90	–	SAD	–	ND
20	M	82	78	SAD	–	3/3

*Apo E* – apolipoprotein E, *APP* – amyloid precursor protein, *EO-FAD* – early-onset familial Alzheimer's disease, *LO-FAD* – late-onset familial Alzheimer's disease, *SAD* – sporadic Alzheimer's disease, *PSEN1* – presenilin 1, *M* – male, *F* – female, *ND* – not determined

98% formic acid for 6 minutes, which enhances  $A\beta$  immunoreactivity.  $A\beta$  was visualised using the streptavidin-biotin horseradish peroxidase procedure with diaminobenzidine as the chromogen. Sections were also stained with haematoxylin.  $A\beta$  deposits were identified in the sections using criteria published by Delaere *et al.* [20]: 1) diffuse deposits were 10-200  $\mu$ m in diameter, lightly stained, irregular in shape, and with diffuse boundaries, 2) primitive deposits were 20-60  $\mu$ m, well demarcated, symmetrical in shape, and strongly stained, and (3) classic deposits were 20-100  $\mu$ m and had a distinct central amyloid core surrounded by a 'corona' of DN [20].

### Morphometric methods

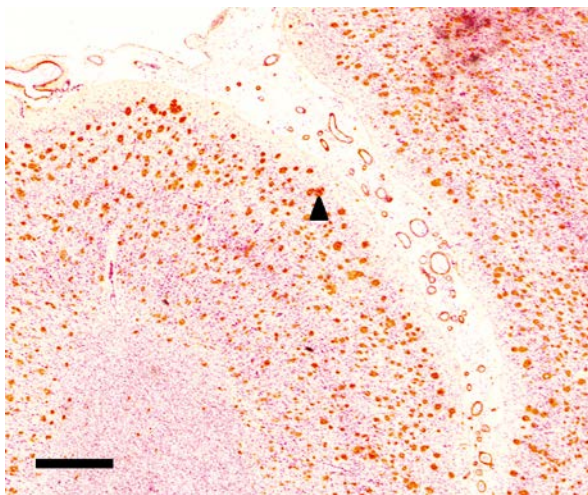
The distribution of the  $A\beta$  deposits in the SFG of each case was studied from the pia mater to white matter using methods described previously [23].

Five traverses from the pia mater to the edge of the white matter were located at random within each gyrus [9]. All deposits were then counted in 200 x 1000  $\mu$ m sample fields arranged contiguously, the larger dimension of the field parallel with the surface of the pia mater. An eye-piece micrometer was used as the sample field and was moved down each traverse one step at a time from the pia mater to the edge of the white matter. Histological features of the section were used to correctly position the field. The mean of the counts from the five traverses was calculated to study variations in density of histological features across each cortical gyrus.

### Data analysis

No attempt was made to locate precisely the boundaries between individual cortical layers. First, the degree of cortical degeneration present in many

gyri made laminar identification difficult. Second, identification was especially difficult in the frontal cortex because it exhibits a heterotypical structure, i.e., six layers cannot always be clearly identified and vary in prominence from case to case. Third, A $\beta$  deposits appeared to exhibit complex patterns of distribution across the cortex rather than being confined to specific layers. Hence, variations in density of A $\beta$  deposits with distance across the cortex were analysed using a polynomial curve-fitting procedure (STATISTICA software, StatSoft Inc., 2300 East 14<sup>th</sup> St, Tulsa, OK, 74104, USA) [2,49]. For each gyrus, polynomials of order 1, 2, 3 up to the 4<sup>th</sup> order were fitted successively to the data. Hence, second-order curves are parabolic, third-order curves are 'S' shaped, and fourth-order curves are double-peaked (bimodal). With each fitted polynomial, the correlation coefficients (Pearson's 'r'), regression coefficients, standard errors of the mean (SEM), values of *t*, and the residual mean square were obtained. At each stage, the reduction in the sums of squares (*SS*) was tested for significance. A polynomial was accepted as the best fit using the procedure described by Snedecor and Cochran [49], viz. when either a non-significant value of *F* was obtained or there was little gain in explained variance. The distributions of the A $\beta$  deposits across the cortex were



**Fig. 1.** Distribution of  $\beta$ -amyloid (A $\beta$ ) peptide deposits (arrowhead) in the superior frontal gyrus (SFG) in a case of early-onset familial Alzheimer's disease (EO-FAD) (*PSEN1* mutation). A $\beta$  deposits occur across the cortex in all layers but with a greater density of larger-sized deposits in the upper layers. A $\beta$  immunohistochemistry, haematoxylin, bar = 0.75 mm.

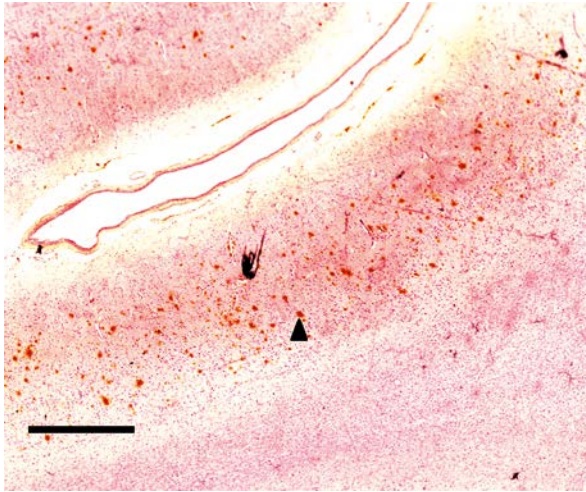
classified initially into three groups: 1) a single density peak was present (unimodal distribution), peak density being located in either upper or lower layers, 2) two density peaks were present (bimodal distribution), density peaks occurring in upper and lower layers, and 3) there was no significant change in density across the cortex, A $\beta$  deposits not being confined to particular layers. Bimodal distributions were then classified further according to whether the density peaks in the upper and lower layers were of similar or different magnitude. To study the effect of *Apo E* genotype, cases were classified into two groups: those not expressing allele  $\epsilon 4$ , i.e., genotypes  $\epsilon 2/3$  and  $\epsilon 3/3$ , and those expressing at least one allele  $\epsilon 4$ , i.e., genotypes  $\epsilon 3/4$  and  $\epsilon 4/4$ . In addition, the location of peak density and the maximum density of deposits were compared in EO-FAD, LO-FAD, and SAD. Hence, the point of maximum density (peak density) was identified for each deposit type for each gyrus while location of the peak was determined as the distance from the pia mater to that of the maximum density of A $\beta$  deposits, expressed as a percentage of the total distance from the pia mater to the edge of the white matter.

## Results

Examples of the distribution of A $\beta$  peptide deposits across the SFG are shown in Figures 1 and 2. In a case of EO-FAD (Fig. 1) (*PSEN1* mutation), A $\beta$  deposits occurred across the cortex with a greater density of larger deposits in the upper layers. By contrast, in a case of SAD (Fig. 2), A $\beta$  deposits occur largely in the lower layers.

Examples of the laminar distribution of the diffuse, primitive, and classic A $\beta$  deposits in the SFG of a single EO-FAD case (case 2, *APP* mutation) is shown in Figure 3. The distribution of the diffuse deposits was fitted by a first-order (linear) regression ( $r = 0.82$ ,  $p < 0.01$ ) consistent with greater densities of diffuse deposits in the upper layers and a linear decrease in density across the cortex from pia mater to white matter. The distribution of the primitive A $\beta$  deposits was fitted by a third-order polynomial ( $r = 0.91$ ,  $p < 0.001$ ) with a large density peak in the upper layers, while the classic deposits were also fitted by a third-order polynomial ( $r = 0.82$ ,  $p < 0.01$ ) with slightly higher densities adjacent to the pia mater and in the lower layers.

A comparison of the laminar distributions is shown in Table II. In FAD, diffuse A $\beta$  deposits exhib-

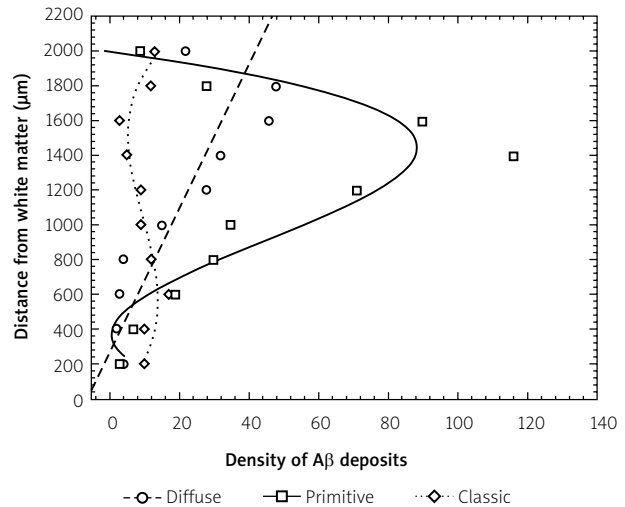


**Fig. 2.** Distribution of  $\beta$ -amyloid ( $A\beta$  deposits) (arrowhead) across the superior frontal gyrus (SFG) in a case of sporadic Alzheimer's disease (SAD).  $A\beta$  deposits occur largely in the lower layers.  $A\beta$  immunohistochemistry, haematoxylin, bar = 0.75 mm.

ited a density peak in the upper layers in 6/10 cases, and the primitive deposits did so in 9/10 cases. The distribution of the classic deposits was more variable; in 6/10 cases there was either a density peak in lower cortex or a bimodal distribution was present with density peaks in upper and lower layers. In SAD, diffuse and primitive  $A\beta$  deposits exhibited a density peak in the upper cortex in 7/10 cases and 9/10 cases respectively. Distribution of the classic  $A\beta$  deposits was more variable, a density peak in the lower layers or a bimodal distribution being present in 5/10 cases. The frequency of the various types of distribution of the diffuse ( $\chi^2 = 3.74$ ,  $p > 0.05$ ), primitive ( $\chi^2 = 0.71$ ,  $p > 0.05$ ), and classic ( $\chi^2 = 11.18$ ,  $p > 0.05$ )  $A\beta$  deposits was similar in EO-FAD, LO-SAD, and SAD.

Comparison of the mean location of maximum density and peak density of deposits among the three groups of cases is shown in Table III. Although there were significant differences in the layers at which peak density occurred among  $A\beta$  deposit subtypes ( $F = 4.44$ ,  $p < 0.01$ ), there were no significant differences among EO-FAD, LO-FAD, or SAD ( $F = 0.89$ ,  $p > 0.05$ ). In addition, there were no significant differences in peak density of  $A\beta$  deposits ( $F = 3.28$ ,  $p > 0.05$ ) among patient groups.

A comparison of the distributions exhibited by the  $A\beta$  deposits in cases classified according to *Apo E*



**Fig. 3.** Examples of the laminar distribution of the diffuse, primitive, and classic  $\beta$ -amyloid ( $A\beta$  deposits) in frontal lobe of a case of early-onset familial Alzheimer's disease (EO-FAD) (case 2, amyloid precursor protein (*APP*) gene mutation).

genotype groups is shown in Table IV. There were no significant differences in distribution of diffuse ( $\chi^2 = 2.55$ ,  $p < 0.05$ ), primitive ( $\chi^2 = 0.003$ ,  $p < 0.05$ ), or classic ( $\chi^2 = 3.41$ ,  $p < 0.05$ )  $A\beta$  deposits in cases expressing *Apo E* genotypes  $\epsilon 2/3$  and  $\epsilon 3/3$ , compared with those expressing genotypes  $\epsilon 3/4$  and  $\epsilon 4/4$ .

## Discussion

The objective of this study was to determine whether genetic factors were associated with a specific pattern of cortical degeneration, as revealed by the deposition of  $A\beta$  deposits in the frontal cortex in AD. The data confirm the need for quantitative assessment of  $A\beta$  deposition in different layers of cortex as deposits often occur over many layers with variation in abundance across the cortex. This study demonstrated: 1) laminar distributions of diffuse, primitive, and classic  $A\beta$  deposit subtypes were essentially similar in EO-FAD, LO-FAD, and SAD, 2) within FAD, laminar distributions were similar in *APP/PSEN1* cases compared with LO-FAD, and 3) laminar distributions were similar in cases expressing *Apo E*  $\epsilon 4$  alleles compared with cases expressing  $\epsilon 2$  or  $\epsilon 3$  alleles.

The data suggest no significant differences in  $A\beta$  deposit density in EO-FAD, LO-FAD, and SAD or when cases were classified according to *Apo E* genotype. Previous quantitative studies comparing SP or  $A\beta$

**Table II.** Comparison of the frequencies of different types of laminar distribution of  $\beta$ -amyloid (A $\beta$ ) deposits in superior frontal gyrus (SFG) in three groups of cases: early-onset familial Alzheimer's disease (EO-FAD), late-onset familial AD (LO-FAD), and sporadic AD (SAD). Data are the number of cases in which a particular type of laminar distribution is present

Group	Deposit type	Type of laminar distribution			
		UL	LL	B	NS
EO-FAD	Diffuse	3	0	1	0
	Primitive	4	0	0	0
	Classic	2	0	2	0
LO-FAD	Diffuse	3	1	1	1
	Primitive	5	0	0	1
	Classic	0	4	0	2
SAD	Diffuse	7	1	0	2
	Primitive	9	0	0	1
	Classic	2	4	1	3

Comparison of frequencies between groups:  $\chi^2$  contingency tables: diffuse deposits  $\chi^2 = 5.09$  (6 DF,  $p > 0.05$ ), primitive deposits  $\chi^2 = 0.74$  (2 DF,  $p > 0.05$ ), classic deposits  $\chi^2 = 11.26$  (6 DF,  $p > 0.05$ )

UL – maximum A $\beta$  deposit density occurred in upper cortical layers, LL – maximum A $\beta$  deposit density occurred in lower cortical layers, B – bimodal distribution with peaks of density in upper and lower layers, NS – no significant change in density of A $\beta$  deposits across the cortex

**Table III.** Comparison of mean peak location, i.e., percentage distance from the pia mater at which maximum density of  $\beta$ -amyloid (A $\beta$ ) deposits occurred expressed as a percentage of the width of the gray matter, and mean peak density, i.e., the actual density of deposits at the peak (mm<sup>2</sup>) in the superior frontal gyrus (SFG) in three groups of cases: early-onset familial Alzheimer's disease (EO-FAD), late-onset familial AD (LO-FAD), and sporadic AD (SAD). Standard errors of the mean (SEM) are in parentheses

Group		A $\beta$ deposit subtype		
		Diffuse	Primitive	Classic
EO-FAD	Peak location	32.5 (12.11)	21.3 (5.98)	40.0 (12.54)
	Peak density	36.5 (6.39)	86.5 (21.85)	12.3 (2.19)
LO-FAD	Peak location	39.5 (9.89)	31.7 (4.88)	51.7 (10.24)
	Peak density	28.0 (5.22)	70.3 (17.84)	8.17 (1.79)
SAD	Peak location	32.0 (9.66)	27.20 (3.69)	55.10 (6.09)
	Peak density	20.8 (3.20)	38.3 (10.76)	10.3 (1.50)

Analysis of variance (ANOVA): 1) Peak location: patient group  $F = 0.89$  ( $p > 0.05$ ), deposit type  $F = 4.44$  ( $p < 0.01$ ), interaction  $F = 0.22$  ( $p > 0.05$ ); 2) Peak density: patient group  $F = 3.28$  ( $p > 0.05$ ), deposit type  $F = 18.73$  ( $p < 0.001$ ), interaction  $F = 1.36$  ( $p > 0.05$ )

deposit abundance in FAD and SAD have been controversial [16,18,24,36]. Hence, no significant differences in severity scores of SP were observed in FAD and SAD [36], and A $\beta$  'load' in the frontal cortex and temporal isocortex was similar in SAD and FAD cases linked to the *APP*<sub>717</sub> mutation [16]. Nevertheless, cultured cells expressing a double mutation in *APP* produced six times more A $\beta$  than normal cells [18]. In addition, other studies have reported increased

amyloid deposition in individuals expressing allele  $\epsilon 4$  [24]. However, it is possible that A $\beta$  deposition could be more widely distributed across the cortical layers of the SFG in FAD, but with similar peak densities.

In the SFG of both SAD and FAD, maximum density of the diffuse and primitive A $\beta$  deposits occurred most frequently in the upper layers. By contrast, the distribution of the classic deposits was more variable, peak densities occurring either in the lower lay-

**Table IV.** Comparison of frequencies of different types of laminar distribution of A $\beta$  deposits in superior frontal gyrus (SFG) of Alzheimer's disease (AD) cases divided into two groups according to apolipoprotein (Apo E) genotype, i.e., those expressing genotypes  $\epsilon$ 2/3 and  $\epsilon$ 3/3 compared with those expressing genotypes  $\epsilon$ 3/4 and  $\epsilon$ 4/4. Data are the number of cases in which a particular type of laminar distribution is present

Group	A $\beta$ deposit subtype	Laminar distribution			
		UL	LL	B	NS
$\epsilon$ 2/3, $\epsilon$ 3/3	Diffuse	8	0	2	1
	Primitive	10	0	0	1
	Classic	3	4	2	2
$\epsilon$ 3/4, $\epsilon$ 4/4	Diffuse	7	0	0	0
	Primitive	7	0	0	0
	Classic	1	5	0	1

Chi-square ( $\chi^2$ ) contingency tables: diffuse deposits  $\chi^2 = 2.29$  (2 DF,  $p > 0.05$ ), primitive deposits  $\chi^2 = 0.006$  (1 DF,  $p > 0.05$ ), classic deposits  $\chi^2 = 2.69$  (3 DF,  $p > 0.05$ )  
 UL – maximum A $\beta$  deposit density occurred in upper cortical layers, LL – maximum A $\beta$  deposit density occurred in lower cortical layers, B – bimodal distribution with peaks of density in upper and lower layers, NS – no significant change in density of A $\beta$  deposits across the cortex

ers, or in both upper and lower layers. Similar results have been reported in studies of the laminar distribution of SP [15,19], Apo E-immunoreactive SP [53], neuritic plaques (NP) [42], and A $\beta$  deposits in AD [2], which are frequently abundant in layers II and III. In addition, in a transgenic mouse model expressing the *APP*<sub>717</sub> mutation, A $\beta$  deposits were most abundant in layers II and III, similar to AD [51]. However, aged dogs often show a different distribution of A $\beta$  deposits to humans, being usually abundant in the deep cortical layers but with evidence of spread to superficial cortical layers with increasing age [43].

Various hypotheses could explain the laminar distribution of A $\beta$  deposits in the SFG in AD. First, mRNA of APP is preferentially expressed by the large pyramidal neurons in layers III and V [13]. Degeneration of these neurons could then result in increased secretion of APP and formation of A $\beta$  deposits within these layers [4]. Second, interleukin-immunoreactive microglia (IL-Mg) have a similar laminar distribution as APP-immunoreactive NP [46]. Hence, the laminar distribution of microglia could be a factor determining the distribution of the A $\beta$  deposits. Third, the laminar distribution of the classic deposits could be spatially related to blood vessels [2,37]. Large blood vessels often exhibit a bimodal distribution in the cortex, whereas smaller capillaries occur at maximum density in the deeper layers [2]. In addition, Akiyama *et al.* [1] found that A $\beta$  deposits accumulated vertically in columns, with blood vessels often occurring perpendicular to the column and pene-

trating its centre. Previous studies suggest, however, that although classic A $\beta$  deposits are clustered around blood vessels in SAD [6], there are fewer spatial associations with blood vessels in FAD [7].

Laminar distributions of A $\beta$  deposits in frontal lobe AD are essentially similar in the FAD and SAD cases examined and similar whether Apo E allele  $\epsilon$ 4 was present or not [31]. In addition, among FAD cases, there was no evidence that a specific type of laminar distribution was influenced by genetic subtype. Hence, neither *APP/PSEN1* mutations nor the presence of Apo E allele  $\epsilon$ 4 uniquely determines A $\beta$  deposition and therefore the pattern of frontal lobe degeneration in AD. Uchihara *et al.* [53] found that Apo E labelled a subset of deposits in lamina III with more Apo E-immunoreactive diffuse deposits in the deeper layers. However, only a proportion of the diffuse deposits were Apo E-immunoreactive, suggesting that Apo E was not involved in the process of cortical degeneration but immunoreactivity was acquired by certain deposits later in the disease. Hence, pathological changes initiated by the various genetic changes in FAD and, by other causes in SAD, appear to follow a parallel course resulting in very similar patterns of cortical degeneration in the SFG.

In conclusion, there were no essential differences in the laminar distribution of the A $\beta$  deposits in the SFG between FAD and SAD, or between different subtypes of FAD. Hence, *APP* and *PSEN1* mutations and the presence of Apo E genotype  $\epsilon$ 4 appear to have little influence on laminar distribution. Although the

mechanism of generating fibrillogenic species of A $\beta$  may differ among disease subtypes, gene expression appears to have little effect on the pattern of degeneration of the frontal lobe in AD.

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## Disclosure

Author reports no conflict of interest.

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