

# Plastic Neuronal Remodeling Is Impaired in Patients with Alzheimer's Disease Carrying Apolipoprotein $\epsilon$ 4 Allele

Thomas Arendt,<sup>1</sup> Cornelia Schindler,<sup>2</sup> Martina K. Brückner,<sup>1</sup> Klaus Eschrich,<sup>2</sup> Volker Bigl,<sup>1</sup> Dyrk Zedlick,<sup>3</sup> and Lena Marcova<sup>4</sup>

<sup>1</sup>Department of Neurochemistry, Paul Flechsig Institute of Brain Research, Leipzig, Germany, <sup>2</sup>Institute of Biochemistry and <sup>3</sup>Hospital for Psychiatry, University of Leipzig, Leipzig, Germany, and <sup>4</sup>Institute of Brain Research, Medical Academy of Russia, Moscow, Russia

A relationship between the apolipoprotein E (apoE) genotype and the risk to develop Alzheimer's disease has been established recently. Apolipoprotein synthesis is implicated in developmental processes and in neuronal repair of the adult nervous system.

In the present study, we investigated the influence of the apolipoprotein polymorphism on the severity of neuronal degeneration and the extent of plastic dendritic remodeling in Alzheimer's disease. Changes in length and arborization of dendrites of Golgi-impregnated neurons in the basal nucleus of Meynert, locus coeruleus, raphe magnus nucleus, medial amygdaloid nucleus, pedunculopontine tegmental nucleus, and substantia nigra were analyzed after three-dimensional reconstruction. Patients with either one or two apoE  $\epsilon$ 4 alleles not only showed a more severe degeneration in all areas investigated than in patients lacking the apoE 4 allele but also re-

vealed significantly less plastic dendritic changes. ApoE  $\epsilon$ 4 allele copy number, furthermore, had a significant effect on the pattern of dendritic arborization. Moreover, the relationship between the intensity of dendritic growth and both the extent of neuronal degeneration and the stage of the disease seen in patients lacking the apoE  $\epsilon$ 4 allele was very weak in the presence of one  $\epsilon$ 4 allele and completely lost in patients homozygous for the  $\epsilon$ 4 allele.

The results provide direct evidence that neuronal reorganization is affected severely in patients with Alzheimer's disease carrying the apoE  $\epsilon$ 4 allele. This impairment of neuronal repair might lead to a more rapid functional decompensation, thereby contributing to an earlier onset and more rapid progression of the disease.

**Key words:** Alzheimer's disease; apolipoprotein E; degeneration; dendritic sprouting; plasticity; regeneration

In the adult CNS affected by age- or disease-related degeneration, mechanisms of compensation and repair are activated in an attempt to counteract functional sequelae of neuronal loss (Scheff et al., 1990; Lapchak et al., 1991; Fritschy and Grzanna, 1992; Lippa et al., 1992; Ramirez and Ulfhake, 1992; Arendt et al., 1995d,e). As a consequence, degenerative events become manifest as a disorder only after exceeding a critical threshold, thereby exhausting the capacity of compensation (Marsden, 1981; Arendt and Bigl, 1987). Under certain conditions, however, these compensational processes might be maladaptive and eventually even contribute to the development and progression of the disease. This seems to be the situation in Alzheimer's disease (AD) (Geddes et al., 1985; Arendt et al., 1986; Butcher and Woolf, 1989), where an aberrant dendritic growth is observed (Scheibel and Tomiyasu, 1978; Ferrer et al., 1983; Arendt et al., 1986, 1991, 1995b,c; Ihara et al., 1988; McKee et al., 1989) instead of the regular dendritic growth seen during aging and in a variety of related degenerative disorders (Buell and Coleman, 1979; Graveland et al., 1985; Coleman and Flood, 1987; Arendt et al., 1995b,c).

The activation of processes involved in the reorganization of membrane components is a major requirement for restructuring the dendritic organization. In the brain, apolipoprotein E (apoE) is the major lipoprotein involved in lipid transport and metabo-

lism (Elshourbagy et al., 1985; Mahley, 1988; Kinoshita et al., 1993; Poirier et al., 1993a). ApoE synthesis is implicated in neuronal growth and in repair after injury of both peripheral and central neurons (Ignatius et al., 1986; Snipes et al., 1986; Poirier et al., 1991a,b).

There are three major isoforms of apoE (E2, E3, and E4), which are products of the three alleles  $\epsilon$ 2,  $\epsilon$ 3, and  $\epsilon$ 4 located at a single gene locus (Mahley, 1988). It has been shown recently that the  $\epsilon$ 4 allele is a major risk factor for late-onset familial AD and probably also for sporadic AD (Corder et al., 1993; Mayeux et al., 1993; Poirier et al., 1993b; Rebeck et al., 1993; Saunders et al., 1993; Strittmatter et al., 1993a). Moreover, AD patients with  $\epsilon$ 4 alleles usually show an earlier age of onset (Corder et al., 1993; Basun et al., 1995; Locke et al., 1995) and a more rapid progression of the disease (Bennett et al., 1995), suggesting a causal link between apoE polymorphism and the development of AD (for review, see Roses et al., 1994).

ApoE is detectable immunohistochemically in senile plaques, neurofibrillary tangles (NFT), and cerebrovascular amyloid in AD (Namba et al., 1991; Wisniewski and Frangione, 1992; Rebeck et al., 1993; Strittmatter et al., 1993a,b; Kida et al., 1994; Benzing and Mufson, 1995). An enhanced burden of both  $\beta$ /A4-amyloid (Schmechel et al., 1993; Strittmatter et al., 1993a,b; Czech et al., 1994; Gearing et al., 1995; Nagy et al., 1995; Ohm et al., 1995; Oyama et al., 1995) and NFT (Nagy et al., 1995; Ohm et al., 1995) was gene dose dependently seen in apoE  $\epsilon$ 4-carriers. Although isoform-specific interactions of apoE with both  $\beta$ /A4-amyloid (Strittmatter et al., 1993a,b; Wisniewski et al., 1993) and tau (Cotton et al., 1994; Strittmatter et al., 1994a,b; Whitson et al., 1994) have been reported, the

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Correspondence should be addressed to Dr. Thomas Arendt, Paul Flechsig Institute, Jahnallee 59, 04109 Leipzig, Germany.

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Table 1. Profile of cases

	Controls	Alzheimer's disease/apolipoprotein E genotype		
		ε3/ε3	ε3/ε4	ε4/ε4
Number of cases	20	34	19	11
Sex (male/female)	9/11	15/19	8/11	5/6
Age in years (±SD)	81 ± 8.1	85 ± 6.6	76 ± 7.3	77 ± 7.9
FAST				
5		6	4	2
6b		5	2	1
6d		6	4	2
6e		5	3	2
7a		5	4	2
7b		7	2	2
PMSI (±SD)	2.2 ± 0.4	2.6 ± 0.7	2.7 ± 0.6	2.7 ± 0.5
Cause of death				
Bronchopneumonia	5	30	17	9
Myocardial infarction	11	—	—	2
Embolism of lung	2	1	—	—
Sepsis	2	3	2	—
Postmortem delay in hr (±SD)	2.8 ± 0.3	3.3 ± 0.5	3.1 ± 0.4	3.0 ± 0.7
Fresh brain weight in gm (±SD)	1282 ± 74	1210 ± 53	1208 ± 64	1164 ± 86

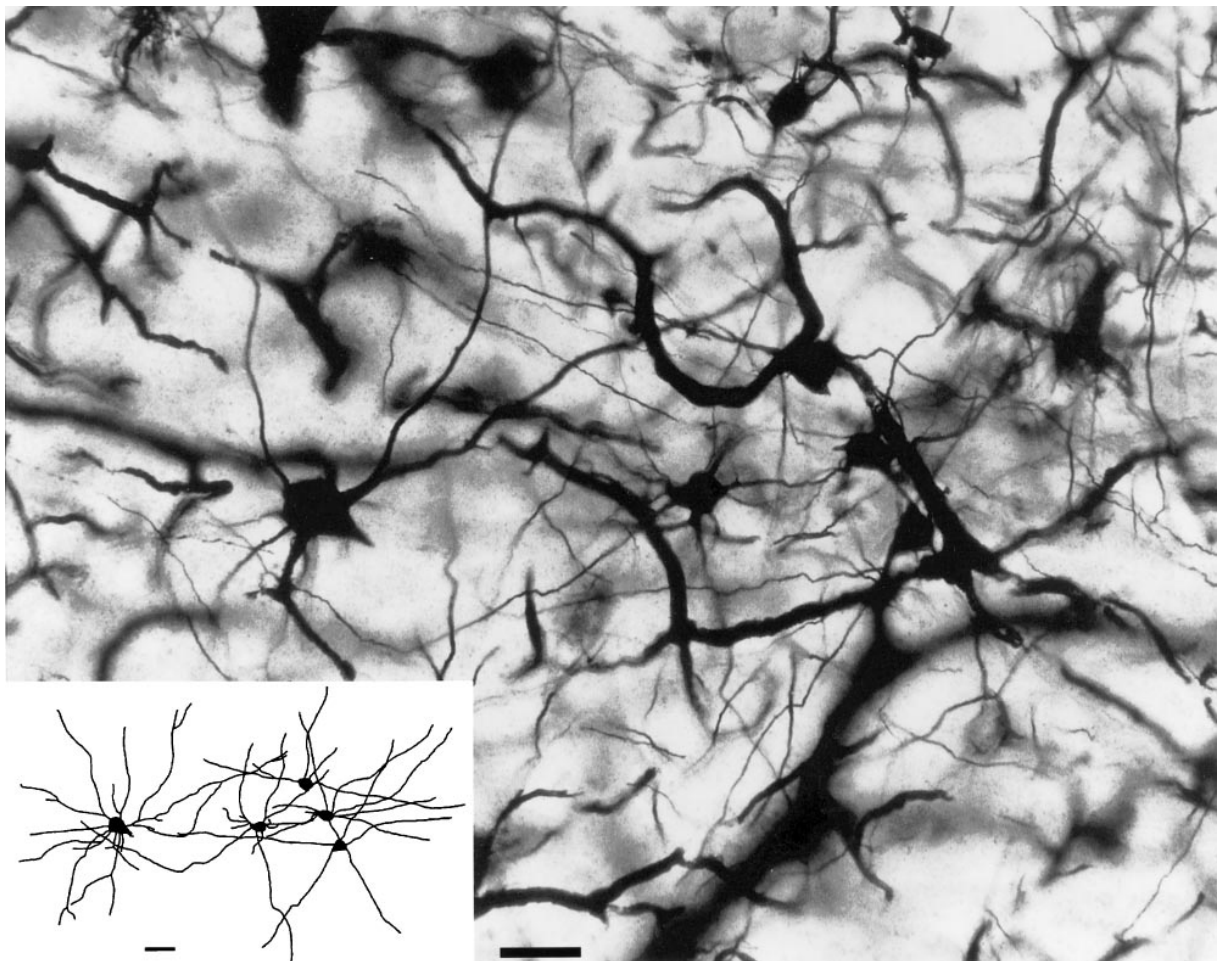
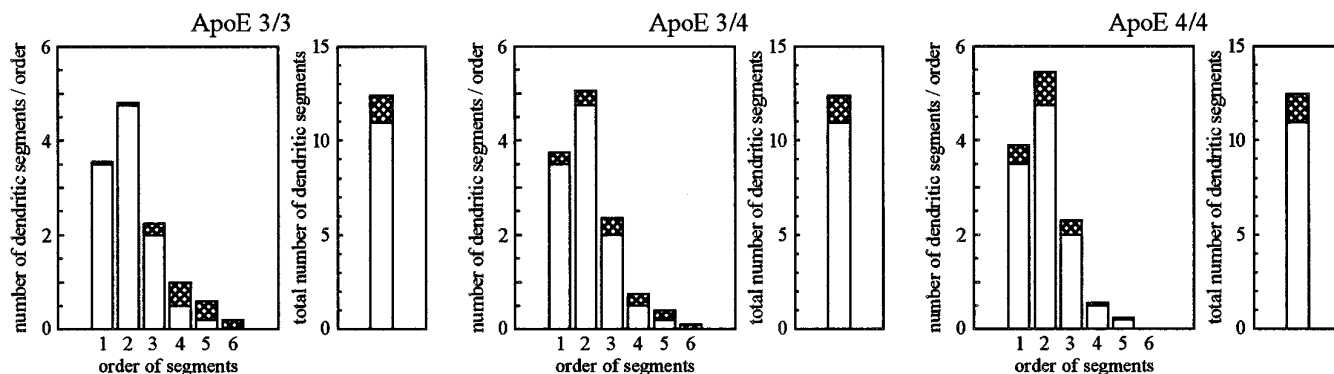
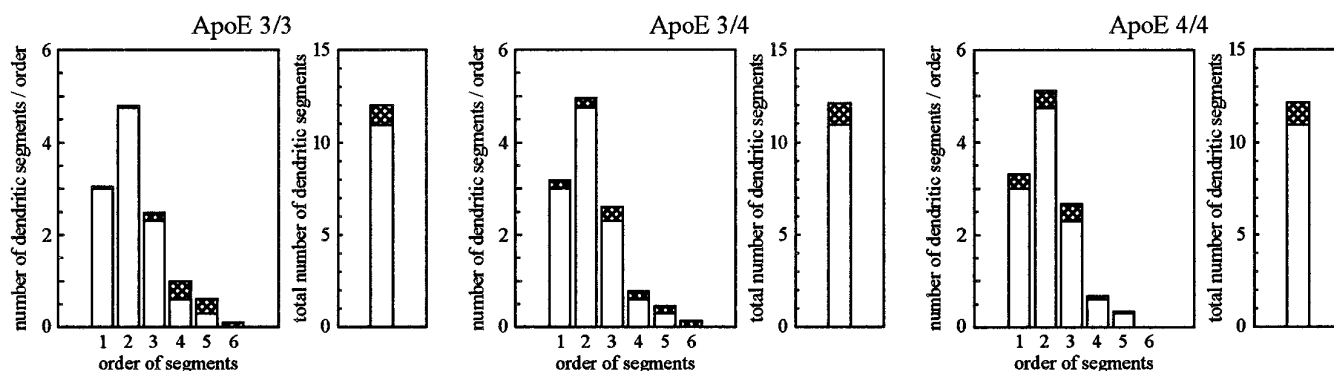


Figure 1. Cluster of Golgi-impregnated reticular neurons in the basal nucleus of Meynert in a patient with AD. Inset, Two-dimensional projection of the three-dimensional image reconstructed from serial sections. Scale bar, 50 μm.

## Nucleus basalis of Meynert



## Locus coeruleus



## Raphé magnus nucleus

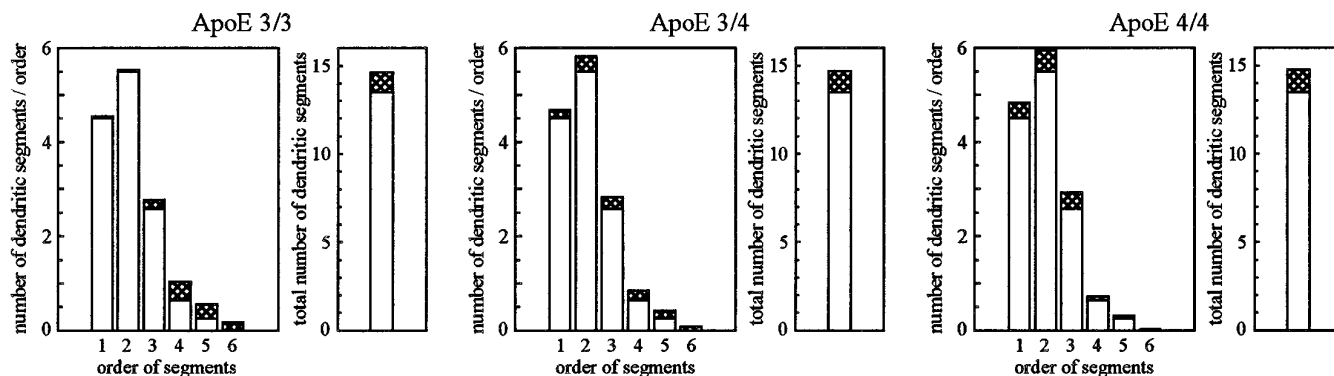


Figure 2. Changes in the total number of dendritic segments and in the frequency distribution of segments obtained for different orders on Golgi-impregnated reticular neurons in different subgroups of patients with AD, as compared with controls (centrifugal system of ordering, i.e., primary segments correspond to order 1; most distal branches correspond to highest segment order). Open columns, Control values; open plus cross-hatched columns, AD patients. Data are mean values. For each case, 300 neurons were analyzed. For group size, see Table 1; for summary of statistical analysis, compare Table 3. Figure 2 continues.

mechanism behind the pathological effects of the apoE polymorphism remains unknown.

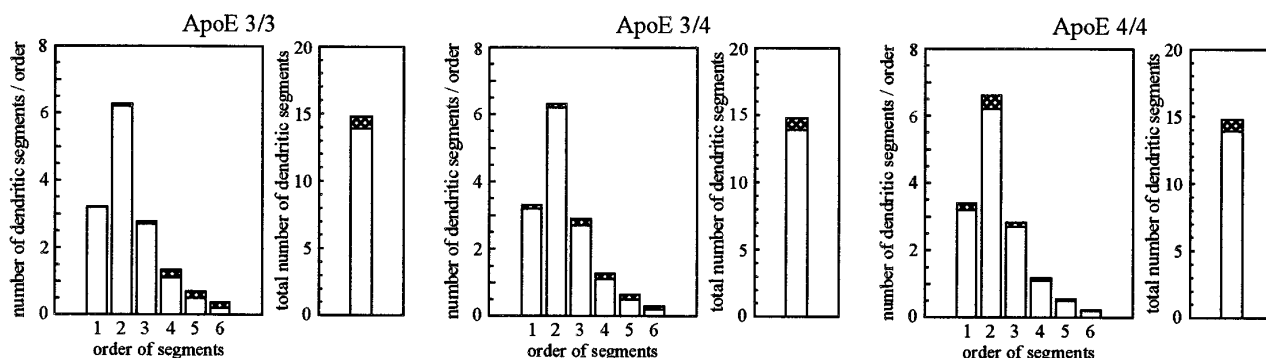
In the present study, we provide direct evidence that dendritic remodeling, which occurs in AD on subcortical neurons and which is likely to represent a response to neurodegenerative changes, is impaired severely in patients carrying the apoE  $\epsilon 4$  allele. This deficiency might be regarded as a failure of plastic neuronal response that could

contribute to a more rapid decompensation, resulting in an earlier onset and a more rapid progression of the disease.

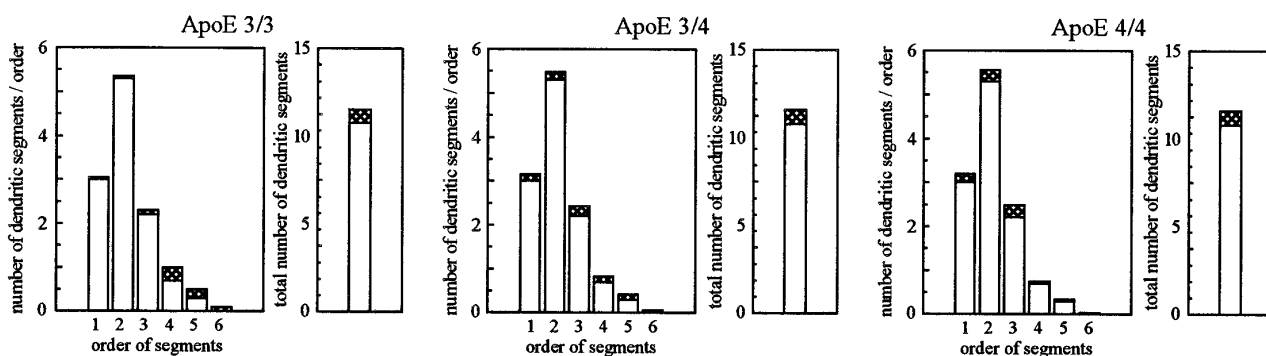
## MATERIALS AND METHODS

**Cases.** Brains were obtained from 64 patients with AD and from 20 age-matched control patients. The profile of cases is summarized in Table 1. The clinical diagnosis of AD was based on the occurrence of significant

## Medial amygdaloid nucleus



## Pedunclopontine tegmental nucleus



## Substantia nigra

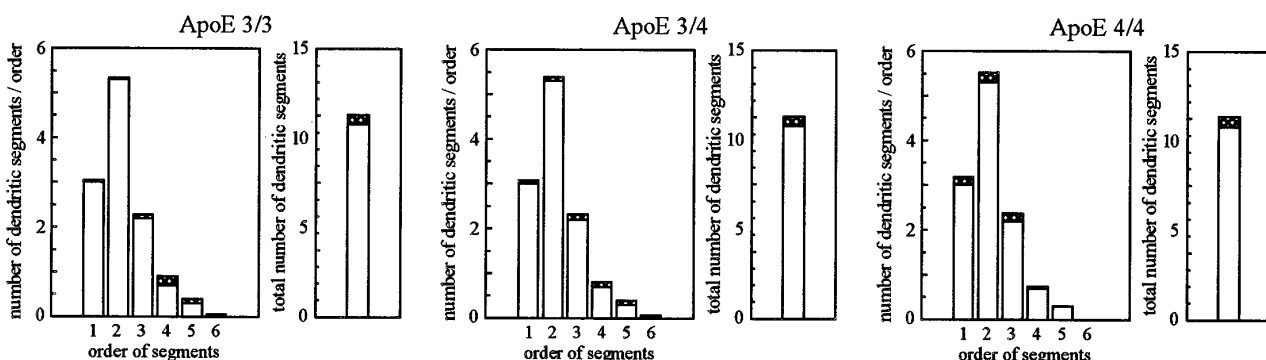


Figure 2 continued.

intellectual dysfunction, i.e., the presence of deficits in at least four aspects of cognitive and social behavior. Other causes of dementia were excluded by medical, psychiatric, and paraclinical examination (American Psychiatric Association, 1987). Each case met the National Institute of Neurological and Communicative Disorders and Stroke (NINCDS) and Alzheimer's Disease and Related Disorders Association (ADRDA) criteria for definite diagnosis of Alzheimer's disease (McKhann et al., 1984), based on the presence of NFTs and neuritic plaques observed in the hippocampal formation and neocortical areas, as recommended (Khataturian, 1985). Clinical severity of the disease was assessed according to the Functional Assessment Stages of Alzheimer's disease (FAST) according to Reisberg et al. (1982). Rating of major stages (5, 6, 7) and its substages (a, b, c, etc.) was based on a regular clinical evaluation during the last month of life with the final examination performed within <2 weeks before death. Because the distinctions between FAST substages are less marked than between major stages, two scales, one including substages and the other consisting only of major stages, were used to

analyze the influence of the clinical stage of AD on both neuronal degeneration and dendritic plasticity. No major differences, however, were obtained using either one of the scales. Therefore, only data obtained with the full scale, including substages, are reported. To minimize the likelihood of an artificial influence by premortem hypoxia and hypovolemia, we matched cases of AD and controls with respect to the Premortem Severity Index (PMSI) of Monfort et al. (1985). Brains used as normal controls were obtained at routine autopsy from patients dying without a history of neuropsychiatric disorder or mental impairment. There had to be clear evidence that the patient was alert, well oriented, and capable of functioning relatively independently shortly before death. No pathological signs were detected by neuropathological examination. None of the control cases carried an apoE  $\epsilon$ 4 allele. The entire procedure of case recruitment, acquisition of the patients' personal data, performing the autopsy, and handling the autopsic material has been approved by the Ethical Committee of Leipzig University.

**Tissue preparation.** Samples of the cerebral cortex were dissected bilat-

**Table 2. Effects of the ApoE genotype on the loss of subcortical neurons and on dendritic growth in AD**

	Control	Alzheimer's disease		
		ApoE 3/3	ApoE 3/4	ApoE 4/4
Basal nucleus of Meynert				
Dendritic length (mm)	2.612 ± 0.049	3.600 ± 0.047***	3.353 ± 0.045***##	3.275 ± 0.036***###
Neuronal density (%)	100	36.3 ± 3.8***	26.6 ± 3.9***	13.4 ± 2.9***##
Locus coeruleus				
Dendritic length (mm)	2.230 ± 0.041	3.021 ± 0.031***	2.838 ± 0.027***###	2.732 ± 0.019***###
Neuronal density (%)	100	49.9 ± 3.5***	33.1 ± 5.3***##	14.2 ± 5.0***###
Raphe magnus nucleus				
Dendritic length (mm)	2.520 ± 0.026	2.982 ± 0.021***	2.922 ± 0.027***	2.855 ± 0.029***###
Neuronal density (%)	100	70.3 ± 2.2***	62.5 ± 8.5***	49.1 ± 3.7***###
Medial amygdaloid nucleus				
Dendritic length (mm)	2.342 ± 0.026	2.776 ± 0.022***	2.730 ± 0.028***	2.654 ± 0.030***##
Neuronal density (%)	100	75.1 ± 2.1***	66.3 ± 2.8***#	44.1 ± 3.7***###
Pedunculopontine tegmental nucleus				
Dendritic length (mm)	2.410 ± 0.024	2.867 ± 0.020***	2.805 ± 0.032***	2.733 ± 0.025***##
Neuronal density (%)	100	77.7 ± 3.8***	58.8 ± 3.4***##	35.4 ± 5.1***###
Substantia nigra				
Dendritic length (mm)	2.155 ± 0.037	2.389 ± 0.030***	2.299 ± 0.027**	2.256 ± 0.008#
Neuronal density (%)	100	87.8 ± 1.3***	77.3 ± 2.7***###	62.6 ± 4.8***###

Data are mean ± SEM. Significantly different from control: \*\*\* $p$  < 0.001; \*\* $p$  < 0.01 (Student's  $t$  test). Significantly different from AD subgroup ApoE 3/3: ### $p$  < 0.001; ## $p$  < 0.01; # $p$  < 0.05 (Student's  $t$  test). For group size, see Table 1.

erally from Brodmann areas 8 and 22. White matter was removed carefully, and specimens were snap-frozen and stored at  $-80^{\circ}\text{C}$ . To prevent artifacts by postmortem delay (Williams et al., 1978; Buell, 1982; De Ruiter, 1983), we immersed the remaining parts of the brains in 4% formaldehyde/0.1 M sodium phosphate buffer adjusted to pH 7.2 within <3.5 hr postmortem. After fixation for ~1 month, cerebellum and brainstem were removed, and brains were cut in the coronal plane into slabs of ~5 mm thickness. One hemisphere was processed further for Golgi impregnation, the other hemisphere for immunohistochemical techniques and Nissl staining.

**ApoE genotyping.** Genomic DNA was isolated by treating 15–25 mg of frozen brain tissue with proteinase K (Sambrook et al., 1989). The digest was deproteinized by phenol/chloroform/isoamyl alcohol extractions, recovered by ethanol precipitation, dried, and resuspended in buffer. A 227 bp sequence of the apoE gene was amplified by PCR using primers and reaction conditions as described (Wenham et al., 1991). Genotyping was performed after digestion with the restriction enzyme *Cfo*I (Wenham et al., 1991) and separation of fragments on a 10% polyacrylamide gel (Hixson and Vernier, 1990).

**Golgi impregnation.** Tissue slabs 5 mm thick were trimmed to blocks varying in size between 3 and 15 cm<sup>2</sup>. Blocks were incubated at room temperature for 8–10 d in a solution of 5% formaldehyde, 3% potassium dichromate, and 12.5% sucrose. Solution was freshly prepared every day. Then tissue was rinsed and kept for a further 3–6 d in 0.75% silver nitrate at room temperature in the dark. After dehydration and celloidin embedding, serial sections 240  $\mu\text{m}$  thick were cut in the coronal plane. Blocks were oriented carefully before cutting to ensure that sectioning was parallel to the frontal reference plane. Sections were mounted between two coverslips to be observable from both sides.

**Histochemistry.** Tissue blocks containing the basal nucleus of Meynert of one hemisphere were immersed in 30% sucrose for cryoprotection and cut in the coronal plane on a freezing microtome at a thickness of 30  $\mu\text{m}$ . Sections were treated for 15 min with 0.3% H<sub>2</sub>O<sub>2</sub> in methanol to destroy endogenous peroxidase and preincubated in 0.3% nonfat dried milk and 0.1% gelatin in 0.1 M PBS for blocking. Free-floating sections were incubated at room temperature overnight in one of the following primary antibodies in 0.1 M PBS also containing 0.01% Triton X-100: rat monoclonal anti-choline acetyltransferase (ChAT; Boehringer Mannheim, Mannheim, Germany; 1:200) or mouse monoclonal anti-NGF-receptor p75 (p75<sup>NGFR</sup>; clone ME 20.4, 1:200). Immunoreaction was detected with peroxidase-labeled anti-rat Ig (Boehringer Mannheim; 1:500) or biotinylated sheep anti-mouse Ig (Amersham, Buckinghamshire, UK; 1:300), the ExtrAvidin-peroxidase complex (Sigma-Aldrich GmbH, Deisen-

hofen, Germany; 1:300) and 0.04% 3,3'-diaminobenzidine/0.015% H<sub>2</sub>O<sub>2</sub> in 0.1 M PBS. For intensification, selected sections were, instead, treated with 0.02% 3,3'-diaminobenzidine, 0.015% H<sub>2</sub>O<sub>2</sub>, and 0.4% nickel ammonium sulfate in 0.5 M Tris-HCl, pH 8. Sections were rinsed, mounted on chrome-alum-coated slides, dehydrated, and coverslipped with Entellan (Merck, Darmstadt, Germany).

**Quantitative analysis of Golgi-impregnated neurons.** A three-dimensional morphometric analysis of Golgi-impregnated reticular neurons in the basal nucleus of Meynert, locus coeruleus, raphe magnus nucleus, medial amygdaloid nucleus, pedunculopontine tegmental nucleus, and substantia nigra was performed as described (Arendt et al., 1995a,c). The morphological delineation of the basal nucleus was based on a detailed cytological and histochemical study (Fischer and Fischer, 1987). Reconstruction of neurons and quantitative measurements were made in three dimensions with reference to a defined Cartesian system of axis by digitizing the neuron directly from serial sections with help of a semiautomatic image analyzing system (Orthoplan-3D, Leitz Wetzlar and analySIS, Münster, Germany). From the digitized image, the total dendritic length per neuron and the number of dendritic segments of each order using the centrifugal ordering system were calculated (Arendt et al., 1986, 1995a). No correction for tissue shrinkage was made. Data acquisition was performed blind to diagnosis.

**Neuronal cell counts.** Numerical neuronal density, defined as the number of nucleolated neurons per tissue volume, was determined after Nissl staining in every tenth section (30  $\mu\text{m}$  thickness) throughout the brain areas listed above with help of an automatic image analyzing system (analySIS). To allow a direct comparison of figures obtained for different areas and different subgroups of patients, we expressed data as percentages of control values. In the Ch4 part (Mesulam et al., 1983) of the basal nucleus of Meynert, cell counts were performed additionally on every 20th section processed for ChAT or p75<sup>NGFR</sup> immunohistochemistry. Total neuronal number was determined by interpolating over the entire structure.

**Activity of ChAT.** Determination of activity of ChAT was performed on homogenates of tissue samples in 0.25 M sucrose containing 0.2% Triton X-100 according to the method of Fonnum (1969) in the modification of Bigl (1975), using [<sup>3</sup>H]acetyl CoA (specific activity 1.5 MBq/ $\mu\text{mol}$ , Amersham) as substrate. Protein content was determined according to Peterson (1977).

**Statistics.** For each case, 300 Golgi-impregnated neurons were analyzed, and data were averaged to determine a case mean. These case means were averaged to obtain group means, the differences of which were analyzed with parametric statistical tests (program SPSS/PC<sup>+</sup>, SPCC, Chicago, IL). One way ANOVA and orthogonal  $t$  tests were used

**Table 3. Statistical summary for the analysis of frequency distribution of dendritic segments in AD compared with controls**

		ApoE 3/3		ApoE 3/4		ApoE 4/4	
	Effect	df	F	df	F	df	F
Basal nucleus of Meynert							
Control	AD	1, 52	72.12***	1, 37	143.34***	1, 29	226.82***
	AD × Lin segm.or.	1, 52	32.34***	1, 37	63.70***	1, 29	108.19***
AD: ApoE 3/4	AD × Lin segm.or.	1, 51	8.29**			1, 28	9.16**
AD: ApoE 4/4	AD × Lin segm.or.	1, 43	21.22***				
Locus coeruleus							
Control	AD	1, 52	43.78***	1, 37	82.10***	1, 29	136.16***
	AD × Lin segm.or.	1, 52	33.10***	1, 37	71.90***	1, 29	112.34***
AD: ApoE 3/4	AD × Lin segm.or.	1, 51	7.56**			1, 28	8.72**
AD: ApoE 4/4	AD × Lin segm.or.	1, 43	18.32***				
Raphe magnus nucleus							
Control	AD	1, 52	45.28***	1, 37	93.10***	1, 29	129.45***
	AD × Lin segm.or.	1, 52	29.34***	1, 37	58.22***	1, 29	89.45***
AD: ApoE 3/4	AD × Lin segm.or.	1, 51	5.16*			1, 28	4.82*
AD: ApoE 4/4	AD × Lin segm.or.	1, 43	8.29**				
Medial amygdaloid nucleus							
Control	AD	1, 52	41.18***	1, 37	83.19***	1, 29	124.12***
	AD × Lin segm.or.	1, 52	23.18***	1, 37	48.22***	1, 29	75.60***
AD: ApoE 3/4	AD × Lin segm.or.	1, 51	4.56*			1, 28	5.28*
AD: ApoE 4/4	AD × Lin segm.or.	1, 43	7.78**				
Pedunculopontine tegmental nucleus							
Control	AD	1, 52	28.92***	1, 37	62.10***	1, 29	89.52***
	AD × Lin segm.or.	1, 52	22.12***	1, 37	43.29***	1, 29	63.17***
AD: ApoE 3/4	AD × Lin segm.or.	1, 51	5.10*			1, 28	6.23*
AD: ApoE 4/4	AD × Lin segm.or.	1, 43	7.56**				
Substantia nigra							
Control	AD	1, 52	21.12***	1, 37	31.19***	1, 29	67.32***
	AD × Lin segm.or.	1, 52	14.20***	1, 37	19.12***	1, 29	38.17***
AD: ApoE 3/4	AD × Lin segm.or.	1, 51	4.78*			1, 28	4.31*
AD: ApoE 4/4	AD × Lin segm.or.	1, 43	7.30**				

Data are *F* values for the comparison of subgroups of patients with AD with the control group and for comparison between AD subgroups obtained by one-way ANOVA. \**p* < 0.05; \*\**p* < 0.01; \*\*\**p* < 0.001; for group size, see Table 1. Lin segm.or., Linear trend of segment order.

to compare individual subgroups of patients with each other and with the control group. Analysis of linear regression was used to analyze the relationship between dendritic growth and neuronal loss, between dendritic growth and clinical stage, and between regenerative capacity and clinical stage of the disease.

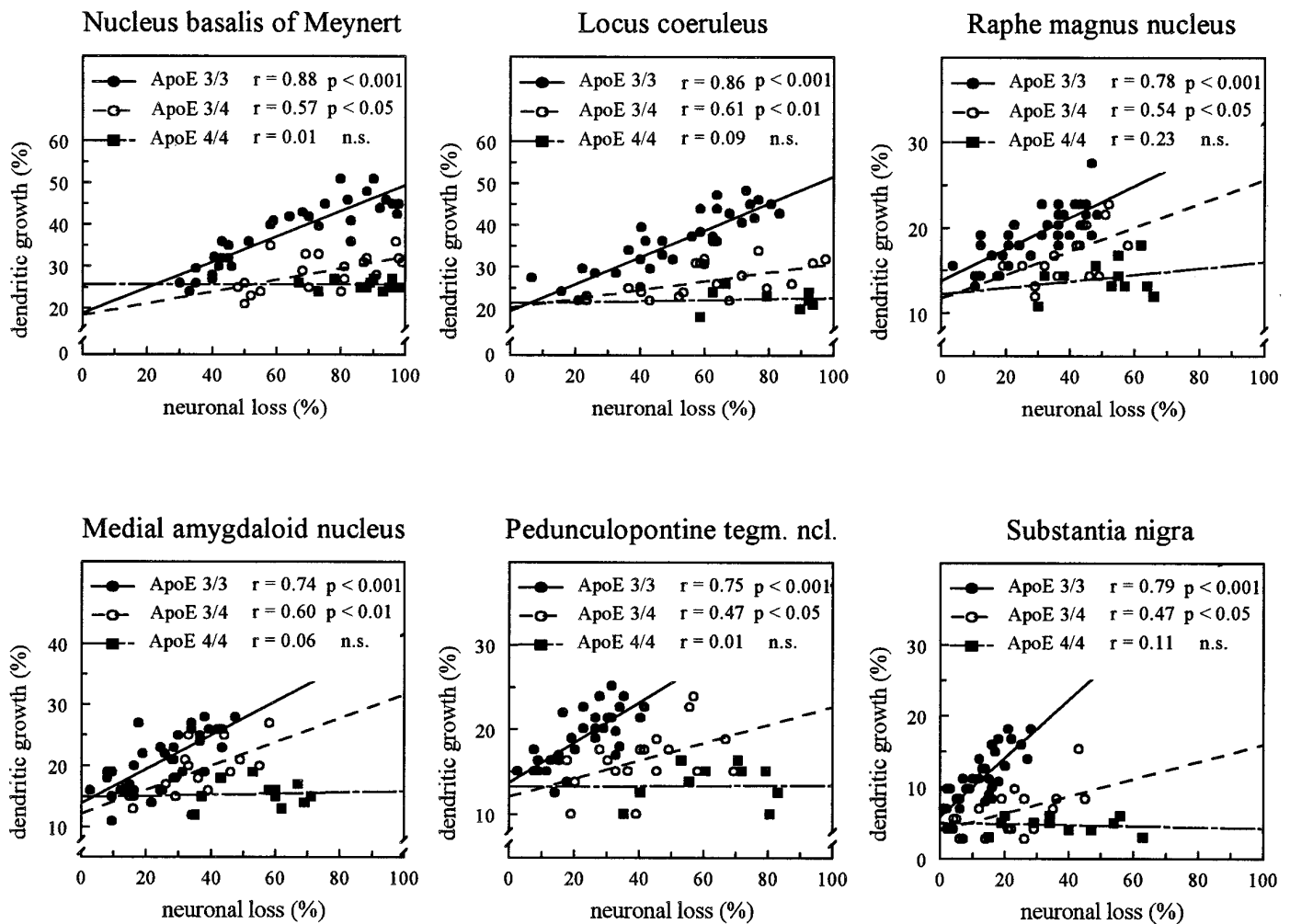
## RESULTS

### Plastic dendritic changes of subcortical neurons

Golgi-impregnated reticular neurons were sampled in the basal nucleus of Meynert (Fig. 1), locus coeruleus, raphe magnus nucleus, medial amygdaloid nucleus, pedunculopontine tegmental nucleus, and substantia nigra. The three-dimensional analysis of the dendritic tree of reticular neurons revealed an increase in dendritic length in patients with AD, as compared with controls in all areas investigated (Table 2). This increase in length of dendrites varied for the different areas. It ranged from ~8% in the substantia nigra to ~26% in the basal nucleus of Meynert when all AD patients were averaged. The intensity of changes in dendritic length within a given area, however, markedly differed for subgroups of patients grouped according to their apoE genotypes. Dendritic changes were most pronounced in patients carrying apoE ε3/3. They were clearly marked less strongly in patients with the genotypes apoE ε3/4 or ε4/4 (Table 2). This influ-

ence of the apoE ε4 allele on the intensity of plastic dendritic changes was present most clearly in the basal nucleus of Meynert and the locus coeruleus but also could be detected in all other areas investigated.

To characterize changes in the pattern of arborization that might accompany the observed process of dendritic growth and to study the distribution of newly formed dendritic segments within the dendritic tree, we analyzed the frequency distribution of all dendritic segments of different orders (Fig. 2). The centrifugal system of ordering segments was used, i.e., primary dendritic segments correspond to segment order 1; the highest order of segments represents most distal branches. Although no subgroup-specific changes were noted for the total number of segments (all *p* > 0.1), subgroups of patients were significantly different with respect to the distribution of newly formed dendritic segments (Table 3). In patients with apoE ε3/3, newly formed segments were mostly of higher order, i.e., more distally localized. Growth processes were approximately equally present in distal and proximal parts of the dendritic tree in patients with apoE ε3/4 and were most frequently localized in proximity to the soma in patients with apoE ε4/4.



**Figure 3.** Relationship between the loss of neurons and the extent of dendritic growth in subcortical brain areas separately analyzed for different subgroups of patients with AD. Analyses of linear regression were performed, and correlation coefficients according to Bravais–Pearson were calculated. For group size, see Table 1.

### Degeneration of neurons

Neuronal degeneration, characterized by formation of NFT and neuronal loss, was observed in all areas investigated. Neuronal loss was most severe in the basal nucleus of Meynert, followed in decreasing order of severity by the locus coeruleus, raphe magnus nucleus, medial amygdaloid nucleus, pedunculopontine tegmental nucleus, and substantia nigra. Again, the extent of changes differed for different subgroups of patients according to their apoE genotype (Table 2). Loss of neurons was more severe in patients carrying one apoE  $\epsilon 4$  allele, as compared with patients with apoE  $\epsilon 3/\epsilon 3$ , and was even higher in patients with apoE  $\epsilon 4/\epsilon 4$ . These differences applied to all areas analyzed in the present study.

### Relationship between degeneration and dendritic reorganization

To investigate whether the process of dendritic reorganization in AD might be related to the extent of neuronal loss, we analyzed the dependency of these two parameters by statistical methods (Fig. 3). A highly significant relationship between dendritic growth and neuronal loss was obtained for patients with apoE  $\epsilon 3/\epsilon 3$  in all areas investigated. The relationship of these two parameters, however, was only marginally significant in patients with the apoE  $\epsilon 3/\epsilon 4$  genotype.

The small increase in dendritic length observed in patients with apoE  $\epsilon 4/\epsilon 4$  was unrelated to the extent of neuronal loss.

To further characterize subgroups of AD patients concerning differences in the dependency of dendritic growth on the extent of neuronal loss, we calculated the individual ratios of the two parameters. This ratio of the relative increase in dendritic length per relative neuronal loss gives an indication of the overall extent of regeneration/repair in the different structures that might occur in response to degeneration of a given extent. This parameter was, therefore, designated as “reparative capacity.” For example, a reparative capacity with a value of “1” would indicate that the newly formed dendritic elements are quantitatively equivalent to those that have been lost by neuronal degeneration. It should be borne in mind, however, that this parameter of reparative capacity solely represents a measure of the mathematical dependency of the two processes, i.e., cell loss and dendritic growth. A strong statistical relationship obtained by this method does not necessarily imply a causal relationship between the two processes. The reparative capacity emerged as a particularly useful parameter with a high potential to discriminate different subgroups of AD patients (Fig. 4). In comparison to patients with AD lacking the apoE  $\epsilon 4$  allele, the value of this parameter

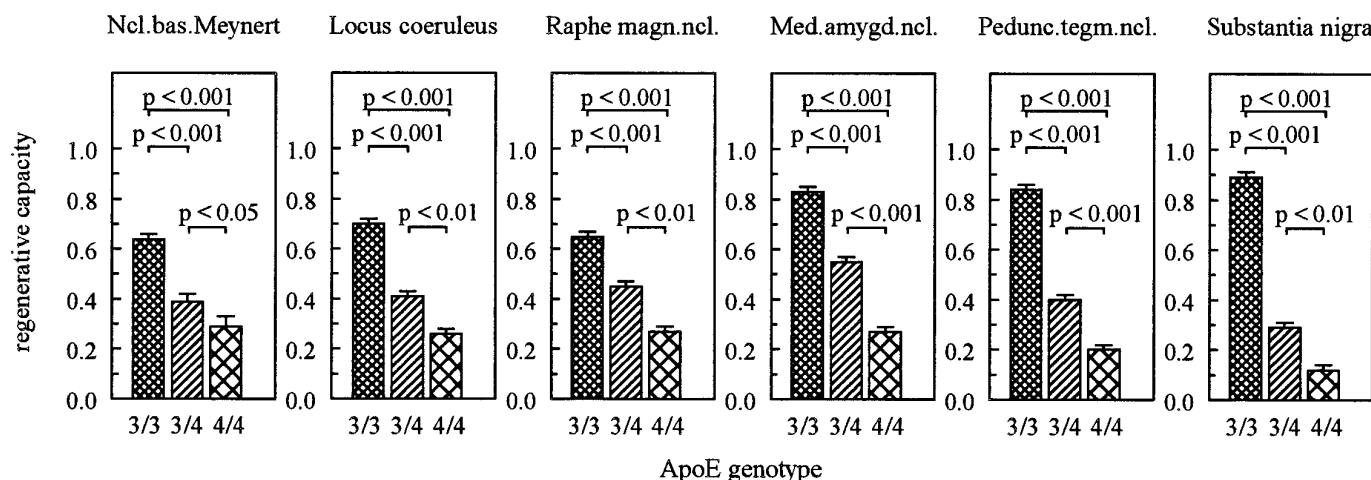


Figure 4. Differences in the reparative capacity of patients with AD grouped according to their apoE genotypes. Student's *t* tests were used to compare individual group means  $\pm$  SEM. For group size, see Table 1.

was reduced by up to 68 and 85% in patients carrying one or two apoE  $\epsilon$ 4 alleles, respectively.

#### Stage-dependent differences of dendritic plasticity

The intensity of dendritic growth observed in patients with AD not only might depend on the extent of neuronal degeneration but also might be influenced by the progression of the disease and might, thus, vary for different disease stages. The influence of the stage of the disease on both the intensity of dendritic growth and the reparative capacity was, therefore, evaluated for each subgroup of AD patients separately (Fig. 5 and Table 4). During early stages of the disease, the extent of dendritic growth was similar for all subgroups of AD patients, independent of its apoE genotype (Fig. 5, *left panels*). The progression of the disease, however, had a significantly different influence on further dendritic changes (Table 4). Whereas in patients with apoE  $\epsilon$ 3/3 dendritic growth continuously increased over the progression of the disease, this increase was only marginally significant in the medial amygdaloid nucleus and insignificantly small in all other areas in patients with apoE  $\epsilon$ 3/4. Disease progression had no influence on dendritic growth in patients with apoE  $\epsilon$ 4/4. Contrary to changes in dendritic length, reparative capacity tended to decline over the progression of the disease (Fig. 5, *right panels*). Again, changes were significant for the apoE  $\epsilon$ 3/3 genotype but only marginally significant or insignificantly small in the presence of one or two apoE  $\epsilon$ 4 alleles (Table 4). Although at early stages of the disease subgroups of patients clearly differed with respect to their reparative capacity, these differences declined considerably during the progression of the disease (Fig. 5).

#### Degeneration of cortical cholinergic axon terminals

Activity of ChAT in the frontal cortex (Brodmann area 8) and temporal cortex (Brodmann area 20) was reduced in all cases with AD. Differences of these changes related to subgroups of patients with different apoE genotypes were pronounced even more clearly than the loss of cholinergic neurons in the basal nucleus of Meynert identified by anti-ChAT and anti-p75<sup>NGFR</sup> immunoreactivity (Table 5). Whereas ChAT activity was reduced by 35–40% in cases with apoE  $\epsilon$ 3/3, reductions of ~70% and even >90% were observed in cases with apoE  $\epsilon$ 3/4 and 4/4, respectively.

Comparing the loss of cholinergic neurons in the basal nucleus giving rise to the cortical cholinergic innervation with changes in

ChAT activity at cortical target sites might give some indication on adaptive changes of cholinergic neurotransmission related to plastic phenomena occurring in remaining neurons. These plastic changes at the sites of cholinergic axon terminals varied among different subgroups of patients, as shown in Figure 6. In patients with apoE  $\epsilon$ 3/4, activity of ChAT at cortical sites was reduced to a similar extent as the number of cholinergic basal forebrain neurons (ratio of ChAT activity to cell number close to 1). Cortical ChAT activity was affected less severely than cell number in the basal nucleus in patients with apoE  $\epsilon$ 3/3 (ratio of ChAT activity to cell number >1), whereas cortical ChAT activity was reduced even more dramatically than neuronal number in the basal forebrain in patients with apoE  $\epsilon$ 4/4 (ratio of ChAT activity to cell number <1).

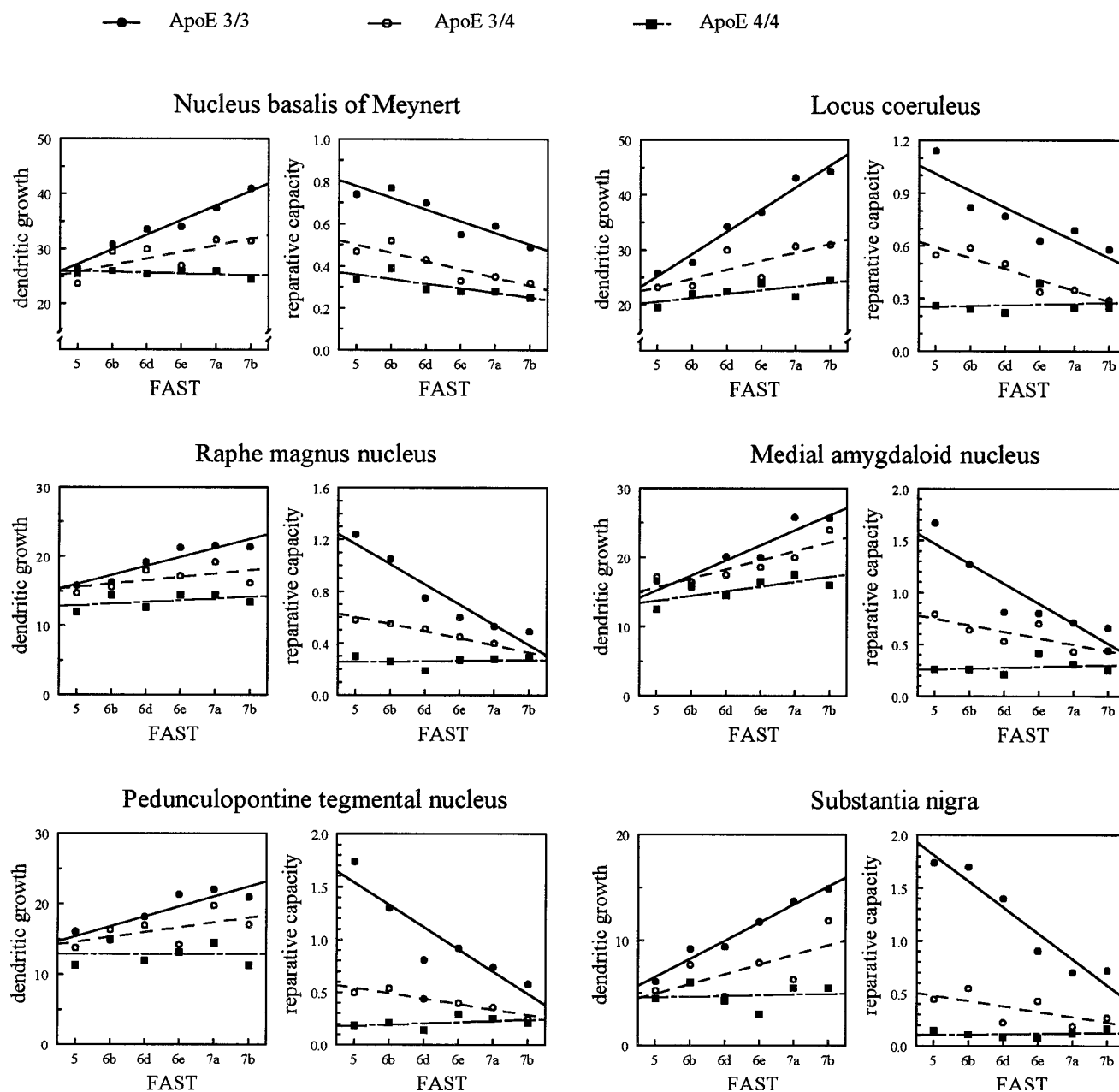
#### DISCUSSION

Chronic neuronal degeneration in a number of disorders related to quite different etiologies, such as AD, post alcoholic Korsakoff's disease, Parkinson's disease, or Huntington's chorea, is accompanied by growth and reorganization of the dendritic tree of certain types of neurons (Graveland et al., 1985; Arendt et al., 1986, 1995b,c; Scott, 1993). These changes have been regarded as an attempt of the nervous system to counteract the functional impairments resulting from degenerative events.

In AD, plastic changes that occur in response to degeneration in cortical and subcortical areas show severe aberrations with respect to their localization, morphological appearance (Scheibel and Tomiyasu, 1978; Scheibel, 1979; Ferrer et al., 1983, 1990; Probst et al., 1983; Arendt et al., 1986, 1995a–c; Arendt and Zvegintseva, 1987; Ihara et al., 1988; Arendt and Brückner, 1991, 1992), and composition of cytoskeletal elements (McKee et al., 1989). The reasons for these aberrancies are unknown.

The results of the present study demonstrate that, in a number of subcortical brain areas, the extent of neuronal degeneration as well as the intensity and pattern of plastic dendritic changes in AD is related to the apoE genotype (For synopsis of the major findings, see Table 6). Because none of the control cases carried an apoE  $\epsilon$ 4 allele, the present study does not allow any conclusion on whether apoE polymorphism might affect neuronal number and branching in patients without AD.





**Figure 5.** Relationship between the dendritic growth and reparative capacity of reticular neurons in subcortical brain areas and the clinical stage of AD, assessed according to FAST. Different subgroups of patients classified according to their apoE genotypes were compared by ANOVA (effect, subgroup  $\times$  linear trend of disease progression). df, apoE 3/3 versus apoE 3/4, 1, 51; apoE 3/3 versus apoE 4/4, 1, 43; apoE 3/4 versus apoE 4/4, 1, 28; for both dendritic growth and reparative capacity, all  $p < 0.01$ . For group size, see Table 1; for analysis of linear regression, compare Table 4.

### ApoE gene dose and degeneration of subcortical neurons

A severe neuronal degeneration was observed in the basal nucleus of Meynert, locus coeruleus, the median raphe nucleus, medial amygdaloid nucleus, pedunculopontine tegmental nucleus, and substantia nigra, thereby confirming a number of earlier studies (Herzog and Kemper, 1980; Arendt et al., 1983; Mann et al., 1983; Shortridge et al., 1985; Yamamoto and Hirano, 1985; German et al., 1987; Jellinger, 1988; Woolf et al., 1989; Esiri et al., 1990; Goto et al., 1990; Aletrino et al., 1992; Förstl et al., 1992; Halliday et al., 1992; Scott et al., 1992). The overall extent of degeneration and the gradual variation of cell loss between different areas are also

in agreement with previous reports (Zweig et al., 1988; Chan-Palay et al., 1992; Arendt et al., 1995c).

In the present study, the severity of degeneration in these areas was found to vary among different patients, depending on their apoE genotype. A gene dosage effect of the apoE  $\epsilon 4$  allele on the severity of degeneration of cholinergic basal forebrain neurons, as revealed in the present study, had been reported previously by Poirier et al. (Poirier, 1994; Poirier et al., 1995) and others (Soininen et al., 1995). The concept has, therefore, been put forward that the integrity of cholinergic neurons may be compromised selectively in apoE  $\epsilon 4$  carriers (Poirier, 1994). The present study demonstrates that effects mediated by the apoE  $\epsilon 4$  allele are

**Table 4.** Analysis of stage dependency of dendritic growth and reparative capacity of different subgroups of AD patients<sup>a</sup>

	ApoE 3/3 dendr.gr.	ApoE 3/4 regen.cap.	ApoE 4/4 dendr.gr.	regen.cap.	dendr.gr.	regen.cap.
Basal nucleus of Meynert	0.98***	0.92**	0.74	0.88*	0.41	0.80
Locus coeruleus	0.98***	0.89*	0.80	0.93**	0.68	0.13
Raphe magnus nucleus	0.93**	0.96**	0.57	0.98***	0.44	0.12
Medial amygdaloid nucleus	0.94**	0.91*	0.88*	0.81	0.72	0.23
Pedunculopontine tegmental nucleus	0.88*	0.92**	0.59	0.46	0.01	0.38
Substantia nigra	0.99***	0.96**	0.66	0.65	0.09	0.18

<sup>a</sup>Analysis of linear regression of dendritic growth (dendr.gr.) and reparative capacity (regen.cap.) on the clinical stage of the disease assessed according to FAST (Reisberg et al., 1982).

The correlation coefficient is significantly different from zero for \*\*\* $p < 0.001$ ; \*\* $p < 0.01$ ; \* $p < 0.05$ . For group size, see Table 1.

**Table 5.** Changes in the number of cholinergic neurons in the basal nucleus of Meynert and the activity of ChAT in the cerebral cortex in different subgroups of patients with AD

	Alzheimer's disease			
	Control	ApoE 3/3	ApoE 3/4	ApoE 4/4
Neuronal number				
ChAT-immunoreactive	178,467 ± 4230	64,834 ± 4852***	47,519 ± 5115***#	23,850 ± 4635***###
p75 <sup>NGFR</sup> -immunoreactive	182,389 ± 4820	66,972 ± 5210***	49,608 ± 4864***#	25,160 ± 4645***###
ChAT activity (nmol/mg protein × h)				
Brodmann area 8	11.9 ± 0.70	7.14 ± 0.52***	3.33 ± 0.34***###	0.83 ± 0.22***###
Brodmann area 20	8.3 ± 0.40	5.47 ± 0.25***	2.40 ± 0.25***###	0.55 ± 0.05***###

Data are mean values ± SEM; for group size, compare Table 1. Significantly different from control: \*\*\* $p < 0.001$  (Student's  $t$  test). Significantly different from AD subgroup ApoE 3/3: ### $p < 0.001$ ; # $p < 0.05$  (Student's  $t$  test). For group size, see Table 1.

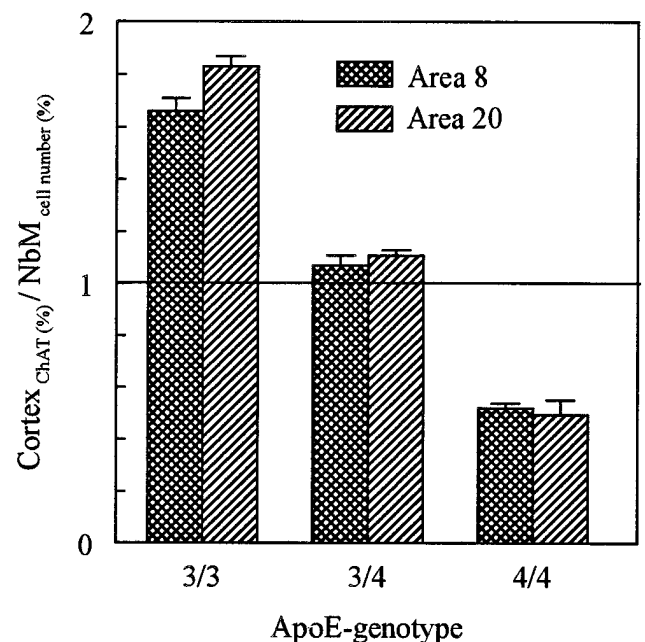
not confined to the cholinergic basal forebrain neurons. Effects related to the apoE-polymorphism might, therefore, be of more global relevance to the process of neuronal degeneration in AD.

### ApoE polymorphism and dendritic plasticity in AD

ApoE  $\epsilon 4$  allele copy number in patients with AD showed an inverse relationship to the extent of plastic neuronal remodeling. In AD patients lacking the apoE  $\epsilon 4$  allele, reactive dendritic growth parallels both the extent of neuronal degeneration and the progression of the disease. Although conclusions of dynamic events from static images need to be drawn with caution, it is tempting to regard this correlative relationship as an indication of a "coupling" between the process of dendritic reorganization and the functional demands of compensating degenerative events. In these patients, newly formed dendritic branches were localized mainly on segments of higher order, resulting in an extensive pattern of growth. A similar pattern of dendritic growth has been established previously for normal aging (Arendt et al., 1995b).

In patients carrying one apoE  $\epsilon 4$  allele, the increase in dendritic length was much less pronounced. Plastic dendritic changes were reduced further in patients homozygous for the  $\epsilon 4$  allele. These effects of the apoE polymorphism on reactive dendritic growth in AD also could be the reason for previous discrepancies between different studies on the extent of dendritic growth in AD (Buell and Coleman, 1979, 1981; Arendt et al., 1986, 1995b,c; Flood et al., 1991).

In ApoE  $\epsilon 4$ -carriers, newly formed dendritic elements were gene dose dependently shifted from distal segments to more proximal parts of the dendritic tree. This different distribution of growth processes resulted in an intensive pattern of growth that already was observed previously in AD (Arendt et al., 1995b). The intensity of growth, furthermore, was related only weakly to both the extent of neuronal loss and the progression of the disease in patients carrying one apoE  $\epsilon 4$  allele. It was completely independent of these two parameters in patients homozygous for the  $\epsilon 4$  allele. These findings



**Figure 6.** Ratio between ChAT activity in the cerebral cortex and number of cholinergic neurons in the basal nucleus of Meynert in patients with AD carrying different apoE genotypes. Values of both ChAT activity and neuronal number (determined on every 20th section processed for anti-ChAT immunocytochemistry) are expressed as percentages of control values. Student's  $t$  tests were used to compare mean values of individual groups; all  $p < 0.001$ . For group size, see Table 1.

**Table 6. Synopsis of the influence of apoE polymorphism on degeneration and dendritic reorganization on subcortical neurons in AD**

	ApoE 3/3	ApoE 3/4	ApoE 4/4
<b>Degeneration</b>			
Neuronal density (depending on brain region) <sup>a</sup>	30–90%	20–80%	10–70%
<b>Dendritic reorganization</b>			
Dendritic length (depending on brain region) <sup>a</sup>	110–140%	100–130%	100–130%
Localization of newly formed dendritic branches	Preferentially distally localized	Intermediate	Preferentially proximally localized
<b>Relationship between degeneration and dendritic reorganization</b>			
Relationship between neuronal loss and dendritic growth	Significantly positively correlated	Marginally significant	Insignificantly small
<b>Reparative capacity</b>			
Ratio: dendritic growth/neuronal loss (depending on brain region) <sup>a</sup>	0.6–0.9	0.3–0.6	0.1–0.3
<b>Stage dependency of dendritic reorganization</b>			
Relationship between disease stage and dendritic growth	Significantly positively correlated	Marginally significant	Insignificantly small
Relationship between disease stage and reparative capacity	Significantly negatively correlated	Marginally significant	Insignificantly small
<b>Degeneration and reorganization of the cortical cholinergic afferentation</b>			
Cortical activity of ChAT	60–70%	20–40%	<10%
Number of ChAT-positive neurons in the basal nucleus of Meynert	30–40%	20–40%	10–20%
Ratio of cortical ChAT (%) versus number (%) of ChAT-positive neurons in the basal nucleus	>1	~1	<1

<sup>a</sup>The regional intensity of both degenerative and plastic dendritic changes varies in the following order: basal nucleus > locus coeruleus > raphe magnus nucleus > medial amygdaloid nucleus > pedunculopontine tegmental nucleus > substantia nigra.

might indicate an “uncoupling” of dendritic growth processes from their functional requirements under these conditions.

The gene dosage effects of the apoE  $\epsilon 4$  allele on plastic neuronal changes observed in the present study were not restricted to dendrites and similarly could be detected on axons. In patients lacking apoE  $\epsilon 4$ , ChAT activity, determined in the frontal or parietal cortex, was reduced less drastically than the number of cholinergic neurons in the basal forebrain, giving rise to the cholinergic cortical innervation. In patients homozygous for  $\epsilon 4$ , on the contrary, loss of ChAT activity was more pronounced than loss of cholinergic neurons, whereas the situation was intermediate for patients carrying one  $\epsilon 4$  allele. These findings imply either an upregulation of ChAT activity in surviving neurons in the absence of the apoE  $\epsilon 4$  allele or compensatory axonal sprouting of surviving neurons. The present data do not allow us to distinguish between these two possibilities but clearly show the impairment of these plastic axonal processes in patients carrying  $\epsilon 4$  allele.

#### **ApoE genotype and stage dependency of reparative capacity**

In the present study, the mean age of patients with apoE  $\epsilon 4$  alleles was ~8 years lower than for patients without  $\epsilon 4$ . This earlier age of onset is a consistent finding associated with apoE  $\epsilon 4/4$  (Corder et al., 1993; Soininen, 1995). Other studies, furthermore, have reported on a more rapid progression of the disease in these patients (Bennett et al., 1995). These findings implicate that patients with apoE  $\epsilon 4$  alleles might reach more advanced stages of the disease at a younger age than patients without  $\epsilon 4$ . Previous studies have indicated that younger AD patients show a more severe degeneration (Bird et al., 1983; Rossor et al., 1984; Perry et al., 1992; Arendt et al., 1995b; Soininen et al., 1995) and a more intense plastic response of subcortical neurons (Arendt et al., 1995c) than older patients. Comparative studies on carriers and

noncarriers of the  $\epsilon 4$  allele might, therefore, be biased by age-related effects. In the present study, two measures were taken to minimize the likelihood of such influence. First, effects related to different apoE genotypes were matched according to the clinical stage of the disease. Second, a parameter, the reparative capacity, was defined, which allowed us to distinguish effects of the  $\epsilon 4$  allele on qualitative, rather than quantitative, grounds. Differences in the reparative capacity related to the presence of the apoE  $\epsilon 4$  allele were independent of the clinical stage of the disease. These results clearly show that the gene dose of the apoE  $\epsilon 4$  allele has an effect on the intensity of reactive dendritic remodeling.

Although dendritic elements continue to grow during the progression of AD in patients lacking the apoE  $\epsilon 4$  allele, reparative capacity continuously declines in these patients. For the most advanced stages, they converge with the low levels of reparative capacity seen in the presence of the apoE  $\epsilon 4$  allele. This observation indicates that, even in patients lacking the apoE  $\epsilon 4$  allele, the presumptive reparative capacity is progressively exhausted. Differences in the mechanism attributed to the apoE-polymorphism might, therefore, be particularly relevant in early stages of the disease.

The present results support the assumption of an involvement of apoE in the repair of central neurons, as suggested by Poirier et al. (1991 a,b, 1993a) and Roses (1994). This proposed mechanism for apoE is supported further by recent observations on apoE knockout mice that show a severe impairment of age and lesion-related plastic synaptic changes (Masliah et al., 1995a,b).

An impairment of the expression of the  $\epsilon 4$  gene and/or the functional properties of the E 4 protein might be related causally to the aberrancies of neuronal repair in AD and might, thus, explain the effect of apoE-polymorphism on the onset and development of AD at the cellular level.

It remains to be determined whether the molecular mechanisms behind these effects are related to genotype-specific alterations of apoE levels in the brain (Blennow et al., 1994; Bertrand et al., 1995), to isoform-specific differences in interactions with cellular proteins that mediate neurotrophic (Nathan et al., 1994; Bellosta et al., 1995; Holtzman et al., 1995) or cytotoxic (Crutcher et al., 1994; Clay et al., 1995) effects, or to molecular interactions with the  $\beta$ /A4-amyloid (Wisniewski and Frangione, 1992; Strittmatter et al., 1993b; Wisniewski et al., 1993; Gallo et al., 1994; Castano et al., 1995; Evans et al., 1995) or microtubule-associated proteins (Cotton et al., 1994; Huang et al., 1994, 1995; Strittmatter et al., 1994a,b; Whitson et al., 1994).

Despite these clear-cut effects of the  $\epsilon$ 4 allele on neuronal plasticity in AD, the plastic neuronal response was not completely absent, even in patients homozygous for the  $\epsilon$ 4 allele. These observations indicate that plastic reorganization under conditions of neurodegeneration is not disturbed completely but appears to be uncoupled from its functional demands. It might, therefore, be hypothesized that effects mediated by the presence of the apoE  $\epsilon$ 4 allele are not the direct cause of the disease. These effects, instead, might lead to a more rapid functional decompensation of a neuronal system under the conditions of a slowly progressing degeneration, which results in a more early onset and more rapid progression of the disease.

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