

**Experimental approaches to study cerebral
amyloidosis in a transgenic mouse model of
Alzheimer's disease**

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Summary

Misfolding, aggregation and the accumulation of proteins in the brain are common characteristics of diverse age-related neurodegenerative diseases. Each of these neurodegenerative diseases is associated with abnormalities in the folding of a different protein leading to protein aggregation and finally to neuronal death. Alzheimer's disease (AD) is one of these protein conformational diseases characterized by two major neuropathological features: extracellular accumulation of amyloid- β (A β) peptide in the form of plaques and intracellular tangles consisting of hyperphosphorylated tau protein. Although the majority of AD cases are sporadic, three genes have been described whose mutations cause early-onset familial AD (FAD). The identification of mutations in these genes has provided new opportunities to explore pathogenic mechanisms using transgenic approaches. Based on the finding that mutations in these genes all lead to elevated levels of A β , new anti-amyloid therapies have been developed to either lower the production of A β or to clear the amyloid peptides.

In the past few years, several groups have generated transgenic mouse models of cerebral amyloidosis that exhibit age related A β deposition similar to AD patients through expression of mutated human amyloid precursor protein (APP). The studies presented herein were done using such a transgenic mouse model, the APP23 mouse, that overexpresses human APP with the Swedish mutation under the control of a neuron specific Thy-1 promotor. APP23 transgenic mice develop cerebral amyloidosis in an age- and region-dependent manner. Plaque formation starts early at 6 months of age and is associated with the typical AD-like pathology including cerebral amyloid angiopathy, neuron loss, glial activation and cognitive impairment.

The purpose of this thesis was to study the mechanism and initiation of amyloid formation as well as the spread of cerebral amyloidosis *in vivo*. The first series of studies were conducted to define the role and contribution of extracellular versus intracellular β -amyloid in plaque formation. To this end, we transplanted embryonic wildtype (wt) and APP23 transgenic (tg) brain tissue into the hippocampus and cortex of both APP23 and wt mice. We observed that APP23 grafts into wt hosts did not develop amyloid deposits up to 20 months post-grafting. In contrast, both tg and wt grafts

into APP23 hosts developed amyloid plaques already 3 months post-grafting. The amyloid deposits in wt grafts were surrounded by neuritic changes and gliosis similar to the amyloid-associated pathology described in APP23 mice as well as in AD patients. These results suggest that the phenotype of the transplanted tissue is strongly influenced by the properties of the host. Moreover, these results provide evidence that diffusion of A β in the extracellular space is important for the spread of A β pathology, that amyloid formation starts extracellularly and that it is the extracellular amyloid that causes neurodegeneration.

The second set of experiments were performed to study the initiation of amyloid deposition and to clarify which factors are involved in the seeding process *in vivo*. Since seeded polymerization of A β has already been demonstrated *in vitro* and *in vivo*, we replicated and advanced these findings by intracerebral injection of diluted brain extract from AD patients and brain extracts from aged APP23 transgenic mice into young pre-depositing APP23 mice. AD and APP23 brain homogenate induced a similar amount of seeded A β deposits in the brain parenchyma and vessel walls four month post-infusion. This seeding was time- and concentration-dependent. In contrast, no seeding was observed when PBS was injected or when the same extract was injected into wt mice. To address whether A β itself is the seeding agent we injected synthetic A β into young APP23 mice. These synthetic A β injections resulted in limited A β deposition compared to that obtained with A β -rich brain extract. Our findings suggest that A β -containing human and mouse brain extracts can induce cerebral amyloidosis *in vivo*, and that A β , in combination with additional factors, initiates amyloid formation.

The third part of the work presented here follows up on our previous finding that diffusion of A β in the extracellular space plays an important role in the spread of cerebral amyloidosis. Therefore, we came up with the hypothesis that amyloid deposition and the accompanied pathophysiology could influence extracellular space (ECS) volume and interstitial fluid (ISF) diffusion properties. By using diffusion weighted magnetic resonance imaging (DWI), we determined the diffusion properties in the brains of young and aged APP23 transgenic mice and control littermates. Our results indicate that fibrillar amyloid formation and the associated gliosis are accompanied by a decrease in the apparent diffusion coefficient (ADC), suggesting that both build a

barrier for interstitial fluid diffusion. Thus, in elderly people, ADC measurements and the assessment of diffusion properties in the ECS could serve as a biomarker to detect pathological events in the brain of AD patients.

In summary, the studies presented herein have increased our understanding of the mechanisms leading to protein aggregation and finally to neurodegeneration in a transgenic mouse model. We have shown that factors other than local A β production, such as diffusion in the extracellular space, are important in determining whether amyloid pathology will occur. Moreover, the results highlight the relevance of extracellular A β to the pathogenesis of the disease. It still remains an open question whether A β itself is sufficient to initiate plaque formation, and if so, what conformational form of A β is required. Together, these studies provide insights into the mechanisms and disease pathways which may lead to AD.

1. Introduction

1.1 Protein misfolding and aggregation in neurodegenerative diseases

The misfolding and aggregation of proteins is one of the major threats for living organisms. The necessary information for a polypeptide chain to fold correctly into a three-dimensional structure is encoded in its primary amino acid sequence (Anfinsen, 1973). Normally, protein folding is assisted by chaperones that help correct folding and remove misfolded and malfunctioning proteins. However, under physiological conditions of high protein concentrations and cellular crowding (van den Berg et al., 1999; Minton, 2000) and especially at high temperatures, many proteins tend to form stable insoluble aggregates (Jaenicke, 1995). Aggregation occurs when folding/unfolding intermediates become trapped in partially misfolded states that successively associate, mainly through hydrophobic interactions, into an oligomeric continuum of increasingly larger, more stable, and less soluble complexes. In the aggregated state, inactive proteins are enriched in anti-parallel β -strands (Fink, 1998). They can appear as amorphous structures like inclusion bodies, or as ordered fibers (straight, unbranched, 10-nm wide fibrils) such as amyloid plaques and prion particles (Prusiner, 1998; McLaurin et al., 2000).

Age-related neurodegenerative diseases such as Alzheimer's disease (AD), Parkinson's Disease (PD), Huntington's Disease (HD), amyotrophic lateral sclerosis (ALS) and transmissible spongiform encephalopathies (TSE) form a pathologically diverse group that can affect thinking, skilled movements, emotion, cognition, memory and other abilities (Table 1). However, despite their differences in clinical symptoms and disease progression, these disorders do share some common features: most of them appear later in life and their pathology is characterized by neuronal loss and synaptic abnormalities (Martin, 1999). The hallmark feature of these conformational disorders is that proteins can fold into an alternative stable conformation, which results in its aggregation and accumulation as fibrillar deposits (Carrell and Lomas, 1997; Dobson, 1999). Insoluble aggregates of disease related proteins are deposited as inclusions or plaques, the characteristics of which are often disease specific. In AD, amyloid plaques are

deposited in the brain parenchyma and vessel walls, composed mainly of the amyloid- β protein (Glennner et al., 1984). In patients with PD, aggregates called Lewy bodies deposit in the substantia nigra (Forno, 1996), with the major constituent being α -synuclein (Spillantini et al., 1997). Intracellular deposits of a polyglutamine-rich version of huntingtin protein are typical for the brains of HD patients (DiFiglia et al., 1997). In ALS, patients have aggregates that are mainly composed of superoxid-dismutase in the cell bodies and axons of motor neurons (Bruijn et al., 1998). Finally, the brains of humans that have been affected by TSE are characterized by aggregates of the prion protein (PrP) (Bolton et al., 1982).

Disease	Clinical features	Protein deposits	Toxic protein	Affected brain regions	Disease gene
Alzheimer's Disease	Progressive dementia	Extracellular plaques Intracellular tangles	Amyloid- β and Tau	Hippocampus, cerebral cortex	APP PS 1 PS 2
Parkinson's Disease	Movement disorder	Lewy bodies	α -Synuclein	Substantia nigra, hypothalamus	α -Synuclein Parkin UCHL1
Huntington's Disease	Dementia, motor and psychiatric problems	Nuclear and cytoplasmic inclusions	Huntingtin	Striatum, cerebral cortex	IT-15
Amyotrophic lateral sclerosis	Movement disorder	Bunina bodies	Superoxide dismutase	Motor cortex, brainstem	SOD1
Transmissible spongiform encephalopathies	Dementia, ataxia, psychiatric problems	Prion plaque	Prion protein	Various regions depending on disease	PRNP

Table 1. Clinical and pathological features of neurodegenerative disorders characterized by aggregation and deposition of abnormal protein (adapted from Soto, 2003).

Although these cerebral aggregates have been recognized as a typical feature of neurodegenerative diseases for many years, neuropathological studies have been unable to determine whether they are directly involved in the pathogenesis of the disease. Analysis of post-mortem tissue shows only a poor correlation between the amyloid deposit load and the severity of clinical symptoms (Terry et al., 1991; Gutekunst et al.,

1999). Moreover, the appearance of protein deposits in clinically normal, healthy, elderly people (Katzman et al., 1988) question the role of protein aggregates in neurodegeneration, even though protein aggregates accumulate mainly in the brain areas that show degeneration. Support for a causal role of protein misfolding in neurodegenerative diseases has come more recently from genetic studies since mutations in the respective fibrillar proteins have been found in AD, PD, HD, ALS and TSE (Hsiao et al., 1990; Goate et al., 1991; Group, 1993; Rosen et al., 1993; Polymeropoulos et al., 1997). Furthermore, the generation of transgenic animal models, which express mutant proteins, has provided good evidence for the contribution of protein misfolding to the disease pathogenesis. Several pathological and clinical features have been observed in transgenic models in which protein aggregates were successfully produced (Hsiao et al., 1990; Gurney et al., 1994; Games et al., 1995; Mangiarini et al., 1996; Davies et al., 1997; Masliah et al., 2000). AD may therefore be seen as one example of many neurodegenerative diseases in which an abnormally folded molecule accumulates in the brain and causes neuronal damage.

1.2 Alzheimer's disease

It is almost a hundred years ago since the German psychiatrist Alois Alzheimer (1864-1915) first described the clinical and pathological symptoms of the disease that now bears his name. At a meeting in 1906, he first presented the case of a 54 year old woman who suffered from progressive dementia. Although his presentation of the typical clinical and morphological features of AD was not considered to be worthy of discussion at this time, it was published one year later (Alzheimer, 1907). The clinical manifestations of the disease consist of progressive memory loss, disordered cognitive function as well as altered behavior including paranoia, delusions, impairments of attention, perception, reasoning and comportsment, and a progressive decline in language function (Price et al., 1993; Morris, 1996). Motor functions remain initially intact but deteriorate in the final phase of the disease leading to a picture resembling motor disorders such as parkinsonism (Selkoe, 2001). Mental functions and activities of daily living become progressively impaired (Forstl and Kurz, 1999). A state of dementia becomes identified when these deficits undermine the capacity for independent living. The onset of the dementia in AD is preceded by a transitional preclinical period of

many years during which relatively isolated memory difficulties exceed those expected on the basis of normal aging, without however, becoming severe enough to interfere with daily living activities (Linn et al., 1995; Fox et al., 1998). Even normal aging does not necessarily imply that memory function has remained intact (Small et al., 1999).

AD is the most common form of dementia in the elderly accounting for over 50% of the typical, late-onset cases of dementia. The estimated number of patients is approximately 20 million worldwide and is expected to keep growing as the current population ages. Epidemiological studies on the prevalence of AD estimate that not more than 10% of the population over 65 years but 25% of the people over 80 years of age are affected with AD (Pfeffer et al., 1987; Bachman et al., 1992; Lautenschlager et al., 1996; Hy and Keller, 2000). Problems with memory and cognition appear during the seventh decade in most individuals with AD, but may appear earlier, particularly in familial cases (FAD)(Albert and Drachman, 2000). The largest proportion of AD cases are sporadic, occurring with no clearly defined etiology. However, it has been known for several decades that AD can occur in a familial form that is autosomal dominant inherited. Estimates of the proportion of Alzheimer cases that are genetically based varied from as low as 5% to as high as 15% (Selkoe, 1994; Lendon et al., 1997). Familial AD (FAD) is clinically and pathologically indistinguishable from sporadic AD, except for the early age of onset (Lehtovirta et al., 1996; Lippa et al., 1996).

1.2.1 Neuropathological hallmarks

The brains of AD patients contain two hallmark pathological features: neurofibrillary tangles (NFT) and neuritic plaques. Classical senile or neuritic plaques are multicellular lesions containing extracellular deposits of amyloid- β protein (A β) that include abundant amyloid fibrils (7-10 nm) intermixed with non-fibrillar forms of the A β peptide. Neuritic plaques are surrounded by variable numbers of dystrophic neurites, both axonal terminals and dendrites. Many such plaques contain activated microglial cells situated near the amyloid core, as well as reactive astrocytes around the periphery of the plaque (Itagaki et al., 1989; Wisniewski et al., 1989). Although many plaques with these features can be found in the AD brain, an even larger number of deposits seem to lack surrounding dystrophic neurites or glia. These lesions are called diffuse or

preamyloid plaques, where the A β occurs in a non-fibrillar, less dense and amorphous form in the neuropil (Tagliavini et al., 1988; Yamaguchi et al., 1988; Yamaguchi et al., 1989). In brain regions that are generally not implicated in the clinical syndrome of AD, such as cerebellum, striatum and thalamus, almost all A β deposits are diffuse with little or no glial and neuritic reactions (Joachim et al., 1989; Ogomori et al., 1989; Yamaguchi et al., 1989). Besides diffuse and compact plaques, extracellular A β deposits often occur in another morphological form in AD, in the walls of cerebral and leptomeningeal blood vessels (Mandybur, 1975; Vinters et al., 1988). According to the amyloid cascade hypothesis, accumulation and deposition of A β in the brain is the primary influence and early process driving AD pathogenesis leading to neuronal dysfunction (Hardy and Higgins, 1992).

Increasing attention is turning towards small oligomeric aggregates that form initially before they assemble into fibrils or plaques (Fig. 1). These oligomeric A β assemblies intermediates are potent neurotoxins and therefore are thought to be the key effectors of neurotoxicity in AD (Klein et al., 2001). Importantly, such oligomeric forms of A β are neurotoxic *in vitro* (Oda et al., 1995; Walsh et al., 1997; Lambert et al., 1998; Hartley et al., 1999; Walsh et al., 1999; Nilsberth et al., 2001) and *in vivo* (Walsh et al., 2002). In transgenic mice expressing the human APP, neurological deficits develop even before amyloid deposits occur (Mucke et al., 2000). It has been shown recently that oligomers of A β even inhibit long-term potentiation *in vivo* (Walsh et al., 2002). Another intermediate in the pathway of A β fibril formation is the protofibril. Protofibrils are short assemblies of 5 nm in diameter and up to 200 nm in length (Harper et al., 1997; Walsh et al., 1997). They are not only important in terms of amyloid fiber formation but also in causing neuronal cell death (Hartley et al., 1999; Walsh et al., 1999). The role of these intermediates in amyloid fiber assembly will be discussed further in section 1.3.2.

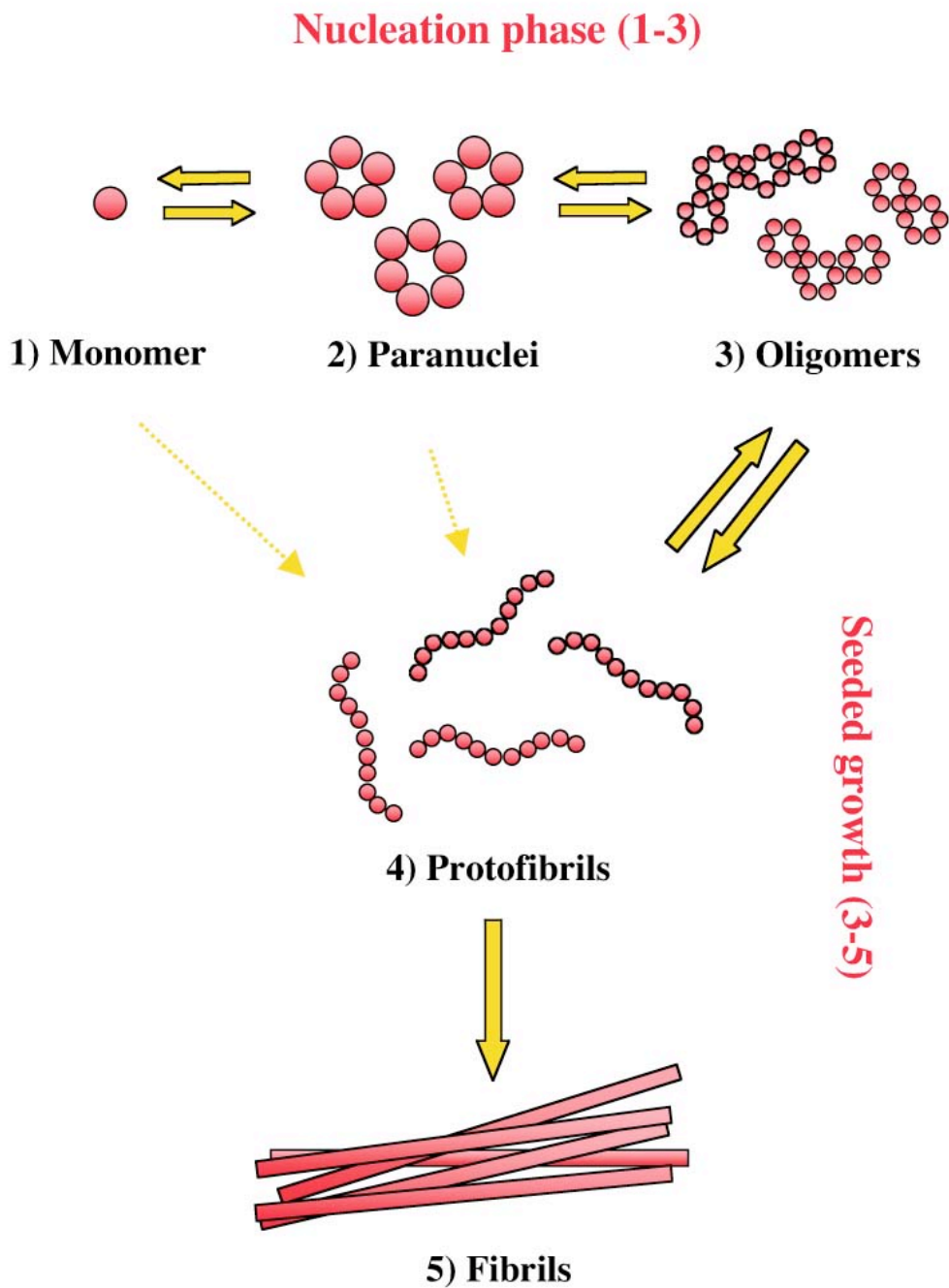


Figure 1. Model of A β assembly via nucleation-dependent polymerization. Monomers form via several steps an unstable oligomeric nucleus. Once the critical protein concentration is reached, seeded growth starts and generates amyloid fibrils via protofibrils. This direct pathway may not be the only one since the addition of monomers or paranuclei to protofibrils could also lead to the maturation into fibrils (adapted from Harper and Lansbury, 1997; Bitan et al., 2003).

Neurofibrillary tangles are intraneuronal cytoplasmic lesions consisting of hyperphosphorylated isoforms of the microtubule-associated protein tau, which assemble into poorly soluble paired helical 10-nm filaments (Goedert, 1998). Six tau isoforms are produced in the adult human brain by alternative mRNA splicing from a single gene located on chromosome 17 (Goedert et al., 1989; Andreadis et al., 1992). The six resulting polypeptides have a molecular mass ranging from 45kDa to 65kDa (Buee et al., 2000). Neurofibrillary tangles generally occur in large numbers in the AD brain, particularly in limbic and paralimbic structures such as the entorhinal cortex, hippocampus and amygdala. Intensive studies have shown that the tau protein, which normally enhances the polymerization of tubulin into microtubules and acts to stabilize these organelles in neurons, becomes excessively phosphorylated which reduces binding to microtubules (Lovestone and Reynolds, 1997). The tau and tangle hypothesis argues that in AD the normal role of tau in stabilizing microtubules is impaired, and in diseased neurons microtubules are replaced by tangles (Gray et al., 1987). The resultant cytoskeletal dysfunction in these neurons eventually leads to the degeneration of dendrites and a loss of synapses at their axonal projection targets. The observation that neurofibrillary tangles form in some cell bodies whose axons terminate in regions containing amyloid-bearing neuritic plaques suggests that neurofibrillary tangle formation in some perikarya and neurites may be related to events associated with amyloid plaque formation in the AD brain (Hyman et al., 1986; Rasool et al., 1986).

In addition to plaques and tangles, the AD brain is characterized by extensive neuronal degeneration. The cortex and hippocampus of AD patients undergo a marked atrophy (Mouton et al., 1998), region specific neuron and synapse loss (Terry et al., 1991; West et al., 1994; Gomez-Isla et al., 1996) and a loss of cholinergic innervation (Coyle et al., 1983). Yet, despite intensive interest in a potential association between the tangles, amyloid plaques, vascular amyloid and neurodegeneration observed in the AD brain, the relationship remains poorly understood.

1.2.2 Genetics of Alzheimer's disease

While the complete etiological picture of AD remains unresolved, the inheritance of predisposing genetic factors appears to play a major role. After age, family history is the second greatest risk factor for AD. To date, three genes have been identified whose mutations cause early-onset familial AD (FAD). These genes are the amyloid precursor protein (APP), presenilin 1 (PS1) and presenilin 2 (PS2) (Schellenberg, 1995). Studies on the amyloid precursor protein (APP) as a genetic determinant of AD have begun with the observation that individuals with Down's syndrome develop the clinical and pathological features of AD (Mann et al., 1985; Mann, 1988). These data pointed to the involvement of chromosome 21 in AD leading to the first genetic linkage discovery between a locus on chromosome 21q and autosomal dominant early-onset familial AD (St George-Hyslop et al., 1987). Sequencing of the APP gene and screening for mutations led to the discovery of several missense mutations in families with early-onset AD (Goate et al., 1991; Hendriks et al., 1992; Mullan et al., 1992). Although the APP mutations account for less than 0.1% of all AD cases (Tanzi and Bertram, 2001), they carry virtually complete penetrance leading to AD between the fourth and seventh decades of life.

Since most early-onset AD families do not have mutations in the APP gene, it was expected that other AD loci might exist. A second locus was linked to chromosome 14, and the gene was later identified and named presenilin 1 (PS1) (Van Broeckhoven et al., 1992; Sherrington et al., 1995). A second gene was found based on its homology to PS1 and mapped on chromosome 1 (Levy-Lahad et al., 1995b; Levy-Lahad et al., 1995a; Rogaev et al., 1995). Mutations in these two genes (PS1 and PS2) are thought to cause up to 80% of familial early-onset AD cases. To date there are more than 120 different mutations identified in the PS1 gene (<http://molgen-www.uia.ac.be/ADMutations>), while only 8 missense mutations have been identified in PS2. Early-onset familial AD mutations in PS1, PS2 and APP lead to an increase of secreted A β 42, the primary component of β -amyloid plaques in the brain (Price and Sisodia, 1998; Haass and De Strooper, 1999).

In addition to the mutations that have been associated with early-onset FAD, other genetic factors have been identified that increase the risk of developing sporadic AD. The $\epsilon 4$ allele of the apolipoprotein E (APOE) gene on chromosome 19 (Olaisen et al., 1982) was first discovered as a major risk factor for AD (Strittmatter et al., 1993a). In humans there are three common alleles of the APOE gene: $\epsilon 2$, $\epsilon 3$ and $\epsilon 4$. APOE normally plays a role in cholesterol transport and lipid metabolism (Mahley, 1988; Vancea et al., 2000). While the $\epsilon 4$ allele is strongly associated with increased neuritic plaques and cerebral amyloid angiopathy in AD (Olichney et al., 1996), the $\epsilon 2$ allele has been shown to have an impact on longevity and may confer protection against AD (Corder et al., 1994). Distinct binding properties to the A β peptide (Strittmatter et al., 1993b) have suggested ways by which APOE might mediate its action. Less than 50% of non-familial AD cases are carriers of the APOE $\epsilon 4$ allele (Corder et al., 1993) suggesting that other susceptibility genes must be involved in the pathogenesis of the disease (Shastri and Giblin, 1999). Most of these genes are located on chromosome 9, 10 and 12 (Okuizumi et al., 1995; Ertekin-Taner et al., 2000; Lambert et al., 2000) and are proteolytic enzymes, plasma proteins, growth factors or membrane receptors such as $\epsilon 2$ -macroglobulin ($\epsilon 2$ -M) (Blacker et al., 1998), low density lipoprotein receptor-related protein (LRP) (Okuizumi et al., 1995), angiotensin converting enzyme (ACE) (Kehoe et al., 1999) and insulin degrading enzyme (IDE) (Bertram et al., 2000; Ertekin-Taner et al., 2000). $\epsilon 2$ -M is thought to mediate the clearance and IDE has been implicated in the degradation of A β peptides (Vekrellis et al., 2000). Thus, all AD genetic factors discovered so far point to an alteration of A β production or clearance as a direct cause of the disease.

1.2.3 APP processing and A β

The isolation of A β from senile plaques and cerebrovascular deposits of AD brains (Glennner and Wong, 1984; Masters et al., 1985) and the cloning of the full-length cDNA of APP (Kang et al., 1987) were the beginning of the modern AD research era. APP is a type-I integral transmembrane glycoprotein that is ubiquitously expressed. Three major isoforms of APP are produced by alternative splicing: the two longer isoforms of 751 and 770 amino acids contain a protease inhibitor domain (Kitaguchi et al., 1988; Ponte et al., 1988; Tanzi et al., 1988), whereas the shorter form of 695 amino

acids lacks this Kunitz domain and is the main isoform expressed in neurons (Kang et al., 1987; Golde et al., 1990). The physiological function of APP in the brain remains unclear, although it has a putative function in transmembrane signal transduction (Nishimoto et al., 1993), calcium regulation (Mattson et al., 1993), cell proliferation (Saitoh et al., 1989), regulation of cell/cell or cell/matrix interactions (Breen et al., 1991), cell-adhesion (Jin et al., 1994), neurite outgrowth (Milward et al., 1992) and synaptogenesis (Mucke et al., 1994). Initial studies of APP knockout mice did not reveal an obvious disease phenotype (Zheng et al., 1995), however, other studies have suggested cognitive deficits (Dawson et al., 1999; Phinney et al., 1999) and impaired synaptic plasticity (Seabrook et al., 1999).

APP can be processed by several different proteases, namely the α -, β - and γ -secretases (Fig. 2). The α -secretase cleavage pathway corresponds to the default secretory pathway and is considered as non-amyloidogenic. α -secretase cleaves within the A β region to produce the soluble α -APPs (Esch et al., 1990; Sisodia, 1992) and the 83-residue COOH-terminal fragment C83. The identity of the α -secretase is unknown, although more than one enzyme may be involved as the preferential cleavage site varies in different cell types (Zhong et al., 1994). TACE, ADAM-10 and MDC-9, which all belong to the family of adamalysins have been postulated as α -secretase candidates (Buxbaum et al., 1998; Koike et al., 1999; Lammich et al., 1999; Lopez-Perez et al., 2001). The β -secretase cleavage corresponds to the alternative cellular pathway that process APP and constitutes the first step in the formation of A β peptides. In contrast to the α -secretase, the β -secretase generates the NH₂-terminus of A β , cleaving APP to produce a soluble version of APP and C99 which remains membrane-bound (Seubert et al., 1993). Two novel aspartyl protease homologous, BACE and BACE-2 were recently identified that cleave APP at the β -secretase sites (Sinha et al., 1999; Vassar et al., 1999). In addition, genetic experiments have demonstrated that BACE is required for A β peptide formation, confirming its identity as the β -secretase gene (Cai et al., 2001; Luo et al., 2001). Both the C83 and C99 fragments remain anchored in the membrane and may become degraded or further processed by γ -secretase to produce p3 from C83 or A β from C99. The γ -secretase cleavage is not sequence specific and can produce a variety of A β peptides. A β 40 is the most common A β peptide, while A β 42 is the

second most common form. The longer the peptide the more insoluble it gets and the more likely it is to form toxic fibrils and aggregates (Jarrett et al., 1993b). The identity of the γ -secretase remains elusive. However, it has been proposed that γ -secretase forms a large complex composed of presenilin, nicastrin, PEN-2, and APH-1, and that the activity of the γ -secretase depends on these proteins (Steiner and Haass, 2000; Yu et al., 2000; Francis et al., 2002; Lee et al., 2002; Steiner et al., 2002). Most recent studies have shown that APP is cleaved by PS-dependent γ -secretase not only in the middle of the transmembrane domain (β -cleavage) but also inside the cytoplasmic membrane boundary (β' -cleavage) (Gu et al., 2001; Sastre et al., 2001; Yu et al., 2001a; Weidemann et al., 2002). However, the functional significance of γ -secretase cleavage remains to be clarified.

Many different pathogenic mutations such as Swedish (Mullan et al., 1992), Flemish (Hendriks et al., 1992), Dutch (Levy et al., 1990), Florida (Eckman et al., 1997) and London (Goate et al., 1991) have been identified in APP, all of which are missense mutations lying at the three secretase cleavage sites or within the A β region, as shown in Figure 2. AD-causing mutations in APP near the β - and β' -secretase cleavage sites all increase A β 42. Those near the β -secretase cleavage site augment β -site proteolysis, leading to elevation of both A β 40 and A β 42 (Citron et al., 1992), whereas those near the β' -site increase the production of A β 42 (Suzuki et al., 1994).

While A β 40 is the major A β product of APP processing and the major component of the amyloid deposits of congophilic angiopathy (Joachim et al., 1988; Miller et al., 1993; Gravina et al., 1995), A β 42/43 has a much greater tendency to aggregate spontaneously into amyloid fibrils (Hilbich et al., 1991; Burdick et al., 1992; Jarrett and Lansbury, 1993) and is the predominant form of A β in senile plaque cores (Roher et al., 1993; Fukumoto et al., 1996). In addition, A β 42 displays enhanced neurotoxicity relative to A β 40 (Younkin, 1995; Selkoe, 1999; Dahlgren et al., 2002). Interestingly, A β 42 levels are reduced in the cerebrospinal fluid (CSF) of AD patients (Motter et al., 1995), which has been attributed to reduced clearance. Taken together, these findings implicate A β 42 in the pathogenesis of AD. The development of antisera specific for A β 40 and for A β 42 enabled plaque evolution and composition to be studied

systematically (Iwatsubo et al., 1994; Gravina et al., 1995). The results are consistent with a nucleation or seeding hypothesis (Jarrett et al., 1993a; Jarrett and Lansbury, 1993), whereby A β 42 forms the nucleus of a plaque initially, leading to the subsequent deposition of soluble A β 40 (Iwatsubo et al., 1994). The seeding model and its role in the pathogenesis of AD will be described in section 1.3.2.

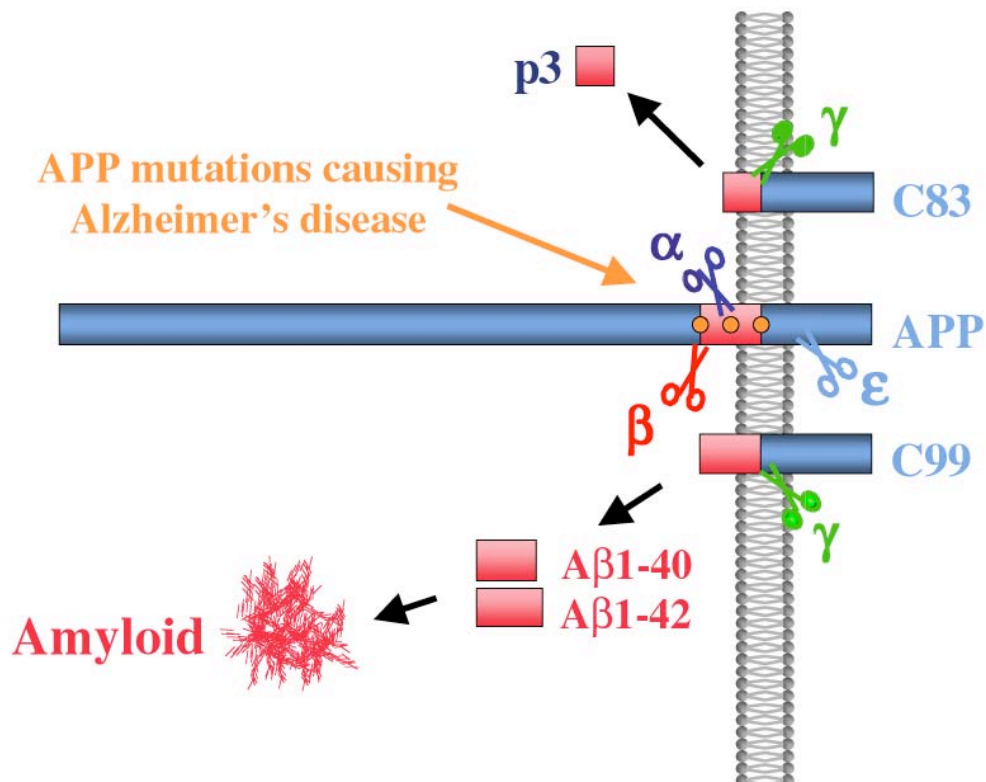


Figure 2. Proteolytic processing of APP. APP is a 770 amino acid type I transmembrane protein which is cleaved by different proteases. The initial cleavage by α -secretase is the most common cleavage, resulting in the release of the large extracellular domain. Since the α -secretase cleavage is in the middle of the A β domain, no A β can be generated. Instead, the remaining fragment (C83) is cleaved by γ -secretase and releases the small peptide p3. Less frequently, APP is cleaved by β -secretase and then in the transmembrane domain either at position 40 or 42 by γ -secretase. This β -secretase pathway is the first step in the formation of A β peptides and amyloid. There are many APP mutations known that cause familial AD. Note that these mutations are localized within the A β domain near the cleavage sites of the secretases.

1.2.4 Mouse models of Alzheimer's disease

The identification of mutations in specific genes has provided new opportunities to explore pathogenic mechanisms using genetically engineered mice. To generate animal models of A β amyloidosis, many groups have produced transgenic mice that express wild-type APP, APP fragments, A β and FAD linked mutant APP and PS1. Although early efforts were disappointing because transgenic mice did not exhibit abnormalities characteristic of AD, more recent work has shown that multiple lines of APP transgenic mice now exist that develop age-related deposition of cerebral amyloid similar to AD (Games et al., 1995; Hsiao et al., 1996; Sturchler-Pierrat et al., 1997). These mice have mild neuron loss, dystrophic neurites, glial responses, learning impairments and deficits in synaptic transmission and/or long-term potentiation (Hsiao et al., 1996; Irizarry et al., 1997; Calhoun et al., 1998; Frautschy et al., 1998; Chapman et al., 1999; Stalder et al., 1999; Bondolfi et al., 2002; Kelly et al., 2003). The disease severity is influenced by the level of transgene expression and the specific mutation. This is in line with the observation that mice expressing both mutant PS1 and mutant APP showed accelerated A β amyloidosis well before APP transgenic mice (Borchelt et al., 1997; Holcomb et al., 1998). APP transgenic mice have also been crossed with mice deficient in APOE. In contrast to the APP/PS1 double transgenic mice, these mice show a remarkable decrease in congophilic plaques, suggesting that APOE promotes the deposition and fibrillization of A β (Bales et al., 1997; Holtzman et al., 2000).

As APP transgenic models failed to develop tau filaments, several research groups have generated human wildtype tau transgenic mice (Gotz et al., 1995; Brion et al., 1999; Ishihara et al., 1999; Spittaels et al., 1999; Probst et al., 2000). However, none of these transgenic mice developed tau filaments. Only one group reported that their human wildtype tau transgenic mice developed tangle like structures at high age (Ishihara et al., 2001). The expression of P301L mutation was more successful in terms of pathology since these tau transgenic mice showed numerous neurofibrillary tangles (Lewis et al., 2000; Gotz et al., 2001b).

However, appropriate mouse models that display the entire neuropathological spectrum of AD have so far not been entirely successful (Wong et al., 2002). In an attempt to

obtain mice with both plaques and tangles, APP transgenic mice were bred with mice expressing the P301L mutant tau. Although the NFT pathology was substantially enhanced, plaque formation was unaffected by the presence of the tau lesions (Lewis et al., 2001). Similarly, tau pathology can be induced by introducing A β 42 fibrils into P301L tau mutant mice (Gotz et al., 2001a). Together, both experiments demonstrated pathological interactions between A β and tau that lead to increased NFT formation. Recently, triple-transgenic mice (PS1, APP and P301L) have been created that develop extracellular A β deposits prior to tangle formation in AD relevant brain regions (Oddo et al., 2003). However, more appropriate models of AD might require co-expression of mutant APP and all six isoforms of wildtype human tau.

In an effort to understand the functions of the AD related genes, researchers have ablated most of them. As mentioned above, homozygous APP $-/-$ mice show only a subtle phenotype (Zheng et al., 1995). PS1 $-/-$ mice fail to survive beyond the early postnatal period, but exhibit severe problems in the development of the axial skeleton and ribs (Shen et al., 1997). The lack of PS1 expression in the forebrain in conditional PS1-targeted mice resulted in decreased A β generation, further establishing that PS1 is critical for β -secretase activity in the brain (Feng et al., 2001; Yu et al., 2001b). BACE1 $-/-$ mice are healthy and have also no obvious phenotype, but secretion of A β peptides is abolished suggesting that BACE1 is the principal β -secretase in neurons (Cai et al., 2001; Luo et al., 2001; Roberds et al., 2001).

1.2.5 Therapeutic approaches

According to the amyloid cascade hypothesis, the accumulation of A β 42 in certain brain regions is an early and critical event in the pathogenesis of AD (Hardy and Selkoe, 2002). Therefore, anti-amyloid strategies are currently being investigated to lower the production of A β (either by inhibiting β - and γ -secretases or by favouring the β -secretase pathway over the γ -secretase pathway), or to clear the amyloid peptides by the use of defibrillating agents or by immunization (Fig. 3).

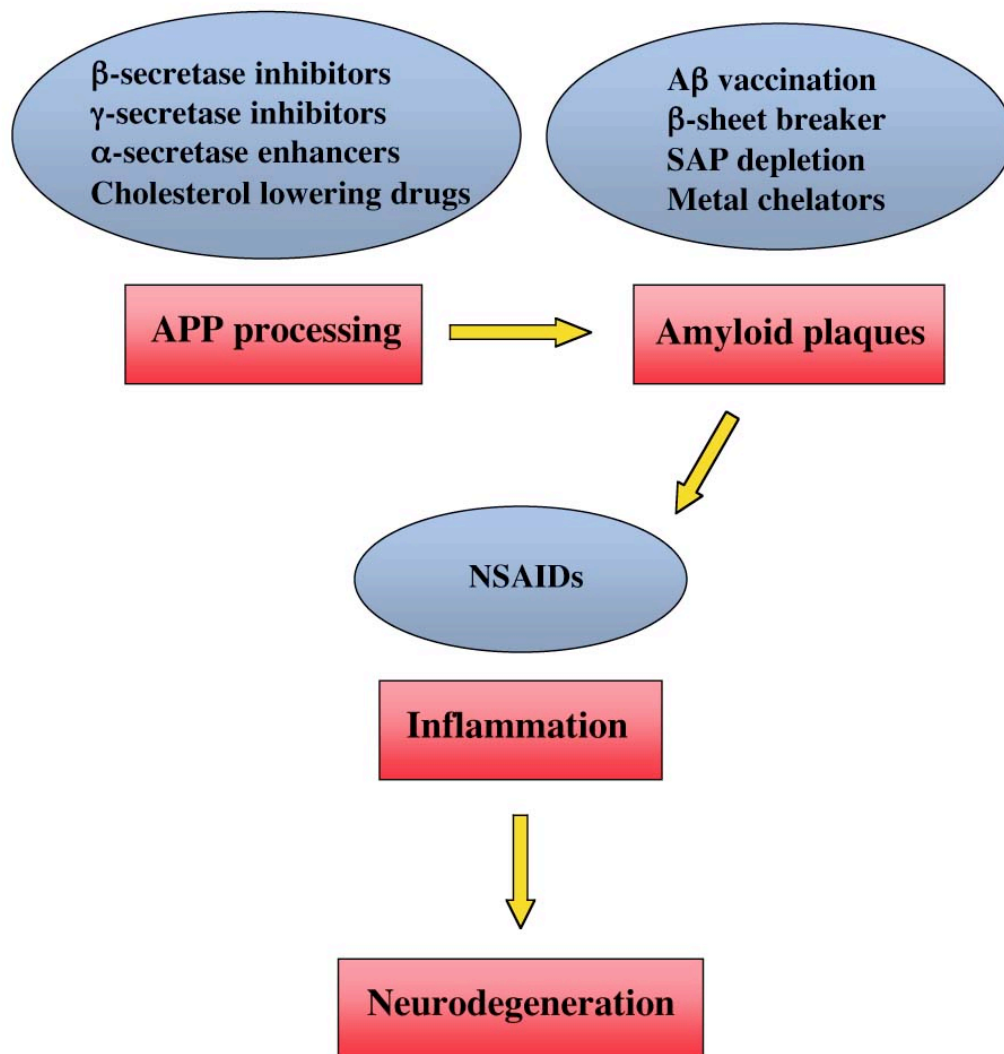


Figure 3. Summary of the main pathological processes and the related therapeutic approaches in AD. These approaches target the pathogenesis of the disease such as APP processing, amyloid plaques and inflammation. The treatment strategies include β - and γ -secretase inhibitors, α -secretase enhancers, cholesterol-lowering drugs and NSAIDs but also amyloid- β -peptide vaccination, β -sheet breakers and metal chelators (adapted from Scarpini et al., 2003).

The β -secretase BACE is thought to be an ideal therapeutic target as it catalyses the first step of $A\beta$ production. Inhibitors described so far are polypeptides that are effective *in vitro* (Sinha et al., 1999; Ghosh et al., 2001), however, these peptidic compounds do not penetrate the blood-brain barrier to a sufficient extent (Citron, 2002). γ -secretase also represents a potential therapeutic target. Potent inhibitors have recently been developed that are effective in preventing the formation of both $A\beta_{40}$ and $A\beta_{42}$ (Wolfe et al.,

1999; Esler et al., 2000; Seiffert et al., 2000; Shearman et al., 2000; Dovey et al., 2001). However, a limitation to the use of β -secretase inhibitors may be that they interfere with Notch signaling and possibly other cellular functions (De Strooper et al., 1999). A recent report that described compounds that inhibit A β formation without affecting Notch cleavage may offer new leads for the design of APP-specific β -secretase inhibitors (Petit et al., 2000). It has also been shown that β -secretase inhibitors reduce amyloid burden in animal models (Dovey et al., 2001). A subset of non-steroidal anti-inflammatory drugs (NSAIDs) are allosteric inhibitors of β -secretase (Weggen et al., 2001). This effect was independent of the intended effect of the NSAIDs, namely cyclooxygenase inhibition. Metabolism by β -secretase can be stimulated by the activation of several cell-surface receptors coupling to protein kinase C (Lee et al., 1995). Estrogens and testosterone were found to promote β -secretase cleavage of APP in cell culture and to reduce A β formation. Thus, hormone replacement therapy may have positive effects in preventing AD. Estradiol treatment is currently being investigated in animal models (Petanceska et al., 2000). *In vitro* and *in vivo* experiments have proposed a molecular link between AD and cholesterol (Simons et al., 1998; Frears et al., 1999; Puglielli et al., 2001). High cholesterol uptake increased A β deposition in transgenic mice and in rabbits (Sparks et al., 1994; Refolo et al., 2000). It has also been shown in animals that a cholesterol synthesis inhibitor reduced plaque formation in transgenic mice (Refolo et al., 2000). Therefore, cholesterol lowering drugs such as simvastatin could be an attractive treatment strategy for AD (Fassbender et al., 2001).

Another approach to reduce amyloid accumulation and toxicity in AD is the use of chemical agents that prevent A β oligomerization and the formation of neurotoxic protofibrils. A variety of compounds including Congo red (Lorenzo and Yankner, 1994), rifampicin (Tomiyama et al., 1996) and benzofurans (Soto, 1999) have been proposed but are not applicable *in vivo* due to their toxicity. The design of short synthetic peptides homologous to the central hydrophobic region of A β has been suggested to disrupt β -sheet stabilisation (Soto, 2001). Administration of this β -sheet breaker reduced the plaque load and ameliorated astrogliosis, microglial activation as well as neurodegeneration in APP transgenic mice (Permanne et al., 2002). Moreover,

this peptide did not induce antibody production and could cross the blood-brain barrier. Serum amyloid P component (SAP) is a glycoprotein that is present in amyloid deposits and seems to protect amyloid fibrils from degradation and clearance (Tennent et al., 1995). Recently, a small compound has been developed that dimerizes SAP, reduces its concentration in the plasma and destabilizes A β deposits *in vivo* (Pepys et al., 2002). Oligosaccharide (Fraser et al., 2001) and metal chelators (Cherny et al., 1999; Cherny et al., 2000) can also inhibit plaque formation and constitute potential plaque busters. The administration of clioquinol to transgenic mice reduced A β deposition (Cherny et al., 2001).

Apart from the prevention of plaque formation, the removal of deposited plaques has become possible through immunotherapeutic approaches. Transgenic mice have been successfully immunized either by active or passive immunization against A β (Schenk et al., 1999; Bard et al., 2000). This vaccination not only prevents the development of A β plaques, neuritic dystrophy and astrogliosis, but also markedly reduces the extent and progression of AD-like pathology. Importantly, further studies revealed that such vaccination can improve learning and memory deficits (Janus et al., 2000; Morgan et al., 2000; Younkin, 2001; Dodart et al., 2002). However, a recent phase IIa clinical trial for an AD vaccine (AN-1792) was stopped when 17 of 360 patients developed severe inflammatory side effects. One of the trial participants developed leptomenigeal infiltrates around amyloid-laden blood vessels with T lymphocytes (Nicoll et al., 2003). In summary, the development of anti-A β therapeutics remains a rational approach to treat AD, based on our current understanding of the earliest features of the disease. Despite all difficulties, current data from various laboratories indicates that A β -targeted therapy is the most likely to be effective in reducing amyloid-related pathology in AD.

1.3 Studying cerebral amyloidosis *in vivo*

1.3.1 Intracerebral grafting

Grafting tissue into the brain as a possible therapy for neurodegenerative disease has been of enormous interest for many scientists in the past and present research. Parkinson's patients especially have benefited from such therapeutic approaches (Bjorklund, 1991; Freed et al., 1992; Lindvall et al., 1992; Freed et al., 1993; Koutouzis et al., 1994). What is often ignored is the enormous benefit grafting techniques can provide basic research with. For many scientists, intracerebral grafting is a tool used to learn more about neuronal plasticity of the adult nervous system, and as a technique to establish principles of neuronal development. To these ends, intracerebral grafting is established as a powerful method to study basic problems in neuroscience (Fig. 4).

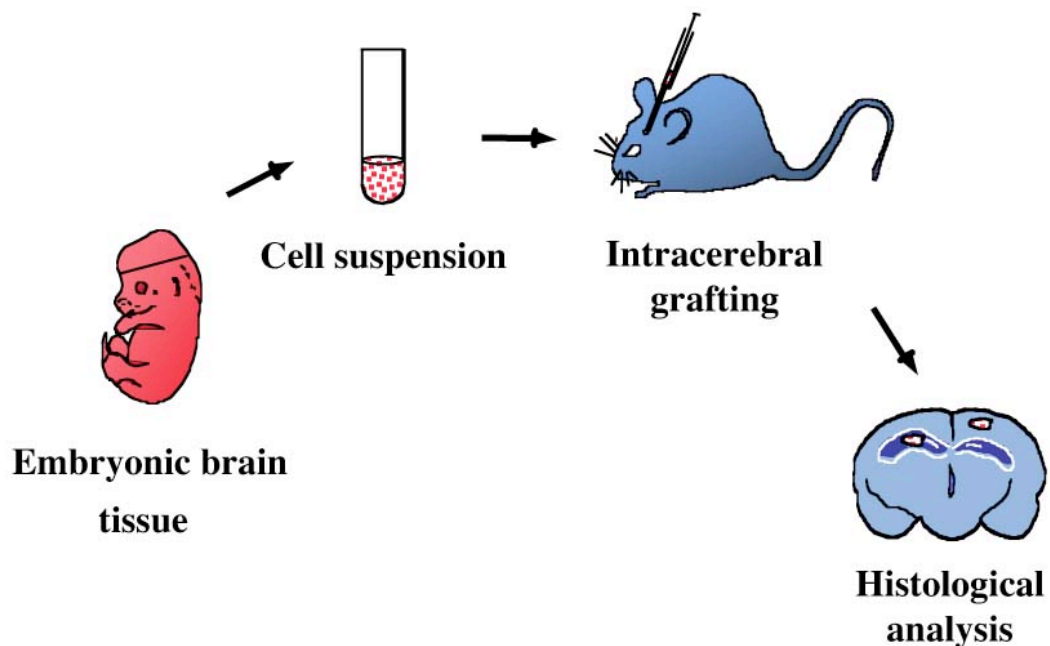


Figure 4. Schematic diagram of the transplantation procedure. Cortical and hippocampal tissue is dissected from E16-17 embryos and then mechanically dissociated into a cell suspension. Several 2.5 μ l aliquots of this cell suspension are stereotactically injected via a microsyringe into the neocortex and hippocampus of the host animal. Histological analysis are performed after different time points.

Many important issues concerning the biological requirements for graft survival and optimal function of the graft have been identified. It has been postulated that for most donor tissues, the younger the cells are, the better their chance of survival (Gage et al., 1983; Brundin et al., 1985; Barker et al., 1995). This is particularly true for neuronal tissue, which explains the focus of attention on fetal cells in intracerebral grafting. An equally important factor is the age of the host brain. While grafts will survive in the adult and aged brain (albeit less well in the aged brain), significantly better survival is observed in the young neonatal brain. Both the limitations of donor age and host may be attributed in part to the availability of trophic factors which can address specifically the survival and growth of grafted tissue independently of many other issues (Gage et al., 1984; Mayer et al., 1993). In order for a graft to survive and integrate in the brain, it must be appropriately vascularized and thereby receive essential nutrients from the host environment (Broadwell et al., 1987; Dusart et al., 1989).

While the above factors can be used to influence the survival of the graft, a major objective is to establish graft function in the host brain. One such function is the interaction with the host brain, which should provide at least partial replacement for missing connections. Alternatively, the graft may serve as an endogenous minipump which secretes a missing transmitter (Horellou et al., 1991; Campbell et al., 1993; Cenci et al., 1994) or a trophic factor, which directly affect the survival and growth of damaged cells (Gage et al., 1984). Finally, the graft can function as a bridge that contributes to neuronal and axonal connections (Zimmer et al., 1986; Zimmer et al., 1988).

Before grafting can be considered as a likely approach to cure Alzheimer's disease, the cause of this neurodegenerative disease needs to be determined. With the availability of experimental animal models which mimic the pathological characteristics of the disease, intracerebral grafting could be used to address the pathophysiology of AD. Therefore, intracerebral grafting became a good tool to study the pathogenesis of plaque formation and the underlying and associated pathology.

1.3.2 Seeding mechanism

Kinetic studies have shown that the aggregation of A β follows a seeding/nucleation mechanism (Jarrett and Lansbury, 1993), which resembles a crystallization process. The critical event is the formation of protein oligomers that act as a nucleus and which direct further growth of aggregates. Nucleation-dependent polymerization involves two stable states, the monomeric protein and the fibril (Harper and Lansbury, 1997). There is first a slow nucleation phase in which the protein forms an ordered oligomeric nucleus and then a growth phase in which the nucleus rapidly grows to form larger polymers. Polymers are not observed until the monomer concentration exceeds a certain level known as the critical concentration. Below this critical concentration, the monomer is the predominant species. At protein concentrations that exceed the critical concentration by a small amount, there is a lag time before polymerization occurs. During this lag time, addition of pre-formed nuclei or seeds will immediately induce fibril formation. This nucleation-dependent polymerization postulates that aggregation is dependent on protein concentration and time.

At least two intermediates have been identified in this process from the native monomeric protein to the fibrillar, fully aggregated structure *in vitro* (Teplow, 1998). The first are soluble oligomers, which have been detected in test tube experiments, in the medium of A β secreting cells and in human cerebro-spinal fluid CSF and brain homogenate (Levine, 1995; Kuo et al., 1996; Lambert et al., 1998; Walsh et al., 2002). The second intermediates are short structures termed protofibrils, which have been studied by electron microscopy and atomic force microscopy (Walsh et al., 1997; Walsh et al., 1999). Protofibrils are 5 nm wide and up to 200 nm long. Protofibrils are the direct precursor of amyloid-like fibrils. Secondary structure studies show that protofibrils have a high β -sheet content like fibrils (Walsh et al., 1999). Evidence indicates that these intermediates, the monomers, and the fibrils are all present in a dynamic equilibrium (Teplow, 1998).

1.3.3 Magnetic resonance imaging

Magnetic resonance imaging (MRI) is a special, non-invasive technique used to image internal structures of the body, particularly the soft tissues. Unlike conventional radiography and computed tomography (CT), which make use of potentially harmful X-rays to generate images, MRI is based on the magnetic spin properties of atoms. In essence, MRI produces a map of hydrogen distribution in the body. Hydrogen is the simplest element known and the most abundant in biological tissues. It will align itself within a strong magnetic field, like the needle of a compass. The earth's magnetic field is not strong enough to keep a person's hydrogen atoms pointing in the same direction, but the superconducting magnet of an MRI machine can. This comprises the "magnetic" part of MRI. Once the hydrogen atoms have been aligned in the magnet, they absorb energy emitted in the form of specific radio wave frequencies, which knocks them back out of the alignment. Thus, the atoms alternately absorb and emit radio wave energy, vibrating back and forth between their resting (magnetized) and their agitated (radio pulse) state. This comprises the "resonance" part of MRI. The equipment records the duration, strength, and source location of the signals emitted by the atoms as they relax and translates the data into an image on a television monitor. The state of hydrogen in diseased tissue differs from healthy tissue of the same type, making MRI particularly good for the diagnosis of many disease processes due to high soft-tissue contrast and discrimination.

Recent advances in MRI technology with high magnetic fields and the introduction of transgenic mouse models of neurodegenerative diseases in MRI studies produced a great interest for identifying alterations in cellular pathology. MRI can not only provide high-resolution anatomical details on the distribution of the pathology but also pathological information through additional parameters such as relaxation times, cerebral blood flow (CBF) and apparent diffusion coefficient (ADC). As the change of ADC in brain tissue was measured in one of the studies presented in this thesis, it will be briefly discussed further. ADC describes the self-diffusion of water molecules, which is characterized by Brownian motion in the restricted environment. In biological systems, most of the diffusive motion of water molecules is restricted by the structures in the microenvironment, including cell membranes. In many pathological conditions, ADC

has been shown to be a very sensitive parameter reflecting changes in the cellular environment including cell membrane disruption and cell swelling (Norris, 2001). A more recent variant of diffusion MRI is diffusion tensor imaging (DTI) (Basser et al., 1994), which is an important method to map and characterize white matter tracts in the brain (Mori et al., 2001). This new technique uses the anisotropy of ADC. When water molecules are unconstrained, as in the CSF, the diffusion is equal in all directions, i.e., isotropic. In contrast, when water molecules are in highly organized tissue structures, such as white matter fiber tracts that restrict the self-diffusion of water, the resulting diffusion is directional or so called anisotropic. DWI determines this anisotropy and plays therefore an important role in the characterization of neuronal disease in humans and in animal models (Kauppinen et al., 1993).

The recent technical advances of *in vivo* MRI, including the availability of higher magnetic fields permitting improved spectral and spatial resolution are of great use to examine morphological changes noninvasively. With the potential of MRI and the availability of a number of transgenic and rodent models of neurodegeneration, we can gain further understanding of the brain pathogenesis of neurodegenerative disorders and the therapeutic strategies, which will eventually benefit humans.

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2. Extracellular amyloid formation and associated pathology in neural grafts

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2.1 Abstract

Amyloid precursor protein (APP) processing and the generation of β -amyloid peptide (A β) are important in the pathogenesis of Alzheimer's disease. While this has been studied extensively at the molecular and cellular levels, much less is known about the mechanisms of amyloid accumulation *in vivo*. We transplanted transgenic APP23 and wildtype B6 embryonic neural cells into the neocortex and hippocampus of both B6 and APP23 mice. APP23 transgenic grafts into wildtype hosts do not develop amyloid deposits up to twenty months post-grafting. In contrast, both transgenic and, surprisingly, wildtype grafts into young transgenic hosts develop amyloid plaques as early as three months post-grafting. Although largely diffuse in nature, some of the amyloid deposits in wildtype grafts were congophilic and were surrounded by neuritic changes and gliosis similar to the amyloid-associated pathology previously described in APP23 transgenic mice. Our results suggest that diffusion of soluble A β in the extracellular space plays an important role in the spread of A β pathology and that extracellular amyloid formation can lead to neurodegeneration.

2.2 Introduction

Alzheimer's disease (AD) is a late onset progressive neurodegenerative disorder that is characterized by aggregation of A β into senile plaques and cerebrovascular amyloid. Genetic studies and transgenic (tg) mice support the view that the production and/or deposition of A β is an early and critical process in AD pathogenesis that triggers a cascade of pathological events leading to neurofibrillary tangle formation, neuroinflammation, synapse and neuron loss, and dementia (Selkoe, 2001; Hardy and Selkoe, 2002). A matter of controversy, however, is whether this cascade is initiated by the accumulation of A β in the extracellular space or by intraneuronal A β generation (Martin et al., 1994; Hartmann et al., 1997; Geula et al., 1998; Gouras et al., 2000; Kane et al., 2000; Walsh et al., 2000; Bayer et al., 2001; D'Andrea et al., 2001; Selkoe, 2001; Wirths et al., 2001).

In an attempt to study mechanisms and early stages of abnormal protein deposition and related neurotoxicity, neural transplantation techniques have previously been applied to study prion encephalopathy (Brandner et al., 1996b). In AD research, similar approaches have been pursued, but no convincing amyloid pathology has been associated with the grafts (Richards et al., 1991; Holtzman et al., 1992; Mantione et al., 1995; Bayer et al., 1996). However, the recent generation of APP tg mice that develop robust cerebral amyloid with aging (Games et al., 1995; Hsiao et al., 1996; Sturchler-Pierrat et al., 1997) has opened new opportunities to study the mechanism of cerebral amyloidosis by using neurografting techniques. To test the hypothesis that transplanted neural tissue would retain its native properties regardless of host genotype, we transplanted wildtype (wt) and APP tg embryonic cortical and hippocampal brain tissue into the brains of both APP tg and wt mice. Our results suggest that the phenotype of the transplanted tissue is strongly influenced by the properties of the host, and suggest a key role of extracellular diffusion of A β in cerebral amyloidogenesis. Moreover, we conclude that extracellular amyloid formation is closely associated with neurodegeneration.

2.3 Results

APP23 grafts in B6 hosts do not develop amyloid up to 20 months post-grafting

Cell suspensions of APP23 tg embryonic cortical and hippocampal tissue were injected into the neocortex and hippocampus, respectively, of young, 3 month-old B6 wt mice. Grafts were analyzed 3-24 months later. All grafts integrated well into the surrounding host tissue and appeared healthy and viable by morphological analysis. Only minimal gliosis was found at the graft-host interface. Nevertheless, grafts could easily be identified in cresyl-violet stained sections due to differences in cytoarchitecture between graft and host tissue (Fig. 1A,D,G). Immunohistochemistry revealed robust and exclusive expression of human APP (hAPP) in the graft, where hAPP was restricted to neurons and their processes, consistent with the neuron-specificity of the Thy-1 promoter used to generate APP23 tg mice (Fig. 1B,E,H). Cellular hAPP expression within the grafts looked very similar to that in adult APP23 mice.

Surprisingly, in light of the observation that amyloid deposition in neocortex and hippocampus of APP23 mice starts at the age of 6 months (Sturchler-Pierrat et al., 1997), no amyloid formation was observed in hippocampal or cortical grafts up to 20 months post-grafting (Fig. 1C,F and Table 1). Beyond 20 months, i.e. when hosts were more than 23 months of age, we identified two out of nine mice that developed massive amyloid in the parenchyma and vasculature within at least one graft (Fig. 1I). Interestingly, some amyloid was found outside the grafts, in particular in vessels. To study whether it is the age of the host or the post-grafting time that determines amyloid deposition in the grafts, we also placed tg grafts in adult and aged wt hosts (Table 1). Again, grafts appeared healthy, viable and, well integrated into the host. However, no amyloid formation was observed in such grafts, suggesting that it is not the age of the host but the time of the grafts residing in the host that determines amyloid formation.

APP expression and A β levels in APP23 grafts

We have tested several hypotheses to explain the delay/lack of amyloid formation in neocortical and hippocampal APP23 tg grafts in wt hosts compared to the cerebral amyloidosis observed in APP23 mice at the age of 6 months. First, we have tested the hypothesis that hAPP expression may be downregulated in grafted neurons compared to

the expression in normal adult APP23 mice (Fig. 2A,B). However, no obvious differences in hAPP expression were found in micropunches taken from neocortical and hippocampal tg grafts compared to micropunches taken from neocortex and hippocampus of normal adult APP23 mice.

Second, we tested the hypothesis that alterations in the processing of hAPP in grafted neurons may lead to a reduction in A β production and/or a shift in the ratio of A β 1-40 to A β 1-42. To this end, A β was immunoprecipitated from micropunches taken from tg grafts and compared to micropunches taken from 4-6 month-old APP23 mice, at an age before these mice develop amyloid deposits. Results revealed similar A β levels in the grafts compared to young APP23 mice with a predominance of A β 1-40 over A β 1-42 (Fig. 2B). Differences of A β 1-40 levels in the grafts compared to young APP23 mice ranged from -31% to +28% (mean -2% \pm 17%; $P > 0.05$). Interestingly, in the oldest transplant analyzed (20 months post-grafting), there was a relative increase of A β 1-42 compared to A β 1-40 (Fig. 2C). This observation is consistent with the result obtained in aged amyloid-depositing APP23 mice. Overall, these results do not support the hypothesis that reduced A β production by the grafted tissue is responsible for the slow onset of amyloid deposition in APP23 grafts in wt mice vs. normal APP23 mice.

Third, we have followed the hypothesis that intracerebral grafting of tg hAPP/A β producing tissue might evoke a humoral immune response in a wt host that has never seen hAPP/A β before (Brandner et al., 1996a), which might prevent amyloid deposition in the transplant (Schenk et al., 1999). However, serum titers of anti-A β antibodies in engrafted mice were not different from serum titers of normal B6 mice (both $< 1:100$). As a positive control we have used sera of APP23 mice passively immunized with A β antibodies with titers of 1:4,000 to 1:20,000 (Pfeifer et al., 2002).

APP23 and B6 grafts in APP23 hosts develop amyloid 3 months post-grafting

To study the influence of the host upon amyloid deposition in the graft, we then intracerebrally grafted cortical and hippocampal APP23 tg and B6 wt control tissue into the cortex and hippocampus, respectively, of 6 month-old APP23 mice (Fig. 3 and Table 2). Again, grafts integrated well into the surrounding host tissue and appeared

healthy and viable by morphological analysis with no evidence of neurodegeneration. Strikingly, after only 3 months post-grafting, tg grafts developed significant amounts of amyloid (Fig. 3A-C). Even more surprising, amyloid deposition was also found in wt cortical and hippocampal grafts 3 months post-grafting (Fig. 3D-I and Table 2). Amyloid deposition occurred throughout the grafts, with the greatest accumulation along the border of the graft. At this age the host tissue has also developed amyloid plaques, but in most animals they were significantly fewer than in the grafts. Semiquantitative analysis of double-stained sections for A β and Congo red indicated that 50% and 12% of the amyloid was compact in tg and wt grafts, respectively (Fig. 3I insert). Immunohistochemical analysis with antibodies specifically to A β _{x-40} and A β _{x-42} revealed the presence of both A β species in the graft with a predominance of A β _{x-40} over A β _{x-42} (Fig. 4A,B). Immunolabeling with an antibody that recognizes mouse but not human A β showed very faint labeling of the amyloid deposits suggesting that minimal amounts of mouse A β may contribute to the amyloid in wt grafts (Fig. 4C), as was also seen for the host tissue amyloid. A β was also immunoprecipitated from micropunches taken from amyloid-bearing wt grafts. Results revealed again both human A β ₁₋₄₀ and A β ₁₋₄₂ in the graft similar to that seen in aged amyloid-depositing APP23 mice (Fig. 4D,E).

Mechanism of amyloid formation in B6 grafts

To study whether the site of the graft placement may influence A β formation in the grafts, we injected hippocampal and cortical wt tissue into either the thalamus or striatum of 6 month-old APP23 mice ($n = 4$ for thalamus and $n = 8$ for striatum). Grafts were analyzed 3 months later. Thalamus and striatum were selected because they markedly differ in amyloid deposition in APP23 mice. The thalamus develops significant amyloid deposition in aged APP23 mice, while amyloid deposition in the striatum is low even in aged APP23 mice (Sturchler-Pierrat et al., 1997; Calhoun et al., 1998). Consistent with the idea that the properties of the surrounding host tissue influence amyloid deposition in the graft, our results revealed that 2 out of 4 mice (50%) developed amyloid in the “thalamic” grafts, while 0 out of 8 mice (0%) developed amyloid in the “striatal” grafts (Fig. 5). To examine whether the influence of the surrounding host tissue may be explained by differences in soluble extracellular A β

levels among the engrafted brain regions, extracellular soluble A β -40 and A β -42 was measured by ELISA in thalamus, striatum, neocortex, and hippocampus of 6 month-old pre-depositing APP23 mice. Results revealed the highest A β levels in hippocampus (A β -40: 2.61 ng/g wet weight; A β -42: 0.41 ng/g) and neocortex (A β -40: 2.42 ng/g; A β -42: 0.42 ng/g), followed by the thalamus (A β -40: 1.94 ng/g; A β -42: 0.32 ng/g). The levels in the striatum (A β -40: 0.71 ng/g; A β -42: 0.15 ng/g) were more than 50% lower compared to these other regions. Consistently, recent microdialysis analysis of A β also revealed significant lower extracellular soluble A β concentrations in striatum compared to hippocampus (Cirrito et al., 2002). Thus, soluble extracellular A β levels in the host may determine amyloid deposition in the graft suggesting that A β is transported extracellularly from the host into the graft. We also tested the hypothesis that the grafts in thalamus and striatum differ in levels of A β -degrading enzymes. However, no evidence of a difference was found using immunohistochemical analysis of insulin-degrading enzyme (Qiu et al., 1998) and neprilysin (Iwata et al., 2000) expression in graft vs. host (results not shown), although expression levels do not necessarily reflect enzyme activity.

We have also addressed several other hypothetical mechanisms to how A β may be transported from the host into the graft. We considered that hAPP/A β is anterogradely transported from host tg neurons into the wt graft. It has been described previously that host axons can penetrate a homotypic transplant, although in very low numbers (Clinton and Ebner, 1988). Moreover, APP is anterogradely transported to synaptic sites where it is released (Lazarov et al., 2002). Thus, we have used Holmes silver staining to visualize processes between host and graft. Overall, fiber density within the graft was much lower compared to the host. Although a few fibers were found to cross the host-graft border, most of the fibers were confined to their host or graft compartments (Fig. 6A). Moreover, immunohistochemistry did not reveal any hAPP-positive fibers that penetrated significantly into wt grafts (Fig. 6B, but see also Fig. 3E,H).

We also considered the possibility that microglia transport APP/A β from the host into the graft. Thus, not the graft *per se* but the lesion-associated gliosis or inflammation may induce amyloid accumulation in wt grafts. In favor of this explanation is the

observation that amyloid was predominantly observed at the border of the graft, an area of a slightly higher density of microglia compared to the rest of the graft. To test this possibility, we performed stab wounds in the neocortex and hippocampus of 6 month-old APP23 mice and analyzed the mice 3 months later. However, in none of lesioned mice ($n=4$) was amyloid found despite appreciable microgliosis around the lesion site, which was more extensive compared to that seen at the host-graft border (Fig. 6C,D). Moreover, we have injected three APP23 mice with wt grafts in which cells were killed by freezing before transplantation. Again, no amyloid accumulation was found although microgliosis was clearly present (not shown). Thus, surgical trauma alone is insufficient to induce β -amyloidosis in APP23 mice.

These results favor the idea that soluble and diffusible A β is transported extracellularly via interstitial fluid from the host into the graft where A β undergoes fibril formation and deposition. To exclude the hypothesis that host-derived soluble extracellular A β is internalized by wt neurons in the graft and that such internalized A β is the nidus and a prerequisite for extracellular amyloid plaque formation, we have used immunohistochemical staining with a variety of antibodies to A β , including antibodies previously suggested to recognize intracellular human A β (Wirhns et al., 2002). However, no evidence for intracellular human A β in the wt graft was found (Fig. 6E,F).

Amyloid-associated pathology in B6 grafts

The observation of amyloid in wt grafts allowed us to study the impact of host-derived extracellular amyloid formation on neurodegeneration. Although most of the amyloid in wt grafts was of the diffuse type and did not induce any notable neuropil changes, some compact and congophilic amyloid plaques were observed (Fig. 3I insert, 7A). Only a few neuronal cells were observed in the area proximal to compact plaques and some of them showed signs of degeneration (Fig. 7B). Moreover, plaques were surrounded by dystrophic synaptophysin-positive boutons (Fig. 7C) and abnormal thickened acetylcholinesterase-positive processes (Fig. 7D). A subpopulation of these distorted neuritic structures were positive for hyperphosphorylated tau (Fig. 7E). Signs of gliosis and neuroinflammation were also evident by clusters of darkly stained complement receptor 3 (CD11b)-positive and ionized calcium binding adaptor molecule 1 (Iba1)-positive microglia cells with hypertrophic processes that cover the amyloid core (Fig. 7F,G). Hypertrophic reactive glial fibrillary acidic protein (GFAP)-positive astrocytes

were also observed around the amyloid deposits (Fig. 7H). Overall, these changes appeared similar to the amyloid-associated degeneration, microgliosis and reactive astrocytosis previously described in amyloid-depositing APP23 mice (Sturchler-Pierrat et al., 1997; Calhoun et al., 1998; Phinney et al., 1999; Stalder et al., 1999; Boncristiano et al., 2002; Bondolfi et al., 2002).

2.4 Discussion

Whether cerebral amyloidosis is initiated by a gradual increase in extracellular A β or by the formation of intracellular A β aggregates with subsequent accumulation in the extracellular space (ECS) is controversial (Martin et al., 1994; Hartmann et al., 1997; Geula et al., 1998; Gouras et al., 2000; Kane et al., 2000; Walsh et al., 2000; Bayer et al., 2001; D'Andrea et al., 2001; Wirths et al., 2001). Our results suggest that local intracellular A β is not a prerequisite for extracellular amyloid deposition and that amyloid deposition is not necessarily confined to the region of A β production. Rather, A β can be transported and/or diffuses considerable distances in ECS of the brain before it aggregates, deposits extracellularly and causes neurodegeneration.

First, we have grafted APP23 tg tissue into wt hosts. Because cerebral amyloidosis in the neocortex and hippocampus of APP23 tg mice appears at six months of age (Sturchler-Pierrat et al., 1997), our initial expectation was that tg neocortical and hippocampal grafts placed in wt hosts would also develop amyloid at about six months post-grafting. However, no A β deposition was observed in the tg grafts up to twenty months post-grafting despite APP/A β production in the tg graft similar to APP23 tg mice. Thus, a reasonable explanation for this lack of or delay in amyloid formation is that extracellular A β in the graft does not reach high enough concentrations for amyloid formation due to movement of extracellular soluble A β out of the graft into the large volume of surrounding host tissue. It has previously been shown that extracellular A β is transported via interstitial fluid to the vasculature where A β is cleared by drainage along the perivascular spaces into the lymph nodes and/or by transport through the blood-brain-barrier into the blood (Weller, 1998; Calhoun et al., 1999; Shibata et al., 2000; DeMattos et al., 2002). That A β is transported from the tg graft into the wt host is consistent with the observation of amyloid outside the tg graft, in particular in vessel walls.

Second, the most compelling evidence that A β is extracellularly transported between host and graft comes from our observation that wt grafts in tg hosts exhibit amyloidosis. The amyloid in wt grafts is largely human A β and thus transgene- and host-derived. Moreover, amyloid in wt graft only develops when grafts are placed in brain areas such

as neocortex, hippocampus, and to some extent thalamus, that have relative high levels of soluble extracellular A β , but not when placed into the striatum, which has significant lower concentrations of extracellular soluble A β . Finally, axonal transport and transport by microglia of A β from the host into the graft did not appear to be significant. Thus, our results argue strongly that soluble A β from the tg host is transported extracellularly via the interstitial fluid from the host into the graft where A β then undergoes deposition. Recent findings have suggested that synaptic dysfunction may precede frank neuronal degeneration in AD and that this dysfunction may be caused by extracellular, diffusible oligomeric assemblies of A β (Selkoe, 2002). Our results would also suggest that such soluble A β aggregates may diffuse over considerable distances in the ECS and therefore may have an unexpected ability to impact synapses distal to the site of A β production.

It could be argued that host-derived human A β may be internalized by neurons within the wt graft (Bi et al., 2002; Nagele et al., 2002) and that such internalized human A β may undergo fibril formation intracellularly prior to being released and providing the nidus for extracellular amyloid formation (Gouras et al., 2000; Walsh et al., 2000; Bayer et al., 2001; D'Andrea et al., 2001; Wirths et al., 2001). However, we found no evidence of intracellular human A β in wt grafts. Additionally, we did not find evidence that the amyloid in wt grafts consisted of significant amounts of mouse A β , which might have originated in neurons within the wt graft. Moreover, we have previously reported no difference in amyloid deposition between APP23 mice on a wt background and APP23 mice on a *App*-null background (Calhoun et al., 1999). Thus, intraneuronal A β does not appear to be a prerequisite for extracellular amyloid formation. Nevertheless, a contributing role of intraneuronal A β to extracellular amyloid deposition cannot be entirely ruled out. The relatively lower amount of compact amyloid in the wt grafts vs. tg grafts (both in tg hosts) may be explained by amyloid formation initiated by an intracellular nidus leading to predominantly compact amyloid plaques while the formation of amyloid in the ECS is largely of the diffuse type. However, these differences may simply be explained by an A β concentration difference given that the tg graft produces A β in addition to that provided by the surrounding host tissue. The unexpected observation that wt grafts often had amyloid deposition prior to the tg host tissue, suggests that the graft provides a particularly favorable environment for

amyloidogenesis. Increased levels of extracellular matrix proteins in the ECS of neural grafts (Gates et al., 1996; Sykova et al., 1999) may act as chaperones for A β fibril formation or promote amyloidogenesis by inhibiting interstitial fluid transport of A β to the vessels and subsequent drainage (Snow et al., 1994; Gates et al., 1996; Weller, 1998; Sykova et al., 1999). Using diffusion-weighted MRI, we have recently shown that amyloidosis in APP23 mice is closely related to a decrease in brain ECS diffusion properties (Mueggler *et al.*, unpublished data). Interestingly however, the induction of extracellular matrix constituents by a stab wound or by the injection of dead cells does not promote amyloid deposition in APP23 tg mice. Thus, in future studies it will be important to characterize the factors that promote amyloidogenesis in the wt grafts. Identifying such amyloid-promoting factors may provide additional targets for amyloid-lowering therapies.

Whether neurodegeneration is related to intracellular A β generation or to its extracellular deposition is key to understanding the pathobiology of AD (Selkoe, 2001). Oligomerization and accumulation of intracellular A β ₁₋₄₂ in AD has been consistently reported (Hartmann et al., 1997; Wild-Bode et al., 1997; Gouras et al., 2000; Walsh et al., 2000; Wirths et al., 2001). Based on these findings, it has been suggested that accumulation of A β ₁₋₄₂ in intracellular compartments may lead to neuronal death (Walsh et al., 2000; D'Andrea et al., 2001; Wirths et al., 2001). However, our findings of amyloid-associated nerve cell degeneration, with dystrophic neuritic and synaptic structures including hyperphosphorylated processes in the vicinity of compact amyloid deposits in wt grafts, suggest that intracellular A β is not a prerequisite for neurodegeneration. In fact, amyloid pathology in wt grafts is very similar to that previously described in APP tg mice, including neuron death in the vicinity of amyloid plaques (Calhoun et al., 1998; Bondolfi et al., 2002; Urbanc et al., 2002), synaptic abnormalities (Phinney et al., 1999), dystrophic cholinergic processes (Wong et al., 1999; Boncristiano et al., 2002), and tau hyperphosphorylated distorted neuritic structures (Sturchler-Pierrat et al., 1997). Moreover, the gliosis and the increase in neuroinflammatory markers was also similar between wt grafts with amyloid and amyloid-depositing APP tg mice (Sturchler-Pierrat et al., 1997; Frautschy et al., 1998; Stalder et al., 1999).

In conclusion, our grafting experiments demonstrate that cerebral β -amyloidosis does not require locally generated intracellular A β to initiate A β deposition. Indeed, transport or diffusion of A β in the ECS of the brain is sufficient and can lead to amyloid pathology at a considerable distance from the site of A β generation. Overall, our findings demonstrate that factors beyond those directly involved in cellular A β production and intracellular A β metabolism are important for amyloidogenesis *in vivo*. The amyloid-associated pathology in wt grafts further suggests that neurodegeneration in AD is dependent, at least partially, upon extracellular amyloid accumulation.

2.5 Materials and Methods

Donor embryos for grafting

Donor mice were C57BL/6 (B6) mice (RCC, Füllinsdorf, Switzerland) and APP23 tg mice (Sturchler-Pierrat et al., 1997). APP23 mice overexpress mutated human APP (hAPP) with the Swedish double mutation under a neuron-specific murine Thy-1 promoter element. All APP23 mice were from generation 9-11 of backcrossing to B6. To produce tg grafts, homozygous male APP23 tg mice were bred with hemizygous female APP23 mice. For wt grafts B6 mice were bred. Pregnant females were staged by vaginal plugs and embryos at E15-17 (crown to rump length: 13-15 mm) were taken for grafting.

Intracerebral grafting

Embryonic neocortices and hippocampi were dissected on ice under semi-sterile conditions in Dulbecco's modified Eagle's Medium (Life Technologies, Basel, Switzerland). All neocortical tissue pieces from the entire litter were pooled. The same procedure was performed for the hippocampal pieces. Cell suspensions were prepared mechanically through repeated pipetting with Pasteur pipettes on ice. Total cell counts and cell viability were determined with a hemocytometer and 0.4% trypan blue stain (0.4%; Fluka, Buchs, Switzerland). Cell number was adjusted to 100,000 cells/ μ l and cell viability was 80%.

Host mice (3-24 month-old, females) were anaesthetized with a mixture of ketamine (10mg/kg body weight) and xylazine (20mg/kg body weight) in saline. Bilateral stereotaxic injections of 2.5 μ l cortical and hippocampal cell suspensions were placed with a Hamilton syringe into the neocortex (AP +1.0mm from bregma, L +/-2.0mm, DV -1.5mm) and hippocampus (AP -2.5mm, L +/-2.0mm, DV -2.5mm), respectively. In selected mice cell suspensions were injected into the striatum (AP +1.0mm, L +/-2.0mm, DV -3mm), and thalamus (AP -2.5mm, L +/-1.5mm, DV -3mm). Injection speed was 1.25 μ l/minute and the needle was kept in place for an additional 2 minutes before it was slowly withdrawn. The surgical area was cleaned with sterile saline and the incision was sutured. As a control, some mice received stab wound lesions, i.e. the needle was lowered in the brain without injection of any cell suspension. All experiments were in compliance with protocols approved by the Animal Care and Use Committee (Veterinäramt, Basel).

Histology and immunohistochemistry

At various time points after grafting host mice were deeply anaesthetized and either perfused transcardially with 0.1 M PBS (pH 7.4), followed by 4% paraformaldehyde in PBS (pH 7.4) or brains were removed and immersion fixed for 2d in 4 % paraformaldehyde in PBS. Brains were then dehydrated overnight in 30% sucrose. After freezing, 20-30 μ m serial coronal sections were cut through the transplants on a freezing-sliding microtome and collected in 0.1 M Tris-buffered saline (pH 7.4).

Sections were stained histologically with cresyl-violet and immunohistochemically according to previously published protocols (Stalder et al., 1999) with the following antibodies: polyclonal antibody NT12 (NT11) to A β (courtesy of P. Paganetti) (Sturchler-Pierrat et al., 1997); mouse monoclonal antibody W0-2 specific to human A β (Wirhth et al., 2002) (courtesy of K. Beyreuther); mouse monoclonal antibody 6E10 specific to hAPP/A β (Signet Pathology Systems, Inc., Dedham, MA, USA); monoclonal antibodies specific to A β x-40 (JRF/cA β 40/10) and A β x-42 (JRF/cA β 42/26) (Mathews et al., 2002); monoclonal antibody specific to rodent/mouse A β that does not cross-react with human A β (JRF7rA β 1-15/2) (Mathews et al., 2002); polyclonal antibody A4CT to the C-terminal 100 amino acid of hAPP (courtesy of K. Beyreuther); mouse monoclonal

antibody 56C6 to neprilysin (CD10; Novocastra Laboratories, Newcastle upon Tyne, U.K); polyclonal affinity-purified antibody IDE-1 to IDE protein (courtesy of D. Selkoe); polyclonal antibody to synaptophysin (Dako, Glostrup, Denmark), mouse monoclonal antibody AT8 which recognizes hyperphosphorylated tau (Innogenetics, Heiden, Germany), rat monoclonal antibody to CD11b (Mac-1; Serotec, Oxford, UK), polyclonal antibody to Iba1 (Ohsawa et al., 2000) (courtesy Y. Imai) and polyclonal antibody to GFAP (Dako). Some sections were double stained with NT12 and Congo Red. Additional sections were stained histochemically for acetylcholinesterase (Boncristiano et al., 2002) and Holmes silver staining.

Assessment of APP and A β in graft and host using Western blotting

Under deep inhalation anesthesia, engrafted mice were sacrificed by decapitation. Brains were removed and sectioned on a vibratome (400 μ m thick slices) in ice-cold PBS. Slices were transferred onto a glass slide on ice. With the aid of a dissecting microscope and a Stoelting micropunch device (Stoelting, Wood Dale, IL, USA), micropunches were taken from the graft. For comparison, micropunches were also taken from normal APP23 and B6 mice. Punches had a diameter of 0.74 mm and thus consisted of 172 μ g wet weight tissue (Palkovits, 1983).

To determine hAPP expression, punched samples were diluted in 30 μ l of homogenization buffer (50 mM Tris, pH 8; 150 mM NaCl; 5 mM EDTA; protease inhibitor cocktail (Roche Diagnostic, Mannheim, Germany) and 90 μ l sample buffer (0.48 M Bistris; 0.21 M Bicine; 1.33% w/v SDS; 20% w/v sucrose; 3.33% v/v 2-mercaptoethanol; 0.0053% w/v bromophenol blue). Subsequently, samples were Dounce-homogenized, sonicated, and subjected to 10% Bicine/Tris 8M Urea SDS-PAGE (Wiltfang et al., 1997). Proteins were transferred onto a PVDF membrane and hAPP was detected using monoclonal mouse 6E10 antibody. The secondary antibody was horseradish peroxidase-conjugated goat anti-mouse IgG (Chemicon, Temecula, CA, USA). Bands were visualized using SuperSignal β (Pierce, Rockford, IL, USA) and developed onto Kodak X-OMAT AR film (Rochester, NY, USA).

To analyze A β levels, tissues of two micropunches were combined and human A β was immunoprecipitated with antibody 6E10 and protein G Sepharose (Sigma, St Louis, MO, USA). Subsequent SDS-PAGE and blotting was done as described above. Synthetic A β 1-40 and 1-42 (Bachem, Bubendorf; Switzerland) was used as control. Different exposures of the films were digitized and band density measurements for A β 40 were made using NIH Image Version 1.61 (NIH, Bethesda, MD, USA). Only bands within the linear range of the film were analyzed.

ELISA to measure soluble extracellular A β in engrafted brain regions

Neocortex, hippocampus, striatum, and thalamus of 6 month-old male pre-depositing APP23 mice ($n = 8$) were dissected on ice. Tissue of four mice was pooled and homogenized in 50mM Tris, pH 8, 150mM NaCl, and 5 mM EDTA and protease inhibitor cocktail (Roche Diagnostic). The homogenate was centrifuged at 107,000g for 1 hr at 4°C and human A β x-40 and A β x-42 levels were determined in the supernatant by previously described sandwich ELISAs (Mathews et al., 2002). In brief, A β was captured with carboxy-terminal monoclonal antibodies that recognize exclusively either A β x-40 (JRF/cA β 40/10) or A β x-42 (JRF/cA β 42/26) and detected with horseradish peroxidase-conjugated JRF/A β tot/17, which was raised against the amino-terminal 16 residues of human A β . All measurements were done in two or more replica wells. ELISA results reported are the mean of the two pooled tissue samples (ng A β per g wet weight), based on standard curves using synthetic A β 1-40 and A β 1-42 peptide standards (American Peptide Co. Sunnyvale, CA).

Serum titers of A β antibodies

At various time points post-grafting (30-510 days), retro-orbital blood samples were collected from selected engrafted mice by using heparin-coated capillary tubes. Samples were centrifuged and the sera were immediately frozen on dry ice. Antibody titers were assessed by ELISA. Serial dilutions of sera were made onto microtiter plates coated with A β 1-40 (50 ng/ml; Bachem). Detection was through alkaline phosphatase-conjugated rabbit anti-mouse IgG (Calbiochem, San Diego, USA) and 4-nitrophenol phosphate disodium salt hexahydrate (Fluka). Titers were defined as dilution yielding 50% of the maximal signal.

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2.7 Figure Captions

Figure 1 Neural grafting of APP23 tissue into B6 hosts. Embryonic APP23 hippocampal tissue was injected into the hippocampus of 3 month-old wt B6 hosts. Mice were analysed at various times post-grafting (see Table 1). Shown are mice analysed 3 months (A-C), 18 months (D-F), and 21 months (G-I) post-grafting. Cresyl violet staining was used to identify the grafts (asterisks in A,D,G). Immunohistochemistry revealed strong hAPP expression restricted to neurons in the grafts (B,E,H). Nevertheless, A β -immunostaining did not reveal any amyloid deposits 3 and 18 months post-grafting (C,F). At 21 months postgrafting massive amyloid deposition in both plaques and vessels was found that was to a great extent congophilic (i, insert). The bulk of the amyloid was confined to the graft, however, some vessels with amyloid (arrow) and some diffuse amyloid (arrowhead) were found outside the graft in the host tissue. Similar results were observed when cortical tg tissue was injected into the cortex of wt mice (not shown). Bars are 250 μ m (A-C) and 400 μ m (D-I).

Figure 2 APP and A β levels in APP23 grafts placed in B6 hosts. (A) Hippocampus of a 15 month-old wt B6 mouse that received an APP23 tg graft at the age of 3 months. A micropunch (p) was taken from the graft (asterisk) and subjected to Western blot analysis. The remaining tissue was immersion-fixed and immunostained for hAPP demonstrating that hAPP expression was exclusively confined to the graft. Bar is 150 μ m. (B) Western blotting of the micropunch taken from the APP23 hippocampal graft shown in (A) with hA β /APP-specific antibody 6E10. In addition micropunches from hippocampus were taken from young 4-6 month-old APP23 mice and control B6 mice for comparison. *Lane 1*: Synthetic A β 40/42. *Lane 2*: punch from the hippocampal APP23 graft. *Lane 3*: punch from B6 hippocampus (CA1 area). *Lane 4*: punch from APP23 hippocampus (CA1 area). Note that hAPP expression and A β levels in the graft (lane 2) were comparable to that in APP23 hippocampus (lane 4). In both graft and young APP23 mouse

A β 1-40 was several fold higher than A β 1-42. (C) Similar analysis as in (B) but for an APP23 cortical graft 20 months post-grafting. For comparison micropunches from neocortex were taken from a 12 month-old amyloid-depositing APP23 mouse and a control B6 mouse. *Lane 1*: Synthetic A β 40/42. *Lane 2*: punch from the cortical APP23 graft. *Lane 3*: punch from B6 neocortex. *Lane 4*: punch from APP23 neocortex.

Figure 3 Neural grafting of APP23 and B6 tissue into APP23 hosts. Embryonic APP23 tg hippocampal tissue was injected into the hippocampus of a 6 month-old APP23 mouse and analyzed three months later (A-C). Cresyl violet staining reveals the graft (asterisk) in the dentate gyrus (A). Immunohistochemistry with an antibody to hAPP reveals robust neuronal expression of hAPP in the graft and host (B). Immunostaining for A β in adjacent sections revealed considerable amyloid deposits in the graft (C). Strikingly, amyloid deposition in the graft was also found when B6 wt hippocampal tissue was injected into the hippocampus of a 6 month-old APP23 mouse and analyzed 3 months later (D-F). In this case no hAPP expression was found in the graft (E). The same was found when embryonic B6 wt cortical tissue was grafted into the neocortex of a 6 month-old APP23 mouse and analyzed 3 months later (G-I). Although the majority of the amyloid was of the diffuse type, some amyloid was compact and Congo red-positive. The insert in (I) shows an adjacent section stained for Congo red and viewed under cross-polarized light. Note, that in most cases amyloid deposition in the graft was more intense compared to the only scattered amyloid plaques in the 9 month-old host tissue (arrows in F and I). Bars are 90 μ m (A-F) and 50 μ m (G-I).

Figure 4 Amyloid in B6 grafts. Immunostaining with antibody JRF/cA β 40/10 specific to A β x-40 (A) and antibody JRF/cA β 42/26 specific to A β 1-42 (B) of a B6 wt graft placed in neocortex of an APP23 host. Antibodies specific to rodent (mouse) A β (JRF7rA β 1-15/2) revealed only very faint labeling of the amyloid (C). Shown are adjacent sections with arrowheads indicating

the same amyloid deposits. Note that the faint labeling of vessels and microglia is due to unspecific labeling of the secondary anti-mouse IgG antibody. (D) A micropunch (p) was taken from a cortical wt graft (asterisk) and subjected to Western blot analysis. The remaining tissue was immersion-fixed and immunostained for hAPP demonstrating that hAPP expression was exclusively confined to the host. (E) Western blotting of the micropunch taken from the wt cortical graft shown in (A) with human A β -specific antibody 6E10. In addition micropunches were taken from the neocortex of a 12 month-old amyloid-depositing APP23 mouse and a control B6 mouse. *Lane 1*: Synthetic A β 40/42. *Lane 2*: punch from the cortical wt graft. *Lane 3*: punch from B6 neocortex. *Lane 4*: punch from APP23 neocortex. Note the presence of both A β 1-40 and A β 1-42 in the wt graft. Bars are 20 μ m (C) and 120 μ m (D).

Figure 5 Neural grafting of B6 tissue into thalamus and striatum of APP23 hosts. (A,B) Embryonic B6 hippocampal tissue was injected into the thalamus of a 6 month-old APP23 mouse and analyzed 3 months later. Cresyl violet staining revealed that the grafts were nicely integrated into the host thalamus and are viable without any signs of neurodegeneration (asterisk in A). Immunostaining for A β in adjacent sections revealed amyloid deposits in the graft (B). In contrast, when embryonic B6 tissue was placed into the striatum of a 6 month-old APP23 mouse, no amyloid formation was observed 3 months later (C,D). Bars are 250 μ m.

Figure 6 Mechanism of amyloid formation in B6 grafts. (A) Holmes silver staining of the graft-host border of a B6 wt graft (asterisk) in a 9 mo-old APP23 tg host. There is a lower density of fibers in the graft compared to the host. Only few fibers cross the host-graft border (dotted line) and most fibers were confined to either the host or the graft. (B) Consistent with the Holmes fiber staining, immunostaining for hAPP reveals many positive neurons and processes in the host tissue, but none in the wt graft. (C) Cortical stab wound in a tg 6 month-old APP23 host shows an appreciable increase in

CD11b-positive microglia around the lesion site 3 months post-surgery (arrowheads). (D) However, immunohistochemistry for A β in an adjacent section does not indicate any amyloid deposition in the vicinity of the lesion. (E) A β immunostaining with antibody NT12 in the neocortex of a 9 month-old APP23 tg host reveals punctate intraneuronal staining. (F) In contrast, no evidence of intraneuronal A β was found in neocortical wt grafts into APP23 hosts. An amyloid plaque in the wt graft is shown by an arrowhead. The same results were observed with monoclonal antibody W0-2 specific to human A β . Bars are 10 μ m (A,B), 100 μ m (C,D), and 15 μ m (E,F).

Figure 7 Amyloid pathology associated with congophilic amyloid plaques in B6 grafts. (A) Compact plaque in B6 wt graft in a 9 month-old APP23 tg host reveals birefringence when viewed under cross-polarized light (double staining for A β and Congo red). (B) Cresyl violet staining demonstrates only a few neuronal cell bodies at the plaque periphery and some of them with a dying phenotype (arrow). (C) Plaques are surrounded and interdigitated by synaptophysin-positive dystrophic boutons (arrows). (D) Acetylcholinesterase-staining reveals abnormal large cholinergic processes in vicinity of the plaque (arrow). (E) Some of these abnormal and distorted processes in the plaque periphery were positive for hyperphosphorylated tau (arrow). (F) CD11b-immunostaining demonstrates hypertrophic microglia clustered around compact amyloid plaques. (G) Amyloid-associated microglia were also intensively positive for Iba1 similar to those observed in amyloid-depositing APP23 mice. (H) GFAP-staining reveals hypertrophic and reactive astrocytes decorating the plaque periphery. Bars are 25 μ m (A,B), 10 μ m (C), 25 μ m (D), 6 μ m (E), 25 μ m (F), 30 μ m (G) and 25 μ m (H).

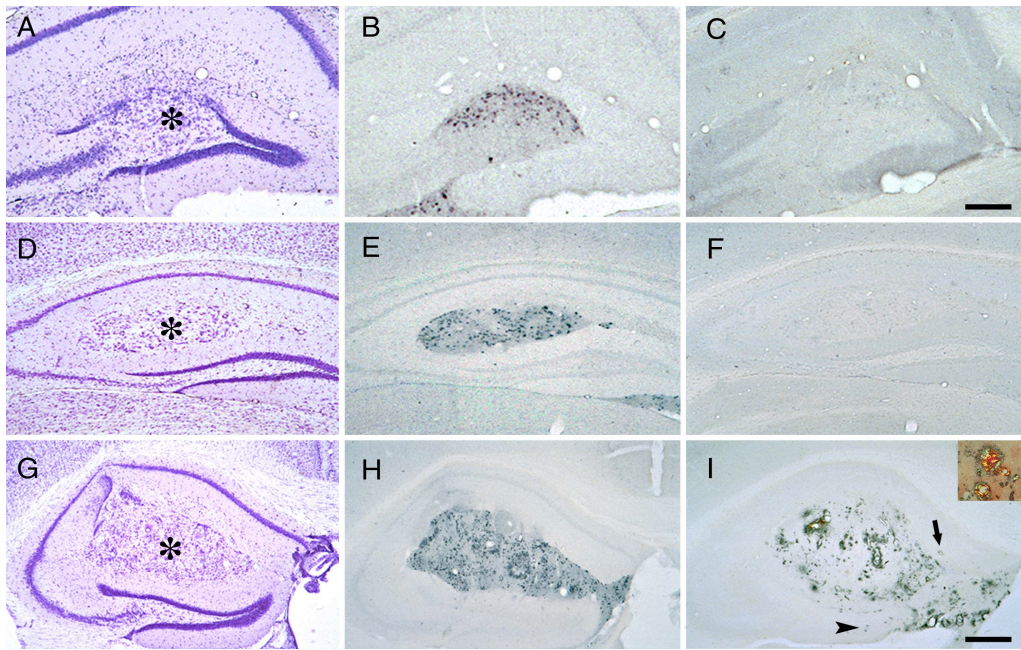


Figure 1

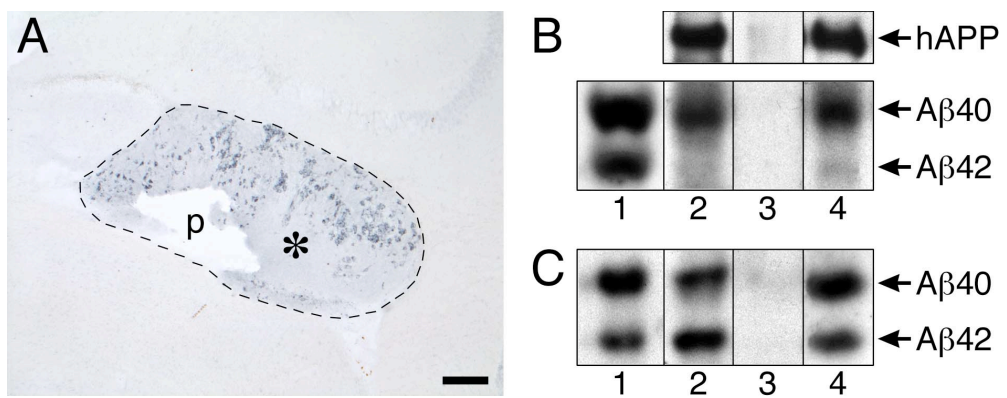


Figure 2

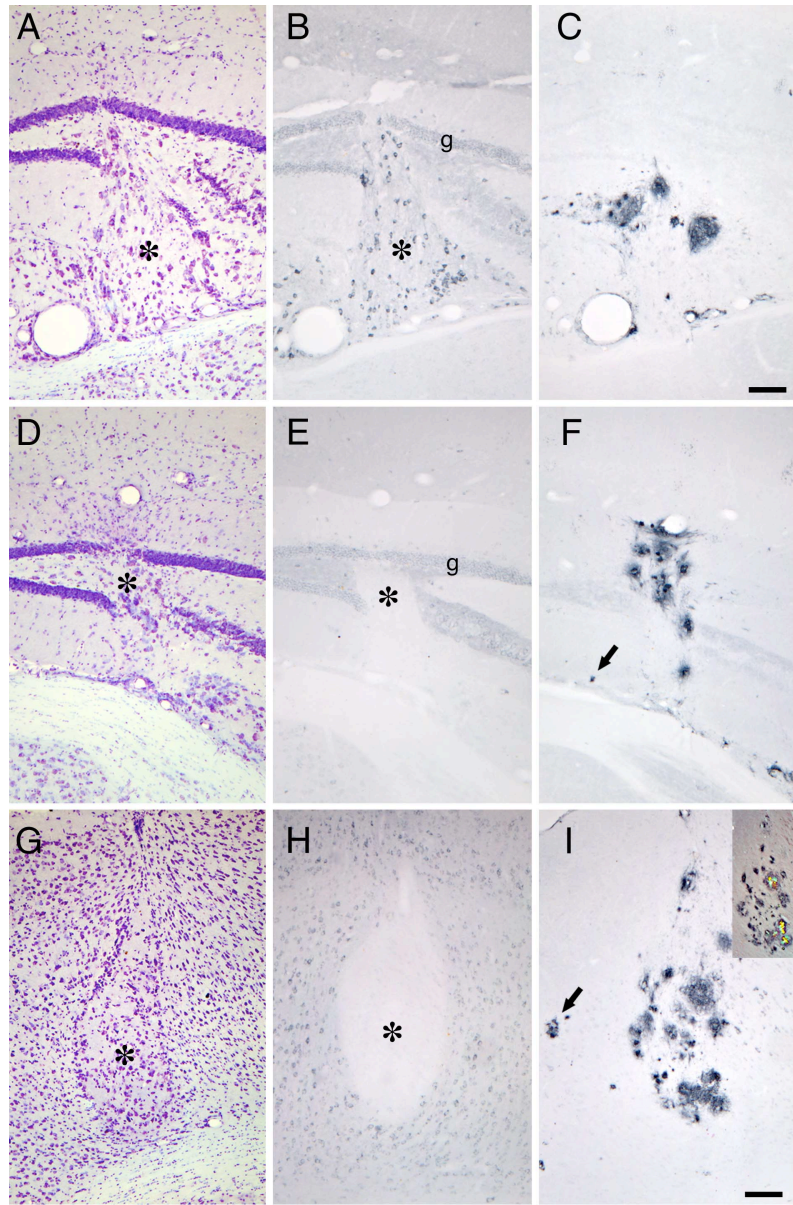


Figure 3

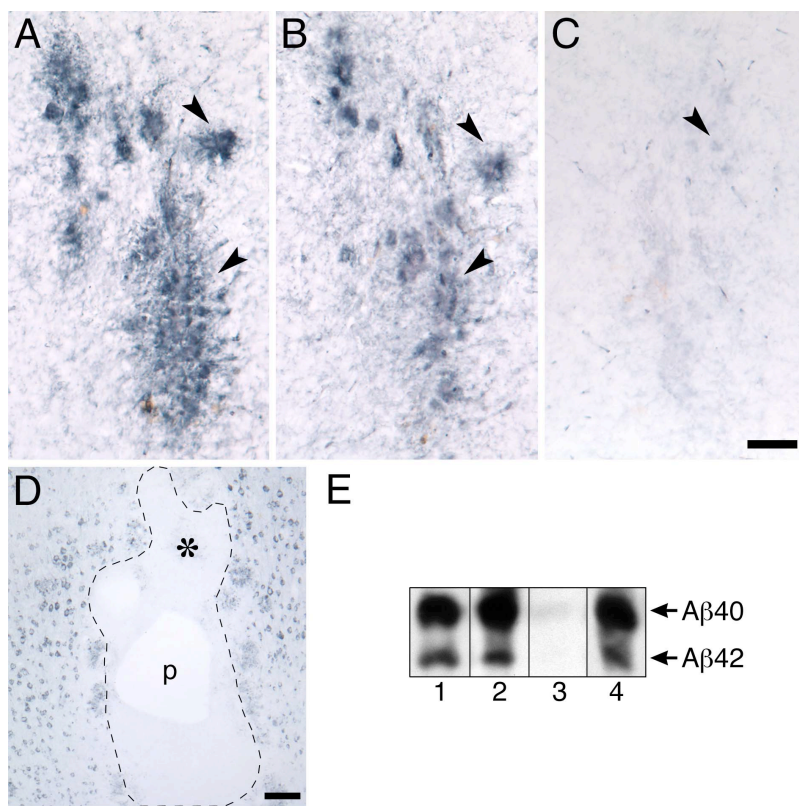


Figure 4

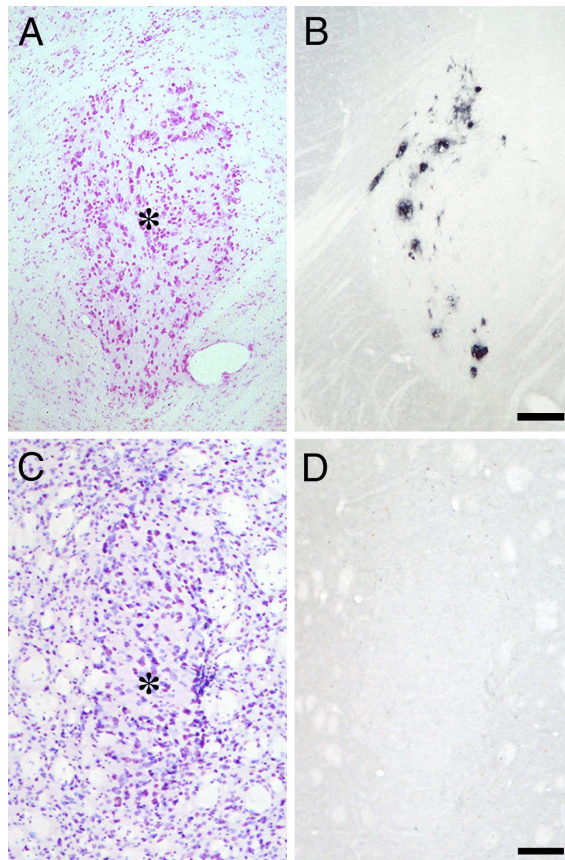


Figure 5

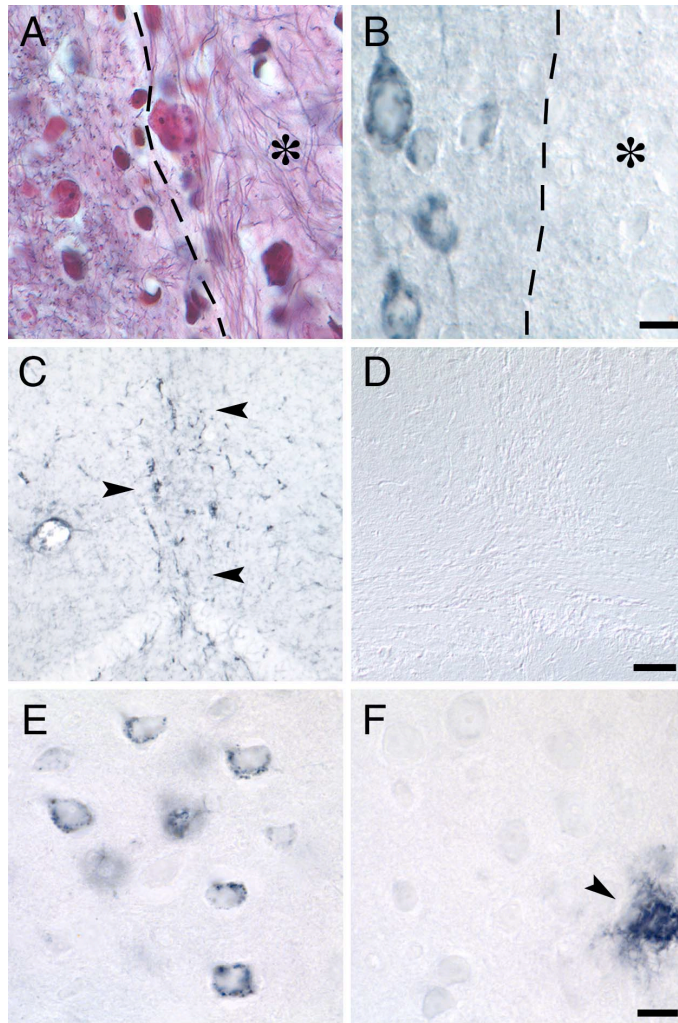


Figure 6

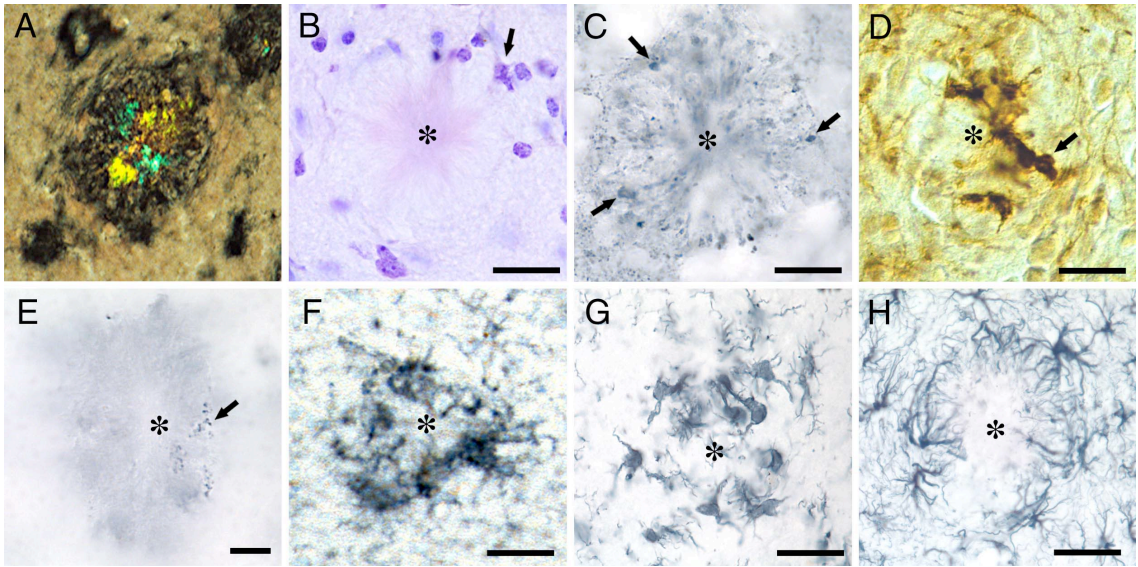


Figure 7

Table 1: *APP23 embryonic tissue transplanted in B6 brain*

Host	Age ^a	Post-grafting interval				
		3 mo	6 mo	12mo	18 mo	> 20 mo
B6	3 mo	0 ^b /12 ^c	0/6	0/6	0/20	2/9
B6	16 mo	0/1	0/2	0/6		
B6	24 mo	0/2	0/5			

^a age of the host at time of transplantation

^b number of mice with amyloid deposits in at least one graft

^c total mice analyzed

Table 2: *APP23 and B6 embryonic tissue transplanted in APP23 brain*

Host	Age ^a	Grafts ^b	
		APP23	B6
APP23	6 mo	7 ^c /7 ^d	10/13

^a age of the host at time of transplantation

^b mice were sacrificed 3 months post-grafting

^c number of mice with amyloid in at least one graft

^d total mice analyzed

3. Seeding of cerebral amyloidosis in APP23 transgenic mice

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3.1 Abstract

The misconformation and self-assembly of specific proteins is well established as a disease process in Alzheimer's disease (AD), but little is known about how protein aggregation is initiated *in vivo*. Seeded polymerization of amyloid beta (A β) has been demonstrated *in vitro* and has recently also been achieved *in vivo* by intracerebral injection of AD brain extract into young amyloid precursor protein (APP) transgenic mice prior to plaque formation. We have confirmed and advanced these findings through studies involving injection of diluted brain extract from AD patients and from aged APP23 transgenic mice into young APP23 mice. Similar to the AD extract, transgenic mouse brain extract induced massive seeded A β deposits in the brain parenchyma and vessel walls four months post-infusion. This seeding was time- and concentration-dependent. These seeded amyloid deposits were mostly diffuse in nature, with some congophilic plaques and vessels. Surprisingly, intracerebral injection of synthetic A β resulted so far in only limited A β seeding deposition compared to that obtained with A β -rich brain extracts. Studies with fresh and aged synthetic A β 40, A β 42 and a mixture of A β 40/42, with levels matching that in AD brain extracts, are ongoing. These studies will unequivocally determine whether A β alone is sufficient to induce seeding, and more specifically, which conformational state of A β has intrinsic properties to act as a seeding agent.

3.2 Introduction

Misfolding, aggregation and abnormal accumulation of proteins is a common feature of many neurodegenerative diseases such as Alzheimer's disease (AD), Parkinson's disease and Creutzfeldt-Jacob disease. Genetic studies support the view that A β fibril formation is a central process in AD pathogenesis (Hardy et al., 1998) and accompanied by neurofibrillary tangle formation, neuron and synapse loss, neuroinflammation, and dementia (Yankner, 1996; Selkoe, 1999; Gotz et al., 2001; Lewis et al., 2001). However, little is known about how protein aggregation and deposition is initiated *in vivo*.

Protein aggregation *in vitro* is dependent on protein concentration and time (Jarrett and Lansbury, 1993; Harper and Lansbury, 1997; Lansbury, 1997). The relatively slow nucleation process of A β can be accelerated by introducing an exogenous seed which then leads to rapid A β aggregation (Koo et al., 1999). Similar to these *in vitro* findings, seeded aggregation may also determine the onset and progression of amyloid formation in amyloid precursor protein (APP) transgenic mice and in AD.

In the past, there have been many attempts to induce AD neuropathology in animal models by injection of A β peptides. Microinjection of fibrillar A β into the cerebral cortex of aged rhesus monkeys resulted in profound neuronal loss and tau phosphorylation (Geula et al., 1998). Moreover, amyloid cores, isolated from AD brains, caused neurodegeneration and neurotoxicity when injected into the cortex and hippocampus of rats (Frautschy et al., 1991; Frautschy et al., 1992). However, none of these studies has focused on the mechanism of A β polymerization *in vivo*. Only recently has it been described that inoculation of dilute AD brain extracts leads to amyloid fibril formation in young Tg2576 mice (Kane et al., 2000; Walker et al., 2002). Nevertheless, it remains to be clarified which factors initiate and are involved in this seeding process. To this end, we have injected diluted brain extracts from AD patients and from aged APP23 transgenic mice into young APP23 mice prior to plaque formation. Our results indicate that A β -containing human and transgenic mouse brain extracts can induce cerebral amyloidosis in APP23 transgenic mice *in vivo*.

3.3 Results

Intracerebral injection of brain extract of aged APP23 mice induces amyloid formation in brains of young APP23 mice similar to that of AD brain extract

To determine whether brain homogenate of aged APP23 mice is as potent in inducing seeding of A β as AD brain homogenate, we injected both brain homogenates into the hippocampus of young, 5-month old APP23 mice. These mice were analyzed 4 months later and, similar to the AD extract, the APP23 brain extract induced robust seeding of β -amyloid deposits, mainly in the hippocampus (Fig. 1A,B). As expected, intracerebral injection of control patient extract induced only few amyloid deposits (Fig. 1C). We observed no seeded amyloid deposits in APP23 transgenic mice injected with wt brain homogenate or PBS (Fig. 1D,E). Infusion of amyloid-containing APP23 brain homogenate into non-transgenic littermates did not produce A β deposition, indicating that the observed immunoreactivity present at 4 months post-infusion was not due to the injected material itself (Fig. 1F).

Seeded amyloid is diffuse and occurs in vessels and plaques

Seeded amyloid exhibits a characteristic distribution pattern along the hippocampal fissure and the granular cell layer of the dentate gyrus, the dorsal lateral geniculate nucleus, in the vasculature of the thalamus and in pial vessels (Fig. 2A-D). In some cases, there was a spread of A β deposits along the corpus callosum (Fig. 2A, right hemisphere). The seeded material in the parenchyma was mainly diffuse in nature (Fig. 2 A,C) with some congophilic compact plaques in the hippocampal area (Fig. 2E) and congo red positive vessels in the thalamus (Fig. 2F). Results revealed that not only tissue near the injection site was affected, but also brain regions away from the infusion site.

Seeding of A β is time- and concentration-dependent

APP23 mouse brain homogenate was injected into 5-month-old APP23 transgenic mice and analyzed after different time points from 1 to 4 months post-infusion. Immunohistological analysis with an antibody against A β revealed no amyloid deposits after 1 month (Fig. 3A). First signs of diffuse seeded amyloid were detectable 2 months

post-infusion (Fig. 3B). Clearly, 3 months after the inoculation of the brain homogenate, the typical seeding pattern was visible (Fig. 3C). The highest number of seeded amyloid plaques was obtained 4 months post-infusion (Fig. 3D). Stereological analysis also confirmed that the amount of A β deposition significantly increases with time (ANOVA, $p < 0.0001$) (Fig. 3E).

To determine if A β concentration in the brain homogenate influences the seeding pattern, we injected different dilutions of the tg brain homogenate (0.5% vs. 10%) and compared the resulting hippocampal amyloid load. Immunohistochemical stainings and stereological analysis both revealed that the 10% extract exceeded seeding obtained with the 0.5% extract by a factor of 2 (Fig. 4A,B,C). Thus, these results clearly demonstrate that seeding of A β in APP23 mice is a time- and concentration-dependent process.

Seeded amyloid contains A β 40 and A β 42

The next set of experiments were designed to determine the nature of the seeded protein deposition. A β was immunoprecipitated from micropunches taken from hippocampus, lateral geniculate nucleus and thalamus out of formalin-fixed prestained tissue (Ikeda et al., 1998) (Fig. 5A-H). These micropunches did not only originate from different brain regions, but they also differed in the type of amyloid they contained. It was clearly visible that the hippocampus showed both, diffuse and compact plaques, in contrast to the only diffuse band in the lateral geniculate nucleus and the mainly congo red positive compact vessels in the thalamus (Fig. 5B,E,H). Western blot analysis revealed the presence of A β in the micropunches (Fig. 5C,F,I). There was a negligible amount of A β 1-42 detectable in the thalamus with a strong predominance of A β 1-40 over A β 1-42 compared to the hippocampus.

Seeding is an extracellular process

Next, we investigated the ultrastructure of A β depositions induced by the infusion of amyloid-containing brain extract. Seeded A β deposits were mostly of the diffuse type but some compact plaques were located along the granular cell layer in the dentate gyrus (Fig. 6A). Ultrastructural analysis revealed that A β deposits were limited to the

extracellular space (ECS) by observation under the electron microscope (Fig. 6B). We found no evidence for intracellular A β , supporting the idea that seeded amyloid aggregates are located extracellularly.

Synthetic A β injections induce only limited seeding properties

To identify the factors that initiated the seeding process in our model, we were wondering whether A β itself may be sufficient to achieve the same degree of seeding seen with A β -containing brain extracts. Therefore, we injected synthetic A β 40 and A β 42 peptides in APP23 transgenic mice and analyzed them 1 day and 4 months post-infusion. After 1 day up to 1 month, we found no immunohistological evidence of the dilute tg brain extract itself (Fig. 3A). In contrast, we observed big aggregates of the injected synthetic A β 40 and A β 42 material, respectively (Fig. 7A,B). After 4 months, a robust seeding pattern was only present in APP23 mice injected with A β -containing brain homogenate (Fig. 1A,B). Synthetic A β 40 and A β 42 injections resulted in only a limited amount of amyloid deposits and failed to induce the same amount of seeding seen with the brain homogenates (Fig. 7C-F).

3.4 Discussion

Accumulation of highly insoluble protein aggregates is a hallmark of AD and other neurodegenerative diseases and considered to be a critical step in disease pathogenesis (Lansbury, 1997; Koo et al., 1999; Walker and LeVine, 2000). The present study was undertaken to characterize the mechanism that leads to amyloid formation in APP transgenic mice in more detail. First, we have confirmed the previous finding (Kane et al., 2000) that intracerebral injection of AD brain extract into APP transgenic mice induces A β deposition. Thus, the AD brain extract provided an extremely potent seeding agent, but the factors that initiated amyloid formation in these mice still remains unknown.

Besides the typical extracellular amyloid deposits and the intracellular neurofibrillary tangles, the AD brain is characterized by a pronounced inflammation with an astroglial and microglial response and the presence of inflammatory cytokines (Itagaki et al., 1989; McGeer et al., 1994; Griffin et al., 1995). Research has shown that inflammation plays a role in plaque formation and it has also been suggested that infectious agents can initiate seeding (Yucesan and Sriram, 2001; Lin et al., 2002). To rule out the latter possibility, we also infused A β -containing tg brain homogenate. Our results revealed robust seeding with both, human and tg animal brain homogenate 4 months post-infusion. The control experiments done in wt control littermates confirmed that it was not the tg injectate itself that was detected 4 month after injection.

The formation of A β fibrils *in vitro* occurs via seeded polymerization (Jarrett and Lansbury, 1993). A β aggregation is dependent upon A β concentration and time and initiated by the formation of an oligomeric nucleus that is thought to be the rate-determining step of amyloid deposition (Lansbury, 1997). We have tested this theory of seeding β -amyloid *in vivo* and found that seeding occurred after a lag phase of 2 months, which increases over time. In addition, plaque formation in our seeding model appears in a concentration-dependent manner. Thus, we show for the first time that the complex processes associated with misfolding and accumulation of β -amyloid protein that emerge from *in vitro* studies are relevant for *in vivo* processes.

Next, we examined the biological properties of the seeded amyloid plaques. APP23 mice develop amyloid plaques and cerebrovascular amyloid throughout the neocortex, hippocampus and to a lesser extent in the thalamus with the majority of cerebrovascular amyloid (Sturchler-Pierrat et al., 1997; Calhoun et al., 1999; Winkler et al., 2001). In the experimental paradigm used here, we always observed a typical seeding pattern with affected brain areas identical to those of aged, untreated APP23 transgenic mice. Thus, not only the hippocampus, but also brain regions further away from the injection site such as thalamus developed seeded plaque formation. This observation is in line with our previous finding that diffusion of A β in the extracellular space is an important parameter for cerebral amyloidosis (Meyer-Luehmann et al., 2003). Alternatively, axonal transport has also been proposed as possible explanation for the spread of seeded A β pathology in the brain (Walker et al., 2002). Electron microscopy of the dentate gyrus revealed A β deposition located between neuron bodies in the extracellular space, indicating that amyloid formation was induced extracellularly. It has been postulated that amyloid deposits in vessels contain A β 40 whereas the longer A β 42 form is the predominant component of parenchymal plaques (Joachim et al., 1988; Gravina et al., 1995; Herzog et al., 2003). Western blot analysis confirmed that the seeded material consisted of both A β peptides, the A β 40 and A β 42 depending on the form of aggregates. Thus, with the injection of A β -containing brain homogenate we induced seeding of both A β species, suggesting that the same initial seeding agent could seed amyloid deposits in the vessels and in brain parenchyma.

Finding out whether A β is sufficient to induce seeding or whether this process requires additional factors, is key to the development of therapeutic strategies against cerebral amyloidosis. Previous attempts to generate consistent amyloid deposition causing neurodegeneration via injection studies have produced inconsistent results (Frautschy et al., 1991; Clemens and Stephenson, 1992; Emre et al., 1992; Frautschy et al., 1992; Games et al., 1992; Kowall et al., 1992; Rush et al., 1992; Stein-Behrens et al., 1992; Waite et al., 1992; Sigurdsson et al., 1997). Most of these studies used single injections of A β peptides into rodent brains, indicating that the method of A β preparation itself like peptide concentration or aggregation state represents a common source of variation. Our data so far favour the possibility of important cofactors because synthetic A β 40 and A β 42 injections resulted in fewer seeded amyloid deposits than those after tg or AD

brain homogenate infusion. We hypothesize that additional plaque enhancing factors are abundant in the brain homogenates since it has been reported that after immunodepletion of A β , similar seeding still occurs (Younkin, 1995). TGF- β , heparan sulfate proteoglycan, apolipoprotein E and A β 43 have been successfully applied in combination with A β 40 to promote plaque formation (Snow et al., 1994; Wisniewski et al., 1994; Frautschy et al., 1996; Stephan et al., 2001). Future studies will help to identify cofactors in the brain homogenate that are required for the initiation of cerebral amyloidosis *in vivo*. In particular, the identification of endogenous chaperone molecules for A β seeding and/or conditions to transform synthetic A β into a seeding agent for cerebral amyloidosis would give new important insights.

Studies with synthetic A β at concentrations equivalent to those found in A β brain extracts are ongoing. In the long run, these studies will hopefully answer the most fundamental question whether A β itself is the seeding factor. Additionally, ongoing analyzes with fresh and aged samples will reveal what conformational state of A β is most effective in inducing seeding. Such results will greatly enhance the understanding of A β aggregation and early amyloidogenesis *in vivo* and will provide new therapeutic targets in the search for compounds that inhibit A β aggregation.

3.5 Materials and Methods

Tissue extracts and synthetic A β

Human tissue for preparation of extracts was derived at autopsy from cortices of 2 patients who had died of confirmed AD and from a neurological normal control. Animal tissue was derived from neocortices of two APP23 transgenic mice and from a wildtype nontransgenic littermate (Table 1). The tissue samples were fresh frozen and stored at -80°C. Pieces of neocortex were homogenized at 10% (w/v) in sterile PBS, vortexed, sonicated for 5 seconds and centrifuged at 3000x g for 5 minutes. The supernatant was immediately frozen (-80°C) and sometimes further diluted 1:20 to a final concentration of 0.5%.

Synthetic A β 40 and A β 42 was prepared by resuspending lyophilized A β 40 and A β 42 (Bachem, Bubendorf, Switzerland) in phosphate-buffered saline (PBS), pH 7.4,

followed by an incubation period of 5 days at 37°C to allow them to aggregate. A β was vortexed, sonicated and injected in the same manner as the brain homogenates. Congo red staining confirmed the formation of fibrils.

Western blot analysis

To determine A β levels in the brain homogenates used in this study, samples were further diluted 1 to 4 in sample buffer (0,48 M Bis-Tris; 0,21 M Bicine; 1,32 % (w/v) SDS; 20% (w/v) sucrose; 3,33 % (v/v) 2-mercaptoethanol; 0,0053% (w/v) bromophenol blue). Samples were then subjected to 10% Bicine-Tris 8M Urea SDS-PAGE (Wiltfang et al., 1997). Synthetic A β 1-40 and 1-42 (Bachem, Bubendorf, Schweiz) were used as controls. Proteins were transferred onto a PVDF membrane and probed with monoclonal antibody 6E10 (dilution 1:500). The secondary antibody was horseradish peroxidase–conjugated goat anti-mouse IgG (Chemicon, Temecula, California). Bands were visualized using SuperSignal (Pierce, Rockford, Illinois) and developed onto Kodak X-OMAT AR film (Rochester, New York).

Stereotaxic surgery

Host mice (5 month-old, males) were anaesthetized with a mixture of ketamine (10 mg/kg body weight) and xylazine (20 mg/kg body weight) in saline. Bilateral stereotaxic injections of 2.5 μ l brain homogenate were placed with a Hamilton syringe into the neocortex (AP -2.5mm from bregma, L +/-2.0mm, DV -1mm) and hippocampus (AP -2.5mm, L +/-2.0mm, DV -2.5mm), respectively. In an additional experiment 10 mice were injected unilaterally into the cortex and hippocampus. Injection speed was 1.25 μ l/minute and the needle was kept in place for an additional 2 minutes before it was slowly withdrawn. The surgical area was cleaned with sterile saline and the incision was sutured. As a control, some mice received stab wound lesions (n=5), i.e. the needle was lowered into the brain without injection of any cell suspension.

Histology and immunohistochemistry

Brains were removed and immersion-fixed for 24 h in 4% paraformaldehyde in PBS, then dehydrated overnight in 30% sucrose. After freezing, 25 μ m serial coronal sections were cut through the brains on a freezing-sliding microtome and collected in 0.1 M Tris-buffered saline (pH 7.4). Sections were stained with cresyl-violet and immunohistochemically according to previously published protocols (Stalder et al., 1999) with the following antibodies: polyclonal antibody NT12 to A β (courtesy of P. Paganetti, Basel, Switzerland) (Schrader-Fischer and Paganetti, 1996); mouse monoclonal antibody 6E10 specific to hAPP/A β (Signet Pathology Systems, Inc., Dedham, MA, USA); polyclonal antibodies specific to A β x-40 (FCA3340) and 42 (FCA3542) (Barelli et al., 1997), rat monoclonal antibody to CD11b (Mac-1; Serotec, Oxford, UK), and mouse monoclonal antibody AT8 which recognizes human and murine tau phosphorylated at S202 and T205 (Goedert et al., 1995). Additional sections were stained with congo red and viewed under cross-polarized light.

Ultrastructural analysis

Some mice (n=3) were deeply anaesthetized and perfused transcardially with fresh 0.1 M phosphate buffered 4% paraformaldehyde (PFA) and 0.5% glutaraldehyde. Brains were postfixed in 4% PFA in 0.1 M phosphate buffer (PB) overnight. The next day they were sectioned on a vibratome and the 40 μ m slices were collected in 0.01 M PBS. Immunohistochemistry was performed as described in a previous publication (Stalder et al., 1999) except that they were treated with Triton and DMSO (5, 10 and 20%) instead of H₂O₂ to perforate the cells. The polyclonal antibody, NT12, served as primary antibody against human A β (Schrader-Fischer and Paganetti, 1996) and the visible reaction was achieved by means of diaminobenzidine (DAB). For the background staining, slices were incubated in 0.5% OsO₄ and 6.8% sucrose in PBS. After washing them with PBS, slices were dehydrated in ethanol and acetone and subsequently incubated overnight in Durcupan® ACM (Fluka Chemie GmbH, Buchs SG, CH) according to the instructions of Fluka. Finally, the slices were embedded between a mould parting compound coated slide and coverslipped before incubation at 60°C for 48 hours. Coverslips were removed by means of a glass-cutter and pieces of brain tissue were excised with a scalpel and subsequently glued on Durcupan-blocs in order to

dissect them on the ultramicrotome. Semithin cuts (1 μm) were collected on Superfrost (gel coated) glass slides and some of them were stained with toluidine blue. Ultrathin cuts (at about 80 nm) were collected on standard 200 mesh copper grids (Provac AG, Balzers, FL). Investigation of the ultrastructure in the ultrathin brain samples was done on a Transmission Electron Microscope (Philips EM 900). The high-tension was set to 50 kV in order to achieve an optimal contrast.

Assessment of A β in different brain regions using Western blotting

Under deep inhalation anesthesia, mice were sacrificed by decapitation. Hippocampi of unilateral injected mice (n=4) were dissected and homogenized at 10% (w/v) in sterile PBS, vortexed and sonicated for 5 seconds. In another group of animals (n=3), brains were removed and sectioned on a vibratome (400 μm thick slices) in ice-cold PBS. Slices were transferred onto a glass slide on ice. With the aid of a dissecting microscope and a Stoelting micropunch device (Stoelting, Wood Dale, IL, USA), micropunches were taken from the hippocampus and thalamus of the injected and uninjected hemisphere, respectively. For comparison, micropunches were also taken from normal APP23 and B6 mice. Punches had a diameter of 0.74 mm and 0.96 mm and thus consisted of 172 μg and 362 μg wet weight tissue (Palkovits, 1983). To analyze A β levels, tissues of five micropunches were combined and human A β was immunoprecipitated with antibody 6E10 and protein G Sepharose (Sigma, St Louis, MO, USA). Subsequent SDS-PAGE and blotting was done as described above. Synthetic A β 1-40 and A β 1-42 (Bachem, Bubendorf; Switzerland) were used as controls.

3.6 References

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3.7 Figure Captions

Figure 1 *Seeding of A β by infusion of AD and Tg brain extract into APP23 mice.* Brain extract was intracerebrally injected into the hippocampus of 5 month-old transgenic APP23 hosts (A-E) and nontransgenic littermates of the same age (F). Mice were analyzed 4 month post-injection. A β -immunoreactivity (A-F) in the hippocampus of APP23 transgenic mice after injection of AD (A) and tg (B) brain extract (10%) reveals similar amyloid deposits. In contrast, only few and no amyloid deposits are detected when brain extract of a control patient (C) and wildtype mouse (D) into APP23 transgenic mice was injected. Similarly, no amyloid is observed, when PBS was injected (E) into APP23 transgenic mice. No amyloid is observed, when APP23 brain extract is injected into a wildtype nontransgenic mouse (F). Scale bar: 350 μ m.

Figure 2 *Regional distribution of seeded amyloid.* (A) A β -immunostaining 4 months after bilateral injection of tg brain extract (10%) into APP23 mice with local distribution of seeded amyloid. Shown at a higher magnification (B-D) are the affected areas indicated by white arrowheads. A β deposits are predominately located in the dentate gyrus, along the hippocampal fissure (B), as a band in the dorsal lateral geniculate nucleus (C) and in vessels in the thalamus (D). Congo red stained amyloid plaques in the hippocampus (E) and congo red positive vessels in the thalamus (F) are shown. Scale bars: 90 μ m (B), 60 μ m (C), 25 μ m (D), 40 μ m (E) and 100 μ m (F).

Figure 3 *Time course study of seeding in APP23 mice.* Transgenic mice were analyzed after 1 to 4 months post-infusion. (A) A β -immunoreactivity is absent 1 month after injection of APP23 brain extract. (B) In contrast, after 2 months seeded amyloid is clearly detectable, distributed along the hippocampal fissure, dentate gyrus and around blood vessels. The amount of A β deposition further increases after 3 and 4 months of incubation (C and D). Scale bar: 350 μ m. (E) Stereological analysis confirms a significant

increase in amyloid load with increasing post-injection time (ANOVA, $p < 0.0001$).

Figure 4 *0.5% and 10% APP23 brain extract injections into APP23 mice. (A) Moderate A β -immunoreactivity in APP23 transgenic mice injected with 0.5% APP23 brain extract compared to A β -immunoreactivity after injections with 10% APP23 brain extract (B). (C) Stereological analysis reveals that 10% APP23 brain extract injections resulted in a significantly higher seeded amyloid load than injections with 0.5 % tg brain extract (ANOVA, $p < 0.01$). Scale bar: 350 μ m.*

Figure 5 *A β levels in different seeded brain regions. (A,D,G) Hippocampus, dorsal lateral geniculate nucleus and thalamus of a 9 month-old APP23 mouse after injection of tg brain extract. Seeding resulted in mostly diffuse amyloid in hippocampus and dorsal lateral geniculate nucleus and vascular amyloid in the thalamus. (B,E,H) Micropunches were taken (asterisks) and subjected to Western blot analysis. Bar is 150 μ m. (C,F,I) Western blotting of micropunches. (C) Lane 1: Synthetic A β 40/42. Lane 2: punch from injected hippocampus of an APP23 mouse. Note the strong A β 1-40 and A β 1-42 band. (F) Lane 1: Synthetic A β 40/42. Lane 2: punch from the dorsal lateral geniculate nucleus of an APP23 mouse. (I) Lane 1: Synthetic A β 40/42. Lane 2: punch from the thalamus of an APP23 mouse. Note the strong A β 1-40 band and the weak A β 1-42 suggesting that A β 1-40 is the predominant peptide in seeded vessels.*

Figure 6 *Ultrastructure of seeded amyloid fibrils in APP23 transgenic mice. (A) Semithin section of the boundary of the granular cell layer (GCL) and the hilus (H) of the dentate gyrus of an APP23 mouse injected with tg brain extract. A β -immunoreactivity seems to be limited to the extracellular space. Amyloid plaques are indicated by asterisks. Counterstaining was achieved by toluidine blue. (B) Ultrastructural analysis reveals that A β (indicated by*

arrowheads) is located between neurons (N). A closer look suggests that A β deposits are located extracellularly. Scale bars: 10 μ m (A) and 1 μ m (B).

Figure 7 *Synthetic A β peptide injections at 1 day and 4 month post-infusion.* One day after the injection of synthetic A β 42 (A) and A β 40 (B) peptide a clot of the injected material in the hippocampus is detectable. (C,D) This clot disappeared and only minimal amount of amyloid deposits in the parenchyma and vessels were observed 4 month post-infusion. (E,F) Immunoreactivity is absent in nontransgenic control mice injected with synthetic A β 40 and A β 42. Scale bar: 350 μ m.

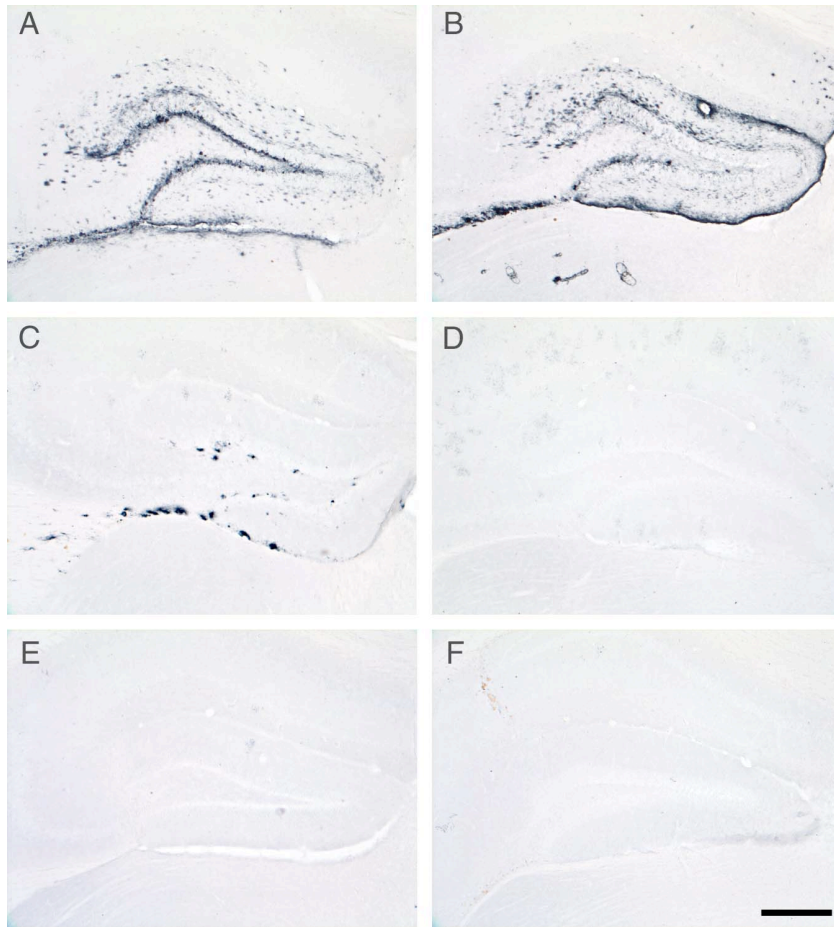


Figure 1

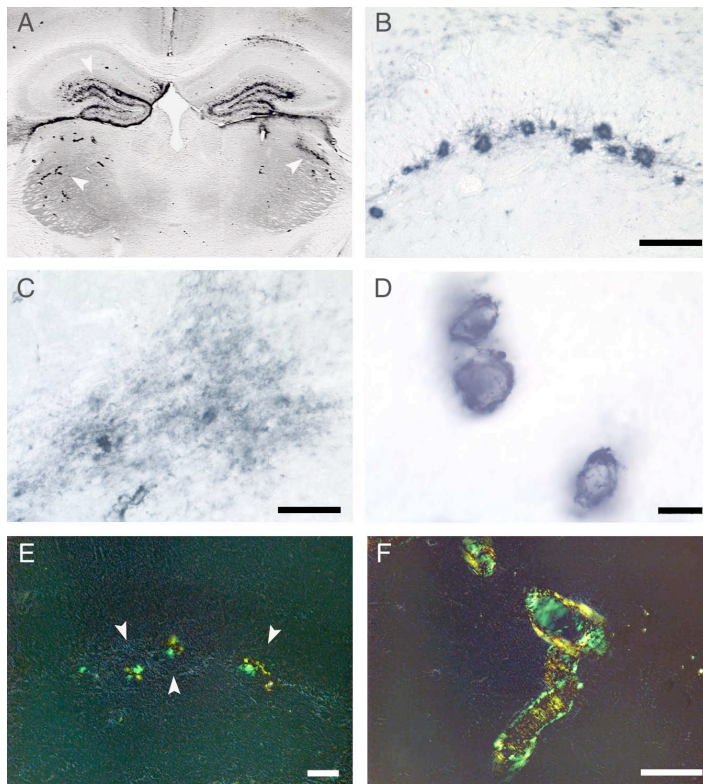


Figure 2

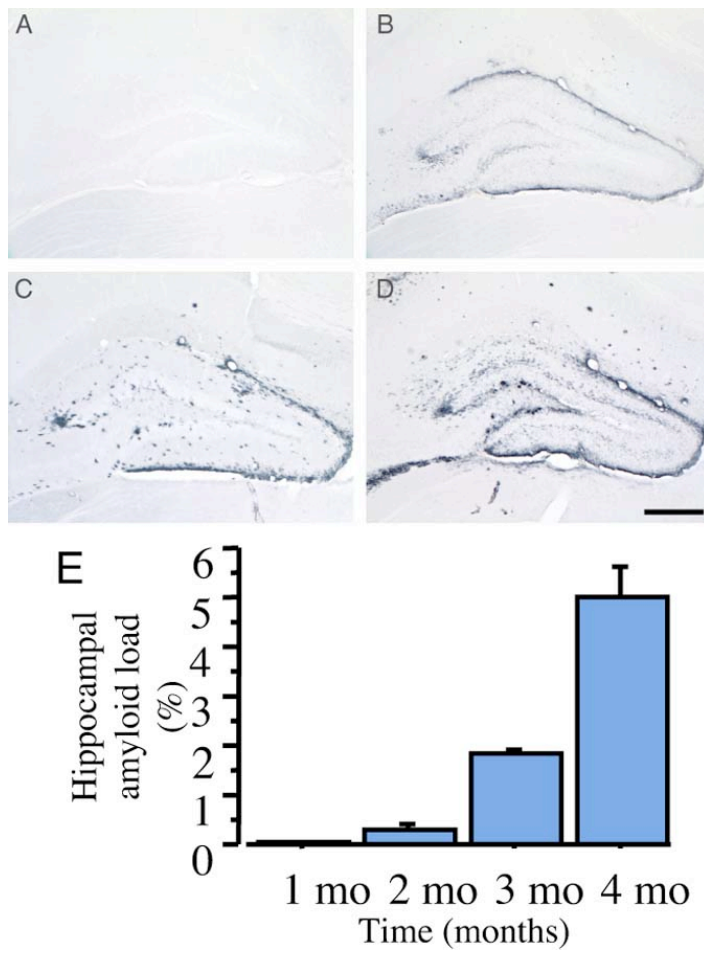


Figure 3

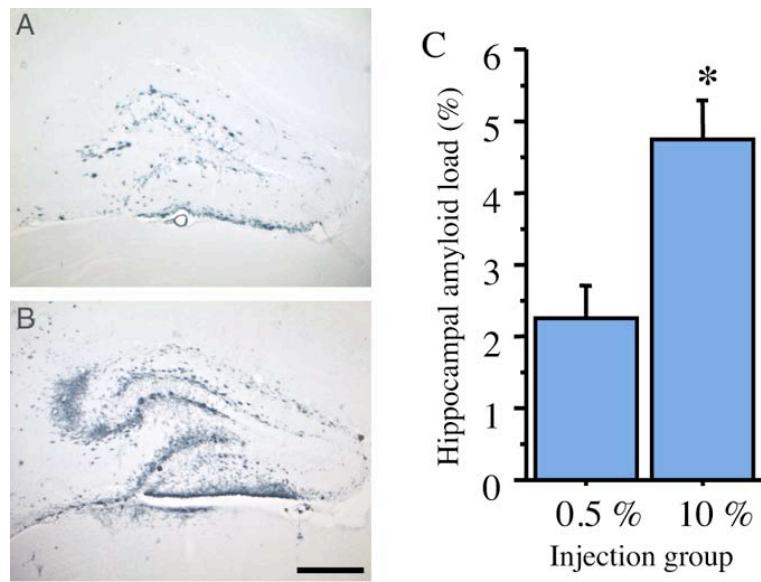


Figure 4

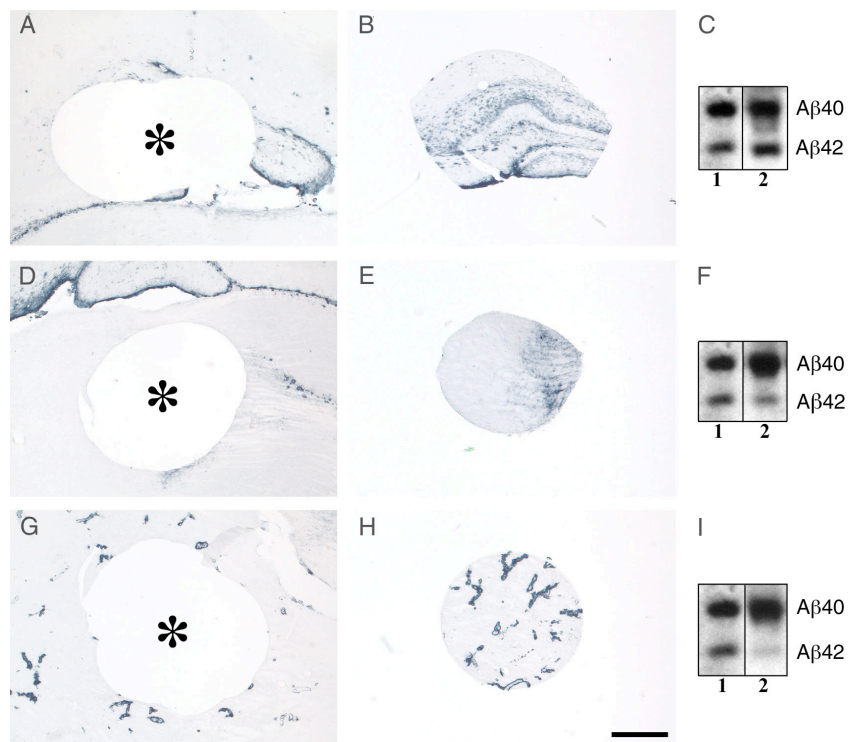


Figure 5

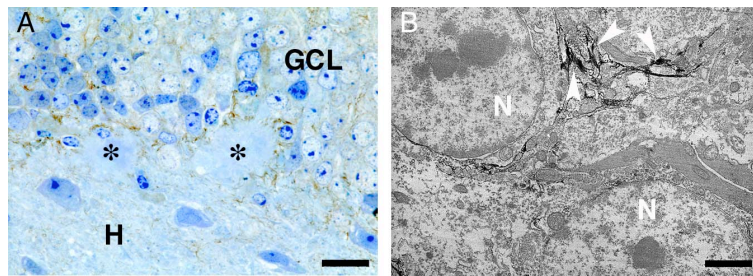


Figure 6

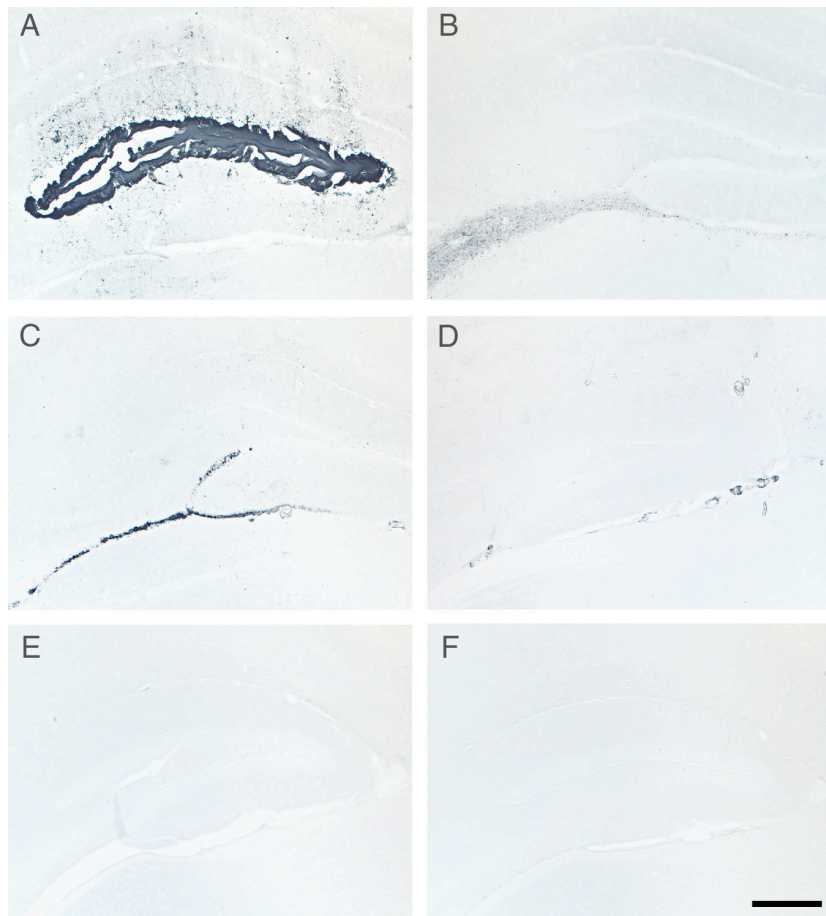


Figure 7

Table 1: *AD and animal brain extracts*

Case	Age	PMD ^a	Sex
AD case #1	85 y	6	f
AD case #2	74 y	7	m
Control #1	95 y	27	f
APP23 #1	26 mo	-	f
APP23 #2	26 mo	-	f
WT #1	24 mo	-	f

^a Postmortem delay (hours)

4. Restricted diffusion in the brain of transgenic mice with cerebral amyloidosis

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4.1 Abstract

A prominent hallmark of Alzheimer's disease (AD) pathology is cerebral amyloidosis. However, it is not clear how extracellular amyloid- β peptide (A β) deposition and amyloid formation compromise brain function and lead to dementia. It has been argued that extracellular amyloid deposition is neurotoxic and/or that soluble A β oligomers impair synaptic function. Amyloid deposits, on the other hand, may affect diffusion properties of the brain interstitium with implications for the transport of endogenous signalling molecules during synaptic and/or extrasynaptic transmission. We have used diffusion weighted magnetic resonance imaging (DWI) to study diffusion properties in brains of young (6-month-old) and aged (25-month-old) APP23 transgenic mice and control littermates. Our results demonstrate that fibrillar amyloid deposits and associated gliosis in brains of aged APP23 transgenic mice are accompanied by a reduction in the apparent diffusion coefficient (ADC). This decrease was most pronounced in neocortical areas with a high percentage of congophilic amyloid and was not significant in the caudate putamen, an area with only modest and diffuse amyloid deposition. These findings suggest that extracellular deposition of fibrillar amyloid and/or associated glial proliferation and hypertrophy cause restrictions to interstitial fluid diffusion. Reduced diffusivity within the interstitial space may impair volume transmission and therefore contribute to the cognitive impairment in AD.

4.2 Introduction

Alzheimer's disease (AD) is a late onset progressive neurodegenerative disorder that is characterized by aggregation of the amyloid- β peptide (A β) into senile plaques and cerebrovascular amyloid deposits (Selkoe, 1999). In the past few years, several transgenic mouse models of cerebral amyloidosis have been generated through overexpression of mutated amyloid precursor protein (APP). These mice develop amyloid plaques predominately in neocortex and hippocampus and share many features with the amyloid deposits in AD brain (Wong et al., 2002). In both AD patients and transgenic mice amyloid deposits are surrounded by activated microglia, and reactive astrocytes (Frautschy et al., 1998; Stalder et al., 1999; Akiyama et al., 2000). Diffusion and drainage of A β throughout the brain interstitium (Meyer-Luehmann et al., 2003) and along perivascular pathways (Calhoun et al., 1999; Weller et al., 2002; Preston et al., 2003) play an important role in the initiation and spread of cerebral amyloidosis. Once deposited, cerebral amyloid and accompanied pathophysiology may influence extracellular space volume and interstitial fluid (ISF) diffusion properties. A reduction in ISF diffusion impairs synaptic and/or extrasynaptic transmission and compromises neuronal function (Nicholson and Sykova, 1998).

The purpose of this study was to assess the apparent diffusion coefficient of water (ADC) in brains of APP23 transgenic mice using diffusion-weighted magnetic resonance imaging (DWI) followed by immunohistochemical analysis of cerebral amyloidosis. DWI is sensitive to microscopic incoherent motion of water molecules and can detect disturbances in ion and water homeostasis by measuring the ADC in the tissue (Moseley et al., 1990; Kauppinen et al., 1993).

4.3 Results

Reduced cortical Apparent Diffusion Coefficient in aged APP23 mice

Changes in ADC values in relation to cerebral amyloidosis were studied in young 6-month-old (pre-depositing) and aged 25-month-old male APP23 mice. We focused on the neocortex, a brain region that exhibits robust age-related congophilic amyloid deposition in aged APP23 mice, and on the caudate putamen, a brain region that develops only few and mostly diffuse amyloid in aged APP23 mice (Sturchler-Pierrat et al., 1997; Calhoun et al., 1998). Three coronal planes were selected and ROI within the caudate putamen and neocortex were defined (Fig. 1).

In caudate putamen ADC-values did not differ among the four groups of mice (Fig. 2). In contrast, ANOVA for the neocortex revealed a significant (age)x(genotype) interaction ($F(3,25) = 7.87, p < 0.001$). Subsequent Tukey post-hoc analysis revealed that aged APP23 mice had significantly reduced cortical ADC value compared to both age-matched wt mice ($p = 0.004$) and young APP23 mice ($p = 0.017$) (Fig. 2).

Visual inspection of pseudo-coloured ADC maps (Fig. 3) confirmed the robust decrease in ADC values in the neocortex. However, a closer analysis indicated that this decrease was mainly confined to the medial and dorso-lateral part of the neocortex that contributed to this decrease. Subregional analysis of neocortical structures (medial, dorso-lateral and ventro-lateral part, Fig. 1) revealed substantial ADC reductions in the medial and dorso-lateral cortex of the aged APP23 mice compared to aged-matched control mice ($p = 0.012$ and $p = 0.001$, respectively). However, ADC-values were not reduced in the ventro-lateral cortex of aged APP23 mice (Table 1).

Amyloid burden and associated gliosis in aged APP23

Following ADC measurements mice were sacrificed and brains analysed for amyloid burden. No amyloid plaques were detected in the 6-month-old male APP23 transgenic mice and in wt control mice (Fig. 4). In the 25-month-old APP23 mice A β -immunostaining revealed prominent amyloidosis throughout the neocortex but only little amyloid in the caudate putamen (Fig. 4D). Quantitative stereological analysis

yielded an amyloid load of 12.6 ± 1.2 % and 3.1 ± 0.7 % in the neocortex and caudate putamen, respectively (Table 2). Combined Congo red and A β -immunostaining revealed that the amyloid in the neocortex consisted mainly of compact plaques with some diffuse amyloid, whereas all the amyloid in the caudate putamen was of the diffuse type (Table 2, Fig. 4; Fig.5 A, B). Interestingly, there was also considerable region-specific variability in diffuse amyloid in neocortex with the ventro-lateral cortex exhibiting more diffuse amyloid compared to the dorso-lateral and medial part of the neocortex (Fig. 4).

Amyloid-associated activation of astrocytes and microglia has been associated with congophilic amyloid deposition (Frautschy et al., 1998; Stalder et al., 1999). Consistently, in the neocortex of aged APP23 mice, abundant hypertrophic GFAP-immunoreactive astrocytes closely associated with A β deposits were found (Fig. 5C), whereas in the caudate putamen only modest activation of astrocytes occurred (Fig. 5D). No such astrocytosis was found in neocortex and caudate putamen of young APP23 mice and wt control mice (Fig. 5E, F). Similarly, numerous Iba1-immunoreactive activated microglia were observed in the vicinity of compact plaques in neocortex of aged APP23 mice (Fig. 5G) while no such microgliosis was seen associated with the diffuse amyloid in the caudate putamen (Fig. 5H). No activated microglial cells were found in young APP23 mice and wt control mice (Fig. 5I, J).

4.4 Discussion

Diffusion-weighted MRI in APP23 mice revealed reduced ADC values in substantial portions of the neocortex of aged 25-month-old transgenic mice when compared to age-matched control mice and to young 6-month-old APP23 mice. In contrast, there was no reduction of ADC values in the caudate putamen of aged APP23 mice compared to age-matched wt control mice or young APP23 mice.

Consistent with previous findings (Sturchler-Pierrat et al., 1997; Calhoun et al., 1998) aged (25-month-old) APP23 mice revealed robust compact A β plaques in the neocortex but only few and diffuse amyloid in the caudate putamen. No amyloid deposition was detected in 6-month-old APP23 mice and in wt control mice. This observation suggests that the ADC reduction in the neocortex of the aged APP23 mice is related to the amyloid deposition in the brain of these mice. However, ADC values were not reduced in all cortical areas containing amyloid. For example, the ventro-lateral part of the neocortex did not show a reduction in ADC although this region exhibited a significant amyloid load. Interestingly, this brain region develops more diffuse amyloid compared to the medial and dorso-lateral parts of the neocortex. Thus, one may argue that not only the amount of amyloid deposition but also its nature (compact versus diffuse) determines diffusivity within cerebral tissue.

ADC values *in vivo* are thought to be affected by the net movement of compartmental water, transmembrane molecular exchange and hence an alteration of viscosity in the intra- and extracellular compartments (Le Bihan, 2003). In addition it has been shown that an increase in the tortuosity of the diffusion paths in the extracellular space leads to a decrease of the extracellular ADC (Norris et al., 1994; van der Toorn et al., 1996; Le Bihan, 2003). Increased tortuosity is a characteristic feature of injured brain tissue (Nicholson and Sykova, 1998; Sykova, 2001). A higher tortuosity may be caused by macromolecules produced by activated glial cells that in turn and may form diffusion barriers and lead to an increased tissue viscosity. In addition, astrogliosis and microgliosis, characterized by glial hypertrophy and an increase in glial number, interposed between the cells, can form diffusion barriers (Roitbak and Sykova, 1999). Recently it has been suggested that increased tortuosity during ischemia, and other

conditions involving cellular swelling, is caused by the formation of dead-space microdomains in the extracellular space. Such dead-end pores occluding the gaps between the cells can be formed by astrocytic expansions occurring during gliosis (Hrabetova et al., 2003).

In APP23 mice a substantial inflammatory reaction is observed and exemplified by activated and hypertrophic astrocytes and microglia (Sturchler-Pierrat et al., 1997; Bornemann et al., 2001). This gliosis occurs in vicinity of congophilic compact but not diffuse amyloid (Stalder et al., 1999). Therefore, we investigated whether ADC changes found in the present study might be attributable to the gliosis and not the amyloid deposition *per se*. Indeed, prominent astrocytosis and microgliosis was observed in the neocortex with virtually no gliosis in the caudate-putamen in the aged APP23 mice. An expected difference in gliosis between the medial and dorso-lateral as compared to ventro-lateral neocortex was less obvious but might have been masked by a ceiling effect due to a prominent gliosis throughout all neocortical areas. Thus, the observed ADC reduction in neocortex of aged APP23 mice might be related to the gliosis associated with congophilic compact amyloid plaques.

ADC measurements in AD patients revealed a loss of diffusion anisotropy in the white matter and an increased diffusivity within the hippocampus (Sandson et al., 1999). This ADC increase has been related to a loss of neurons as it correlates with the brain atrophy observed in the hippocampus (Sandson et al., 1999). However, other studies reported similar ADC values in the hippocampus of AD patients compared to controls (Hanyu et al., 1998; Bozzao et al., 2001). These studies suggest that an increase in water diffusivity due to neuron loss is compensated for by the gliosis that accompanies neurodegeneration and the accumulation of senile plaques and tangles. Thus, the present observation of a significant reduction of diffusion in APP23 mice may be explained by the only modest neuron loss but prominent gliosis in the neocortex of these mice (Stalder et al., 1999; Bondolfi et al., 2002).

In summary we have shown that aged 25-month-old APP23 mice exhibit reduced cortical ADC values. This reduction goes in parallel with the formation of compact amyloid plaques and might be attributable to the pronounced gliosis associated with the

amyloid deposits. Alterations of ISF diffusion properties may change synaptic and/or extrasynaptic (volume) transmission and compromise neuronal function (Zoli et al., 1999). Thus, reduced ISF diffusion may contribute to the cognitive deficits previously reported in these mice (Lalonde et al., 2002; Kelly et al., 2003) and to the AD cognitive impairment that has primarily been attributed to neuron and synaptic loss.

4.5 Materials and Methods

Animals

APP23 mice express human mutated APP₇₅₁ under the control of the murine Thy-1 promoter and exhibit the age-related deposition of cerebral amyloid (Sturchler-Pierrat et al., 1997). Mice have been generated on a B6D2 background and have subsequently been backcrossed for more than 8 generations to B6. For the present experiments 6-month-old (n=8) and 25-month-old (n=10) male APP23 mice and age-matched wild-type (wt) control mice were used (n=5 and n=6, respectively).

Animal preparation

All animal experiments were carried out in strict adherence to the Swiss Law for Animal Protection. For MRI experiments animals were anesthetized using an initial dose of 3 % isoflurane (Abbott, Cham, Switzerland) in air:O₂ (2:1), intubated with a tube made from polyethylene (PE; ID/OD: 0.4/0.8 mm) and artificially ventilated using a ventilator for small animals (KTR 3, Alfos Electronics, Biel-Benken, Switzerland). The animals were positioned in a cradle made from Plexiglas and kept anesthetized with 1.4 % isoflurane in air:O₂ (2:1). The animals were paralysed with 10 mg/kg (intravenously, i.v.) gallamine triethiodide (Aldrich, Milwaukee, USA) in saline (3 mg/ml). Blood carbon dioxide levels were monitored transcutaneously (PtcCO₂) during the experiment using a paediatric monitoring device (TCM3, Radiometer Copenhagen) as described in previous studies (Mueggler et al., 2001). Body temperature was maintained at 36.5 ± 1°C using warm air, regulated by a rectal temperature probe (MLT415, ADInstruments, Hastings, UK).

MRI protocol

DWI experiments were carried out on a Biospec system equipped with a PharmaScan 70/16 magnet (Bruker Medical System, Ettlingen, Germany) equipped with actively shielded gradient coils (maximum gradient strength: 200 mT/m; rise time <150 μ s). The radiofrequency probe was a birdcage resonator of 28 mm inner diameter. A spin-echo sequence with echo time TE=50 ms and repetition time TR=1325 ms comprising a diffusion-sensitizing pulsed gradient pair according to Stejskal and Tanner (1965) was used for DWI (Stejskal and Tanner, 1965). Seven 1 mm thick coronal slices, separated by a 0.6 mm inter-slice gap, were imaged with a field-of-view of 2.56 x 2.56 cm² and a matrix size of 128 x 64. For the determination of the apparent diffusion coefficient (ADC) of water, diffusion-encoding was applied along slice gradient direction (for coronal images caudal to rostral) with five b-values covering a range from 0 to 2000 s/mm². ADC values are determined by the volume fraction of the various tissue compartments and by the water diffusion coefficient within each of these compartments, which by itself depends on fluid viscosity and tortuosity λ , which is a measure for the steric hindrance of diffusion within a compartment ($\lambda^2 = \text{free diffusion coefficient}/\text{apparent diffusion coefficient}$).

Tissue preparation and immunohistochemistry

One day after the DW-MRI experiments mice were sacrificed by decapitation under deep inhalation anaesthesia. Brains were removed and post fixed for 4d in 4% paraformaldehyde then dehydrated overnight in 30% sucrose. After freezing, serial coronal sections of 25 μ m thickness were cut on a freezing-sliding microtome throughout the brains. Sections were stained immunohistochemically according to previously published protocols (Stalder et al., 1999) by using the avidin-biotin-peroxidase complex method with SG blue as chromogen (Vector Laboratories, Burlingame, CA). The following antibodies were used: polyclonal antibody NT12 to A β (Sturchler-Pierrat et al., 1997); polyclonal antibody to glial fibrillary acidic protein (GFAP; Dako, Glostrup, Denmark); and polyclonal antibody to ionized calcium binding adaptor molecule 1 (Iba1; courtesy of Y. Imai, Tokyo, Japan) (Ohsawa et al., 2000). Some sections were double stained with NT12/GFAP and NT12/ Iba1 antibodies using SG blue and DAB as chromogens.

Stereological assessment of amyloid load

Stereological techniques were used to estimate the amyloid load in the neocortex and caudate putamen. To this end every 24th section throughout the brain was immunostained with NT12 yielding 10-12 sections for the neocortex and 4-5 section for the caudate putamen. The amyloid load was determined by calculating the area fraction occupied by amyloid in two-dimensional disectors on a single focal plane (20X objective; 0.45 numerical aperture (Calhoun et al., 1998)). One set of sections was double stained for Congo red and NT12 and the percentage of congophilic (compact) amyloid was estimated. Stereological analysis was performed with the aid of Stereologer™ software and a motorized x-y-z stage coupled to a video-microscopy system (Systems Planning and Analysis, Inc., Alexandria, VA). Region definitions of neocortex and caudate putamen were based upon mouse brain atlas (Franklin and Paxinos, 1997).

Data analysis

Quantitative ADC maps were calculated on a pixel-by-pixel basis by linear regression analysis using the model function:

$$\ln(S(b)/S_0) = -b \cdot ADC \quad [1]$$

where $S(b)$ refers to the pixel intensity as a function of b (the diffusion-sensitizing gradient value), S_0 refers to the signal intensity of the pixel with zero diffusion gradient amplitude ($b=0$).

Analysis was carried out using in-house developed imaging analysis software (Biomap v3.1). To assess ADC values from different brain structures regions-of-interest (ROIs) were defined (Fig. 1).

Statistics

The ADC values of both hemispheres measured in the DWI experiments were averaged to yield one value per region. Data in text and figures are expressed as mean values \pm standard deviation (SD), unless stated otherwise. Multiple comparisons were evaluated by the analysis of variance (ANOVA) and subsequent post-hoc group comparison

(Tukey test). Probability values of less than 0.05 were considered as statistically significant.

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4.6 References

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4.7 Figure Captions

- Figure 1 Coronal sections through mouse brain (1.6, 0, -1.6 mm anterior to Bregma) with regions-of-interest (ROI) selected for quantitative data analysis indicated. ROI 1 was defined for the lateral and medial caudate putamen. ROI 2 was defined for neocortex. In a subsequent analysis ROI 2 was further subdivided into medial (2a) cortex; dorso-lateral cortex (2b), and ventro-lateral cortex (2c).
- Figure 2 Apparent diffusion coefficient (ADC) of water in neocortex (ROI 1) and caudate putamen (ROI 2) of 6- and 25-month-old APP23 mice and age-matched littermates (wt). Data reveal a significant reduction of cortical ADC in aged APP23 animals. Values are given as mean \pm SD. *= $p < 0.05$; **= $p < 0.01$.
- Figure 3 Colour-coded representative ADC maps of a coronal brain section at the level of Bregma for a 6- (A) and 25-month-old APP23 mouse (D) and age-matched controls (6 months (B), 25 months (C)). ADC values given as $10^{-6} \text{ cm}^2 \text{ s}^{-1}$. A significant ADC decline, colour-coded in blue, is detectable in the medial and dorso-lateral cortex of aged APP23 mice. In the dorso-lateral cortex of the 6-month-old APP23 mouse few areas with reduced ADC are visible.
- Figure 4 Histological analysis of A β deposition in APP23 and wt mice. A β immunostaining with the NT12 antibody shows no amyloid deposits in young wt mice (A), young APP23 transgenic mice (B) and aged wt mice (C), while aged APP23 transgenic mice displayed massive amyloid deposits predominantly in neocortex (D). Histological analysis of A β deposition in APP23 and wt mice. A β immunostaining with the NT12 antibody shows no amyloid deposits in young wt mice (A), young APP23 transgenic mice (B)

and aged wt mice (C), while aged APP23 transgenic mice displayed massive amyloid deposits predominantly in neocortex (D).

Figure 5 Activation of plaque-associated astro- and microglia in 25-month-old APP23 transgenic mice. In aged APP23 neocortex massive compact amyloid plaque formation was observed (A) whereas in the caudate putamen only diffuse plaques was found (B). A β related astrocytosis was extensive around plaques in the neocortex (C) compared to the weakly GFAP-positive astroglia in the caudate putamen (D) and in wt control mice (E,F). A β deposits in the neocortex of 25-month-old APP23 mice were also associated with activated and Iba1-immunoreactive microglia (G). In contrast no activation of microglia was observed in the caudate putamen (H) and neocortex and caudate putamen of wt control mice (I, J). Bars are 100 μ m (A,B) and 60 μ m (C-J).

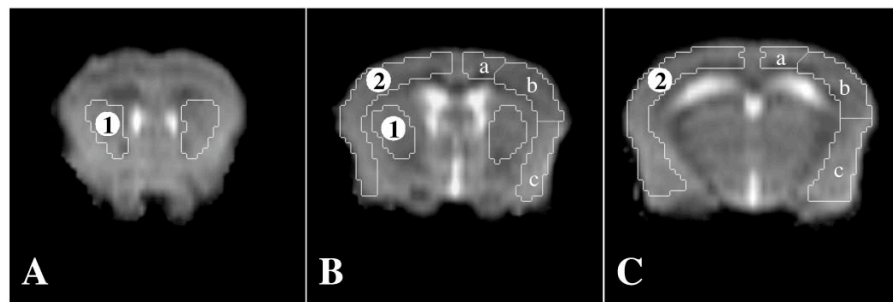


Figure 1

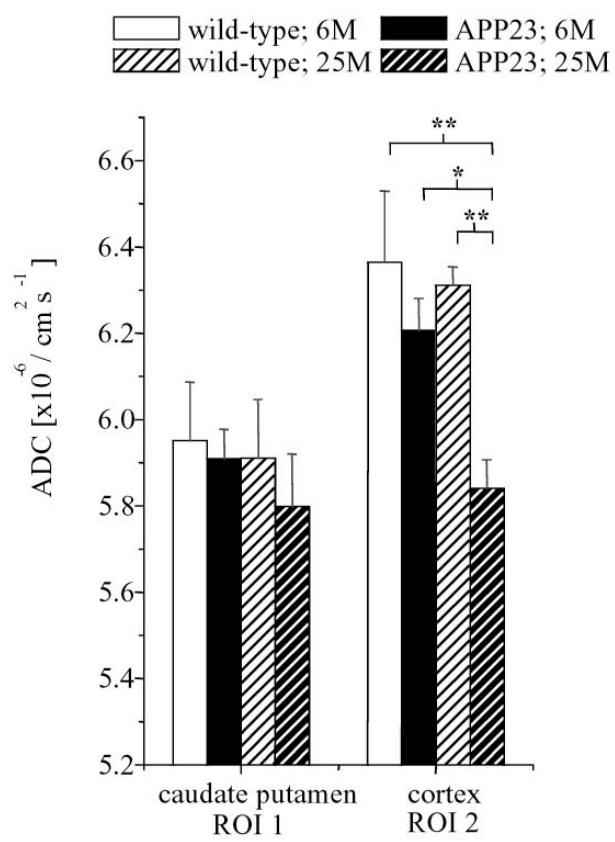


Figure 2

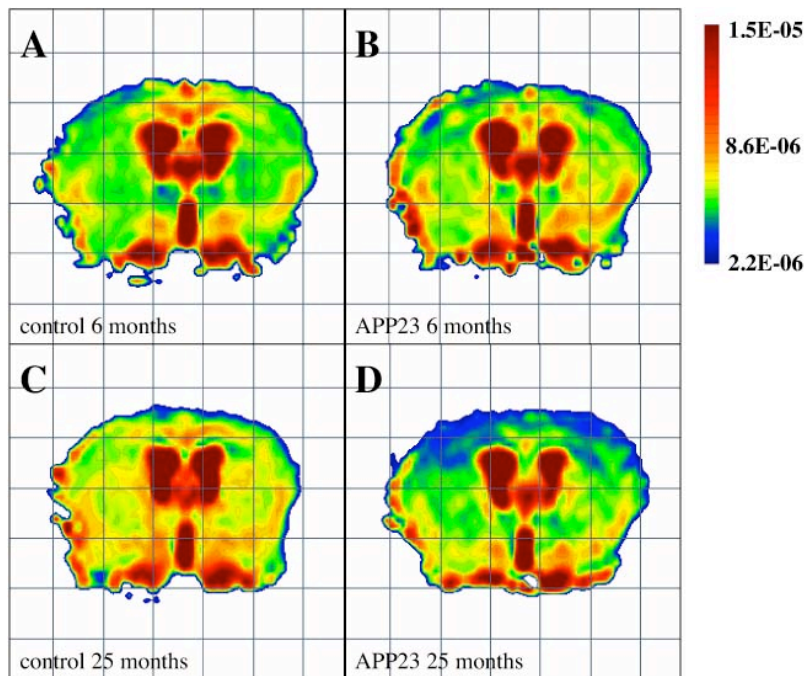


Figure 3

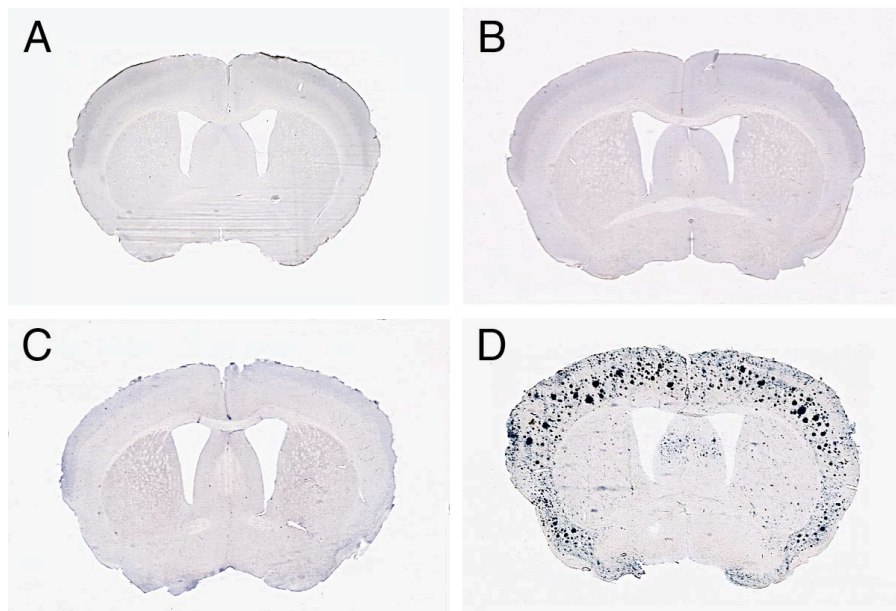


Figure 4

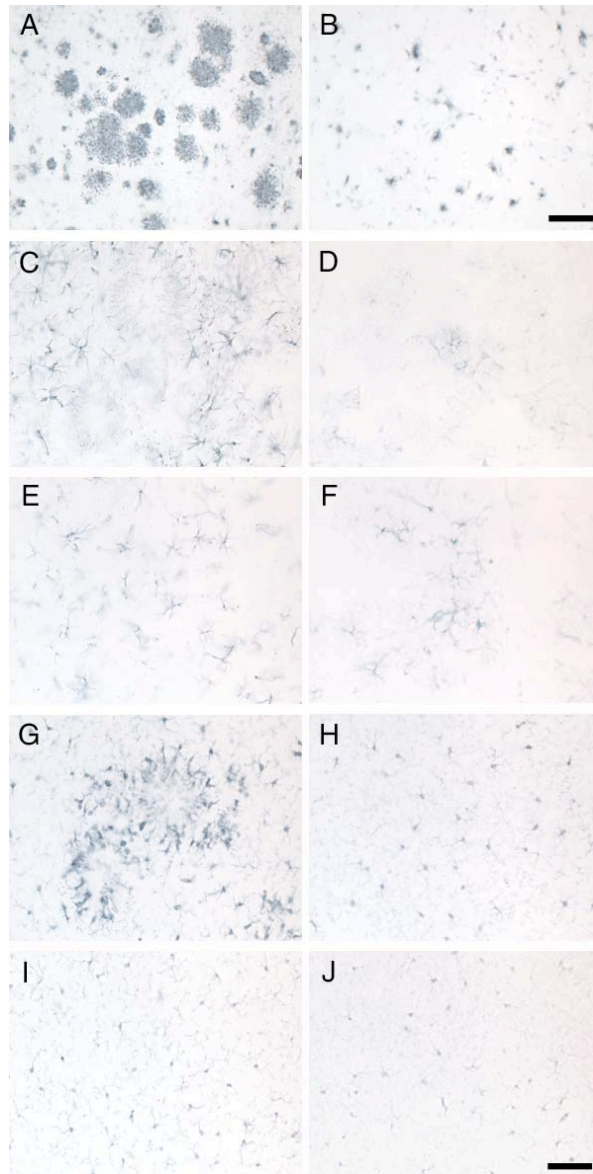


Figure 5

Table 1 ADC values for different brain regions of wild-type control and APP23 mice

region analyzed	6 months				25 months			
	wildtype N=5		APP23 N=8		wildtype N=6		APP23 N=10	
	mean	SD	mean	SD	mean	SD	mean	SD
striatum	5.95	0.3	5.9	0.19	5.91	0.57	5.79	0.43
ventro-lateral cortex	6.96	0.43	6.91	0.22	6.88	0.22	6.86	0.34
dorso-lateral cortex	5.61	0.45	5.47	0.25	5.8	0.21	5.09 ^{§, &}	0.33
medial cortex	6.52	0.54	6.23	0.38	6.25	0.29	5.57 ^{§, &, *}	0.37

Significant differences ($p < 0.05$) are indicated (for p values see text). Aged APP23 compared to 6-month-old control[§], 25-month-old control[&] and 6-month-old APP23* (ANOVA Tukey test). Diffusion coefficient calculated as $\times 10^{-6} \text{ cm}^2 \text{ s}^{-1}$

Table 2 Amyloid load in 25-month-old APP23 mice

region analyzed		cortex		caudate putamen	
		mean [%]	SEM	mean [%]	SEM
total amyloid load		12.6	1.2	3.2	0.7
composed of	A. compact plaques	92.8	1.1	0	
	B. diffuse plaques	7.2	1.1	100	

5. Conclusion

This thesis aims at understanding some of the underlying mechanisms of plaque formation and associated neurodegeneration in cerebral amyloidosis. Although the generation of A β and the processing of its precursor APP has been studied extensively at the molecular and cellular level, it still remains a mystery how neuronally derived A β propagates and forms fibrillary amyloid deposition *in vivo* and what factors influence this pathological process. The availability of transgenic mice that reflect the neuropathological spectrum of AD enabled neuroscientists to explore the relationship between the pathological features, biochemical abnormalities, and memory dysfunctions occurring in the human AD brain.

In this thesis, several methods have been applied in order to identify the mechanisms of cerebral A β accumulation *in vivo*. As a first attempt to investigate early stages of abnormal protein deposition and related neurotoxicity, we have used neural transplantation techniques. These experiments have shown that neuronally produced A β diffuses over considerable distances in the extracellular space of the brain before it aggregates, deposits extracellularly and finally causes neurodegeneration. There has been a long debate in the field about the origin of plaques and a longstanding question in AD research is whether amyloid formation occurs intra- or extracellularly. Our experiments clearly indicate that cerebral amyloidosis does not require locally generated intracellular A β to initiate A β deposition. We conclude that the properties of the host influence plaque formation in the transplant and that extracellular amyloid can cause neurodegeneration in the absence of intracellular A β production.

The second part of the present study has focused on the identification of factors which initiate or are involved in the seeding process and that induce A β fibril formation in APP23 transgenic mice and in AD patients. Preliminary results indicate that cerebral amyloidosis can be induced by the infusion of A β -containing brain extract into APP23 transgenic mice, and that this seeded polymerization of A β is both time- and concentration-dependent. Interestingly, the infusion of synthetic A β peptide has so far failed to induce amyloid seeding. Future experiments will have to examine whether A β itself is the seeding factor, and in particular, what A β conformation is important to

initiate plaque formation *in vivo*. The identification of the seeding agent and cofactors would open new therapeutic strategies to prevent cerebral A β deposition in diseases such as AD.

The findings presented above highlight the importance of transport or diffusion of soluble A β in neural tissue as a major factor implicated in amyloid formation and neurotoxicity. In order to characterize A β diffusion properties in more detail, non-invasive diffusion-weighted MRI measurements have been applied to APP23 transgenic mice in the final part of this thesis. The results strongly suggest that amyloid deposition and the associated gliosis build a hindrance in the brain tissue leading to decreased interstitial fluid diffusion properties, which may finally change synaptic and extrasynaptic transmission. Diffusion measurements by MRI would therefore be a valuable tool to detect early alterations in the diffusion properties in AD patients.

What can we learn from these studies? The discovery of the underlying origin of cerebral amyloidosis is important in view of the increasing relevance of protein misfolding diseases for the aging society. Further progress on the key question about the mechanism of plaque formation and associated neurodegeneration in cerebral amyloidosis discussed in this thesis may also shed some light on other protein misfolding diseases. Uncovering some of the fundamental mechanisms of cerebral amyloidosis will give us the opportunity to develop new therapeutic strategies in the future to cure these devastating diseases.

6. Abbreviations

A β	β -amyloid
AD	Alzheimer's disease
ADC	apparent diffusion coefficient
ALS	amyotrophic lateral sklerosis
APP	amyloid precursor protein
APOE	apolipoprotein E
CBF	cerebrovascular blood flow
CSF	cerebrospinal fluid
CAA	cerebral amyloid angiopathy
DWI	diffusion weighted magnetic resonance imaging
ECS	extracellular space
FAD	familial Alzheimer's disease
HD	Huntington's disease
IDE	insulin degrading enzyme
ISF	interstitial fluid
MRI	magnetic resonance imaging
NFT	neurofibrillary tangle
NSAID	non-steroidal anti-inflammatory drugs
PD	Parkinson's disease
PrP	prion protein
PS	presenilin
SAP	serum amyloid P component
TSE	transmissible spongiform encephalopathy

7. Curriculum Vitae

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8. Bibliography

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Invited Presentations

1. **Meyer-Luehmann, M.** : Extracellular amyloid formation causes neurodegeneration. 1st Symposia of the Hertie-Institute of Clinical Brain Research, Tübingen, 2003.

2. **Meyer-Luehmann, M.** : Mechanism of cerebral amyloidosis. Departmental lecture series, Institute of Anatomy, Basel, 2003.

3. **Meyer-Luehmann, M.** : Initiation and spread of cerebral amyloidosis in transgenic mice. Alzheimer meeting Eltem, Neurex, Basel, 2003.

During my studies and PhD thesis, lectures and seminars of the following persons were taken: S. Arber, Y.-A. Barde, B. Bettler, H.-R. Brenner, M. Burger, P. Caroni, A. Gratwohl, N. Hynes, M. Jucker, J. Kapfhammer, A. Lüthi, A. Lüthi, A. Matus, T. Meier, A. Merlo, H. Moch, D. Monard, C. Moroni, D. Oertli, P. Piguet, U. Müller, H. Reichert, C. Rochlitz, M. Rudin, M. Rüegg, N. Schaeren-Wiemers, G. Spagnoli and E. Stöckli.