Network: Systematic Gene Identification and Functional Analyses in Common CNS Disorders

Project: Transgenic Mouse Models to Study Family Forms of Cerebral Amyloid Angiopathy

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Introduction

In normal aging and Alzheimer's disease (AD), Abeta deposition occurs in both parenchymal amyloid plaques and in vessels (cerebral amyloid angiopathy, CAA). However, CAA can also occur in the absence of parenchymal amyloid plaques. For example, patients with hereditary cerebral hemorrhage with amyloidosis-Dutch type (HCHWA-D) develop a severe form of Aβ-CAA but very few amyloid plaques and no neurofibrillary tangles (Levy *et al.*, 1990). These patients suffer recurrent intracerebral hemorrhages leading to death between the ages 45 and 55. HCHWA-D is caused by a point mutation Glu->Gln at codon 693 of the amyloid-beta precursor protein (APP) (amino acid 22 of Aβ)(Levy *et al* 1990).

Although the most common form of cerebral amyloidosis is of the Abeta-type, there are other proteins which have been linked to severe familial forms of cerebral amyloidosis. Hereditary cerebral hemorrhage with amyloidosis Icelandictype (HCHWA-I), is caused by a point mutation Leu->Gln at position 68 of cystatin C (Ghiso et al., 1986; Levy et al., 1989). HCHWA-I patients suffer the first often fatal hemorrhagic stroke often in their twenties (Olafsson and Grubb, 1999). The amyloid in these patients (ACys) consists of mutated cystatin with an N-terminal truncation of 10 amino acids. Familial British Dementia (FBD) and Familial Danish Dementia (FDD) are autosomal dominant disorders with death occurring at 40-60 years of age (Vidal et al., 1999, 2000). The hallmark lesion in both diseases is cerebral amyloidosis with predominant CAA. In addition the patients develop neurofibrillary tangles (Vidal et al., 1999, 2000). Both diseases are caused by mutations in the recently discovered BRI gene. In FBD a stop codon mutation generates a longer open reading frame while in FDD a 10-nucleotide duplication immediately before the stop codon produces a frame-shift mutation. Both mutations generate 277-amino acid long precursor proteins, ABriPP and ADanPP, that are 11 amino acids longer compared to the 266-amino acid-long wildtype protein (BriPP). In both FBD and FDD, furin-cleavage releases a 34-amino acid carboxy-terminal of the mutant precursor protein to generate the amyloidogenic ABri and ADan peptides, respectively, which are then deposited as amyloid in the brain (Vidal et al., 1999, 2000; Kim et al., 1999).

Although normal aging, AD, HCHWA-D/HCHWA-I, and FDB/FDD share cerebral CAA as common pathology, there are a puzzling variety of additional neuropathological lesions and clinical phenotypes. It is well established that CAA is a risk factor for spontaneous and often fatal hemorrhagic stroke. However, it has also become clear that CAA affects cognition independent of strokes: FBD and FDD patients do not suffer hemorrhage but are demented (Vidal et al., 1999, 2000). Dementia in HCHWA-D is associated with CAA independent of hemorrhagic strokes (Natte et al., 2001). Moreover, CAA in normal aging and AD has recently been linked to cognitive impairment (Pfeifer et al., 2002a) and cerebral hemorrhage following anti-A_β-immunotherapy, one of the currently followed approaches to cure AD (Pfeifer et al., 2002b). Thus, mechanisms by which these amyloidogenic proteins are deposited in the vessel wall and lead to stroke and/or dementia remain puzzling.

Project Status

Over the last five years we have studied APP transgenic mouse models that develop amyloid plaques and CAA. We



have also demonstrated that amyloid plaque formation in these mice leads to region-specific neuron death, synaptic and cholinergic dysfunction, and neuroinflammation (Stürchler-Pierrat *et al.*, 1997; Calhoun *et al.*, 1998; Phinney *et al.*, 1999; Stalder *et al.*, 1999; Bondolfi *et al.*, 2002; Boncristiano *et al.*, 2002). Furthermore, we have demonstrated that CAA in these mice leads to a loss of vascular smooth muscle cells, aneurysmal vasodilatation, vessel wall rupture, and recurrent hemorrhages (Calhoun *et al.*, 1999; Winkler *et al.*, 2001; Pfeifer *et al.*, 2002b).

To study the significance of CAA proper, we have developed new APP transgenic mice harboring the HCHWA-D mutation under a neuron-specific Thy-1 promoter (APPDutch mice). These mice develop extensive CAA with only very few parenchymal deposits (Fig. 1). CAA in these mice leads to severe CAA, smooth muscle cell degeneration, cerebral hemorrhages, and neuro-inflammation. In contrast, neuronal overexpression of wild-type human APP (APPwt mice) results in predominantly parenchymal amyloidosis. In HCHWA-D and APPDutch mice the A_β40:42 ratio is significantly higher than in AD and APPwt mice, and significant wild-type human or murine A_{β40}, respectively, is co-deposited with the ABDutch. Genetically shifting the ratio of A_βDutch40:42 towards A_βDutch42 by crossing APPDutch mice with mutated presenilin 1 transgenic mice redistributes the amyloid pathology from the vasculature to the parenchyma. The understanding that different $A\beta$ species can drive amyloid pathology in different cerebral compartments has implications for current anti-amyloid therapeutic strategies This work has recently been published (Herzig et al., 2004).

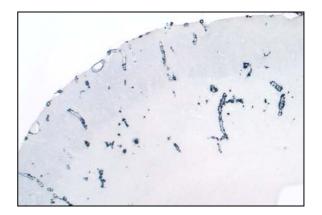


Fig 1: Cerebral amyloid angiopathy in an aged APPDutch transgenic mouse.

To further understand the mechanism by which amyloidogenic proteins are deposited in the vessel wall and lead to stroke and/or dementia, we have also generated cystatin C transgenic mice with the HCHWA-I mutation (Cystatin C_{L680} tg mice) and ABriPP transgenic mice with the FBD mutation. For all these transgenic mouse lines the Thy-1 promoter has been used and mice have been generated on a pure B6 background. Unfortunately, the present Cystatin C_{L680} tg mice and ABriPP tg mice do not develop cerebral amyloid deposits at least until 24 months of age.



Outlook

The overall aim of our proposal is to elucidate the pathobiological mechanism leading to CAA and to provide a basis for therapeutic strategies. To this end we suggested (a) to generate various transgenic mouse models of familial forms of CAA, and (b) to use state-of-the-art morphological, biochemical, and gene expression and protein profiling to study mechanism how CAA leads to stroke and neurodegeneration.

While such analyses of transgenic mice are ongoing for APPDutch mice, we have started new efforts to generate additional Cystatin C and AbriPP/ADanPP transgenic mice using new refined molecular constructs. To understand the pathomechanism of British and Danish dementia we have also initiated studies aiming at generating *Bri*-null mice. We anticipate that these new efforts and the current analysis of APPDutch mice will greatly contribute to the understanding of the pathophysiology and therapy of CAA.

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