CEREBROSPINAL FLUID RESEARCH

# RESEARCH

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# Effects of cerebrovascular disease on amyloid precursor protein metabolites in cerebrospinal fluid

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# Abstract

**Background:** Alzheimer's disease (AD) and cerebrovascular disease (CVD) including chronic small vessel disease of the brain (SVD) are the most frequent causes of dementia. AD is associated with metabolism of amyloid precursor protein (APP) and low levels of amyloid- $\beta$  peptide (A $\beta$ ) X-42 in the cerebrospinal fluid (CSF). CVD and SVD are established risk factors for AD, brain white matter lesions (WML) are established surrogate markers for SVD and are also associated with reduced CSF A $\beta$ X-42.

A cohort survey was performed to examine whether SVD or acute CVD affects APP metabolism and to explore a potential association between WML and APP metabolism in two groups; cognitively impaired patients, subjective and mild (SCI and MCI) and stroke patients. Through measurements of CSF APP metabolite levels in patients with a wide range of WML volumes, this study aimed to determine how SVD influences APP metabolism.

**Methods:** Sixty-three patients were included: 37 with subjective cognitive impairment (SCI) or mild cognitive impairment (MCI) without stroke, and 26 after acute stroke. Chronic and acute WML volume and infarct volume were determined by magnetic resonance imaging (MRI) post-scan processing, and CSF levels of  $\alpha$ - and  $\beta$ -cleaved soluble APP (sAPP- $\alpha$  and sAPP- $\beta$ , A $\beta$ X-38, A $\beta$ X-40 and A $\beta$ X-42) were determined. The Mann-Whitney test was used to compare the patient groups. Chronic and acute WML volumes, infarct volume, age, and sex were used as predictors for CSF biomarker levels in linear regression analysis.

**Results:** CSF levels of sAPP- $\alpha$  and sAPP- $\beta$  were strongly correlated (r = 0.95, p < 0.001) and lower levels of these biomarkers were found in the stroke group than in the SCI/MCI group; median sAPP- $\alpha$  499.5 vs. 698.0 ng/mL (p < 0.001), sAPP- $\beta$  258.0 vs. 329.0 ng/mL (p < 0.005). CSF levels of sAPP- $\alpha$ , sAPP- $\beta$ , A $\beta$ X-38, A $\beta$ X-40 and A $\beta$ X-42 were inversely correlated with chronic WML volume ( $p \le 0.005$ ;  $p \le 0.01$ ;  $p \le 0.05$ ;  $p \le 0.05$ ;  $p \le 0.05$  respectively), but not with acute WML or infarct volumes.

**Conclusions:** Lower CSF levels of sAPP- $\alpha$  and sAPP- $\beta$  in the stroke group than in the SCI/MCI group and an inverse correlation with chronic WML indicate that ischemia lowers the levels of CSF sAPP metabolites and suggests that APP axonal transport or metabolism may be affected in SVD of the brain.

# Background

Alzheimer's disease (AD) and cerebrovascular disease (CVD) are the most frequent causes of dementia. Familial AD is associated with metabolism of the transmembrane amyloid precursor protein (APP) and mutations in the APP gene [1,2], while less is known about the etiology of sporadic AD [3]. However, findings in

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histopathology [4] and molecular imaging [5] imply that amyloid metabolism is also involved in sporadic cases. After fast axonal transport of APP to synaptic terminals [6],  $\alpha$ - or  $\beta$ -secretase cleaves the protein into soluble APP (sAPP- $\alpha$  or sAPP- $\beta$ ) and C-terminal fragments ( $\alpha$ CTFs and  $\beta$ CTFs) [7]. Subsequent cleavage of  $\beta$ CTFs (by  $\gamma$ -secretase) yields amyloid  $\beta$  (A $\beta$ ) peptides X-38, X-40 and X-42 [8]. A $\beta$ X-42 is prone to deposition in amyloid plaques [9], and an association between low



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levels of A $\beta$ X-42 in cerebrospinal fluid (CSF) and presence of amyloid plaques has been shown both in molecular imaging [5] and post-mortem histopathological studies [10]. Low CSF A $\beta$ X-42 is also a predictor of AD [11]. APP and sAPP- $\alpha$  are important factors for neurite outgrowth [12] and neuronal plasticity and memory [13,14].

We have found similar CSF levels of sAPP- $\alpha$  and sAPP- $\beta$  in sporadic AD and mild cognitive impairment (MCI) [15] to control subjects [16]. Subjective cognitive impairment (SCI) [17] is a pre-MCI stage characterized by subjectively impaired cognition which is not demonstrable with objective screening tests. SCI [18], MCI, white matter lesions (WML) [19] and stroke [20,21] all increase the risk of dementia and AD. WML are established surrogate markers of chronic small vessel disease of the brain (SVD) [22,23] and are frequently seen on T2-weighted magnetic resonance imaging (MRI) scans of individuals with and without dementia [24], and are present in increased amount in AD [25]. They are associated with reduced CSF levels of  $A\beta X$ -42 in various diseases [26-28], and an inverse correlation between WML volume and CSF level of sAPP- $\alpha$  and sAPP- $\beta$  has been demonstrated in non-demented elderly people [29]. Experimental stroke [30] and ischemia [31] lead to an increased production of APP, upregulation of  $\beta$ -secretase activity [32], and an accumulation of A $\beta$  peptides and APP around ischemic WMLs [31]. Axonal transport is impeded both by WML and APP metabolites [33,34]. Thus, APP over-expression may then impede axonal transport, also of APP, and impair neuronal plasticity and survival [35]. In ischemic conditions, CSF levels of APP metabolites may be influenced by increased gene expression, impeded axonal transport and deposition in plaques (which mainly contributes to reduced levels of CSF A $\beta$ X-42) [10]. In the case of impeded transport, low cortical levels of APP [14] and sAPP- $\alpha$  [13] may also contribute to cognitive decline.

In this study, a hypothetical association between volume of WML and CSF concentration of sAPP metabolites (sAPP- $\alpha$  and sAPP- $\beta$ ) in cognitively impaired patients was tested by comparing CSF levels of these metabolites in patients with SCI or MCI to those in post-stroke patients and to MRI-based quantitative measures of brain ischemia (WML) in both patient groups. Through measurement of APP metabolite levels in patients with a wide range of WML volumes, this study aimed to determine how SVD influences APP metabolism.

# Methods

Patients with SCI and MCI were recruited from a university-hospital based memory clinic between September 2005 and December 2007. Inclusion criteria were age 40-79, established SCI or MCI for at least 6 months, Global Deterioration Scale [36] score 2 or 3 (scores 4 and higher are per definition dementia) as determined from a clinical interview, Clinical Dementia Rating [37]  $\leq$  0.5 and results of screening tests performed at time of inclusion (mini-mental state examination MMSE [38]; Stepwise comparative status analysis parameters 13-20 [39]; fluency, interference and numeral-letter items from the Iflex [40]; and Cognistat [41]). Exclusion criteria were impaired activities of daily living, established psychiatric disorder, cancer, drug abuse, solvent exposure or anoxic brain damage. Thirty-seven patients fulfilled all criteria, and successfully underwent MRI and lumbar puncture.

Stroke patients were recruited from a university-hospital based stroke unit during 2007. Inclusion criteria for these patients were: cortical and lacunar supratentorial infarctions, classified by MRI, between 40 and 79 years of age, and cognitive complaints but MMSE score >23, no severe problems of language and visual/auditory neglect. Exclusion criteria were a history of medical or psychiatric disorder including depression. Twenty-six patients fulfilled all criteria, and successfully underwent MRI and lumbar puncture. Table 1 presents sex, age and MMSE scores for each patient group. All patients gave their written consent, and the regional ethics committee approved the study.

# MRI

MRI scans were from two sites (site 1: 10 patients, all SCI/ MCI; site 2: 53 patients). Site 1: Siemens Symphony 1.5 T (Siemens, Erlangen, Germany) with a conventional quadrature head coil. Two 3 D magnetization-prepared gradient echo (MP-RAGE), T1-weighted sequences in succession (TR/TE/TI/FA = 2730 ms/3.19 ms/1100 ms/15°, matrix =  $256 \times 192$ ), 128 sagittal slices, thickness = 1.33 mm, inplane resolution of  $1.0 \times 1.33$  mm. Site 2: Siemens Espree 1.5 T using two 3 D MP-RAGE, T1-weighted sequences in succession (TR/TE/TI/FA = 2400/3.65/1000/8°, matrix =  $240 \times 192$ ), 160 sagittal slices, thickness = 1.2 mm, inplane resolution of 1.0 mm × 1.2 mm.

For the SCI/MCI patients MRI was performed after inclusion, whereas it was performed 3 months after the stroke for stroke patients. Acute stroke localization was determined during the acute stage hospitalization. At

#### **Table 1 Patient characteristics**

Variable		Stroke	SCI/MCI
Sex	Men (total)	20 (26)	20 (37)
Age	Median	66.0	60.4
	Range	42-78	43-77
MMSE	Median	29.0	28.0
	Range	23-30	23-30

SCI: subjective cognitive impairment, MCI: mild cognitive impairment, MMSE: mini-mental state examination

this stage, diffusion-weighted MRI was also performed if infarct localization was not evident from cerebral computed tomography and clinical examination. WML, including white matter hyperintensities associated with current or previous infarction(s), were quantified with a semi-automated method in the Nordic ICE clinical image processing and analysis software application (NordicNeuroLab AS, Norway). In the T2-weighted fluid attenuated inversion recovery (FLAIR) images, pixel values in white matter higher than two standard deviations (SD) above mean pixel value were defined as WML (Fig 1). White matter hyperintensities associated with current or previous infarctions were classified as acute or chronic WML. The former was also considered likely to represent acute post-stroke changes, possibly stroke penumbra volume and subtracted from total WML volume in post-stroke patients to determine the volume of chronic WML used for further analysis.

Using the FLAIR-weighted images, hypointensities classified as current or previous infarctions were quantified by manually assigning a freehand region of interest to the area. The areas of each slice were added and multiplied with slice thickness to obtain total volumes. Images from one MCI patient were excluded due to motion artifacts.

# CSF

CSF samples were collected by lumbar puncture through the L3/L4 or L4/L5 intervertebral-space. The lumbar puncture was performed consecutively after inclusion in

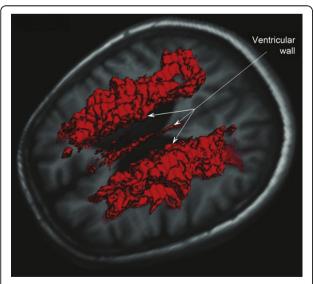


Figure 1 Visualization of chronic white matter lesions obtained by MRI. The hyper-intense areas in the FLAIR images (in red) are overlain onto the T1 images and shown in three dimensions. The ventricular walls (arrows), adjacent periventricular and subcortical hyper-intensities are visualized.

the SCI/MCI group or, in the stroke group (7-10 days post stroke). CSF was collected in a polypropylene tube and centrifuged at 2,000 × g at +4°C for 10 min. The supernatant was removed, gently mixed to avoid possible gradient effects, and stored within one hour at -80°C, pending biochemical analyses.

CSF concentrations of sAPP- $\alpha$  and sAPP- $\beta$ , and A $\beta$ X-38, A $\beta$ X-40 and A $\beta$ X-42 were determined using the MSD° sAPP- $\alpha$ /sAPP- $\beta$  Multiplex Assay and MSD° A $\beta$  Triplex Assay as described by the manufacturer (Meso Scale Discovery, Gaithersburg, MD, USA). Coefficients of variation were < 10% for all analyses. All analyses from the MCI and stroke groups were performed in the same batch.

#### Statistics

The statistics software package PASW 18 (SPSS Inc, USA) was used for statistical analysis. Linear regression was used to regress out age and sex, providing standardized residuals for further use. Due to skewed data, the Mann-Whitney test was used on these residuals to compare for group differences.

Pearson correlation coefficients between the CSF variables and the WML volumes were determined. To assess predictors for APP metabolites, scanner and sex were linearly regressed out from chronic WML and acute WML, again providing standardized residuals for further use. Consecutively, age, acute WML volume and chronic WML volume were entered as independent variables for all APP metabolite values. The different APP metabolites were sequentially entered as dependent variables in linear regression analysis.

#### Results

After regressing out age and sex from the biomarkers, the median CSF sAPP- $\alpha$  and sAPP- $\beta$  levels were significantly lower (p < 0.001 for sAPP- $\alpha$  and p < 0.005 for sAPP- $\beta$ ) in the post-stroke patients than in the SCI/MCI patients. There were no significant differences in CSF levels of A $\beta$  X-38, X-40 and X-42 between the two groups. The volume of chronic WML was higher in the stroke group than in the SCI/MCI group, but this difference was not significant (Table 2). The volume of acute post-stroke changes (infarct volume and surrounding WML halo) did not correlate with levels of CSF APP metabolites (data not shown). sAPP- $\alpha$  and sAPP- $\beta$  levels were linearly related in both the stroke (r = 0.942, p < 0.001) and the SCI/MCI (r = 0.955, p < 0.001) groups (Fig 2).

Most patients with high chronic WML volumes had low CSF sAPP- $\alpha$  (Fig 3), and a clear negative correlation between the two variables was seen (r = -0.36, p < 0.01). A similar relation was seen between chronic WML volumes and CSF sAPP- $\beta$  (r = -0.33, p < 0.05). In the regression analyses, chronic WML volume was a significant predictor for all examined CSF APP metabolites;

Table 2 CSF concentrations of APP metabolites in two patient groups

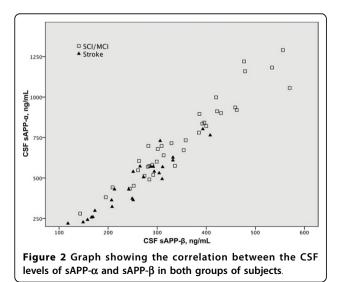
Variable	Stroke median, (SD)	SCI/MCI median, (SD)	p
CSF sAPP- $\alpha$ ng/mL	499.5 (168.9)	698.0 (248.4)	< 0.001
CSF sAPP- $\beta$ ng/mL	258.0 (74.2)	329.0 (101.4)	<0.005
CSF A <mark>β</mark> X-38 pg/L	1048.0 (576.9)	1141.0 (810.1)	n.s.
CSF Aβ X-40 pg/L	5989.0 (1777.6)	6323.0 (2209.3)	n.s.
CSF AβX-42 pg/L	485.5 (167.5)	483.0 (227.7)	n.s.
Chronic WML cubic mm	5051.8 (11569.0)	971.0 (6105.8)	n.s.

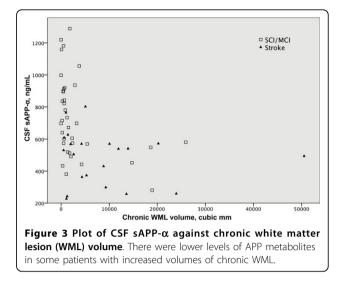
Group differences after age and sex have been corrected for by linear regression. CSF sAPP- $\alpha$  and sAPP- $\beta$  are significantly lower in the stroke group as compared to the SCI/MCI group. n.s.: not significant.

 $p \le 0.005$  for sAPP- $\alpha$ ,  $p \le 0.01$  for sAPP- $\beta$  and A $\beta$ X-38 levels, and  $p \le 0.05$  for A $\beta$ X-40 and A $\beta$ X-42 (Table 3). When repeating this analysis separately for the post-stroke and SCI/MCI groups, chronic WML still predicted sAPP- $\alpha$  and sAPP- $\beta$  in the SCI/MCI group ( $p \le 0.005$  for both sAPP- $\alpha$  and sAPP- $\beta$ ), but not in the post-stroke group. Age was not significantly related to sAPP, but significantly predicted higher levels of A $\beta$ X-38 ( $p \le 0.01$ ) and A $\beta$ X-40 ( $p \le 0.05$ ). Sex and scanner site were not significantly related to any of the metabolites (data not shown).

# Discussion

We have shown that CSF levels of sAPP metabolites are lower in post-stroke patients than in SCI/MCI patients. There was an inverse relation between chronic, but not acute WML volume and all examined CSF APP metabolites across the groups. This suggests that the reduction is associated with the severity of chronic, but not acute ischemic disease.





One limitation of our study is that it did not include a healthy control group. It may therefore be argued that an alternative interpretation of our results would be that there was an increase of APP metabolite levels in the SCI/MCI group instead of a decrease in the post-stroke group. At least two pieces of evidence argue against this interpretation. Firstly, we did not see any significant difference in CSF levels of sAPP between controls and MCI patients in an earlier study from our laboratory [16]. Unfortunately however, the results from this study, although employing the same sAPP assay, cannot be directly compared to those in the present investigation due to batch-to-batch variation in sAPP concentration between different kits. Secondly, the inverse correlation of CSF sAPP levels with WML volume seen in the present study, as well as in an earlier investigation [29], suggest that subcortical changes are associated with sAPP reductions. Similar results have been reported in

Table 3 Chronic and acute white matter lesion volumes as predictors for APP metabolites in all patients.

Dependent Variables	es Independent variables p (beta)			
	cWML	aWML	Age	
CSF sAPP-α	≤0.005	n.s.	n.s.	
	(-0.396)			
CSF sAPP <b>-β</b>	≤0.01	n.s.	n.s.	
	(-0.369)			
CSF AβX-38	≤0.01	n.s.	≤0.01	
	(-0.359)		(0.371)	
CSF AβX-40	≤0.05	n.s.	≤0.05	
	(-0.357)		(0.309)	
CSF AβX-42	≤0.05	n.s.	n.s.	
	(-0.273)			

Chronic WML significantly predicts CSF levels of APP metabolites. cWML = chronic WML, aWML = acute WML. other diseases that are characterized by white matter changes, including multiple sclerosis [42] and dementia in acquired immunodeficiency syndrome [43]. It should be noted that Lewczuk and colleagues recently reported higher CSF concentrations of sAPP in patients with cognitive impairment (CI) *and* other neurochemical CSF findings characteristic of AD, than in patients with CI *without* CSF findings characteristic of AD [44]. Importantly, however, this study did not include a healthy control group and it is not unlikely that a significant number of the cognitively-impaired individuals without AD-like CSF biomarker changes in fact suffered from chronic cerebrovascular disease.

Our finding that increased chronic WML volume predicts lower sAPP metabolites is however clear-cut, and is most easily interpreted in favor of reduced sAPP metabolites in the stroke group. When repeating this analysis separately in the post-stroke and SCI/MCI groups, the prediction was not, however, significant in the stroke group, but was still significant within the SCI/MCI group. As CVD both increases APP production and impedes transport, this suggests that the net negative effects on APP levels occur at mild to moderate WML levels. Also, the stroke group was somewhat smaller than the SCI/MCI group, and this might explain why the prediction was not significant within the stroke group alone. In accord with earlier findings [16], sAPP- $\alpha$  and sAPP- $\beta$  levels were very tightly correlated, indicating that the mechanism for the reduction lays upstream of  $\alpha$ - and  $\beta$ -secretase activity. Energy-dependent fast axonal transport of APP [45] may well suffer in chronic ischemic brain disease affecting white matter tracts, resulting in reduced axonal transport of the precursor protein [34], reduced substrates for the secretases resulting in the observed reduction in metabolite levels in the CSF. Alternatively, ischemia may have an effect on earlier stages, e.g. reduce APP gene expression, but observations in experimental ischemia suggest that this is less likely [30,31].

Chronic WML volume predicts levels of both sAPP metabolites and A $\beta$  peptides, but there is a significant difference between the post-stroke and the SCI/MCI groups for sAPP, suggesting that for the amyloid  $\beta$  peptides there is more extensive interplay with other factors. In a previous study [46], acute stroke was not shown to have a significant short term effect on levels of A $\beta$ X-42. However, as human A $\beta$  clearance rates are close to 10% per hour [47]; changes temporally related to acute stroke onset are not expected to be detected in the present study.

#### Conclusions

In the patient group as a whole, there was a strong correlation between CSF sAPP- $\alpha$  and sAPP- $\beta$  concentrations.

An inverse relationship was demonstrated between the volume of chronic WML and CSF APP metabolites (sAPP- $\alpha$ , sAPP- $\beta$ , A $\beta$  X-38, X-40 and X-42) in both stroke patients and SCI/MCI patients. In addition, there were lower levels of CSF sAPP- $\alpha$  and sAPP- $\beta$  in the stroke group when compared to the SCI/MCI-group. This suggests that ischemia influences APP metabolism probably through inhibition of fast axonal transport of APP. If confirmed, the present results implicate new mechanisms for reduction of CSF APP metabolites, including CSF A $\beta$ X-42, which is a predictor for development of AD.

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#### Authors' contributions

PS determined the WML volumes and performed the statistical analyses, participated in the collection of data and in drafting of the manuscript. KB participated in the immunological analyses and the conception of the study. HZ participated in the immunological analyses and the conception of the study. RG participated in the collection of data. LR participated in the immunological analyses and the study. LJ participated in the collection of data. TF participated in the conception of the study, its design and coordination and helped to draft the manuscript. All authors read and approved the final manuscript.

#### **Competing interests**

The authors declare that they have no competing interests.

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