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CSF Neurofilament Proteins Levels are Elevated in Sporadic Creutzfeldt-Jakob Disease

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Abstract. In this study we investigated the cerebrospinal fluid (CSF) levels of neurofilament light (NFL) and heavy chain (NFHp35), total tau (t-tau), and glial fibrillary acidic protein (GFAP) to detect disease specific profiles in sporadic Creutzfeldt Jakob disease (sCJD) patients and Alzheimer's disease (AD) patients. CSF levels of NFL, NFHp35, t-tau, and GFAP of 23 sCJD patients and 55 AD patients were analyzed and compared to non-demented controls. Median NFL, NFHp35, GFAP, and t-tau levels were significantly increased in sCJD patients and AD patients versus controls ($p < 0.0001$ in all). NFL, NFHp35, and t-tau levels were significantly increased in sCJD patients versus AD patients ($p < 0.005$), but GFAP concentrations did not differ between sCJD and AD. The results suggest that neuroaxonal damage, reflected by higher CSF levels of NFL, NFHp35, and t-tau, is more pronounced in the pathophysiology of sCJD than in AD. The comparable CSF GFAP concentrations suggest that astroglial damage or astrogliosis is equally pronounced in the pathophysiology of AD and sCJD. Prospective studies are needed to determine whether NFL and NFHp35 may be additional tools in the differential diagnosis of rapidly progressive dementias.

Keywords: Alzheimer's disease, cerebrospinal fluid, Creutzfeldt-Jakob disease, diagnosis, prion disease

INTRODUCTION

Alzheimer's disease (AD) is a devastating neurodegenerative disorder characterized by widespread loss of cortical neurons and their connections resulting in memory loss and cognitive decline. Sporadic

Creutzfeldt-Jakob disease (sCJD) is a rare, fatal neurodegenerative disorder, belonging to the spongiform encephalopathies which is clinically characterized by a rapidly progressive dementia with visual or cerebellar signs, (extra)pyramidal symptoms, and ultimately akinetic mutism.

In AD, the typical cerebrospinal fluid (CSF) pattern includes elevated total tau (t-tau) and hyperphosphorylated tau (p-tau) proteins, whereas the amyloid- β_{42} protein ($A\beta_{42}$) concentration is decreased [1]. In sCJD, detection of 14-3-3 protein is highly sensitive and is part of the WHO diagnostic criteria for sCJD [2], and

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strongly elevated levels of t-tau protein are highly specific. Other abnormal proteins in CSF of sCJD patients include elevated S100B and neuron-specific enolase (NSE) and decreased $A\beta_{42}$ [3,4]. However, CSF t-tau levels are usually much higher in sCJD than in AD and $A\beta_{42}$ levels are usually much lower in AD than in sCJD.

Neurofilament (NF) proteins are neuronal cytoskeleton proteins. They play an important role in neuronal structure and are mainly located in large myelinated neurons. NF-proteins are highly phosphorylated and are composed of 3 subunits defined by their molecular weight: 68 kDa (NF-light chain, NFL), 160 kDa (NF-medium chain, NFM), and 200 kDa (NF-heavy chain, NFH). Elevated CSF levels of NF-proteins have been found in vascular dementia, AD, and frontotemporal dementia (FTD) and it was demonstrated that CSF NFL analysis may help to distinguish FTD from AD [5–7]. NF-protein analysis in sCJD has not been reported so far, although neuropathological reports on neurofilament inclusions in sCJD suggest neurofilament levels could be increased in sCJD [8]. In addition, since the concentration of the glial protein S100B is significantly increased in CSF of sCJD patients and because of the reactive astrocytosis in sCJD, it can be speculated that concentrations of the astrocyte marker glial fibrillary acidic protein (GFAP) will also be increased in CSF of sCJD patients.

The aim of the present study was to identify whether markers of neuronal damage (NFL, NFHp35, t-tau) and glial damage (GFAP) are abnormal in CSF of sCJD and AD patients and to determine a disease specific profile.

MATERIALS AND METHODS

Patients

This retrospective study included 23 sCJD patients and 55 AD patients identified through the CSF databases of the Radboud University Nijmegen Medical Centre and the Born Bunge Institute (University of Antwerp) between January 1998 and January 2005. Inclusion into the study was dependent upon the availability of CSF for additional analyses and of the availability of sufficient clinical information to make a proper diagnosis. Only patients with a probable diagnosis according to standard diagnostic criteria were included. The sCJD patients fulfilled the WHO criteria for probable sCJD and a diagnosis of probable AD was made according to the revised NINCDS-ADRDA criteria [2,9]. The

diagnostic examination included physical, neurological and mental status examination, screening laboratory tests, routine CSF analysis, and MRI or CT of the brain. The sCJD patients also underwent EEG. Twenty probable sCJD patients had converted to definite sCJD by the time of data analysis. In the remaining 3 probable sCJD patients, postmortem neuropathological examination had not been allowed by the family. Two of them were tested for 14-3-3 and were positive; the other one had not been analyzed for 14-3-3, but had a typical EEG (with periodic sharp wave complexes) and died after a total disease duration of 14 months. None of the sCJD patients survived more than 2 years after having been diagnosed with sCJD. The sCJD patients from the Belgian center ($n = 15$) were also included in a previous study [10]; the other sCJD patients were not published previously.

The study contained some missing data, since the archived CSF was not sufficient for all analyses. The sCJD patients ($n = 23$) had all been tested for NFHp35 and t-tau, whereas 21 patients had been tested for NFL or 14-3-3 and 18 for GFAP. All AD patients ($n = 55$) had been tested for NFL and t-tau, whereas 52 had been tested for NFHp35, 51 for GFAP, and 40 for 14-3-3. Disease duration at the time of lumbar puncture was also recorded and available in all patients.

For determination of CSF biomarker concentration in a control population, non-demented controls were included with ages between 50–85 years who were referred to the Department of Neurology of the RUNMC and underwent a lumbar puncture in a diagnostic work-up, but who, after extensive examination, turned out not to have a neurological disease, and who had normal CSF cell count, hemoglobin, bilirubin, total protein, lactate, glucose and no oligoclonal IgG bands. For NFHp35, 21 controls (mean age 57 yr, SD 7.9) were used. For NFL, 23 controls (mean age 57.6 yr, SD 7.9) were used, for t-tau 80 controls (mean age 60.2 yr, SD 8.8), and for GFAP 31 controls (mean age 58.7 yr, SD 8.4) were used.

CSF analysis

CSF was obtained by lumbar puncture, collected in polypropylene tubes, and transferred to the hospital laboratory within 30 min. Lumbar punctures were performed after informed consent was obtained from the patient, or from his or her legal representative. After routine investigations, CSF was centrifuged (860 g, 5 min), aliquoted and stored at -80°C until analysis. The concentrations of t-tau and GFAP were analyzed

Table 1
Patient characteristics and concentrations of CSF biomarkers

| | sCJD | AD | CONT | p-value |
|--------------------|---------------------|--------------------|---------------------------------|--|
| Number of patients | 23 | 55 | 130 | |
| Gender (m/f) | 10/13 | 23/32 | | N.S. ¹ |
| *Age (yrs) | 68.4 (8.2) | 68.7 (8.8) | 59.2 (8.4) | N.S. ¹ ; < 0.0001 ^{2,3} |
| #Duration (months) | 5 (2–9.5) | 24 (18–40) | N.A. | < 0.0001 ¹ |
| *MMSE | N.A. | 19 (5.5) | N.A. | N.A. |
| #NFHp35 (ng/L) | 579 (420–796) | 117 (77–165) | 75 (61–88) (<i>n</i> = 21) | < 0.0001 ^{1,2} ; 0.0007 ³ |
| #NFL (ng/L) | 78.0 (32.6–123.3) | 7.2 (6.0–18.1) | 5.20 (0–19.7) (<i>n</i> = 23) | < 0.0001 ^{1,2} ; 0.044 ³ |
| #GFAP (μg/L) | 2.4 (1.3–6.1) | 2.9 (2.3–3.7) | 1.39 (0.6–2.2) (<i>n</i> = 31) | 0.054 ¹ ; 0.0037 ² ; < 0.0001 ³ |
| #Tau (ng/l) | 6438 (2400–15230) | 593 (388–795) | 179 (136–253) (<i>n</i> = 80) | < 0.0001 ^{1,2,3} |
| §14-3-3 | 20 (<i>n</i> = 21) | 5 (<i>n</i> = 40) | N.A. | < 0.0001 ¹ |

Abbreviations: sCJD, sporadic Creutzfeldt-Jakob disease; AD, Alzheimer's disease; CONT, non-demented controls; NFL, neurofilament light chain; NFHp35, phosphorylated neurofilament heavy chain; GFAP, glial fibrillary acidic protein; Tau, tau-protein.

#Median (25th–75th percentile) are shown. *Mean (standard deviation) are shown. §Number of positive tests is shown (total number of tests). N.S., not significant; N.A., not applicable, *n*: number. ¹sCJD compared to AD, ²sCJD compared to CONT, ³AD compared to CONT. p-values for NFHp35 and NFL; Mann Whitney-tests; p-values for GFAP and t-tau: t-tests.

routinely every two weeks, i.e., at a maximum storage time at -80°C of two weeks after sample collection. GFAP concentrations were analyzed by a homemade sandwich ELISA as described earlier [11]. The inter-assay CV% is 8.6% (*n* = 16; at a concentration of $4.5\ \mu\text{g/L}$). T-tau concentrations in CSF were analyzed by using the Innostest hTau assay (Innogenetics, Ghent, Belgium). The inter-assay CV% is 6.9% (*n* = 20; at a concentration of $263\ \text{ng/L}$). Previously described sandwich ELISAs were used to determine NFL and NFHp35 levels [12,13]. Both NFL and NFHp35 were retrospectively analyzed in the patient groups after the selection of the patient groups was completed. The inter-assay CV for NFL was 3.9% at a concentration of $124\ \text{ng/l}$ (*n* = 8), 9.1% at a concentration of $60\ \text{ng/l}$ (*n* = 14), and 17.8% at a concentration of $26\ \text{ng/l}$ (*n* = 15). For NFHp35, the interassay CV was 18% at a concentration of $35\ \text{ng/L}$ (*n* = 18). The presence of the 14-3-3 protein in CSF was routinely determined every two weeks by electrophoresis and immunoblotting. In brief, CSF proteins were separated by SDS-PAGE electrophoresis. Proteins were transferred to nitrocellulose membranes, after which the membranes were successively incubated overnight with Odyssey blocking buffer (LI-COR Biosciences, Lincoln, NE, USA), anti-14-3-3 β antibody (Santa Cruz Biotech, 2 h) and Alexa Fluor680-labeled goat anti-rabbit antibodies (1 h). Incubation steps were each followed by extensive washing in PBS. Blots were scanned by using Odyssey infrared imaging system (LI-COR Biosciences), according to manufacturer's instructions. Total protein concentrations were measured by using the Lowry method.

Statistical analysis

Kolmogorov-Smirnov normality tests showed non-Gaussian distribution of CSF NFHp35 in the AD and control groups and of NFL in the AD group. Therefore, data of NFHp35, NFL, t-tau, and GFAP are all presented as median and p25–p75 range, and non-parametric Mann-Whitney tests were used to analyze non-Gaussian distributed CSF data, whereas t-tests were used to analyze Gaussian distributed CSF data. Adjustments for age, gender, disease duration and total protein concentrations were performed by using linear regression. Correlations between CSF and clinical parameters were analyzed using the Spearman test for non-Gaussian distributed variables and the Pearson test for Gaussian distributed variables; in case of CSF proteins a partial correlation analysis (by group) was used. Graphpad Prism (San Diego, CA) and SPSS16.0 software were used for statistical analysis.

RESULTS

Baseline characteristics and biochemical analyses are summarized in Table 1. All sCJD patients clinically suffered from a (severe) dementia syndrome. Abnormal triphasic waves on EEG analysis were observed in 14 patients, absent in 8 patients and could not be unequivocally determined in one patient. Myoclonic jerks were observed in 15 patients. Seven sCJD patients had pyramidal symptoms (spasticity and extensor plantar responses), 9 patients had extrapyramidal symptoms (hypo/bradykinesia, chorea or rigidity), 16 patients had a cerebellar syndrome (ataxia) and in only

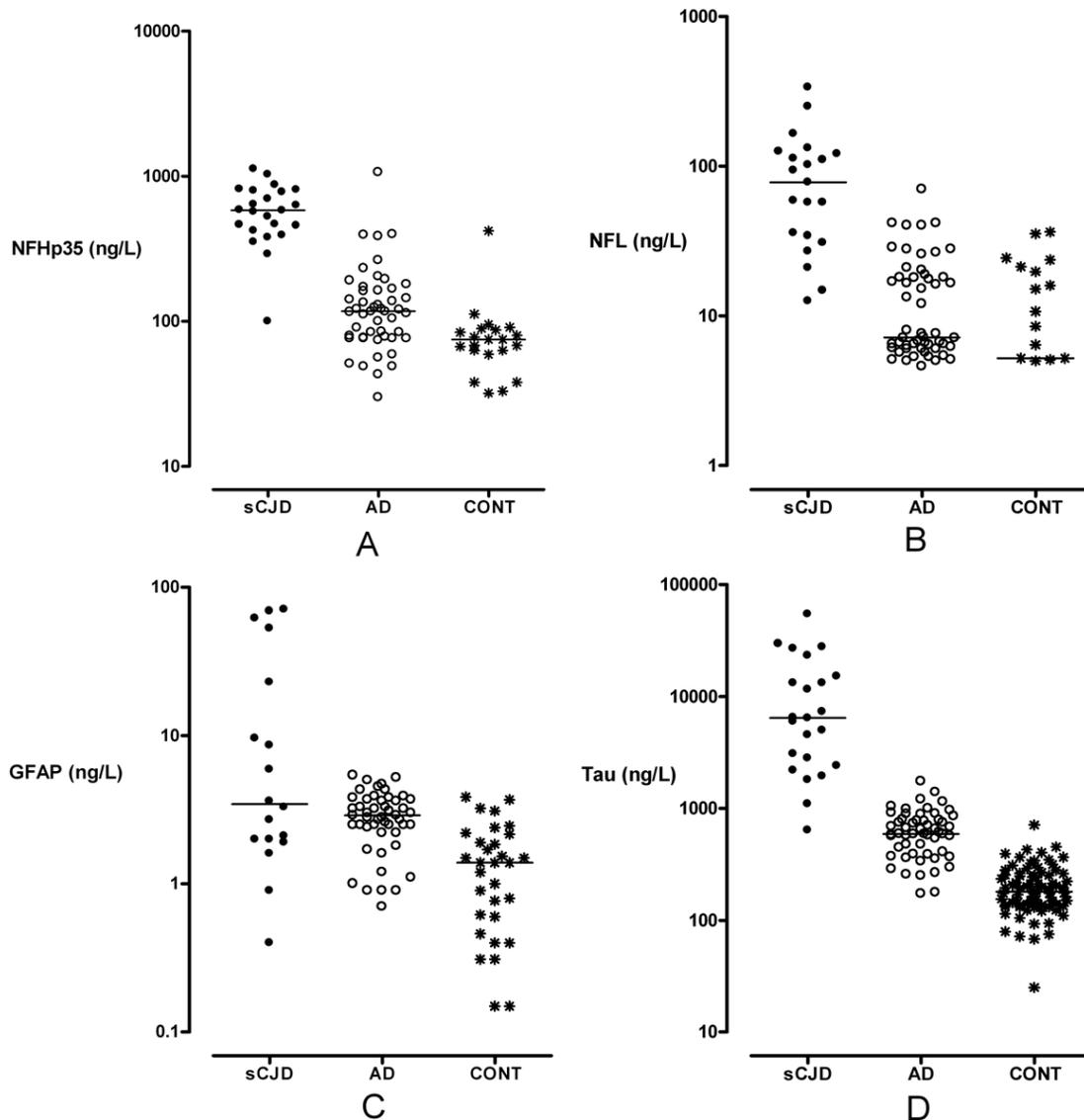


Fig. 1. Concentrations of NFHp35 (A), NFL (B), GFAP (C), and tau (D) per patient group. Horizontal lines indicate median levels. Abbreviations: sCJD, sporadic Creutzfeldt-Jakob disease; AD, Alzheimer's disease; CONT, controls.

7 sCJD patients typical abnormalities on MRI, e.g., bilateral hyper intense signal on T2 weighted images predominantly affecting the caudate nucleus and putamen, were observed.

Levels of CSF NFL, NFHp35, and t-tau were significantly increased in sCJD versus controls and sCJD versus AD patients (Table 1; Fig. 1a, b, d), whereas total protein concentrations were similar in AD (mean 513 mg/L) and sCJD (mean 495 mg/L; $p = 0.75$). Similar results were obtained after adjustment of the data for age, gender, disease duration and total protein concentrations. Also, similar p-values were obtained

if we log-transformed the results for NFL, NFHp35, t-tau, and GFAP (to normalize the data distribution for NFL and NFHp35), either with or without adjustment for age, gender, and disease duration. GFAP levels were higher in either sCJD or AD versus controls (Table 1; Fig. 1c), but did not differ significantly between sCJD and AD patients (Table 1; Fig. 1c). The differences in GFAP results between AD and controls, but not between sCJD and controls, remained significant after adjustment for age, gender, and disease duration. Furthermore, the differences between sCJD and AD remained significant in those subgroups (sCJD, $n = 17$;

AD, $n = 44$) for whom all CSF data of NFL, NFHp35, t-tau, and GFAP were available. The test for the presence of the 14-3-3 protein in CSF by immunoblotting was positive in 20 cases with sCJD and 5 cases with AD ($p < 0.0001$).

No significant correlation between CSF markers and either age or disease duration was found in sCJD patients. In AD patients, we found a weak correlation between NFL concentration and age ($r = 0.297$; $p = 0.022$), but not with disease duration. No other correlations were found between the various CSF markers and clinical parameters in either sCJD or AD patients. Correlations between the CSF proteins are shown in Table 3; significant correlations were observed between NFH, NFL, and GFAP, between GFAP and t-tau, but not between t-tau and either NFL or NFH.

The results of the ROC analysis to discriminate AD from sCJD are shown in Table 2. NFL, NFHp35 and tau analysis all resulted in a high area under the ROC curve and high likelihood ratio. Binary (multivariate) logistic regression analysis selected NFL, NFHp35, tau and duration as independent contributors for differentiation between sCJD and AD according to the model: $Y = 33.68 - 0.465 * NFL - 0.15 * NFHp35 - 0.026 * tau + 1.98 * duration$ (months). The ROC area under curve of this equation was 0.998 and therefore almost perfectly discriminated sCJD from AD patients.

DISCUSSION

In this study, we found significantly elevated concentrations in CSF of NFHp35, NFL, t-tau, and GFAP in sCJD and AD compared to controls. Concentrations of CSF NFL, NFHp35, and t-tau were also significantly elevated in sCJD compared to AD patients, but GFAP levels did not differ between these two patient groups.

Both human and animal studies have shown that NF-proteins play an important role in the axonal environment. These highly phosphorylated proteins are essential in maintaining neuronal cytoskeletal plasticity by influencing axonal caliber and axonal transport [14]. Neuropathological examination in sCJD has shown numerous abnormal neurites labeled by NF-protein immunocytochemistry [8]. In previous studies it has been shown that elevated NFL levels are associated with axonal and white matter damage in neurological diseases such as multiple sclerosis, FTD, and multiple system atrophy [7,13,15]. NF-proteins may be regarded as biomarkers for degeneration of large, myelinated axons. Our data suggest that these axons are either more

subject to degeneration in sCJD than AD or release NF-proteins during the course of this disease, which may lead to the highly elevated CSF concentrations. A close observation of the NFL data in the AD group revealed a potential bimodal distribution of the data; we did not, however, find a correlation with sampling date or storage time which may explain this difference.

T-tau concentration was also strongly elevated in sCJD relative to AD. T-tau is a microtubule associated protein mainly localized in neuronal axons where it has a function in maintaining the normal cytoskeleton, by stabilizing microtubules that are essential for axonal transport. Thus, the high concentration of t-tau in CSF of sCJD patients also suggests a more compromised axonal function in this disorder. Our data are in line with previous studies, in which strongly elevated t-tau concentrations were reported in CSF of sCJD patients [4, 10,16]. The concentrations of both NFL and t-tau were increased by a factor 10 in sCJD compared to AD, suggesting, at first sight, that similar pathophysiological mechanisms may lead to release of these axonal proteins into the CSF of sCJD patients. However, we did not find a correlation between NFL and t-tau levels, suggesting that independent mechanisms contribute to the release of multiple axonal proteins in the extracellular space and, eventually, CSF. Furthermore, in previous studies we found that NFL and t-tau independently contributed to the differentiation of MSA from idiopathic Parkinson's disease [13]. Thus, although both NF and t-tau proteins are both located in axons, these observations strongly suggest that different pathophysiological mechanisms lead to the release of these proteins into the extracellular space. This hypothesis is strengthened by the observation that the concentrations of NFHp35 in sCJD were elevated by a factor 5 compared to AD and also did not correlate to t-tau levels. The respective mechanisms that lead to these differences remain to be elucidated.

We did not find a significant difference in CSF GFAP concentration between sCJD and AD and the GFAP concentration in either group was only marginally, albeit significantly, increased compared to controls. GFAP is a cytoskeletal protein of astrocytes which is important for astrocyte motility and shape. GFAP expression is increased during reactive astrocytosis, a process that is associated with senile plaque formation in AD brains and that is a common feature of sCJD as well [17,18]. In a previous study, it was described that CSF GFAP concentrations were higher in AD than in sCJD [19]. Our data do not support this observation, but suggests that the degree of astrocytosis is compa-

Table 2
ROC analysis of the CSF biomarkers NFL, NFHp35, t-tau, and 14-3-3

| CSF parameter | Cut-off value | Sens % | Spec % | PPV % | NPV % | Likelihood ratio | AUC (95% CI) |
|---------------|---------------|--------|--------|-------|-------|------------------|------------------|
| NFL (ng/l) | > 29.8 | 81 | 91 | 77 | 93 | 4.8 | 0.93 (0.87–0.99) |
| NFHp35 (ng/l) | > 276.5 | 90 | 96 | 82 | 98 | 20.8 | 0.95 (0.88–1.0) |
| t-tau (ng/l) | > 1072 | 93 | 96 | 85 | 98 | 21.7 | 0.98 (0.94–1.0) |
| 14-3-3 | N.A. | 95 | 88 | 80 | 97 | 7.9 | 0.91 (0.83–0.99) |

Abbreviations: NFL, neurofilament light chain; NFHp35, phosphorylated neurofilament heavy chain; t-tau, total tau protein; Sens, sensitivity; Spec, Specificity; PPV, positive predictive value; NPV, Negative predictive value; AUC, area under curve. N.A., not applicable.

Table 3
Partial correlation analysis of the CSF proteins NFH, NFL, GFAP, and t-tau

| | NFH | NFL | t-tau |
|-------|-----------------------------|-----------------------------|-----------------------------|
| NFL | $R = 0.473$ $p < 0.0001$ | | |
| t-tau | $R = 0.189$ $p = 0.147$ | $R = 0.077$ $p = 0.558$ | |
| GFAP | $R = 0.350$ $p = 0.006$ | $R = 0.449$ $p < 0.0001$ | $R = 0.684$ $p < 0.0001$ |

R = partial regression coefficient.

rable in sCJD and AD. Interestingly, our correlation analysis demonstrated a correlation between t-tau and GFAP, suggesting that reactive astrogliosis is linked to axonal degeneration. However, given the magnitude of increase in CSF biomarkers, our observations showed that in particular neuro-axonal degeneration contributes to the pathophysiological mechanism of sCJD, rather than astrocyte degeneration. GFAP expression is limited to astrocytes, whereas the protein S100B is a more general marker for the glial cell population in the brain, also including oligodendrocytes and Schwann cells.

It has been demonstrated previously that the CSF concentrations of S100B are increased in a majority of the sCJD patients, but is less pronounced in AD [3]. Therefore, it seems likely that degeneration of oligodendrocytes or Schwann cells, rather than that of astrocytes, contributes to the elevated S100B concentration in sCJD CSF.

The typical findings in CSF of sCJD patients can be summarized by elevated concentrations of the proteins 14-3-3, t-tau, NSE and S100B. In particular, the diagnostic accuracy of the testing for the presence in CSF of the 14-3-3 protein, as part of the WHO diagnostic criteria, has been extensively studied. Reported sensitivities to detect sCJD on the basis of 14-3-3 analysis vary between 83% and 100% with a specificity of 67%–100% [16,20–28]. Our data fit in with these numbers. A recent autopsy-based study on 280 consecutive cases clinically suspected of sCJD in the Netherlands, however, demonstrated a sensitivity of only 74% and specificity of 79%, demonstrating the limitations of this biochemical assay [29].

Sensitivity for t-tau varied between 86% and 98% with a specificity of 75–97% [16,20–22,25]. Sensitivity to detect sCJD is generally lower for NSE (79%–85%), with specificities ranging from 85%–92% [22, 23,28]. Finally, sensitivity to detect sCJD by using S100B quantification in CSF varied between 84% and 98% and specificity widely ranges from 29% to 90% [3, 23,25]. In the light of these findings the combined sensitivity and specificity we report here for t-tau, NFL and NFHp35 are very well comparable with other biomarkers. In the case of t-tau NFHp35 these numbers are at the high end of reported ranges as are the likelihood ratios and area under the curves.

It should be noted, however, that the timing of the lumbar puncture relative to the onset of clinical symptoms may affect the diagnostic accuracy of several biomarkers. It has been reported that the sensitivity for 14-3-3, t-tau, and S100B is higher in patients with a lumbar puncture taken within 6 weeks after the clinical symptoms compared to a delayed lumbar puncture [25]. Contrasting reports have also been published, however [30]. Finally, the M/V polymorphism at position in the PRNP gene may also slightly affect the diagnostic accuracy of the respective biomarkers [25, 30]. One of the limitations of our study is its retrospective, unblinded nature which might have confounded the results by indication. The missing values might have introduced skewing of data as well; however, given the major differences we observed in the concentrations of CSF NFL, NFHp35, and t-tau, this effect will be of minor importance. In conclusion, we found a disease specific CSF profile of sCJD in which the very high concentrations of NFL, NFHp35, and t-tau reflect a more important role of neuroaxonal damage in sCJD compared to AD. In contrast, astroglial damage seems to play an equally important role in either sCJD or AD. Moreover, this study is the first to evaluate analysis of NF proteins in sCJD-patients. It demonstrated that levels of NFL and NFHp35 are significantly and strongly increased in sCJD versus AD and controls. The ability to measure the neurofilament levels quantitatively with

an ELISA makes this analysis potentially interesting for diagnostic purposes and has advantages over the qualitative assessment of 14-3-3 protein in CSF, which is performed by immunoblotting. Assessment of the potential of NF protein analysis as an additional diagnostic value in rapidly progressive dementias awaits future prospective studies.

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