RESEARCH ARTICLE

Ratio and index of Neurofilament light chain indicate its origin in Guillain-Barré Syndrome

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Abstract

Objective: Neurofilament light chain (NfL) has been established as a biomarker of axonal damage in many diseases of the central nervous system (CNS). Increased levels of serum NfL (sNfL) can derive as well from damage in the peripheral nervous system (PNS) as from CNS, but little is known about the quantities contributing to sNfL. Peripheral nerve damage may be reflected by an increase in sNfL levels, while the NfL CSF/serum ratio and NfL index decreases. Methods: We collected serum and cerebrospinal fluid (CSF) from 21 Guillain-Barré Syndrome (GBS) patients and measured NfL in serum and CSF and compared them with 19 neurologically healthy controls. Results: In general, NfL in CSF and serum was significantly higher in GBS patients. Serum NfL was higher in GBS patients admitted to the intensive care unit (P = 0.02). Controls had a mean CSF/serum NfL ratio of 26.7 (ranging from 5.8 to 69.5) indicating a central origin of NfL. Three GBS patients had a similar range (23.9 to 42.7, mean 33.3) all of them with demyelinating pathology in the PNS. Eighteen GBS patients with axonal or mixed axonal-demyelinating pathology showed significantly lower CSF/serum ratios (0.02–12.2, mean 4.4), indicative of a peripheral origin of NfL. When applying the NfL index subdivisions remain the same. Interpretation: These results demonstrate that the PNS is a relevant contributor to sNfL levels and that the distribution can be identified by a lowered NfL CSF/serum ratio of NfL index. Furthermore, acute or subacute polyneuropathies are likely confounding factors in interpreting sNfL levels in CNS diseases.

Introduction

Neurofilaments are cytoskeletal proteins of exclusively neuronal origin. Neurofilament light chain (NfL) levels in CSF have been established as biomarker of neuronal damage in numerous neurological diseases of the CNS.¹–³

With the advent of the single molecule assay (SIMOA) technique NfL could also be quantitated reliably at low levels in serum, confirming its capacity to reflect disease activity in the CNS.⁴,⁵ Implicitly, NfL in blood was considered of being of CNS origin when measured in terms of CNS disorders not taking into account the possibility of being derived from the PNS also. However, axons of the PNS hence release NfL upon damage.⁶ Accordingly, elevated sNfL levels have already been shown in various vasculitic and hereditary neuropathies.⁶,⁷ GBS separates

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from other neuropathies as it affects intrathecal and extrathecal parts of the peripheral nervous system. Accordingly, increased levels of NfL were found both in CSF,8,9 and serum or plasma.10 However, the question to which extent intrathecal/central and extrathecal/peripheral nerve pathology contribute to the neurofilament levels in serum has not been addressed although of upmost importance for correct interpretation of sNfL levels.

We hypothesized that increased sNfL in GBS results to a relevant degree from extrathecal axonal damage, which may be reflected by a decrease in the NfL CSF/serum ratio and the NfL index. Furthermore, these two parameters render possible to calculate the portions of NfL in every disease and in a consequence to assess the origin of the NfL levels.

Methods

Cohorts

We performed a retrospective analysis of CSF and serum samples from 21 patients suffering from GBS and 19 non-neurological disease controls who underwent lumbar puncture (LP) to exclude diseases such as cerebral hemorrhage or neuroborreliosis and without signs of polyneuropathy or neurodegeneration having normal CSF parameters. At every lumbar puncture CSF and serum were drawn in one session. We included 21 GBS patients who had been diagnosed and treated at the Department of Neurology, University Hospital Magdeburg, Germany. Six patients had clinical follow-up data with longitudinally sampled CSF and serum available. GBS was diagnosed according to the criteria presenting with at least bilateral or flaccid weakness of limbs accompanied by decreased or absent tendon reflexes, CSF cell count under 50 cells/µl, monophasic course and time between onset-nadir of 12 h to 28 days.11

This study included GBS patients with Brighton collaboration criteria 2 with some GBS patients fulfilling Brighton Criteria 1.12 Assessment of disease severity was performed with the GBS Disability Scale (GBSDS).13

Laboratory procedures

Patients were tested for presence of ganglioside antibodies using a standard immunoblot (EUROLINE, DL-1130-1601-2 G/M comprising Anti-Hu, Anti-Ri, Anti-Yo, Anti-Zic4, Anti-DNER, Anti-Recoverin, Anti-GAD65, Anti-SOX1, Anti-Titin, Anti-Ma2, Anti-Amphiphysin, EURO-IMMUN, Lübeck, Germany), had a CSF analysis. Every patient received a standard electrophysiological procedure including measuring the motor and sensory nerve conduction velocities and f-wave latency measurements in at least N. tibialis and N. peroneaus in one leg. Additionally, most patients received the same program on N. medianus and N. ulnaris at least at one arm. Furthermore, a minimum of one electromyographic measurement from the M. Tibialis anterior was added in each patient. Axonal motor GBS subtype required next to pathological nerve conduction velocities present f-waves without pathological latency.11 Electrophysiological classification (axonal, demyelinating, or mixed type neuropathy) was performed in all GBS patients. We used the electrophysiological GBS criteria according to Hadden14 before 2015 and then changed to the electrophysiological criteria according to Rajabally.15

This diagnostical workup enabled us to categorize according to the Brighton Criteria.

NfL measurements

sNfL measurements were performed with SIMOA at the University Hospital Basel, Switzerland, by an in-house assay as described.4 NfL in CSF was measured with the ELISA using the similar antibody pair (Umandiagnostics®, Umeå, Sweden) at the University Hospital Magdeburg.

Statistics

Statistical analyses were performed with GraphPad Prism. We used a one-way ANOVA with Bonferroni correction, for correlation analysis Spearman rho, and considered P < 0.05 as significant. We calculated an NfL index using CSF-NfL/sNfL divided by the Albumin ratio.

The study was approved by the ethics committee of the University Hospital Magdeburg, Germany (approval number 22/19). Every patient gave written informed consent.

Results

Clinical presentation, standard CSF evaluation, and electrophysiological evaluation of GBS patients

Table 1 shows that in the mean all patients demonstrated an increased albumin ratio; 24% (5/21) had serum anti-ganglioside antibodies. Guillain-Barré Disability Score (GBSDS) ranged from 2 to 5 (Fig. 1a).

Based on electrophysiological work-up, 67% (14/21) of patients showed a mixed neuropathy pattern; 9% (2/21) were classified as purely axonal and 24% (5/21) as purely demyelinating neuropathies. The electrophysiological workup was applied according to standardized methods. All patients were treated with either intravenous immunoglobulins or with plasma exchange at first. Six
patients had to be admitted to the Intensive Care Unit (ICU).

**NfL levels in CSF and serum: biomarker for GBS and clinical course**

Serum NfL levels were significantly higher in every GBS patient (mean = 2603, 25.1 to 14935 pg/mL) than in controls (mean = 50.7, 11.1 to 84.9 pg/mL, \( P < 0.01 \), Table 1). Accordingly, CSF-NfL levels (mean = 2871.6, 397.4 to 50.7 pg/mL) were ranging more than 5-fold higher versus CSF-Controls (mean = 15854, 1115 pg/mL, \( P = 0.02 \), Table 1). Serum-NfL, but not CSF-NfL was significantly higher in patients who were admitted to ICU compared to patients who remained on the ward (\( P = 0.02 \) vs. \( P = 0.16 \), Fig. 1B).

**NfL levels in CSF and serum: categorisation into high vs low CSF/serum ratio groups**

The NfL CSF/serum ratio in controls with NfL originating predominantly from CNS structures ranged from 5.9 to 69.8 (mean = 26.7) The margin to its lowest quartile (12.8) was used as a cut-off value indicating the intra- or extrathecal origin of sNfL, respectively. This allowed us to categorize patients into a NfL-ratio\textsubscript{high} \((n = 3, \text{mean} = 33.23, \text{range} 24.00-42.65)\) versus a NfL-ratio\textsubscript{low} \((n = 18, \text{mean} = 3.95, 0.03-10.52, \text{Fig. 2A})\).

Serum-NfL and CSF/serum NfL ratio were strongly correlated (\( P < 0.001 \)), but not CSF-NfL with the CSF/serum NfL ratio (\( P = 0.44 \)). The GBSDS correlated well with CSF-NfL (\( P = 0.005 \)), sNfL (\( P = 0.02 \)), and the Qalb (\( P = 0.04 \)), but not with the NfL CSF/serum ratio (\( P = 0.41 \), Table 1).

**Clinical data show particular differences between both cohorts**

Nerve conduction studies showed either axonal or mixed patterns in the NfL-ratio\textsubscript{low}, while in the NfL-ratio\textsubscript{high} cohorts only demyelinating patterns were found (Fig. 2B). GBSDS was slightly higher in the NfL-ratio\textsubscript{low} group (mean = 3.6, range 2–5) than in the NfL-ratio\textsubscript{high} group (mean = 3.3, range from 2 to 4). The mean albumin ratio scored slightly higher in the NfL-ratio\textsubscript{low} (17.11 pg/mL) than in NfL-ratio\textsubscript{high} (13.77 pg/mL) group, but this difference did not reach levels of significance (\( P = 0.74 \)).

A major difference was time from disease onset to lumbar puncture with NfL-ratio\textsubscript{high} at 3.7 days and in the NfL-ratio\textsubscript{low} at 17.1 days.

**Ratios in follow-up patients**

Six GBS patients had clinical follow-up data. The mean time interval between lumbar punctures was 8.8 days for five patients and one patient having a second spinal tap 127 days after disease onset. Serum and CSF NfL levels increased in most patients with CSF/serum NfL ratios staying within the low or high ratio groups irrespective of GBSDS (Fig. 2C).

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### Table 1. Statistical values and analyses from patients and related measurements of NfL and albumin levels

|                    | GBS-ratio\textsubscript{low} | GBS-ratio\textsubscript{high} | Controls |
|--------------------|-------------------------------|-------------------------------|----------|
| n                  | 18                            | 3                             | 19       |
| Age (f:m)          | 58.4 (6:12)                   | 60.7 (0:3)                    | 64.7 (9:10) |
| Time to LP [days]  | 17.1                          | 3.7                           | n/a      |
| Mean CSF cell count| 2.2                           | 1                             | 1.2      |
| CSF/serum Albumin ratio (Qalb) | 17.1            | 13.8                          | 6.1      |
| Mean serum NfL [pg/mL] | 2871.6                     | 397.4                         | 50.7     |
| Mean CSF NfL [pg/mL] | 5939.5                       | 15854                         | 1115     |
| NfL CSF/Serum ratio | 3.96                         | 33.32                         | 26.7     |
| NfL-index          | 0.4                           | 3.15                          | 4.97     |
| Ganglioside antibody positive | 4                          | 1                             | n/a      |
| GBS disability score | 3.6                          | 3.1                           | n/a      |
| Electrophysiology: axonal/demyelinating | 16/14                       | 0/3                           | n/a      |
| Electrophysiology: F-waves measurable? | 0/18                       | 1/3                           | n/a      |
| sNfL/CSF-NfL       | 0.003*/0.07                   | 0.005*/0.004*                 | n/a      |
| sNfL/CSF-NfL       | 0.023*/0.048*                 | 0.333/0.333                   | n/a      |
| Spearman correlation GBSDS P-value | 0.099/0.443               | -0.001*/-0.001*               | n/a      |
| Spearman correlation CSF/Serum ratio P-value | 0.581/0.003*               | 0.666/0.666                   | n/a      |

*ANOVA NfL levels versus controls 0.003*/0.07 0.005*/0.004* n/a
Spearman correlation GBSDS P-value 0.023*/0.048* 0.333/0.333 n/a
Spearman correlation Qalb P-value 0.099/0.443 -0.001*/-0.001* n/a
Spearman correlation CSF/Serum ratio P-value 0.581/0.003* 0.666/0.666 n/a
Applying the NfL index results in almost similar results

Albumin ratio and sNfL did not correlate in GBS patients overall ($P = 0.76$) and not in the NfL-ratio\textsubscript{low} group ($P = 0.44$) but in the group NfL-ratio\textsubscript{high} ($P < 0.01$). CSF-NfL and Qalb in NfL-ratio\textsubscript{high} also correlated well ($P < 0.01$) but not in the NfL-ratio\textsubscript{low} (Table 1).

NfL index in controls was 4.97 (ranged from 0.78 to 18.27) with lowest quartile at 2.8. Using the NfL index elicited slightly different subdivisions depending on the cut-off used. Applying the lowest quartile from the NfL index obtained from controls would leave only one patient in the intrathecal NfL group. Simply using the NfL index produced a gap (Fig. 2D) separating four GBS patients from the other (NfL index range from 1.77 to 5.83) comprising the four patients with the highest CSF/Serum NfL ratio (ratio ranging from 10.95 to 42.65) and having a mixed demyelinating-axonal PNS pathology being inside the high NfL index group (Fig. 2D). The parameter “time disease onset to LP” would remain highly different between NfL index high versus NfL index low group ($t = 6.25$ days to $t = 17.8$ days).

Mean NfL index in NfL-ratio\textsubscript{high} was at 3.15 (1.78–4.83) and not different from that of controls ($P = 0.34$). In NfL-ratio\textsubscript{low} mean NfL index was 0.4 (ranging from 0.00581 to 2.14) with significant difference to the controls ($P < 0.01$). Comparison of the interquartile range for the

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**Figure 1.** (A) Split up of the GBS patients per GBS Disability score 2 to 5 showing NfL CSF/serum ratio, Serum-NfL and CSF-NfL levels, and Albumin quotient per score. (B) CSF-NfL, Serum-NfL and its ratio separated for GBS patients who had been admitted to the ICU and not. Significant difference is marked with *.
C/S NfL ratio or the NfL index with the controls also showed highly significant differences (Kolmogorov-Smirnoff $P < 0.001$, Fig. 3).

**Discussion**

NfL ratio and NfL index are markers to determine the origin of NfL. This is important to assess the contribution of the different NfL sources to the sNfL level. We could show in this study that NfL of peripheral origin can be a significant contributor of overall sNfL levels. In general, sNfL is a biomarker for GBS and is also a marker for disease severity as seen before for CSF-NfL.

Guillain-Barré Syndrome is a dynamic, acute polyneuropathy causing damage in PNS with intrathecal structures involved. Edema in the nerve root are probably causing the phenomena of cyto-albumin dissociation because proteins with at least the size of 70 kDa as albumin cannot dissociate toward the blood compartment. If this hypothesis is true, NfL having a
molecular weight of 68 kDa may also not dissociate from and in the blood compartment explaining the high NfL levels in the CSF and serum in follow-up examinations. The retrograde neurodegeneration involving the long peripheral nerves during the course of the disease may also contribute to the disproportionate increase in sNfL but this mechanism comes into significance later on. Parallel to this phenomenon axonal damage of the peripheral nerves including at the neuromuscular junction according to the underlying pathology of the GBS may cause an “NfL flood” into the serum releasing NfL into the blood compartment probably keeping the Nfl ratio constant at first. These mechanisms may explain the difference in both groups in time from disease onset to lumbar puncture and may explain the increase in total amount of NfL in CSF and serum over time in the follow-up cohort. One more mechanism may also contribute to the increase of sNFL. Due to the damaged blood–nerve-barrier at the nerve root NfL passes into the serum. This diffusion may be hampered due to the edema in the nerve root. The contribution of this mechanism to the overall sNFL and vice versa is not quantifiable with our data. In total several mechanisms are possibly contributing to the results as seen here.

The precise half-life of NfL in serum is unknown but could be another explanation for the increase of NfL levels overall. Due to its resistance to proteases NfL has an estimated half-life of several months in blood. This resistance could result in an additive effect increasing Nfl levels over time.18

The albumin ratio in the Nfl-ratio<sub>low</sub> group was slightly higher than in the Nfl-ratio<sub>high</sub> group. In favor of this, serum and CSF levels of NfL and Qalb correlated in the Nfl-ratio<sub>high</sub> group ($P < 0.001$), while this was not the case for the Nfl-ratio<sub>low</sub> group (Table 1).

Applying the Nfl index produces highly significant differences between controls and GBS patients pointing at a peripheral origin of sNfL in most GBS patients but not in controls. The Nfl-ratio<sub>high</sub> patients showed a Nfl index similar to that of controls again suggesting that the larger part of NfL should derive from within the CSF space. In contrast, the Nfl-ratio<sub>low</sub> group had a mean Nfl index far lower than in controls, suggesting that a major quantity is released from extrathecal nerve tissue.

Another factor in favor for sNfL as a biomarker is its correlation with the severity of the disease, which is very heterogeneous and dynamic in GBS. Here, we do find a correlation with GBDS and meaning that the higher the GBDS the more NfL is released, predominantly from PNS into the serum. Furthermore, sNfL and not CSF-NfL is higher in patients admitted to the ICU confirming the biomarker status at least for sNfL.

**Conclusion**

The existence of NfL with peripheral origin in blood changes the way we look at sNfL. One suggestion is to measure the Nfl ratio or index to assure the biomarker quality of sNfL for a CNS disease. Another suggestion is to rule out every patient having subacute or acute damage to peripheral nerves due to, for example, contact sports,
medication with potentially neurotoxic agents or diabetic and other inflammatory polyneuropathies.

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**Author Contributions**

PK contributed to design and conceptualization of the study, major role in the acquisition of data, analysis or interpretation of the data, drafting or revising the manuscript for intellectual content. JK contributed to analysis or interpretation of the data drafting or revising the manuscript for intellectual content major role in the acquisition of data. ED contributed to drafting or revising the manuscript for intellectual content major role in the acquisition of data, analysis or interpretation of the data. BS contributed to major role in the acquisition of data, drafting or revising the manuscript for intellectual content, analysis or interpretation of the data.

**Conflict of Interest**

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