Neurofilaments as Biomarkers for Amyotrophic Lateral Sclerosis: A Systematic Review and Meta-Analysis

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Abstract

Background

To allow early diagnosis and monitoring of disease progression, there is a need for biomarkers in amyotrophic lateral sclerosis (ALS). Neurofilaments (NF) are emerging protein biomarkers in other neurological diseases, and are of possible use in ALS.

Objective

The aim of this study is to evaluate the utility of NF levels as blood or cerebrospinal fluid (CSF) biomarker in patients with ALS.

Methods

A systematic search of Pubmed, Embase and Scopus was performed. Methodological quality assessment was applied to refine the final search results. Meta-analysis of the data was performed.

Results

Level of NF heavy chain and light chains were significantly elevated in the CSF of ALS patients compared to healthy controls/controls without parenchymal central nervous system (CNS) involvement and ALS mimic disease patients. NF light chain level in CSF was higher in ALS patients than in neurological patients with CNS involvement (SMD = 1.352, P = 0.01). NF light chain concentration in blood was higher in ALS patients than healthy controls/controls without CNS involvement (SMD = 1.448, P < 0.0001). NF heavy chain levels in CSF were negatively correlated disease duration and ALSFRS-R (r = -0.447, P < 0.0001; r = -0.486, P < 0.0001). NF light chain levels in CSF were negatively correlated with disease duration (r = -0.273, P = 0.011).
Conclusion
NF heavy and light chain levels have potential use as a marker of neural degeneration in ALS, but are not specific for the disease, and are more likely to be used as measures of disease progression.

Introduction
Neurofilaments (NF) are intermediate filaments that are major components of neuronal cytoskeleton. NF can be divided according to the observed molecular weight into NF light chain (68 kda), NF intermediate chain (160 kda) and NF heavy chain (205kda) [1]. The three different NFs share a conserved alpha-helical rod domain, but differ in the head and tail domains [2]. NFM and NFH are always phosphorylated [3]. NFL is susceptible to protease degradation, while NFH, which is phosphorylated, can resist protein degradation [4]. Damage to axons of the central nervous system (CNS) or peripheral nervous system (PNS) could release the NF, which would then appear in the cerebrospinal fluid (CSF) and the bloodstream, where NF can be detected with techniques such as enzyme linked immunosorbent assay (ELISA), western blot and mass spectrometry [5, 6].

Amyotrophic lateral sclerosis (ALS) is a fatal neurodegenerative disease, which involves progressive loss of both upper and lower motor neurons. The incidence of ALS is about 1–2 per 100,000 [7]. Almost 90% of the cases of ALS are sporadic and 5–10% of the cases are familial [8]. Generally, the median survival of patients from symptom onset is about 2–3 years [9] and the cause of the death is respiratory failure. However, there is heterogeneity of disease survival, and approximately 10% of patients survive for more than 10 years [10]. Analysis of spinal cord tissue has suggested abnormal NFH subunit accumulation in neuronal perikarya and spheroids in ALS patients compared with control [11].

There is a need for a biomarker in ALS, for use in diagnosis, prognosis and in clinical trials. Katz et al. defined biomarkers as a "objectively measured and evaluated parameters for indication of pathological processes, disease progression or response to pharmacological interventions" [12]. Various bodily fluids have been used for biomarkers, including blood, CSF, urine and saliva. Among these biofluids, blood and CSF have differing advantages. CSF is expected to contain proteins from degenerating neurons because of its direct contact with the CNS, so is suitable for the study of CNS disease. Blood samples have a less invasive process of collection, which is much more acceptable for patients. Biomarkers that enter the CSF will eventually drain into the veins, so blood samples will include protein released from degenerating neurons. The disadvantage of studying blood is its complexity, as blood contains many different proteins.

The aim of this systematic review and meta-analysis is to investigate whether NF levels in blood or CSF could be a reliable biomarker for amyotrophic lateral sclerosis, either in distinguishing patients from controls or as markers monitoring disease progression or predicting prognosis. The preferred reporting items for systematic review and meta-analysis (PRISMA) checklist was shown in the S1 File [13].

Methods
1. Search strategy and key words
A systematic search was conducted in Pubmed, Embase, Scopus and Medline, with the latest date of search being 20th May 2016.
The search strategy in Pubmed is as follows:

(((((((((("Blood"[Mesh]) OR ((("Cerebrospinal Fluid"[Mesh]) OR "Cerebrospinal Fluid") OR "Cerebrospinal Fluids"))) OR blood)) AND "Amyotrophic Lateral Sclerosis"[Mesh]) AND neurofilament) AND "Humans"[Mesh]) AND English[lang])) AND ((((("Biomarkers"[Mesh]) OR biomarker))))

The search strategy in Embase is as follows:

'amyotrophic lateral sclerosis'/exp AND ('biological marker'/exp OR biomarker OR biomarkers) AND ('neurofilament'/exp OR neurofilament OR neurofilaments) AND ('blood'/exp OR 'cerebrospinal fluid'/exp OR 'cerebrospinal fluid' OR 'cerebrospinal fluids') AND 'human'/
de

The search strategy in Scopus is as follows:

(TITLE-ABS-KEY("Amyotrophic Lateral Sclerosis")and ((TITLE-ABS-KEY(blood) OR TITLE-ABS-KEY("Cerebrospinal Fluid")OR TITLE-ABS-KEY("Cerebrospinal Fluids")) and ((TITLE-ABS-KEY(biomarker) OR TITLE-ABS-KEY(biomarkers)) and ((TITLE-ABS-KEY(neurofilament) OR TITLE-ABS-KEY(neurofilaments)))AND(LIMIT-TO(EXACTKEYWORD,"Human")) AND (LIMIT-TO(LANGUAGE,"English"))

2. Selection criteria

The inclusion criteria are as follows:

1. The study must be carried out in humans more than 18 years of age.
2. All the subjects with ALS must be diagnosed with the EL Escorial criteria or revised El Escorial [14] or Awaji [15] criteria.
3. A demographic description of the patients must be provided.
4. The study must use NF as a biomarker in the blood or CSF to differentiate ALS patients from a control group or to distinguish subgroups of ALS patients.
5. The study must provide clear description of the time point and method for collection of blood and CSF.
6. The study must provide a full description of the method used to measure NF levels.
7. The study must be the original paper rather than abstracts, posters or reviews.

The exclusion criteria are as follows:

1. The subjects were younger than 18 years.
2. The ALS subjects had additional neurological disorders or pathologic changes such as brain tumour, epilepsy or brain injury.
3. The number of subjects was less than 5.
4. Studies without healthy controls or without disease control.
5. The study was not published in English.
6. Studies with missing data such as details about the demographic information and details of method used for NF detection.
7. The papers are abstracts or posters or reviews
3. Assessment of methodological quality

A total of 20 papers were selected for the final systematic review. The original extracted data on NF levels is summarized in S1, S2, S3 and S4 Datasets. We used the QUADAS-2 (Quality Assessment of Diagnostic Accuracy Studies) criteria to assess the 20 papers selected for systematic review[16]. This tool is made up of 4 key domains which assessed the risk of bias covering patient selection, index test, reference standard and flow and timing respectively. The first three domains were also assessed the concerns in terms of applicability. The assessment was done independently by two authors (ZX and PM). If there was disagreement, the third author RH was consulted to resolve the disagreement. The final results of quality assessment of the 20 papers and the proportion of high quality and low quality papers were summarized in S1 Table.

4. Data synthesis and statistics

We attempted to obtain the original data by contacting the authors of the selected papers, but this was not available. Comprehensive meta-analysis software V 2.0 (Biostat, USA) was used to perform the final data combination and meta-analysis. When the mean value, standard deviation (SD), correlation coefficient and size of cohort were not available, a series of formulas, as described by Hozo et al. and Wan et al. [17, 18] was utilized to estimate the sample mean and standard deviation from the published sample size, median, range or inter-quartile range. If the median was the only value provided, the study could not be included into the meta-analysis.

The studies used a variety of disease controls. To deal with this heterogeneity, we subdivided the disease controls into three groups; 1) healthy control/controls without parenchymal involvement of the CNS, 2) ALS mimics and 3) controls with neurological disease with parenchymal CNS involvement. The controls without parenchymal CNS involvement had a range of conditions including Guillain-Barré syndrome, tension headache, back pain, normal pressure hydrocephalus, facial palsy and polyneuropathy.

When doing the meta-analysis, as required by the Cochrane handbook for systematic reviews of Interventions[19], we used standardized mean difference to combine the overall effect on the basis of the mean± SD value from each study. We used Fisher's Z test to combine the overall effect on the basis of the correlation coefficient and sample size. P value less than 0.05 was considered as significant. If the I² statistics of the heterogeneity of the studies was less than 40%, the fixed effect meta-analysis model was chosen. If the I² statistics was more than 50%, the random effect model was applied [19].

The results of the meta-analysis of NF concentration difference between the ALS group and the control groups are illustrated as forest plots that show the standardized mean difference (SMD) between the two groups. The results of the meta-analysis of correlations between NF and disease duration or progression are also illustrated as forest plots that show the correlation coefficient between the two variables.

Results

1. Study characteristics

The total number of papers obtained from the search was 49. The flow diagram based on PRISMA is shown in Fig 1[13]. S2 Table shows the details of 20 papers assessed as being suitable for the systematic review. The details of the controls in every paper are also summarized in the S2 Table. The 20 papers are categorized according to the biofluids that were sampled. The summarized results are shown in the S3 Table. There were more studies of NFH than NFL.
Because the NF heavy chain is phosphorylated, all the detection methods were targeted to phosphorylated NFH (pNFH). The terms pNFH and NFH are used interchangeably in the literature. The summary of the results of the meta-analysis are shown in Table 1.
2. NFH levels in CSF

**ALS versus healthy controls/controls without parenchymal CNS involvement.** In CSF, the NFH meta-analysis between ALS patients and healthy controls/controls without parenchymal CNS involvement was based on 5 studies [20–24], all using ELISA, including 443 ALS patients and 267 healthy controls (Fig 2). The level of NFH in ALS patients is significantly higher than that of the controls (SMD = 1.018, \( P < 0.0001 \)). Another paper used Western blot analysis and identified one unique pNFH in the CSF of ALS patients, which is absent in the CSF of healthy controls [25].

### Table 1. Summary of the meta-analyses.

|                  | ALS v HC/non CNS disease | ALS v mimic disease | ALS v CNS disease |
|------------------|--------------------------|---------------------|-------------------|
|                  | No of studies | Total no of subjects (patients/controls) | P value of meta-analysis | No of studies | Total no of subjects (patients/controls) | P value of meta-analysis | No of studies | Total no of subjects (patients/controls) | P value of meta-analysis |
| NFH in CSF       | 4 studies (5 cohorts)   | 443/267             | P < 0.0001        | 2 studies     | 251/100                         | P = 0.013                       | 6 studies (11 cohorts) | 468/329                     | P = 0.075                       |
| NFH in blood     | 2 studies       | 117/78              | P = 0.057         |               |                                |                                 |                       |                            |                                 |
| NFL in CSF       | 6 studies (7 cohorts) | 463/214             | P < 0.0001        | 2 studies     | 250/99                          | P < 0.0001                      | 5 studies     | 398/405                     | P = 0.001                       |
| NFL in blood     | 3 studies (5 cohorts) | 202/277             | P < 0.0001        |               |                                |                                 |                       |                            |                                 |

doi:10.1371/journal.pone.0164625.t001

![Fig 2. NFH CSF ALS-healthy](image)

Meta-analysis of NFH levels in CSF between ALS patients and healthy controls/controls without parenchymal CNS involvement. The random effect model was used. There was a highly significant difference between the two groups (\( P < 0.0001 \)).

doi:10.1371/journal.pone.0164625.g002
ALS versus ALS mimics disease. In CSF, the NFH meta-analysis between ALS patients and ALS mimic disease controls was based on 2 studies [24, 26] including 251 patients and 100 ALS mimic disease controls on basis of ELISA technology (Fig 3). The standardized mean difference is significant (SMD = 0.796, P = 0.013).

ALS versus other neurological diseases with CNS involvement. The NFH meta-analysis between ALS patients and patients with other neurological disease with CNS involvement was based on 6 studies [20–22, 24, 27, 28] including 329 disease controls and 468 ALS patients (Fig 4). There was no significant difference between the two groups (P = 0.075).

Higher levels of pNFH<sub>SMI35</sub> were reported in patients with upper motor neuron dominant disease than in those with typical ALS [20]. Higher levels were also reported in symptomatic carriers of causative mutations than asymptomatic carriers [23]. Another paper suggested that the ratio of CSF pNFH/complement C3 could be used to discriminate the ALS patients and disease control [29].

3. NFH levels in Blood

ALS versus healthy controls/controls without parenchymal CNS involvement. The result for the NFH meta-analysis between healthy controls/controls without parenchymal CNS involvement and ALS patients in blood was based on 2 studies [30, 31] including 78 healthy controls and 117 patients (Fig 5). The concentration of NFH in the ALS patients group was not significantly increased relative to the healthy controls/without CNS involvement (P = 0.057).

Another paper [29] also found an increased level of NFH in ALS compared to healthy controls and other neurological disease, but there was insufficient data for inclusion in the meta-analysis.

There was no significant difference in levels of NFH between patients treated with Riluzole and those without [30]. Bulbar onset patients were reported to have elevated concentration of NFH in plasma compared with spinal onset patients [32]. There is correlation of pNFH between CSF and blood levels of NFH [29]. Two papers reported that the concentration of NFH in the blood showed an initial rise but later the levels fell [30, 32].

4. NFL levels in CSF

ALS versus healthy controls/controls without parenchymal CNS involvement. The NFL meta-analysis between ALS patients and healthy controls/controls without parenchymal CNS involvement was based on 2 studies [24, 26] including 251 patients and 100 ALS mimic disease controls on basis of ELISA technology (Fig 3). The standardized mean difference is significant (SMD = 0.796, P = 0.013).
involvement in CSF was based on 6 studies [23, 24, 33–36] including 463 ALS patients and 214 healthy controls (Fig 6). The standardized mean difference was significantly increased in comparison with the controls (SMD = 1.627, P < 0.0001).

**ALS versus ALS mimics disease.** The NFL meta-analysis between ALS patients and mimic disease control was based on 2 studies [24, 26] including 250 ALS patients and 99 ALS mimic disease patients (Fig 7). The level of NFL in ALS mimic disease was significantly lower than the ALS patients group (SMD = 0.742, P < 0.0001).

**ALS versus other neurological diseases with CNS involvement.** The meta-analysis of ALS patients and disease control with CNS involvement in CSF samples was based on 5 studies [24, 33–35, 37] including 398 ALS patients and 405 disease controls. A significant elevation of NFL was found in the ALS patients (SMD = 1.625, P = 0.001).

No difference of NFL level in the CSF was found between different phenotypes and no correlation was found between NFL level in the CSF and EMG features of lower motor neuron denervation [37]. The level of NFL in the CSF of symptomatic ALS gene mutation carriers was significantly greater than that of asymptomatic ALS mutation carriers [23] and patients carrying a SOD1 mutation have a lower concentration of NFL than those without [34].

### 5. NFL levels in blood

**ALS versus healthy controls/controls without parenchymal CNS involvement.** In blood, the NFL meta-analysis between ALS patients and healthy controls without CNS involvement was based on 3 studies [23, 35, 36] including 277 healthy controls and 202 ALS patients (Fig 9). The standardized mean difference between the two groups was statistically
Significant (SMD = 1.448, P < 0.0001). The papers used ELISA except for one which used electrochemiluminescence (ECL) based immune assay [35].

Symptomatic ALS gene mutation carriers had significantly increased blood levels of NFL compared to asymptomatic carriers of causative mutations.[23]. There were no differences in the blood NFL level between ALS patients treated with Riluzole and those without [30]. Three papers found that there was a high correlation of NFL levels between CSF and serum and/or plasma [23, 35, 36].

### NFH Blood ALS-healthy

Meta-analysis of NFH levels in blood between ALS patients and healthy controls/controls without parenchymal CNS disease. The random effect model was applied. There is no significant difference between the two groups (P = 0.068).

Fig 5. NFH Blood ALS-healthy. Meta-analysis of NFH levels in blood between ALS patients and healthy controls/controls without parenchymal CNS disease. The random effect model was applied. There is no significant difference between the two groups (P = 0.068).

doi:10.1371/journal.pone.0164625.g005

### NFL CSF ALS-healthy

Meta-analysis of NFL levels in CSF between ALS patients and healthy controls/controls without parenchymal CNS disease. The random effect model was applied. There is significant difference between the two groups (P < 0.0001).

Fig 6. NFL CSF ALS-healthy. Meta-analysis of NFL levels in CSF between ALS patients and healthy controls/controls without parenchymal CNS disease. The random effect model was applied. There is significant difference between the two groups (P < 0.0001).

doi:10.1371/journal.pone.0164625.g006

| Study name     | Std diff in means | Lower limit | Upper limit | Z-Value | p-Value |
|----------------|-------------------|-------------|-------------|---------|---------|
| McCombe 2015   | 0.529             | 0.201       | 0.858       | 3.161   | 0.002   |
| Boylan 2009    | 1.699             | 0.957       | 2.441       | 4.489   | 0.000   |
| Overall effect | 1.065             | -0.077      | 2.207       | 1.827   | 0.068   |

Test for heterogeneity: $\chi^2=7.990; df=1$ (P=0.005); $I^2=87.484\%$

| Study name     | Std diff in means | Lower limit | Upper limit | Z-Value | p-Value |
|----------------|-------------------|-------------|-------------|---------|---------|
| Zetterberg 2007| 1.346             | 0.929       | 1.763       | 6.327   | 0.000   |
| Rosengren 1996 | 1.932             | 1.164       | 2.699       | 4.933   | 0.000   |
| Weydt (ULM) 2016| 2.014            | 1.233       | 2.794       | 5.056   | 0.000   |
| Weydt (UMEA) 2016| 2.060           | 1.416       | 2.704       | 6.268   | 0.000   |
| Gaiottino 2013 | 1.687             | 1.083       | 2.290       | 5.481   | 0.000   |
| Steinacker 2015| 0.696             | 0.411       | 0.982       | 4.778   | 0.000   |
| Lu (Oxford) 2015| 1.990             | 1.339       | 2.641       | 5.988   | 0.000   |
| Overall effect | 1.627             | 1.139       | 2.114       | 6.541   | 0.000   |

Test for heterogeneity: $\chi^2=34.254; df=6$ (P=0.000); $I^2=82.484\%$
6. Correlation with the measures of disease

In the papers selected for analysis, there have been attempts to determine whether levels of NF are related to the measures of disease. This has been done by examining the relationship of NF with the disease duration, disease progression measured by revised ALS functional rating scale (ALSFRS-R) \[38\] and survival time. There was considerable heterogeneity among the statistic approaches used in the different papers. So only limited meta-analysis was possible.

![Fig 7. NFL ALS-mimic. Meta-analysis of NFL levels in CSF between ALS patients and ALS mimics. Because the I² = 0, the fixed effect model was applied. There is significant difference between the two groups (P<0.0001).](#)
doi:10.1371/journal.pone.0164625.g007

Fig 7. NFL ALS-mimic. Meta-analysis of NFL levels in CSF between ALS patients and ALS mimics. Because the I² = 0, the fixed effect model was applied. There is significant difference between the two groups (P<0.0001).

doi:10.1371/journal.pone.0164625.g007

### Test for heterogeneity: $\chi^2=0.036; df = 1 (P=0.850); I^2=0%$

| Study name  | Std diff in means | Lower limit | Upper limit | Z-Value | p-Value |
|-------------|-------------------|-------------|-------------|---------|---------|
| Reijn 2009  | 0.683             | 0.025       | 1.341       | 2.034   | 0.042   |
| Steinacker 2015 | 0.751         | 0.494       | 1.008       | 5.727   | 0.000   |
| Overall effect | 0.742           | 0.502       | 0.981       | 6.074   | 0.000   |

| -1.00 | -0.50 | 0.00 | 0.50 | 1.00 |
|-------|-------|------|------|------|
| NFL higher in control | NFL higher in ALS |

6. Correlation with the measures of disease

In the papers selected for analysis, there have been attempts to determine whether levels of NF are related to the measures of disease. This has been done by examining the relationship of NF with the disease duration, disease progression measured by revised ALS functional rating scale (ALSFRS-R) \[38\] and survival time. There was considerable heterogeneity among the statistic approaches used in the different papers. So only limited meta-analysis was possible.

![Fig 8. NFL CSF ALS-CNS disease. Meta-analysis of NFL levels in CSF between ALS patients and other neurological disease with CNS involvement. The random effect model was applied. There is significant difference between the two groups (P = 0.001).](#)
doi:10.1371/journal.pone.0164625.g008

Fig 8. NFL CSF ALS-CNS disease. Meta-analysis of NFL levels in CSF between ALS patients and other neurological disease with CNS involvement. The random effect model was applied. There is significant difference between the two groups (P = 0.001).

doi:10.1371/journal.pone.0164625.g008

| Study name  | Std diff in means | Lower limit | Upper limit | Z-Value | p-Value |
|-------------|-------------------|-------------|-------------|---------|---------|
| Tortelli 2012 | 1.312             | 0.835       | 1.789       | 5.395   | 0.000   |
| Zetterberg 2007 | 0.774           | 0.507       | 1.041       | 5.681   | 0.000   |
| Rosengren 1996 | 1.156             | 0.272       | 2.040       | 2.564   | 0.010   |
| Gaiottino 2013 | 2.957             | 2.460       | 3.455       | 11.658  | 0.000   |
| Steinacker 2015 | 0.607          | 0.305       | 0.909       | 3.943   | 0.000   |
| Overall effect | 1.352             | 0.582       | 2.121       | 3.443   | 0.001   |

| -4.00 | -2.00 | 0.00 | 2.00 | 4.00 |
|-------|-------|------|------|------|
| NFL higher in control | NFL higher in ALS |

### Test for heterogeneity: $\chi^2=70.132; df = 4 (P=0.000); I^2=94.296%$
Correlation with disease duration. The extracted results and statistic methods of the correlation with disease duration were summarised in S4 Table. The majority of papers showed a negative correlation of disease duration with NF levels. There was heterogeneity in the statistical methods. However, a meta-analysis was possible for correlation of CSF NFH and NFL levels with disease duration. These are shown in Figs 10 and 11 respectively.

The correlation meta-analysis between CSF NFH and disease duration was based on 2 studies [23, 24] including 516 patients. The overall correlation coefficient was significant ($r = -0.447, P < 0.0001$). The correlation meta-analysis between CSF NFL and disease duration was based on 4 studies [23, 24, 34, 37] including 622 patients. The overall correlation coefficient was significant ($r = -0.486, P < 0.0001$).

### Table 1: Correlation of CSF NFH and NFL with disease duration

| Study name         | Std diff in means | Lower limit | Upper limit | Z-Value | p-Value |
|--------------------|-------------------|-------------|-------------|---------|---------|
| Gaiottino 2013     | 1.122             | 0.739       | 1.505       | 5.746   | 0.000   |
| Lu (Oxford) 2015   | 1.606             | 1.141       | 2.071       | 6.769   | 0.000   |
| Lu (London) 2015   | 1.462             | 1.066       | 1.859       | 7.232   | 0.000   |
| Weydt (ULM) 2016   | 1.838             | 1.080       | 2.597       | 4.751   | 0.000   |
| Weydt (UMEA) 2016  | 1.723             | 1.109       | 2.337       | 5.503   | 0.000   |
| Overall effect     | 1.448             | 1.236       | 1.661       | 13.378  | 0.000   |

Test for heterogeneity: $\chi^2 = 5.204; df = 4 (P = 0.285); I^2 = 20.382\%$

Fig 9. NFL blood ALS-healthy. Meta-analysis of NFL levels in blood between ALS patients and healthy controls/controls without parenchymal CNS disease. The fixed effect model was applied. There is significant difference between the two groups ($P < 0.0001$).

doi:10.1371/journal.pone.0164625.g009

### Table 2: Correlation of CSF NFH and NFL with disease duration

| Study name          | Correlation | Lower limit | Upper limit | Z-Value | p-Value |
|---------------------|-------------|-------------|-------------|---------|---------|
| Steinacker 2015     | -0.430      | -0.502      | -0.352      | -9.778  | 0.000   |
| Weydt 2016          | -0.565      | -0.715      | -0.365      | -4.875  | 0.000   |
| Overall effect      | -0.447      | -0.513      | -0.374      | -10.849 | 0.000   |

Test for heterogeneity: $\chi^2 = 1.670; df = 1 (P = 0.195); I^2 = 40.125\%$

Fig 10. CSF NFH and disease duration. Meta-analysis of correlation between CSF NFH and disease duration. The fixed effect model was applied. There is significant negative correlation between two variables ($P < 0.0001$).

doi:10.1371/journal.pone.0164625.g010
Correlation with disease duration. The extracted results and statistic methods of the correlation with disease progression were summarised in S4 Table. The papers used different statistic method to find the correlation with ALSFRS-R or the rate of change of ALSFRS-R. Cross sectional correlation of NF levels with ALSFRS-R scores and comparison between patients with rapid progression and ones with slow progression were also mentioned in the papers. A meta-analysis was possible for correlation of CSF NFH and NFL levels with ALSFRS-R score. These are shown in Figs 12 and 13 respectively.

The correlation meta-analysis between CSF NFH and ALSFRS-R score was based on 3 studies [23, 24, 32] including 536 patients. The overall correlation coefficient was significant (r = -0.273, P = 0.011). The correlation meta-analysis between CSF NFL and ALSFRS-R was based on 3 studies [23, 24, 37] including 543 patients. The overall correlation coefficient was not significant (r = -0.447, P = 0.156). One study suggested using the ratio CSF pNFHSMI35/CSF NFL.

| Study name     | Statistics for each study | Correlation and 95% CI |
|----------------|---------------------------|------------------------|
|                | Correlation | Lower limit | Upper limit | Z-Value | p-Value |                  |
| Tortelli 2012  | -0.550      | -0.742      | -0.275      | -3.606   | 0.000   |                  |
| Zetterberg 2007| -0.518      | -0.663      | -0.335      | -5.001   | 0.000   |                  |
| Steinacker 2015| -0.376      | -0.452      | -0.294      | -8.406   | 0.000   |                  |
| Wedigt 2016    | -0.606      | -0.755      | -0.397      | -4.868   | 0.000   |                  |
| Overall effect | -0.486      | -0.598      | -0.356      | -6.574   | 0.000   |                  |

Test for heterogeneity: $\chi^2=6.524; df=3 (P=0.089); I^2=54.018\%$

Fig 11. CSF NFL and disease duration. Meta-analysis of correlation between CSF NFL and disease duration. The random effect model was applied. There is significant negative correlation between two variables (P<0.0001).

doi:10.1371/journal.pone.0164625.g011

Correlation with disease duration. The extracted results and statistic methods of the correlation with disease progression were summarised in S4 Table. The papers used different statistic method to find the correlation with ALSFRS-R or the rate of change of ALSFRS-R. Cross sectional correlation of NF levels with ALSFRS-R scores and comparison between patients with rapid progression and ones with slow progression were also mentioned in the papers. A meta-analysis was possible for correlation of CSF NFH and NFL levels with ALSFRS-R score. These are shown in Figs 12 and 13 respectively.

The correlation meta-analysis between CSF NFH and ALSFRS-R score was based on 3 studies [23, 24, 32] including 536 patients. The overall correlation coefficient was significant (r = -0.273, P = 0.011). The correlation meta-analysis between CSF NFL and ALSFRS-R was based on 3 studies [23, 24, 37] including 543 patients. The overall correlation coefficient was not significant (r = -0.447, P = 0.156). One study suggested using the ratio CSF pNFHSMI35/CSF NFL.

| Study name     | Statistics for each study | Correlation and 95% CI |
|----------------|---------------------------|------------------------|
|                | Correlation | Lower limit | Upper limit | Z-Value | p-Value |                  |
| Boylan 2012    | -0.590      | -0.819      | -0.200      | -2.794   | 0.005   |                  |
| Steinacker 2015| -0.163      | -0.251      | -0.072      | -3.497   | 0.000   |                  |
| Weydt 2016     | -0.285      | -0.501      | -0.036      | -2.232   | 0.026   |                  |
| Overall effect | -0.279      | -0.468      | -0.065      | -2.540   | 0.011   |                  |

Test for heterogeneity: $\chi^2=4.940; df=2 (P=0.085); I^2=59.515\%$

Fig 12. CSF NFH and ALSFRS-R. Meta-analysis of correlation between CSF NFH and ALSFRS-R. The random effect model was applied. There is significant negative correlation between two variables (P = 0.011).

doi:10.1371/journal.pone.0164625.g012
sAPP $\alpha-\beta$ to monitor disease progression [21]. Another paper discovered that the CSF NFL was correlated with spreading from localized to generalized weakness [39].

**Correlation with survival time or prognosis.** The extracted results and statistic methods of the correlation with disease progression were summarised in S4 Table. Due to the considerable heterogeneity of statistic method. The meta-analysis of correlation between NF and survival time was impossible. Several papers found that the level of NF or the rate of rise of NF was negatively correlated with survival time or increasing odds ratio of death in both CSF and blood[24, 30, 32, 34, 36, 37]. Only one paper pointed out that no correlation was found between cumulative mortality and base line pNFH[31].

7. Sensitivity and Specificity of NF

Some of the papers provided and used receiver operating characteristic (ROC) to analyse the sensitivity and sensitivity of the NF levels in distinguishing ALS from the various control groups. The extracted results of ROC were shown in the S5 Table. Different optimal cut-off values were used among the papers, the sensitivity of ROC analysis ranges from 0.71 to 0.91. The specificity of ROC analysis ranges from 0.64 to 0.95. Area under curve (AUC) ranges from 0.77 to 0.9987. The ROC analysis on blood NF was very few.

**Discussion**

Here we have analysed the studies of NFH and NFL as biomarkers of ALS. The studies used different controls, so we grouped the controls into healthy controls/without parenchymal CNS disease, ALS mimic diseases and CNS parenchymal diseases. In CSF, both NFL and NFH levels were significantly greater in ALS than in healthy controls/patients without parenchymal CNS disease, while for blood, NFL levels showed a highly significant difference. For NFH, the number of subjects was lower than for CSF, and the difference was not statistically significant. Comparing ALS with other CNS disease, and comparing ALS with ALS mimic disease, only CSF was studied. CSF NFL and NFH levels were significantly greater in ALS than ALS mimic disease. CSF NFL levels, but not CSF NFH levels, were significantly greater in ALS than the other CNS diseases that were tested.

Before doing the meta-analysis, we attempted to get all the original data from the author of every paper. However it was not possible due to the confidentiality or other various reasons.
from the authors. So we used a series of formulas to infer and calculate the mean and SD from the published sample size, median, range or inter-quartile range. These formulas were widely acknowledged and used in other meta-analysis [40, 41].

The finding that NFH levels are higher in ALS than in healthy controls/non CNS parenchymal disease is consistent with the view that damage to axons releases NF. There was a significant difference between CSF NFL and the CNS diseases used as controls, but it must be noted that elevation of NF levels is not specific for ALS and has been reported in a number of other CNS diseases [42, 43] [2, 44–47]. For example, levels of NF in subarachnoid haemorrhage were much higher than in ALS patients [22] and similar results were also observed in the other studies [48, 49].

In the meta-analysis, we included GBS data in the control group without parenchymal CNS involvement. However, two researchers reported that the NFH or NFL in the CSF of GBS patients was higher than in the other neurological disease such as AD and PD with CNS involvement but less than ALS patients [22, 35]. NF are found in peripheral as well as central nervous system axons, and can also be released by peripheral axonal degeneration.

Given that raised NF levels are not specific for ALS, the most useful role of NF levels is likely to be in monitoring duration, progression or prognosis of disease. So we did meta-analyses about the correlation between CSF NFH or CSF NFL with ALSFRS-R score and disease duration. The final results showed that CSF NFH had a significant negative association with both disease duration and progression. Significant correlation was also found between CSF NFL and disease duration. Conversely no significant relationship was found between CSF NFL and disease progression. Worse prognosis or high odds ratio of death was also closely related with high baseline NF in most of the papers. However the NFL role in these fields still need to be verified and replicated with larger cohort by using same or standardized measurement scale in these three fields. There were some attempts to correlate NF levels with disease phenotype, but these were few.

To advance the field of NF as a biomarker, there needs to be standardization of the techniques that are used. In the papers reviewed here, ELISA, electrochemiluminescence (ECL) based immunoassay and western blot were used, with ELISA being used most commonly. One paper pointed out the traditional ELISA kit method was more sensitive than the ECL based immunoassay [50]. The versions of ELISA used in these studies were more sensitive than earlier methods [6, 51, 52]. The minimum limit of NFL ELISA kit used by the relevant papers ranges from 30 ng/ml to 125ng/ml. The phenomenon of the "hook effect" is an issue with measurements of NFH. This effect occurs with neurofilaments aggregation when blood samples are not diluted. [53], preventing the accurate quantification by immunoassay. This phenomenon was not observed inNFL in blood or with either NFL or NFH in CSF. As to the specificity and sensitivity of NF in the ALS patients, they not only depended on the detection methods as mentioned above, but also depends on the optimal cut-off and what other diseases to be differentiated from ALS. Neurological disease with more axon injury involvement was less easy to differentiate in comparison with healthy people.

In summary, a series of meta-analyses have shown that the NFH and NFL in CSF sample can be used to discriminate the ALS patients from the healthy people, and ALS mimics disease. NFL chain in CSF and blood sample can also be used to differentiate the ALS patients from the other neurological disease and healthy people respectively. There is promising evidence that NF levels could be used to monitor disease progression but this needs further work.

Supporting Information

S1 Dataset. CSF NFH original extracted data.
(XLSX)
S2 Dataset. CSF NFL original extracted data.
(XLSX)

S3 Dataset. Blood NFH original extracted data.
(XLSX)

S4 Dataset. Blood NFL original extracted data.
(XLSX)

S1 File. PRISMA checklist.
(DOC)

S1 Table. Results of Quadas-2 quality assessment.
(DOCX)

S2 Table. Summarized details of the papers.
(DOCX)

S3 Table. Summary of the papers according to the biofluid.
(DOCX)

S4 Table. Summary of studies correlated with disease duration, progression and survival time.
(DOCX)

S5 Table. Summary of specificity and sensitivity of the papers.
(DOCX)

Acknowledgments
We would like to thank Nicky Foxlee for providing the meta-analysis help.

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References

1. Dong DL, Xu ZS, Chevrier MR, Cotter RJ, Cleveland DW, Hart GW. Glycosylation of mammalian neurofilaments. Localization of multiple O-linked N-acetylglucosamine moieties on neurofilament polypeptides L and M. The Journal of biological chemistry. 1993; 268(22):16679–87. PMID: 8344946.

2. Teunissen CE, Khalil M. Neurofilaments as biomarkers in multiple sclerosis. Multiple sclerosis. 2012; 18(5):552–6. doi: 10.1177/1352458512443092 PMID: 22492131.

3. Nixon RA, Sihaq RK. Neurofilament phosphorylation: a new look at regulation and function. Trends in neurosciences. 1991; 14(11):501–6. Epub 1991/11/01. doi: 10.1016/0166-2236(91)90062-y PMID: 1726767.

4. Goldstein ME, Sternberger NH, Sternberger LA. Phosphorylation protects neurofilaments against proteolysis. Journal of neuroimmunology. 1987; 14(2):149–60. doi: 10.1016/0165-5728(87)90049-x PMID: 3029175.

5. Gnanapavan S, Grant D, Morant S, Furby J, Hayton T, Teunissen CE, et al. Biomarker report from the phase II lamotrigine trial in secondary progressive MS—neurofilament as a surrogate of disease progression. PloS one. 2013; 8(8):e70019. Epub 2013/08/13. doi: 10.1371/journal.pone.0070019 PMID: 23936370; PubMed Central PMCID: PMC3731296.

6. Patzold A. Neurofilament phosphoforms: surrogate markers for axonal injury, degeneration and loss. Journal of the neurological sciences. 2005; 233(1–2):183–98. Epub 2005/05/18. doi: 10.1016/j.jns.2005.03.015 PMID: 15896809.

7. Chio A, Logroscino G, Traynor BJ, Collins J, Simeone JC, Goldstein LA, et al. Global epidemiology of amyotrophic lateral sclerosis: a systematic review of the published literature. Neuroepidemiology. 2013; 41(2):118–30. doi: 10.1159/000351153 PMID: 23860588; PubMed Central PMCID: PMC4049265.

8. Blasco H, Guennoc AM, Veyrat-Durebeex C, Gordon PH, Andres CR, Camu W, et al. Amyotrophic lateral sclerosis: a hormonal condition? Amyotrophic lateral sclerosis: official publication of the World Federation of Neurology Research Group on Motor Neurons. 2012; 13(6):585–8. doi: 10.3109/17482968.2012.706303 PMID: 22873563.

9. Kiernan MC, Vucic S, Cheah BC, Turner MR, Eisen A, Hardiman O, et al. Amyotrophic lateral sclerosis. Lancet. 2011; 377(9769):942–55. doi: 10.1016/S0140-6736(10)61156-7 PMID: 21296405.

10. Talbot K. Amyotrophic lateral sclerosis: cell vulnerability or system vulnerability? Journal of anatomy. 2014; 224(1):45–51. doi: 10.1111/joa.12107 PMID: 24010870; PubMed Central PMCID: PMC3867886.

11. Mendonca DM, Chimelli L, Martinez AM. Quantitative evidence for neurofilament heavy subunit aggregation in motor neurons of spinal cords of patients with amyotrophic lateral sclerosis. Brazilian journal of medical and biological research Revista brasileira de pesquisas medicas e biologicas / Sociedade Brasileira de Biosciencia [et al]. 2005; 38(6):925–33. /S0100-879X2005000600015. PMID: 15933787. doi: 10.1590/s0100-879x2005000600015.

12. Katz R. Biomarkers and surrogate markers: an FDA perspective. NeuroRx: the journal of the American Society for Experimental NeuroTherapeutics. 2004; 1(2):189–95. doi: 10.1602/neurorx.1.2.189 PMID: 15717019; PubMed Central PMCID: PMC534924.

13. Moher D, Liberati A, Tetzlaff J, Altman DG, Group P. Preferred reporting items for systematic reviews and meta-analyses: the PRISMA statement. PLoS medicine. 2009; 6(7):e1000097. doi: 10.1371/journal.pmed.1000097 PMID: 19621072; PubMed Central PMCID: PMC2707599.

14. Brooks BR, Miller RG, Swash M, Munsat TL, World Federation of Neurology Research Group on Motor Neuron D. El Escorial revisited: revised criteria for the diagnosis of amyotrophic lateral sclerosis. Amyotrophic lateral sclerosis and other motor neuron disorders: official publication of the World Federation of Neurology, Research Group on Motor Neuron Diseases. 2000; 1(5):293–9. doi: 10.1080/1468020030079536 PMID: 11464847.

15. de Carvalho M, Dengler R, Eisen A, England JD, Kaji R, Kimura J, et al. Electrodiagnostic criteria for diagnosis of ALS. Clinical neurophysiology: official journal of the International Federation of Clinical Neurophysiology. 2008; 119(3):497–503. doi: 10.1016/j.clinph.2007.09.143 PMID: 18164224.

16. Whiting PF, Rutjes AWS, Westwood ME, Mallett S, Deeks JJ, Reitsma JB, et al. QUADAS-2: A Revised Tool for the Quality Assessment of Diagnostic Accuracy Studies. Annals of Internal medicine. 2011; 155(8):529–36. doi: 10.7326/0003-4819-155-8-201110180-00009 PMID: 22007046.

17. Wan X, Wang W, Liu J, Tong T. Estimating the sample mean and standard deviation from the sample size, median, range and/or interquartile range. BMC medical research methodology. 2014; 14:135. doi: 10.1186/1471-2288-14-135 PMID: 25524443; PubMed Central PMCID: PMC4383202.

18. Hozo SP, Djulbegovic B, Hozo I. Estimating the mean and variance from the median, range, and the size of a sample. BMC medical research methodology. 2005; 5:13. doi: 10.1186/1471-2288-5-13 PMID: 15840177; PubMed Central PMCID: PMC1097734.
19. Higgins JPT, Green S, Cochrane C. Cochrane handbook for systematic reviews of interventions. Chichester, England; Hoboken, NJ: Wiley-Blackwell; 2008.

20. Breitenechle J, Petzold A, Sussman SD, Ludolph AC, Tumani H. Axonal damage markers in cerebrospinal fluid are increased in ALS. Neurology. 2006; 66(6):852–6. doi: 10.1212/01.wnl.0000203120.85850.54 PMID: 16567701.

21. Steinacker P, Fang L, Kuhle J, Petzold A, Tumani H, Ludolph AC, et al. Soluble beta-amyloid precursor protein is related to disease progression in amyotrophic lateral sclerosis. PloS one. 2011; 6(8):e23600. Epub 2011/08/23. doi: 10.1371/journal.pone.0023600 PMID: 2185182; PubMed Central PMCID: PMCPMC3156148.

22. Kuhle J, Regeniter A, Leppert D, Meiling M, Kappos L, Lindberg RL, et al. A highly sensitive electrochemiluminescence immunoassay for the neurofilament heavy chain protein. Journal of neuroimmunology. 2010; 220(1–2):114–9. doi: 10.1016/j.jneuroim.2010.01.004 PMID: 21198418.

23. Weydt P, Oeckl P, Huss A, Muller K, Volk AE, Kuhle J, et al. Neurofilament levels as biomarkers in asymptomatic and symptomatic familial amyotrophic lateral sclerosis. Annals of neurology. 2016; 79(1):152–8. Epub 2015/11/04. doi: 10.1002/ana.24552 PMID: 26528863.

24. Steinacker P, Feneberg E, Weishaupt J, Breitenechle J, Tumani H, Andersen PM, et al. Neurofilaments in the diagnosis of motoneuron diseases: a prospective study on 455 patients. Journal of neurology, neurosurgery, and psychiatry. 2016; 87(1):12–20. doi: 10.1136/jnnp-2015-311387 PMID: 26296871.

25. Mendonca DM, Martins SC, Higashi R, Muscara MN, Neto VM, Chimelli L, et al. Neurofilament heavy subunit in cerebrospinal fluid: a biomarker of amyotrophic lateral sclerosis? Amyotrophic lateral sclerosis: official publication of the World Federation of Neurology Research Group on Motor Neuron Diseases. 2011; 12(2):144–7. doi: 10.3109/17482968.2010.542002 PMID: 21198418.

26. Reijn TS, Abdo WF, Schelhaas HJ, Verbeek MM. CSF neurofilament protein analysis in the differential diagnosis of ALS. Journal of neurology. 2009; 256(4):615–9. doi: 10.1007/s00415-009-0131-z PMID: 19296046.

27. Goncalves M, Tillack L, de Carvalho M, Pinto S, Conradt HS, Costa J. Phosphorylated neurofilament heavy chain and N-glycomics from the cerebrospinal fluid in amyotrophic lateral sclerosis. Clinica chimica acta; international journal of clinical chemistry. 2015; 438:342–9. doi: 10.1016/j.cca.2014.09.011 PMID: 25261856.

28. Lehnert S, Costa J, de Carvalho M, Kirby J, Kuzma-Kozakiewicz M, Morelli C, et al. Multicentre quality control evaluation of different biomarker candidates for amyotrophic lateral sclerosis. Amyotrophic lateral sclerosis & frontotemporal degeneration. 2014; 15(5–6):344–50. doi: 10.3109/21678421.2014.884592 PMID: 24575871.

29. Ganesalingam J, An J, Shaw CE, Shaw G, Lacomis D, Bowser R. Combination of phosphorylated neurofilament heavy chain and complement C3 as CSF biomarkers for ALS. Journal of the neurological sciences. 2015; 353(1–2):122–9. doi: 10.1016/j.jns.2015.04.032 PMID: 25958264.

30. McCombe PA, Pfluger C, Singh P, Lim CY, Airey C, Henderson RD. Serial measurements of phosphorylated neurofilament-heavy in the serum of subjects with amyotrophic lateral sclerosis. Journal of the neurological sciences. 2015; 353(1–2):122–9. doi: 10.1016/j.jns.2015.04.032 PMID: 25058264.

31. Boylan KB, Glass JD, Crook JE, Yang C, Desaro P, et al. Phosphorylated neurofilament heavy subunit (pNF-H) in peripheral blood: a potential prognostic biomarker in amyotrophic lateral sclerosis. Journal of neurology, neurosurgery, and psychiatry. 2013; 84(4):467–72. doi: 10.1136/jnnp-2012-303768 PMID: 23117489.

32. Zetterberg H, Jacobsson J, Rosen G, Blennow K, Andersson PM, et al. Increased phosphorylation of neurofilament light chain in cerebrospinal fluid of patients with amyotrophic lateral sclerosis and other neurodegenerative diseases. Journal of neurochemistry. 1996; 67(5):2013–8. doi: 10.1046/j.1471-4159.1996.67052013.x PMID: 8863508.

33. Zetterberg H, Jacobsson J, Rosengren L, Blennow K, Andersson PM. Cerebrospinal fluid phosphorylated neurofilament light levels in amyotrophic lateral sclerosis: impact of SOD1 genotype. European journal of neurology. 2007; 14(12):1329–33. Epub 2007/10/02. doi: 10.1111/j.1468-1331.2007.01972.x PMID: 17903209.
36. Lu CH, Macdonald-Wallis C, Gray E, Pearce N, Petzold A, Norgren N, et al. Neurofilament light chain: A prognostic biomarker in amyotrophic lateral sclerosis. Neurology. 2015; 84(22):2247–57. doi: 10.1212/WNL.0000000000001642 PMID: 25934855; PubMed Central PMCID: PMC4456658.

37. Tortelli R, Ruggieri M, Cortese R, D’Errico E, Capozzo R, Leo A, et al. Elevated cerebrospinal fluid neurofilament light levels in patients with amyotrophic lateral sclerosis: a possible marker of disease severity and progression. European journal of neurology. 2012; 19(12):1561–7. doi: 10.1111/j.1468-1331.2012.03777.x PMID: 22680408.

38. Cedarbaum JM, Stambler N, Malta E, Fuller C, Hill D, Thurmond B, et al. The ALSFRS-R: a revised ALS functional rating scale that incorporates assessments of respiratory function. BDNF ALS Study Group (Phase III). Journal of the neurological sciences. 1999; 169(1–2):13–21. Epub 1999/12/14. doi: 10.1016/s0022-510x(99)00210-5 PMID: 10540002.

39. Tortelli R, Copetti M, Ruggieri M, Cortese R, Capozzo R, Leo A, et al. Cerebrospinal fluid neurofilament light chain: marker of progression to generalized amyotrophic lateral sclerosis. European journal of neurology. 2015; 22(1):215–8. doi: 10.1111/ene.12421 PMID: 24750431.

40. Rysz-Gorzynska M, Giuba-Brzozka A, Sahebkar A, Serban MC, Mikhailidis DP, Ursoniu S, et al. Efficacy of Statin Therapy in Pulmonary Arterial Hypertension: A Systematic Review and Meta-Analysis. Scientific reports. 2016; 6:30060. Epub 2016/07/23. doi: 10.1038/srep30060 PMID: 27444125; PubMed Central PMCID: PMCPMC4957081.

41. Su Z, Zhu L, Wu J, Zhao R, Ji HL. Systematic review and meta-analysis of nasal potential difference in hypoxia-induced lung injury. Scientific reports. 2016; 6:30780. Epub 2016/08/05. doi: 10.1038/srep30780 PMID: 27489696; PubMed Central PMCID: PMCPMC4973283.

42. Pijnenburg YA, Verwey NA, van der Flier WM, Scheltens P, Teunissen CE. Discriminative and prognostic potential of cerebrospinal fluid phosphoTau/tau ratio and neurofilaments for frontotemporal dementia subtypes. Alzheimer's & dementia (Amsterdam, Netherlands). 2015; 1(4):505–12. Epub 2015/05/31. doi: 10.1016/j.dadm.2015.11.001 PMID: 27239528; PubMed Central PMCID: PMCPMC4879490.

43. Siedler DG, Chuah MI, Kirkcaldie MT, Vickers JC, King AE. Diffuse axonal injury in brain trauma: insights from alterations in neurofilaments. Frontiers in cellular neuroscience. 2014; 8:429. Epub 2015/01/08. doi: 10.3389/fncel.2014.00429 PMID: 25565963; PubMed Central PMCID: PMCPMC4269130.

44. Gresle MM, Butzkueven H, Shaw G. Neurofilament proteins as body fluid biomarkers of neurodegeneration in multiple sclerosis. Multiple sclerosis international. 2011; 2011:315406. Epub 2011/11/19. doi: 10.1155/2011/315406 PMID: 22096635; PubMed Central PMCID: PMCPMC3195785.

45. Petzold A. The prognostic value of CSF neurofilaments in multiple sclerosis at 15-year follow-up. Journal of neurology, neurosurgery, and psychiatry. 2015; 86(12):1388–90. Epub 2015/01/27. doi: 10.1136/jnnp-2014-309827 PMID: 25616604.

46. Cai JY, Lu C, Chen MH, Ba HJ, Chen XD, Lin JH, et al. Predictive value of phosphorylated axonal neurofilament subunit H for clinical outcome in patients with acute intracerebral hemorrhage. Clinica chimica acta; international journal of clinical chemistry. 2013; 424:182–6. Epub 2013/07/03. doi: 10.1016/j.cca.2013.06.019 PMID: 23810564.

47. Nylen K, Csajbok LZ, Ost M, Rashid A, Karlsson JE, Blennow K, et al. CSF -neurofilament correlates with outcome after aneurysmal subarachnoid hemorrhage. Neuroscience letters. 2006; 404(1–2):132–6. Epub 2006/06/30. doi: 10.1016/j.neulet.2006.05.029 PMID: 16806706.

48. Lewis SB, Wolper RA, Miraila L, Yang C, Shaw G. Detection of phosphorylated NF-H in the cerebrospinal fluid and blood of aneurysmal subarachnoid hemorrhage patients. Journal of cerebral blood flow and metabolism: official journal of the International Society of Cerebral Blood Flow and Metabolism. 2008; 28(6):1261–71. Epub 2008/03/06. doi: 10.1038/jcbfm.2008.12 PMID: 18197931.

49. Petzold A, Keir G, Green AJ, Giovannoni G, Thompson EJ. A specific ELISA for measuring neurofilament heavy chain phosphoforms. Journal of immunological methods. 2003; 278(1–2):179–90. Epub 2003/09/06. doi: 10.1016/s0022-1759(03)00189-3 PMID: 12957406.

50. Neurofilament light chain: a prognostic biomarker in amyotrophic lateral sclerosis. Neurology. 2015; 85(10):921. doi: 10.1212/WNL.0000000000001986 PMID: 26346831.

51. Shaw G, Yang C, Ellis R, Anderson K, Parker Mickle J, Scheff S, et al. Hyperphosphorylated neurofilament NF-H is a serum biomarker of axonal injury. Biochemical and biophysical research communications. 2005; 336(4):1268–77. Epub 2005/09/24. doi: 10.1016/j.bbrc.2005.08.252 PMID: 16176808.

52. Rosengren LE, Wikkelso C, Hagberg L. A sensitive ELISA for glial fibrillary acidic protein: application in CSF of adults. Journal of neuroscience methods. 1994; 51(2):197–204. doi: 10.1016/0165-0270(94)90011-6 PMID: 8051950.

53. Lu CH, Kalmar B, Malaspina A, Greensmith L, Petzold A. A method to solubilise protein aggregates for immunobassy quantification which overcomes the neurofilament "hook" effect. Journal of neuroscience methods. 2011; 195(2):143–50. doi: 10.1016/j.jneurneuro.2010.11.026 PMID: 21134399.