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Serum neurofilament light chain levels as a marker of upper motor neuron degeneration in patients with Amyotrophic Lateral Sclerosis

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Aims: Amyotrophic lateral sclerosis (ALS) is the most common motor neuron degeneration disease with a diagnostic delay of about 1 year after symptoms onset. In ALS, blood neurofilament light chain (NfL) levels are elevated, but it is not entirely clear what drives this increase and what the diagnostic performance of serum NfL is in terms of predictive values and likelihood ratios. The aims of this study were to further explore the prognostic and diagnostic performances of serum NfL to discriminate between patients with ALS and ALS mimics, and to investigate the relationship between serum NfL with motor neuron degeneration. **Methods**: The diagnostic performances of serum NfL were based on a cohort of 149 serum samples of patients with ALS, 19 serum samples of patients with a disease mimicking ALS and 82 serum samples of disease control patients. The serum NfL levels were correlated with the number of regions (thoracic, bulbar, upper limb and lower limb) displaying upper and/or lower motor neuron degeneration. The prognostic performances of serum NfL were investigated based on a Cox regression analysis. **Results**: The associated predictive values and likelihood ratio to discriminate patients with ALS and ALS mimics were established. Serum NfL was associated with motor neuron degeneration driven by upper motor neuron (UMN) degeneration and was independently associated with survival in patients with ALS. **Conclusions**: Altogether, these findings suggest that elevated serum NfL levels in ALS are driven by UMN degeneration and the disease progression rate and are independently associated with survival at time of diagnosis.

Keywords: amyotrophic lateral sclerosis, neurofilament light chain, motor neuron degeneration

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Introduction

Lou Gehrig's disease or amyotrophic lateral sclerosis (ALS) is characterized by a degeneration of upper motor neurons (UMN) in the motor cortex and lower motor neurons (LMN) in the brain stem and spinal cord, resulting in a progressive muscle weakness [1,2]. The diagnosis of ALS is currently supported by a clinical examination and an electromyography (EMG), requiring the degeneration of both UMN and LMN, the evidence of disease spreading to other regions and the exclusion of another aetiology [1,3-5]. The diagnosis of ALS is made on average 10-12 months after onset of first symptoms, when patients are already well advanced in their ALS disease trajectory [6]. Thus, an earlier diagnosis of ALS is desirable for early initiation of therapy and for inclusion of patients in clinical trials. Diagnostic biomarkers for ALS in early disease stages might shorten the diagnostic delay. We and others showed that the cerebrospinal fluid (CSF) levels of neurofilament light chain (NfL) and phosphorylated neurofilament heavy chain (pNfH) in patients with ALS are significantly higher in comparison with the CSF levels of ALS mimics, displaying good predictive values towards the diagnosis at the early or conventional diagnostic stage [6-9]. Furthermore, we showed that CSF NfL and pNfH levels correlated with the extent of UMN and LMN degeneration [6,7]. In line with these findings, we and others described that the serum concentrations of pNfH in patients with ALS were higher than in patients with an ALS mimic at the conventional diagnostic stage [7]. A similar change for NfL in blood of patients with ALS has recently been reported [8]. Nevertheless, we did not observe any correlation between pNfH in blood with UMN or LMN loss in ALS, matching our finding of pNfH in blood being less performant than pNfH in CSF to discriminate between ALS and ALS mimics [7]. It is unclear yet to what extent NfL in blood of patients with ALS correlates with UMN and/or LMN degeneration, and if the raised levels of NfL in blood are useful to discriminate between patients with ALS and ALS mimics in terms of predictive values and likelihood ratios.

The objective of this study was to evaluate the diagnostic performances of NfL in blood and whether NfL levels in blood correlate with the hallmark features of ALS, such as the extent of UMN and LMN degeneration, the disease progression rate and survival.

Materials and methods

We sampled peripheral blood in 5-ml BD Vacutainer[®] serum separation tubes (catalogue #:367955; Becton Dickinson and Company, Franklin Lakes, NJ, USA) of patients referred to the University hospital of Leuven neuromuscular reference centre (NMRC) with complaints suspicious of ALS (including muscle weakness or speech/swallowing/breathing difficulties), and hence with a differential diagnosis including ALS, between April 2014 and August 2017. All participants signed an informed consent. Patients were followed up and eventually received the diagnosis of (i) ALS (n = 149)(indicating 56% of patients diagnosed at the NMRC at time of study inclusion) or of (ii) an ALS mimic (n = 19), including patients presenting with symptoms reminiscent of ALS but for whom ALS was ruled out upon follow-up by an expert neurologist at the NMRC (PVD, KGC). The diagnosis of ALS was based on the Awaji and revised El Escorial criteria [3,5,10] (Table 1). Among those patients, 15 were C9orf72 mutation carriers and 15 were diagnosed with frontotemporal dementia (FTD). In addition, 11 patients were diagnosed with primary lateral sclerosis (PLS) and six patients were diagnosed with progressive muscular atrophy (PMA) (Table 2). Moreover, we also included patients whose blood was sampled at the diagnostic phase and who eventually received a diagnosis of Guillain-Barré syndrome (GBS), chronic inflammatory demyelinating polyneuropathy (CIDP) or hereditary spastic paraplegia (HSP) by an expert neurologist at the NMRC.

Symptom duration in patients with ALS was defined as the time difference between estimated first symptom (muscle weakness or speech/swallowing/ onset breathing difficulties) and blood sampling. The disease progression rate, expressed in points/month, was calculated as the difference between the maximum score of 48 and the ALS functional rating scale revised (ALSFRS-R) score the closest to blood sampling, divided by the symptom duration (in months) at date of ALSFRS-R score. Hundred and twenty-four patients with ALS with a delay between blood sampling and ALSFRS-R within 3 months scoring (median: 0.1 month; range: 0-3 months) were included in the correlation analysis between the serum NfL levels and the disease progression rate. Patients with a disease

Table 1. Clinical characteristics of the patients with ALS

	Table	1. ((Continued)	
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	Median (IQR; range)		Median (IQR; range)
<i>n</i> (m/f)	149 (94/55)	Mean survival after sampling in mor	ths (95% CI)
Age at blood sampling (years)	62 (15.4; 29–93)	Low tertile NfL (<115 pg/ml)	40.1 (37.36-42.8)
Age at estimated symptom onset	60 (14.8; 25-84)	Mid-tertile NfL (115 pg/	23.6 (19.78-27.44)
(years)†		$ml \le x \le 235 \text{ pg/ml}$	
Site of onset (bulbar/spinal/ generalized/unknown)†	34/105/4/6	High tertile NfL (>235 pg/ml)	17.3 (13.41–21.1)
Diagnostic delay (months)‡	10.6 (8.6; 3-107)	95% CI, 95th percentile confidence	e interval: ALS, amyotrophic
Symptoms duration at blood sampling (months)†	13.7 (12.5; 2–240)	lateral sclerosis; ALSFRS-R, ALS fu score; IQR, interquartile range; Nfl	nctional rating scale revised
ALSFRS-R score closest to date of	37 (12.0; 11-47)	FVC, forced vital capacity; LMN, low	
blood sampling§		motor neuron	er motor neuron, owny, upper
Disease progression rate (points/ month)¶	0.672 (0.75; 0.058–5.00)	†Six patients with unknown estimat site of onset.	ed date of symptoms onset or
FVC (%)††	92 (32.5; 25–142)	Diagnostic delay could not be calcu	lated in six natients who had
Serum NfL (pg/ml)	179 (164.6; 0.3-1141)	an unknown date of symptoms on	-
		unknown date of diagnosis.	iset and one patient nat an
Serum NfL (pg/ml)		§ALSFRS-R was not performed in sev	ven patients.
1 Region with UMN	106 (145; 25–370)	The second secon	
neurodegeneration $(n = 14)$		R not performed in seven patient	
2 Regions with UMN	188 (136; 12–768)	unknown date of symptoms onset.	L
neurodegeneration $(n = 31)$	251 (155, (2, 1141)	††FVC was unavailable for 17 patient	its.
3 Regions with UMN	251 (155; 62–1141)		
neurodegeneration $(n = 55)$ Serum NfL (pg/ml)			
1 Region with LMN	191 (169.2; 12-781)	the Awaji and revised El Escori	al criteria. UMN involve-
neurodegeneration $(n = 33)$	191 (109.2, 12-781)	ment in the thoracic region is	not assessed routinely at
2 Regions with LMN	221 (281.7; 41-694)	our NMRC. The number of reg	
neurodegeneration ($n = 21$)	221 (201.7, 11 091)	-	-
3 Regions with LMN	225 (161.4; 38–1141)	ation was assessed clinically a	nd by EMG in the bulbar
neurodegeneration $(n = 46)$	(,)	and thoracic regions and in th	e upper and lower limbs
Serum NfL (pg/ml)		according to the revised El Eso	corial and Awaji criteria.
1 Region with LMN and UMN	119 (175.9; 12-781)	One hundred patients with AL	· · · · · · · · · · · · · · · · · · ·
neurodegeneration $(n = 34)$		-	
2 Regions with LMN and UMN	232 (168.9; 41-768)	of 3 months between EMG or	
neurodegeneration $(n = 47)$		blood sampling were includ	led in the correlation
3 Regions with LMN and UMN	268 (192.6; 110–1141)	between serum NfL and the nu	umber of regions affected
neurodegeneration $(n = 19)$		by neurodegeneration. The for	-
Serum NfL (pg/ml)			
Slow disease progression rate	112 (81.4; 0.3–295)	score the closest to blood ta	
(n = 33)		spirometry in 132 patients w	ith ALS. A total of 113
Intermediate disease progression	188 (152.9; 12–554)	patients with ALS having a p	ossible minimum 1-year
rate $(n = 61)$		survival follow-up after blood	-
Fast disease progression rate $(x - 20)$	255 (215.1; 62–1141)	_	
(n = 30)		ment were included in the su	

progression rate below the 25th percentile (<0.372) were considered as slow progressors, patients with a disease progression rate above the 75th percentile (>1.121) were considered as fast progressors and patients with a disease progression rate between or equal to 0.372 and 1.121 were identified as intermediate progressors. The number of regions affected by UMN degeneration was assessed clinically in the bulbar, upper limb and lower limb regions, according to

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sampling between December 2016 and 2017). Patients with ALS with missing data or a delay between blood sampling and ALSRFS-R, EMG or FVC assessment greater than 3 months were excluded from the multivariate Cox regression analysis. The CSF levels of NfL were assessed with an enzyme-linked immunosorbent kit (UmanDiagnostics, Umea, Sweden) as previously published [6]. The serum NfL levels were quantified with an electrochemiluminescent assay as previously described [11]. Patients were then stratified in tertiles according to their serum NfL levels: (i) lowest tertile

Cohort group	Diagnosis	<i>n</i> (m/f)	Age (years)	Serum NfL (pg/ml)
ALS mimics	Chronic radiculopathy/polyradiculopathy	7 (6/1)	54 (6; 38–71)	29 (71; 21–152)
	Brachial plexopathy	2 (2/0)	44 (7; 37-51)	537 (516; 20-1053)
	Myasthenia Gravis	2(1/1)	73 (1; 72–74)	80 (13; 67–93)
	Cervical stenosis	1(1/0)	63	61
	Fewdon syndrome [12]	1(0/1)	25	21
	Hirayama syndrome	1(1/0)	23	6
	IBM	1(1/0)	80	74
	Suspected AMAN	1(1/0)	71	42
	MMN	1(0/1)	28	15
	Motor neuropathy	1(0/1)	57	20
	Brainstem vascular lesions	1(0/1)	72	19
Disease controls	GBS	48 (26/22)	53 (30.8; 7-86)	123 (280.6; 22-9045)
	CIDP	20 (17/3)	59 (12.5; 31-75)	101 (117.5; 29-2863)
	HSP	14 (8/6)	46 (22.2; 25-84)	37 (32.3; 8–639)

Table 2. The serum levels of NfL in ALS mimics and controls

Data expressed as the median (IQR; range).

ALS, amyotrophic lateral sclerosis; AMAN, acute motor axonal neuropathy; CIDP, chronic inflammatory demyelinating polyneuropathy; GBS, Guillain–Barré syndrome; HSP, hereditary spastic paraplegia; IBM, inclusion body myositis; IQR, interquartile range; MMN, multifocal motor neuropathy; Nfl, neurofilament light chain.

serum NfL<115 pg/ml; (ii) mid-tertile serum between or equal to 115 pg/ml and 235 pg/ml and (iii) highest tertile serum NfL >235 pg/ml. Longitudinal serum samples of 16 patients with ALS were available (mean follow-up = 15.5 months; range = 3.8-32.0 months). The different time points for a given patient were assessed on the same run to avoid analytical deviation.

Statistical analysis

Normality of the data and residuals were assessed with the D'Agostino & Pearson's test. Correlations between the serum levels of NfL with the CSF levels NfL, symptom duration at blood sampling or the disease progression rate were assessed with the Spearman's rank-order correlation, with a spearman's coefficient correlation (Spearman's p) at a significance level of 5%. Comparisons between the serum levels of NfL with the disease progression rate, the extent of motor neuron degeneration in patients with ALS or with the different diagnosis were assessed with the Kruskal-Wallis test corrected for multiple comparisons with the post hoc Dunn's test, at a significance level of 5%. Logarithmic transformation of the serum levels of NfL limited the right skewed distribution of the serum NfL levels, allowing the analysis of covariance. Optimal receiver operating characteristics (ROC) curves were generated to calculate the area under the curve (AUC) and the highest Youden's index was used to estimate the optimal cut-offs for which

sensitivity, specificity, positive and negative likelihood ratio (+LR and -LR, respectively) and positive and negative predictive values (PPV and NPV, respectively) were estimated. For the comparison between slow and fast progressors the predictive values were calculated based on the prevalence obtained by Gomeni & Fava [13]. For the comparison between ALS and ALS mimics, the predictive values were calculated based on the prevalence of ALS and ALS mimics in our data set. The effect of age and gender on the AUC was assessed in a multivariate model using logistic regression and an adjusted AUC was reported, based on the predictive probabilities, in the case of significant differences with the AUC obtained with univariate model using the serum NfL values. The Cox proportional-hazard regression analysis was performed via the backward method with a P-value threshold of 0.157, with the time to event being the time from blood sampling to death [14]. Based on a recent study of prognostic factors in ALS [15], the following parameters were included as survival variables into the multivariate Cox regression: the tertiles of the serum levels of NfL, the disease progression rate stratified as slow, intermediate and fast, the number of regions with UMN and LMN degeneration, age at blood sampling, the FVC, gender, the diagnostic delay, the C9orf72 status, FTD status and the site of onset. The exclusion criteria for the Cox regression were a follow-up shorter than a year (sampling between December 2016 and 2017) and, for the multivariate Cox regression, a time difference among the date of ALSFRS-R, EMG and FVC with the date of blood sampling greater than 3 months. The Kaplan-Meier analysis was performed only on the tertiles of the serum levels of NfL and on the full data set (n = 149). Statistical testing was performed with GraphPad Prism (v.7.01; GraphPad Software Inc, San Diego, CA, USA), Kaplan-Meier, ROC curve, logistic regression and analysis of covariance were performed in MedCalc (v.17.9; MedCalc Software bvba, Ostend, Belgium) and univariate and multivariate Cox proportional-hazard regressions were performed in SPSS version 25 (SPSS, Chicago. IL. USA). *P < 0.05.**P < 0.01.***P < 0.001 and ****P < 0.0001 were considered significant.

Results

The demographics of the patients with ALS and the different control cohorts are listed in Tables 1 and 2 respectively. Serum NfL strongly correlated with the CSF NfL as assessed on 97 paired samples of patients with ALS (Spearman's $\rho = 0.79$, P < 0.0001) (Figure 1a). The median time differences between blood sampling and lumbar puncture was less than 24 h (95 pairs of 97 were sampled on the same day).

Serum NfL are correlated with symptom duration and the disease progression rate

In serum of patients with ALS, NfL correlated negatively with symptom duration at time of blood sampling (Spearman's $\rho = -0.39$, P < 0.0001) (Figure 1b). The median symptom duration to blood sampling was 13.7 months [interquartile range (IQR) = 12.5 months; range = 2– 240 months]. Fast progressors and slow progressors displayed a short and long symptom duration at blood sampling respectively (median symptom duration fast progressors = 7.3 months; IQR = 4.7 months; range = 3.4–31.0 months, median symptom duration slow

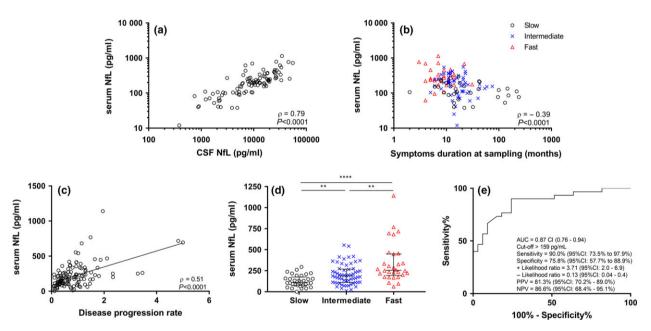


Figure 1. Correlations among the serum levels of NfL, symptom duration and the disease progression rate. (a) Correlation between the serum and CSF levels of NfL in 97 paired ALS samples (Spearman's $\rho = 0.79$; P < 0.0001). (b) Negative correlation between the symptom duration at blood sampling and the serum NfL levels of patients with ALS stratified according to their disease progression in slow (black circles), intermediate (blue crosses) or fast progressors (red triangles) (Spearman's $\rho = -0.39$; P < 0.0001). (c) Positive correlation between the disease progression rate and the serum levels of NfL (Spearman's $\rho = 0.51$; P < 0.0001). (d) Serum NfL levels in patients with ALS stratified according to their disease progression rate in slow (black circles), intermediate (blue crosses) and fast progressors (red triangles). Kruskal–Wallis test corrected for multiple comparison with the Dunn's *post hoc* test. (e) ROC curve to discriminate patients with ALS displaying slow and fast disease progressions. **P < 0.01 and ****P < 0.0001 were considered as statistically significant. + Likelihood ratio, positive likelihood ratio; –Likelihood ratio, negative likelihood ratio; ALS, amyotrophic lateral sclerosis; AUC, area under the curve; CSF, cerebrospinal fluid; NfL, neurofilament light chain; ROC curve, receiver operating characteristics curve; PPV, positive predictive value; NPV, negative predictive value.

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progressors = 17.3 months; IQR = 31.2; range = 2.4-239.6 months), suggesting that the enrichment of fast disease progressors within the cohort of patients with a short disease duration from blood sampling drives the inverse correlation between NfL levels and symptom duration (Figure 1b). Serum NfL levels in patients with ALS correlated with the disease progression rate (Spearman's $\rho = 0.51$, P < 0.0001) (Figure 1c). The median time difference between date of ALSFRS-R scoring and date of blood sampling was 2.5 days (IQR = 25.3 days; ranged from 0 to 90 days). Patients with ALS with a fast disease progression rate (n = 30) displayed significantly higher levels of serum NfL than patients with a slow (n = 33) (median = 255 pg/ ml vs. 112 pg/ml, respectively, P < 0.0001) or intermediate disease progression rate (n = 61) (255 pg/ml vs. 188 pg/ml, P = 0.0097) (Figure 1d and Table 1). The serum levels of NfL were also significantly higher in patients with ALS displaying an intermediate progression rate as compared with slow progressors (188 pg/ml vs. 112 pg/ml. respectively, P = 0.0059) (Figure 1d and Table 1). The AUC to discriminate patients between fast and slow disease progressors was 0.87 [95th percentile confidence interval (95% CI): 0.76-0.94; P < 0.0001 at an optimal cut-off of 159 pg/ml. At this cut-off, the diagnostic accuracy of the serum levels of NfL had a sensitivity of 90.0% (95% CI: 73.5–97.9) and a specificity of 75.8% (95% CI: 57.7–88.9) with a +LR of 3.71 (95% CI: 2.0-6.9), a -LR of 0.13 (95% CI: 0.04-0.4), a PPV of 81.3% (95% CI: 70.2-89.0) and a NPV of 86.6% (95% CI: 68.4–95.1) (Figure 1e).

Serum NfL as a marker of UMN degeneration in ALS

Serum NfL in ALS was increased as a function of the number of regions affected by both UMN and LMN degeneration (1 region vs. 2 regions: 119 pg/ml vs. 232 pg/ml, P = 0.027; 1 region vs. 3 regions: 119 pg/ml vs. 268 pg/ml, P = 0.0021) (Figure 2a). In particular, the levels of serum NfL were significantly increased between patients with 1 or 3 regions affected by UMN degeneration (1 region vs. 3 regions: 106 pg/ml vs. 251 pg/ml, P = 0.0081) (Figure 2b). However, the serum NfL levels were relatively stable with increasing number of regions affected by LMN degeneration as assessed with EMG, applying the Awaji criteria (1 vs. 2 regions: 191 pg/ml vs. 225 pg/ml) and revised El Escorial criteria (Figures 2c and S1,

respectively). When including these parameters in an analysis of covariance together with the disease progression rate, the age at blood sampling, gender, the site of onset and the symptom duration at blood sampling as covariates, serum NfL was still elevated in patients with ALS displaying 3 regions with UMN degeneration compared with patients with ALS with only one region with UMN degeneration (adjusted means log10 serum NfL; 1 vs. 3 regions: 2.063 vs. 2.348, P = 0.0062). Based on a heat map analysis, the serum levels of NfL increase together with the number of regions with UMN degeneration and the disease progression rate (Figure 3a). The levels of serum NfL increase with the disease progression rate, with the former less dependent on the number of regions affected with LMN degeneration (Figure 3b). The serum levels of NfL correlated with the disease progression rate when 1 or 2 regions are affected by UMN and LMN degeneration but not when 3 regions are affected by both UMN and LMN degeneration (Figure 3c).

The serum levels of NfL are elevated in patients with ALS

The serum NfL levels were about 6 times higher in serum of patients with ALS than in patients with an ALS mimic (median serum NfL levels 179 pg/ml vs. 29 pg/ml; P < 0.0001) (Figure 4a). The associated AUC was 0.85 (95% CI: 0.79–0.90; P < 0.0001), generating an optimal cut-off of 93 pg/ml resulting in a sensitivity of 79.2% (95% CI: 71.8-85.4) and a specificity of 84.2% (95% CI: 60.4-96.6), with a +LR of 5.02 (95% CI: 1.8-14.2), a -LR of 0.25 (95% CI: 0.2-0.4), a PPV of 97.5% (95% CI: 93.3-99.1) and a NPV of 34.0% (95% CI: 26.3-42.7) (Figure 4b). At a fixed sensitivity of 95%, the serum levels of NfL had a specificity of 52.6% (95% CI: 5.3-79.0) to discriminate patients with ALS from ALS mimic patients. At a fixed specificity of 95% the serum levels of NfL had a sensitivity of 0.67% (95% CI: 0.0-61.1). As ALS is commonly associated in the differential diagnosis of patients with a cervical spondylotic myelopathy, we retrospectively retrieved an additional set of 16 serum samples of patients with a cervical stenosis/myelopathy. These patients, however, did not present as a genuine ALS mimic at our NMRC. The vast majority of these patients suffered

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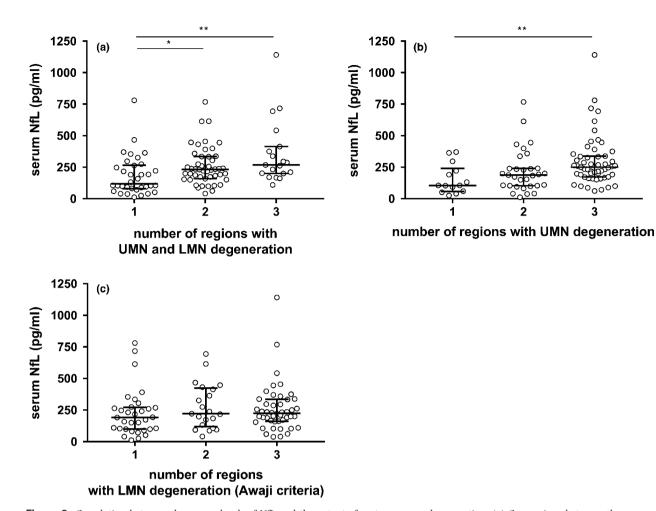


Figure 2. Correlation between the serum levels of NfL and the extent of motor neuron degeneration. (a) Comparison between the serum NfL levels in patients with ALS according to the number of regions with both UMN and LMN degeneration. (b) Comparison between the serum NfL levels in patients with ALS according to the number of regions with UMN degeneration only. (c) Comparison between the serum NfL levels in patients with ALS according to the number of regions with LMN degeneration only according to the Awaji criteria. Kruskal-Wallis corrected for multiple comparison with the Dunn's post hoc test. ALS, amyotrophic lateral sclerosis; LMN, lower motor neuron; NfL, neurofilament light chain; UMN, upper motor neuron.

from a moderate-to-severe cervical myelopathy, with among them high levels of serum NfL (Figure S2). However, the patient with cervical stenosis mimicking ALS presented with a cervical herniated disc (C5–C6) without a clear documented myelopathy, having low levels of serum NfL (Table 2).

The serum levels of NfL were significantly higher in patients with ALS as compared with patients diagnosed with PLS but not with patients with PMA (median serum NfL levels 179 pg/ml vs. 34 pg/ml (P = 0.0008) and 179 pg/ml vs. 71 pg/ml, respectively) (Figure 4a). However, variable levels of serum NfL were observed (range = 13-1185 pg/ml) in patients with PMA. The serum NfL levels correlated with the disease progression rate in patients with PMA but not in patients with PLS (Spearman's $\rho = 0.94$, P = 0.017;Spearman's $\rho = 0.32$, P = 0.56) (Table S1 and Figure S3). The serum levels of NfL in patients with ALS were significantly higher as compared to HSP, but not to GBS or CIDP (median serum NfL levels 179 pg/ml vs. 37 pg/ ml, P = 0.0002; 179 pg/ml vs. 123 pg/ml and vs. 101 pg/ml, respectively) (Figure 4a), which confirms that serum NfL is not specific to ALS. Indeed, ROC curve analysis revealed an AUC of 0.58 (95% CI: 0.51-0.64, P = 0.097), at an optimal cut-off of 139 pg/ml, between patients with ALS and disease

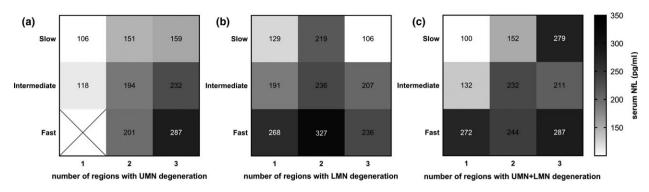


Figure 3. Heat map of the correlations among the disease progression rate, the extent of motor neuron degeneration and the serum levels of NfL. (a) Heat map among the disease progression rate, the number of regions with UMN degeneration and the serum levels of NfL. (b) Heat map among the disease progression rate, the number of regions with LMN degeneration (Awaji criteria) and the serum levels of NfL. (c) Heat map among the disease progression rate, the number of regions displaying both UMN and LMN degeneration and the serum NfL levels. Cell labels are expressed in pg/ml and represent the median for a given combination. The colour gradient corresponds to the median levels of serum NfL, the darker the colour gets, the higher the levels of serum NfL are. A cross ((a) fast progressor with 1 region with UMN degeneration) indicates that no patients with ALS had this combination. LMN, lower motor neuron; NfL, neurofilament light chain; UMN, upper motor neuron.

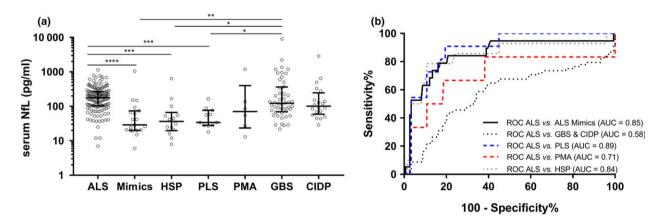


Figure 4. Serum levels of NfL in patients with ALS, ALS mimics and controls. (a) Scatter plots of the serum NfL levels in patients with ALS, disease mimics and in disease controls (median serum NfL levels; ALS vs. ALS Mimics: 179 pg/ml vs. 29 pg/ml P < 0.0001; ALS vs. PLS: 179 pg/ml vs. 34 pg/ml P = 0.0008; ALS vs. HSP: 179 pg/ml vs. 37 pg/ml P = 0.0002; GBS vs. ALS mimics: 123 pg/ml vs. 29 pg/ml vs. 29 pg/ml P = 0.0016; GBS vs. PLS: 123 pg/ml vs. 34 pg/ml P = 0.022; GBS vs. HSP: 123 pg/ml vs. 37 pg/ml P = 0.011). Kruskal–Wallis corrected for multiple comparison with the Dunn's *post hoc* test. (b) ROC curves to distinguish between patients with ALS vs. ALS mimics (solid black line; AUC = 0.85, P < 0.0001), vs. GBS and CIDP (dotted black line; AUC = 0.58, P = 0.097), vs. PLS (dashed blue line; AUC = 0.89, P < 0.0001), vs. PMA (dashed red line; AUC = 0.71, P = 0.13) and vs. HSP (dotted grey line: AUC = 0.84, P < 0.0001). *P < 0.05, **P < 0.01, ***P < 0.001 and ****P < 0.0001 were considered as statistically significant. ALS, amyotrophic lateral sclerosis; AUC, area under the curve; CIDP, chronic inflammatory demyelinating polyneuropathy; GBS, Guillain–Barré syndrome; HSP, hereditary spastic paraplegia; NfL, neurofilament light chain; PLS, primary lateral sclerosis; PMA, progressive muscular atrophy; ROC curve, receiver operating characteristic curve.

controls (GBS and CIDP) with a poor specificity of 63.2% (95% CI: 50.7–74.6) (Figure 4b). The AUC to discriminate between patients with ALS and with HSP was 0.84 (95% CI: 0.78–0.90; P < 0.0001) at an optimal cut-off of 55 pg/ml, with a sensitivity of 89.3% (95% CI: 83.1–93.7), a specificity of 78.6% (95% CI: 49.2–95.3), a +LR of 4.2 (95% CI: 1.5–11.4) and a

-LR of 0.14 (95% CI: 0.1–0.2) (Figure 4b). The AUC to discriminate patients with ALS with patients with PLS was 0.89 (95% CI: 0.83–0.93, P < 0.0001), at an optimal cut-off of 88 pg/ml, resulting to a sensitivity of 80.5% (95% CI: 73.3–86.6), a specificity of 90.9% (95% CI: 58.7–99.8), a +LR of 8.9 (95% CI: 1.4–57.5) and a –LR of 0.2 (95% CI: 0.1–0.3) (Figure 4b). The

AUC to distinguish patients with ALS and patients with PMA was 0.71 (95% CI: 0.63–0.78, P = 0.13), at an optimal cut-off of 86 pg/ml, resulting to a sensitivity of 81.2% (95% CI: 74.0–87.1), a specificity of 66.7% (95% CI: 22.3–95.7), a +LR of 2.4 (95% CI: 0.8–7.6) and a –LR of 0.3 (95% CI: 0.1–0.5) (Figure 4B).

Serum levels of NfL are stable at follow-up

In a subgroup of 16 patients with ALS, longitudinal samples were available. The serum levels of NfL remained relatively stable over time, although changes (mainly an increase) could be seen especially within the first 20 months after onset of disease, irrespective of the disease progression rate (Figure S4).

Serum NfL levels are associated with survival

Based on a Kaplan-Meier analysis, the survival curve of patients with ALS belonging to the different tertiles of serum NfL levels was significantly different (log-rank test P < 0.0001). Univariate Cox proportional-hazards regressions identified serum NfL, the disease progression rate, the number of regions with LMN degeneration, the number of regions with both UMN and LMN degeneration, the diagnostic delay, the FVC and FTD as predictors of the mortality hazard ratio (HR) in patients with ALS (Table 3). All the above-mentioned parameters were entered into a multivariate Cox model, except the number of regions with LMN degeneration to avoid circularity with the number of regions with UMN and LMN degeneration. Gender, the C9orf72 status and age were also entered into the multivariate model. On a total of 52 patients with ALS entered into the multivariate Cox regression model, 26 patients reached the end-point (death) while 26 patients did not (censored values). Among the tested covariates the serum NfL levels, the site of onset and the FVC were significantly associated with survival (Table 3). Serum NfL levels in the mid and high tertile were associated with an increased HR of 4.5 and 5.3 as compared to patients with ALS having NfL levels in the lowest tertile [HR = 4.47 (95% CI: 1.08-18.63), P = 0.04; HR = 5.34 (95% CI: 1.39–20.56), P = 0.015] (Table 3 and Figure 5). Patients with ALS with a spinal onset were associated with an increased HR of 3.7 times compared with patients with a bulbar onset [HR = 3.73 (95% CI: 1.16-12.01), P = 0.027] (Table 3). Higher score at the FVC was strongly associated with a better prognosis in patients with ALS [HR = 0.97 (95% CI: 0.95-0.99), P = 0.005] (Table 3).

Discussion

In this study, we described a strong correlation between CSF and serum NfL levels in patients with ALS in line with previous reports on among others Alzheimer's disease and multiple sclerosis [16,17]. The strong relationship between the CSF and serum levels of NfL supports the theory of a common source for NfL originating from the central nervous system (CNS) [16,18]. We also showed that serum NfL levels rise as a function of the number of regions affected by UMN degeneration. This is in agreement with our previous findings where we observed a better correlation of the CSF levels of NfL with the extent of UMN degeneration than with the extent of LMN degeneration [6]. In line with this, the serum NfL levels were lower in the majority of patients with PMA, as reported before [19]. Importantly, our findings show that the serum levels of NfL increase with the extent of UMN degeneration when correcting for the disease progression rate, the site of onset, the age at blood sampling, gender or the symptom duration at time of blood sampling.

The elevation of NfL is not specific to motor neuron degeneration, as there is evidence that the source of serum NfL also originate from cortical and spinal neuronal damages [16,18]. In a mouse model where neuronal injury was induced by cranial irradiation, increase in serum NfL was observed, independently of the blood-brain barrier permeability [20]. The CNS origin of elevated levels of NfL in CSF and in serum is further confirmed by the correlation between rising levels of serum and CSF NfL with activation of neurodegeneration in an inducible mouse model of neurodegeneration [21]. Furthermore, elevated levels of serum and CSF NfL have been reported in other neurodegenerative diseases mainly affecting the cerebral cortex such as Alzheimer's disease [16], FTD [22] and Huntington's disease [23]. Intriguingly, it is not clear yet how NfL is shed into the peripheral blood in ALS. Brureau and colleagues have identified in CSF of Thy-Tau22 mice a 10-kDa fragment of NfL but not the full-length NfL that is detected with a commercial enzyme-linked immunosorbent assay using the same pair of antibodies as in our study [21]. This might suggest that studies in human body fluids also detect short fragments of NfL

			Univariate Cox regressio	n	Multivariate Cox regres (n = 52)	ssion
			HR (95% CI)	P-value	HR (95% CI)	P-value
Serum NfL†*‡	$n = 113^{\$}$	Low tertile	Ref	_	Ref	_
		Mid tertile	6.92 (2.31-20.73)	0.001	4.47 (1.08-18.63)	0.04
		High tertile	10.49 (3.57-30.88)	< 0.0001	5.34 (1.39-20.56)	0.015
Disease progression rate†‡	n = 106	Slow	Ref	_	Ref	_
		Intermediate	3.38 (1.16-9.82)	0.026	1.24 (0.30-5.17)	0.77
		Fast	9.51 (3.19-28.37)	< 0.0001	1.05 (0.20-5.63)	0.95
UMN†	n = 106	1 Region	Ref	_		
		2 Regions	0.47 (0.19-1.20)	0.113		
		3 Regions	1.21 (0.55-2.68)	0.637		
LMN†	n = 106	1 Region	Ref	_		
		2 Regions	2.15 (0.82-5.66)	0.12		
		3 Regions	2.73 (1.19-6.28)	0.018		
UMN + LMN†‡	n = 106	1 Region	Ref	_	Ref	_
		2 Regions	1.34 (0.65-2.77)	0.424	0.94 (0.26-3.42)	0.93
		3 Regions	4.51 (2.02-10.09)	< 0.0001	2.38 (0.73-7.81)	0.15
Diagnostic delay‡	n = 106	Continuous	0.95 (0.91-0.996)	0.035	0.92 (0.85-1.01)	0.067
Forced vital capacity‡	n = 102	Continuous	0.97 (0.96-0.98)	< 0.0001	0.97 (0.95-0.99)	0.005
C9orf72 status†‡	n = 105	Noncarriers	Ref	_	Ref	_
		Carriers	2.0 (0.88-4.52)	0.097	0.50 (0.12-2.05)	0.34
FTD†"‡	n = 113	Non-FTD	Ref	_	Ref	_
		FTD	2.34 (1.04-5.26)	0.039	1.04 (0.23-4.77)	0.96
Age at onset	n = 107	Continuous	1.01 (0.99-1.04)	0.307		
Age at sampling‡	n = 113	Continuous	1.01 (0.99-1.04)	0.408	1.01 (0.96-1.06)	0.72
Site of onset†‡¶	n = 104	Bulbar	Ref	_	Ref	_
		Spinal	0.64 (0.34-1.20)	0.161	3.73 (1.16-12.01)	0.027
Gender†‡	n = 113	Male	Ref	_	Ref	
		Female	1.18 (0.65-2.17)	0.59	0.82 (0.28-2.42)	0.73

Table 3. Univariate and multivariate Cox regression analysis of possible clinical predictors of survival in patients with ALS

95% CI, 95th percentile confidence interval; ALS, amyotrophic lateral sclerosis; FTD, frontotemporal dementia; HR, hazard ratio; LMN, lower motor neuron; NfL, neurofilament light chain; Ref, reference; UMN, upper motor neuron.

†Univariate analysis based on categorical values.

 \ddagger Predictors entered into the multivariate model.

ALS were excluded from the survival analysis because possible follow-up was shorter than 1 year (Samples from December 2016 and December 2017). Univariate and Multivariate Cox regressions were performed with the backward method with a threshold to inclusion in the model of P < 0.157.

¶Only spinal and bulbar onset were entered in the univariate and multivariate analysis.

that might give clues on which mechanisms are involved in the shedding of NfL from brain into blood.

Our findings are also in line with the correlation between serum NfL and the number of regions displaying a pathological brisk reflex and with white matter fibres degeneration of the superior longitudinal fasciculi and cortical spinal tract [24]. This might indicate that NfL is shed upon damage of long axons of the UMN. This is further supported by the rise of serum and CSF NfL in other diseases damaging the central nerve axons such as multiple sclerosis [25,26].

Intriguingly, serum NfL was relatively low in patients with PLS and HSP, which are two UMN syndromes. The low levels of serum NfL in PLS patients are in agreement with previous findings [8]. A reason for this lack of serum NfL elevation may be the slow speed of UMN degeneration in these disorders. In ALS, we confirmed that serum NfL was correlated with the disease progression rate, as shown previously [27]. Furthermore, the serum NfL of patients with PMA positively correlated with the disease progression rate, and a tendency was seen for patients with PLS, which requires to be confirmed on larger cohorts of patients with PLS and PMA. The relatively high levels of serum NfL in GBS and CIDP, both immune-mediated demyelinating disorders, confirmed the higher levels of NfL obtained in the CSF of

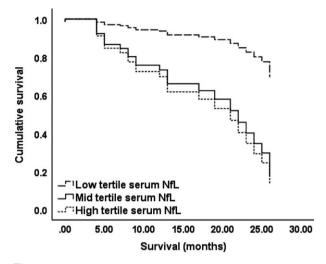


Figure 5. Multivariate Cox proportional-hazards regression survival curves of the patients with ALS according to their serum NfL levels. The serum NfL levels were stratified as low tertile (<115 pg/ml), mid-tertile (115 pg/ml $\leq x \leq 235$ pg/ml) and high tertile (>235 pg/ml). The disease progression, the tertile levels of serum NfL, the number of regions with UMN and LMN degeneration, the diagnostic delay, the age at sampling, the FVC score, the site of onset, the C9orf72 status, FTD and gender were included in the multivariate Cox model. Patients with ALS displaying a time difference between blood sampling and ALSFRS-R or EMG or FVC > 3 months and a follow-up between blood sampling and date of analysis of <1 year were excluded (excluding sampling between December 2016 and 2017). ALS, amyotrophic lateral sclerosis; ALSFRS-R, ALS functional rating scale revised; EMG, electromyography; FTD, frontotemporal dementia; FVC, forced vital capacity; LMN, lower motor neuron; NfL, neurofilament light chain; UMN, upper motor neuron.

these disorders [6,11]. Interestingly, Gaiottino and colleagues showed that the levels of serum and CSF NfL correlate in patients with GBS, suggesting a CNS origin. However, the correlation was less strong in patients with GBS than observed in patients with ALS, suggesting that serum NfL may also partly reflect demyelinating processes of the peripheral nerves in GBS [11].

Altogether, these findings are complementary and support the idea that both pNfH and NfL reflect distinct pathophysiological processes in ALS, with serum and CSF NfL levels rather reflecting UMN degeneration while CSF levels of pNfH are more specific to the ALS pathology. This is supported by our findings confirming that a rise in serum NfL is also seen in other diseases affecting the brain, and by the fact that CSF NfL, unlike CSF pNfH, is a less specific diagnostic biomarker for ALS [6]. Nevertheless, we now showed that the serum NfL levels discriminate patients with ALS and ALS mimics with a PPV of 97.5%, a NPV of 34%, a +LR of 5.02 and a -LR of 0.25.

We further showed that the serum NfL levels are significantly decreased with symptoms duration to the same extent than serum pNfH levels are, as recently described [7]. This decrease, observed in a cross-sectional measurement design is driven by an enrichment of fast progressors that were sampled much earlier in the disease continuum as compared with the slow progressors. Indeed, we described that, longitudinally, the serum levels of NfL levels are relatively stable or slightly increased in a subset of patients with ALS. These results are in accordance with previous reported results for NfL in CSF and serum [6,27,28]. It is of interest to explore whether longitudinal measurements of serum NfL might predict onset of disease in presymptomatic individuals, including asymptomatic carriers of a hexanucleotide repeat expansion in the C9orf72 gene.

Based on univariate Cox proportional-hazards regression we showed that the disease progression rate, the serum NfL levels, the number of regions with LMN degeneration, the number of regions with UMN and LMN degeneration, the diagnostic delay, the FVC and FTD were associated with an increased HR in patients with ALS. Notably, when investigating the levels of serum NfL, independently to the other clinical parameters, patients with serum NfL levels in the highest tertile had a HR rate to mortality 10 times greater than patients with serum NfL levels in the lowest tertile.

Furthermore, in a multivariate model, only the serum NfL levels, the site of onset and the FVC were identified as significant independent predictors of survival in patients with ALS when entered together with the other covariates. Our findings confirm the previous findings that serum NfL at time of diagnosis is a strong predictor of survival [27].

Our study has some limitations: (i) We did not investigate the correlation between the pattern of white matter degeneration in patients with ALS and PLS with the serum levels of NfL. Further studies investigating the relationship between blood and CSF levels of neurofilaments (both for pNfH and for NfL) and brain neurodegeneration (white matter and/or gray matter degeneration) would give further valuable insights on the pathophysiological processes reflected by the neurofilaments. (ii) The low serum NfL levels detected in patients with PLS would need to be confirmed on a larger cohort to validate the involvement of the disease progression rate on the serum concentrations of NfL in patients with PLS. (iii) Follow-up assessments of the serum levels of NfL of asymptomatic ALS mutation carriers would give further insight on the dynamics of the neurofilament prior to symptom onset. (iv) The survival was not controlled for possible comorbidities that could influence the survival after sampling.

Conclusion

In this study, we showed that serum NfL levels were increased as a function of the disease progression rate in ALS. We also described that serum NfL correlates with the extent of motor neuron degeneration mainly driven by UMN degeneration. In contrast with the disease progression rate and the extent of motor neuron disease, the serum NfL levels were found to be strong independent predictors of survival. Our findings warrant confirmation by large multicentre studies especially with respect to increase in the number of ALS mimics, including patients with cervical stenosis/myelopathy mimicking ALS.

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Author contributions

Concept and design of the study: BG, PVD and KP; Drafting the manuscript: BG, PVD and KP;

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Interpretation and statistical analysis: BG, PVD, MDS and KP; All co-authors revised the manuscript for intellectual content.

Data accessibility

The data are available from the corresponding author on reasonable request.

Conflict of interest

None declared.

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Supporting information

Additional Supporting Information may be found in the online version of this article at the publisher's web-site:

Figure S1. Comparison of the serum neurofilament light chain (NfL) levels according to the number of

regions with lower motor neuron (LMN) degeneration only.

Figure S2. Samples of patients with cervical stenosis/ myelopathy that have been retrospectively retrieved from hospital's serum biobank.

Figure S3. Correlation between the disease progression rate and the serum neurofilament light chain (NfL) levels in patients with progressive muscular atrophy (PMA) and primary lateral sclerosis (PLS).

Figure S4. Longitudinal assessment of the serum levels of neurofilament light chain (NfL) as a function of time after blood sampling in patients with amyotrophic lateral sclerosis (ALS).

Table S1. Clinical parameters of the primary lateralsclerosis (PLS) and progressive muscular atrophy(PMA) cohorts.

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