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A prospective observational cohort study

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Complement Profiles in Patients with Amyotrophic Lateral Sclerosis: A Prospective Observational Cohort Study

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Background: The complement system has been suggested to be involved in the pathophysiology of amyotrophic lateral sclerosis (ALS), a progressive motor neuron disease. In the present study, we compared levels of selected complement markers to clinical outcome in ALS patients.

Methods: This observational, explorative cohort study included 92 ALS patients, 61 neurological controls (NCs) admitted for suspected aneurysmal subarachnoid haemorrhage, and 96 neurologically healthy controls (NHCs). Peripheral blood and cerebrospinal fluid (CSF) were obtained for the measurement of ficolin-1, −2, and −3; collectin-11, MBL, MASP-3, MAP-1, C4, C3, PTX-3, and complement activation products C4c, C3bc, and sC5b-9. We recorded clinical outcomes of ALS patients for 24 to 48 months after inclusion in order to analyse the effects of the complement markers on survival time.

Results: Compared with both control groups, ALS patients exhibited increased collectin-11, C4 and sC5b-9 in plasma, as well as increased ficolin-3 in CSF. Ficolin-2 was significantly decreased in plasma of the ALS patients compared with NHCs, but not with NCs. The concentration of collectin-11, C3 and C3bc correlated negatively with the revised ALS functional rating scale (ALSFRS-R). No association was found between levels of complement markers and survival as estimated by hazard ratios.

Conclusion: ALS patients exhibit aberrant expression of selected mediators of the lectin complement pathway as well as increased activation of the terminal complement pathway, corroborating the notion that the complement system might be involved in the pathophysiology of ALS.

Keywords: amyotrophic lateral sclerosis, innate immunity, complement, lectin pathway, cerebrospinal fluid, pathophysiology, observational cohort study

Background

Amyotrophic lateral sclerosis (ALS) is a rare, rapidly progressing disease with a mean life expectancy of 30 months from the time of diagnosis.1 The hallmark of this disease is continual degeneration of the motor neurons, which leads to progressive dysfunction of extremity, trunk and bulbar skeletal muscles.

Several findings have suggested an essential role of the complement system in the pathophysiology of ALS.2 From being regarded as part of the host defence, recent research has shown a role for the complement system in development processes and homeostatic mechanisms of different tissues,3 including the brain, where complement components are crucial for neuronal migration and synaptic remodeling,4 and for
regulating memory. Moreover, a dysregulated complement system is associated with the pathophysiology of many systemic inflammatory diseases such as sepsis, systemic lupus erythematosus, rheumatoid arthritis, anti-neutrophilic cytoplasmic autoantibodies (ANCA) vasculitis, inflammatory myopathies, and microangiopathies such as atypical hemolytic uremic syndrome. Increased complement activity also seems to be implicated in disorders of neurodevelopmental origin, such as schizophrenia, autism spectrum disorder, and Rett syndrome, as well as the neuroinflammatory diseases, neuro-myelitis optica and myasthenia gravis.

Dysfunctional complement activation also seems to be present during the progression of ALS in the surroundings of the diseased motor neurons in the central nervous system (CNS) as well as systemically. Several studies based on animal ALS models suggest that the complement system plays a significant pathophysiological role during the initial phase of ALS, as activated complement proteins appear in the surroundings of the neuromuscular junctions (NMJs) concomitantly with the first pathological changes that appear in the NMJs of the failing motor neurons. A terminal complement inhibitor, ravulizumab, is currently being tested on ALS patients in a Phase III trial (ClinicalTrials.gov Identifier: NCT04248465).

Recently, Xu et al conducted a mass spectrometry analysis of plasma from ALS patients and reported upregulation not only of the complement pathway in general but also ficolin-3 expression. The latter finding validated a previous observation by the same research group and indicates a contribution of the lectin pathway of complement in the pathophysiology of ALS. Lectin pathway components are present in the CNS of healthy individuals and appear to be involved in normal embryogenesis, but have so far not been studied in conjunction with ALS. In the present study, we measured levels of selected markers of the lectin complement pathway as well as complement activation products in peripheral blood and cerebrospinal fluid (CSF) from ALS and non-ALS patients. We hypothesised that these mediators are upregulated both intrathecally and systemically during ALS and that more extreme changes are associated with more severe disease and shorter survival times.

**Materials and Methods**

This observational, exploratory cohort study, which is reported according to the STROBE guidelines, was approved by the Committees on Health Research Ethics in the Capital Region of Denmark (Approval number H-16017145) and the Danish Data Protection Agency (File number 2012-58-0004), registered with Clinicaltrials.gov (NCT02869048) and conducted in accordance with the Declaration of Helsinki. After receiving written and verbal information, all participants gave their written consent prior to enrollment.

In accordance with the Global Data Protection Regulation of the European Union, data are not publicly available but may be available upon reasonable request if a mutual data sharing agreement is approved by the relevant authorities.

**Participants**

As previously described, we included adults (> 18 years) into the following groups:

1. ALS cohort: Patients who were referred to or followed at five large Danish motor-neuron disease outpatient clinics if they fulfilled the El Escorial Revised criteria for probable or definite ALS.
2. Neurologically healthy controls (NHCs): Healthy patients (American Society of Anesthesiologists score, 1–2) scheduled for planned orthopaedic surgery in spinal anaesthesia at a large elective surgical centre were included.
3. Non-ALS neurological controls (NCs): Patients who underwent emergency admission to hospital because of suspected aneurysmal subarachnoid haemorrhage (SAH), and who underwent a lumbar puncture following computed tomography (CT) scanning of the brain; were included if the SAH diagnosis was refuted.

For both control groups (NHCs and NCs), patients with a history or symptoms of motor neuron disease, acute or chronic inflammatory disease, or autoimmune disease were excluded.

**Samples**

Peripheral venous blood and CSF were sampled at inclusion sites (Figure 1). The samples were immediately centrifuged (2000 g). Samples of EDTA plasma, serum, and CSF were kept at −20 °C during transportation and stored at −80 °C until analysis.

The level of the complement lectin pathway components ficolin-1, ficolin-2, ficolin-3, collectin-11, mannose-binding lectin (MBL), MBL/ficolin/collectin-associated serine protease 3 (MASP-3), MBL/ficolin/collectin associated protein...
1 (MAP-1), the acute phase marker pentraxin-3 (PTX-3), the complement components C4 and C3 and the activation products C4c, C3bc and sC5b-9 were measured in plasma and serum as well as in CSF as described below.

Complement Measurements

Complement related proteins ficolin-1, −2, and −3; collectin-11, MBL, MASP-3, MAP-1, PTX-3, and complement activation products C4c, C3bc, and sC5b-9 were measured in EDTA plasma and CSF samples by in-house sandwich enzyme-linked immunosorbent assays (ELISA). The measurements were conducted as previously described.30–41 The antibodies used for each of the ELISA analyses are listed in Supplementary Table 1. The serum concentrations of C3 and C4 were estimated using an automated turbidimetric protein analyzer (SPAPLUS®, The Binding Site group LDT, Birmingham, UK). An overview of the performed analyses is found in Table 1.

Statistical Analyses

The ALS group and the two control groups (NCs and NHCs) were compared using Bonferroni-corrected, stepwise three-way analysis of variance (ANOVA). For this comparison, analysis testing for differences was done, using the following covariates: subject category (ALS, NC, NHC), sex, and age. Data which did not follow a Gaussian distribution were log- or square-root transformed. To further analyse the significant differences between the ALS patients, the NHCs, and the NCs in the subject category, Bonferroni corrected, one-way ANOVA and t-testing were used. Missing observations were omitted from the statistical analyses on a listwise basis and were assumed to be missing at random.

The potential association between complement marker levels and clinical severity of ALS was assessed by a univariate regression model (Pearson’s correlation). For clinical severity, the revised ALS functional rating scale (ALSFRS-R) was used. ALSFRS-R quantifies the level of daily living activities in ALS.42

To evaluate the association between complement activation and survival, markers for which the level in ALS patients differed from both or one of the control groups were analyzed as continuous variables in a univariate Cox’ proportional hazards regression model. Moreover, we calculated the median survival time of the ALS cohort as we defined the primary end point as either death or the date for the initiation of invasive mechanical ventilatory support at censor date (15th of June 2020). ALS patients that had reached primary end point (invasive mechanical ventilation) prior to the test date were not included in the survival analysis.

Results

Patients

Between June 2016 and June 2018, 109 patients were screened for enrollment into the ALS cohort, of whom 92 were enrolled; we excluded 17 because the ALS diagnosis was dismissed. Of 63 potential NCs, two were excluded due to autoimmune disease and neuroinfection, respectively. Finally, of 101 possible NHCs, five were excluded due to the cancellation of surgery.

Plasma and Serum Levels in ALS Patients, NCs, and NHCs

According to the ANOVA, the levels of ficolin-2, collectin-11, C4, C3bc, sC5b-9 and the C3bc/C3 index varied significantly between patients with ALS, NCs,
Table 1 Complement Markers and Mediators

<table>
<thead>
<tr>
<th>Complement Component</th>
<th>Sample</th>
<th>Transf.</th>
<th>Outl.</th>
<th>Obs.</th>
<th>ALS</th>
<th>NC</th>
<th>NHC</th>
<th>P*</th>
</tr>
</thead>
<tbody>
<tr>
<td>Ficolin-1 (ng/mL)</td>
<td>EDTA plasma</td>
<td>Log</td>
<td>None</td>
<td>249</td>
<td>242 (219–268)</td>
<td>246 (217–279)</td>
<td>253 (229–280)</td>
<td>0.83</td>
</tr>
<tr>
<td>Ficolin-2 (µg/mL)</td>
<td>EDTA plasma</td>
<td>Log</td>
<td>None</td>
<td>249</td>
<td>4.21 (3.77–4.70)</td>
<td>4.03 (3.51–4.63)</td>
<td>5.12 (4.58–5.72)</td>
<td>0.01</td>
</tr>
<tr>
<td>Ficolin-3 (µg/mL)</td>
<td>EDTA plasma</td>
<td>Log</td>
<td>None</td>
<td>249</td>
<td>22.8 (21.1–24.7)</td>
<td>22.4 (20.3–24.7)</td>
<td>23.0 (21.3–24.9)</td>
<td>0.92</td>
</tr>
<tr>
<td>Collectin-11 (ng/mL)</td>
<td>EDTA plasma</td>
<td>Log</td>
<td>1</td>
<td>249</td>
<td>209 (196–223)</td>
<td>192 (177–208)</td>
<td>176 (165–188)</td>
<td>3.9 × 10⁻⁴</td>
</tr>
<tr>
<td>MB (ng/mL)</td>
<td>EDTA plasma</td>
<td>Sqrt</td>
<td>None</td>
<td>249</td>
<td>949 (784–1130)</td>
<td>886 (689–1108)</td>
<td>843 (688–1014)</td>
<td>0.69</td>
</tr>
<tr>
<td>MASP-3 (ng/mL)</td>
<td>EDTA plasma</td>
<td>Log</td>
<td>None</td>
<td>249</td>
<td>5227 (4986–5480)</td>
<td>534 (5039–5674)</td>
<td>5081 (4846–5327)</td>
<td>0.39</td>
</tr>
<tr>
<td>MAP-1 (ng/mL)</td>
<td>EDTA plasma</td>
<td>Log</td>
<td>None</td>
<td>249</td>
<td>226 (207–247)</td>
<td>218 (195–244)</td>
<td>219 (200–240)</td>
<td>0.85</td>
</tr>
<tr>
<td>PTX-(ng/mL)</td>
<td>EDTA plasma</td>
<td>None</td>
<td>None</td>
<td>248</td>
<td>4.80 (3.40–6.30)</td>
<td>3.10 (1.40–4.80)</td>
<td>3.22 (1.92–4.52)</td>
<td>0.18</td>
</tr>
<tr>
<td>C (mg/mL)</td>
<td>Serum</td>
<td>None</td>
<td>None</td>
<td>249</td>
<td>0.23 (0.22–0.24)</td>
<td>0.21 (0.19–0.23)</td>
<td>0.19 (0.18–0.20)</td>
<td>2.1 × 10⁻⁴</td>
</tr>
<tr>
<td>C4 (AU)</td>
<td>EDTA plasma</td>
<td>Log</td>
<td>None</td>
<td>249</td>
<td>1.3 (1.24–1.52)</td>
<td>1.3 (1.20–1.55)</td>
<td>1.5 (1.36–1.66)</td>
<td>0.36</td>
</tr>
<tr>
<td>C3 (mg/mL)</td>
<td>Serum</td>
<td>Log</td>
<td>None</td>
<td>248</td>
<td>1.0 (0.99–1.07)</td>
<td>1.0 (0.99–1.09)</td>
<td>1.0 (0.97–1.05)</td>
<td>0.62</td>
</tr>
<tr>
<td>C3bc (AU)</td>
<td>EDTA plasma</td>
<td>Log</td>
<td>None</td>
<td>249</td>
<td>2.97 (2.67–3.30)</td>
<td>4.59 (4.02–5.25)</td>
<td>3.30 (2.97–3.67)</td>
<td>5.1 × 10⁻⁷</td>
</tr>
<tr>
<td>C3bc/C3</td>
<td>Index</td>
<td>Log</td>
<td>None</td>
<td>248</td>
<td>2.89 (2.61–3.20)</td>
<td>4.41 (3.88–5.01)</td>
<td>3.26 (2.94–3.61)</td>
<td>5.1 × 10⁻⁷</td>
</tr>
<tr>
<td>sC5b-(AU)</td>
<td>EDTA plasma</td>
<td>Log</td>
<td>1</td>
<td>249</td>
<td>0.66 (0.61–0.71)</td>
<td>0.58 (0.53–0.64)</td>
<td>0.58 (0.54–0.63)</td>
<td>0.017</td>
</tr>
<tr>
<td>Ficolin-3 (µg/mL)</td>
<td>CSF</td>
<td>Log</td>
<td>None</td>
<td>239ᵃ</td>
<td>0.005 (0.0052–0.0067)</td>
<td>0.004 (0.0038–0.0052)</td>
<td>0.0049 (0.0043–0.0056)</td>
<td>0.013</td>
</tr>
<tr>
<td>C3bc (AU)</td>
<td>CSF</td>
<td>Log</td>
<td>2</td>
<td>233ᵇ</td>
<td>33.4 (27.8–40.2)</td>
<td>53.6 (42.5–67.6)</td>
<td>43.3 (36.0–52.1)</td>
<td>0.0013</td>
</tr>
<tr>
<td>C4c (AU)</td>
<td>CSF</td>
<td>Sqrt</td>
<td>None</td>
<td>235ᶜ</td>
<td>3.78 (3.28–4.32)</td>
<td>3.08 (2.52–3.70)</td>
<td>3.11 (2.66–3.60)</td>
<td>0.36</td>
</tr>
<tr>
<td>sC5b-9 (AU)</td>
<td>CSF</td>
<td>Data</td>
<td>None</td>
<td>236ᵈ</td>
<td>0.0014 (0.0000–0.0030)</td>
<td>0.001 (0.0000–0.0030)</td>
<td>0.000 (0.0000–0.0020)</td>
<td>0.56</td>
</tr>
</tbody>
</table>

Notes: Values are mean (95% confidence interval). Eighty-two ALS patients donated CSF. Six samples failed during the analysis process and therefore no results are available for six observations in this assay. Four samples failed during the analysis process and therefore no results were available for four observations in this assay. Three samples failed during the analysis process and therefore no results were available for four observations in this assay. All P values are Bonferroni-corrected (p-value < 0.017 to discard the null hypothesis).

Abbreviations: AU, arbitrary unit; NCs, neurologically healthy controls; NHCs, neurologically healthy controls; Log, logarithmically transformed; Sqrt, square root transformed; Transf, transformations; Outl, outliers; Obs, observations; CSF, cerebrospinal fluid.
Table 2 Demographic Data

<table>
<thead>
<tr>
<th></th>
<th>ALS Patients (n=92)</th>
<th>Neurological Controls (n=61)</th>
<th>Neurologically Healthy Controls (n=96)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Age at onset of symptoms (years)</td>
<td>62.5 (±11.7)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Age at study date (years)</td>
<td>64.6 (±11.0)</td>
<td>49.6 (±15.5)</td>
<td>46.9 (±11.4)</td>
</tr>
<tr>
<td>Time from symptom onset to study date (months)</td>
<td>25.0 (±32.8)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Time from symptom onset to diagnosis (months)</td>
<td>16.5 (±12.7)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Male sex</td>
<td>52 (56%)</td>
<td>28 (46%)</td>
<td>62 (65%)</td>
</tr>
<tr>
<td>Familial ALS</td>
<td>2 (2%)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Subtypes</td>
<td>Bulbar: 26 (28%)</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>Spinal: 59 (64%)</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>Trunical: 1 (1%)</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>Mixed: 6 (7%)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>ALSFRS-R score at test date (points)</td>
<td>34.5 (±9.5)</td>
<td>26.7 (±9.5)</td>
<td>28.6 (±9.4)</td>
</tr>
<tr>
<td>Bulbar motor score</td>
<td>8.9 (±2.8)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Fine motor score</td>
<td>8.1 (±3.2)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Coarse motor score</td>
<td>8.0 (±3.0)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Respiratory score</td>
<td>9.3 (±4.1)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Ventilatory support (on test date)</td>
<td>16</td>
<td>7</td>
<td></td>
</tr>
<tr>
<td>Invasive mechanical ventilation</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Non-invasive ventilation</td>
<td>11</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Percutaneous endoscopic gastrostomy (on test date)</td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

Note: Values are mean (SD) or numbers (percentage).

Abbreviations: ALSFRS-R, the revised amyotrophic lateral sclerosis functional rating scale; fALS, familial amyotrophic lateral sclerosis.

and NHCs in plasma and serum samples (Figure 2 and Table 1). Following Bonferroni-corrected t-tests, ficolin-2 levels remained lower in ALS patients and NCs compared to NHCs, with no difference between ALS patients and NCs. Moreover, ALS patients had higher collectin-11, C4, and sC5b-9 levels than NHCs, but not compared to NCs. Finally, C3bc and consequently C3bc/C3 index were markedly higher in NCs than in ALS patients and NHCs, with no difference between the latter groups.

There were no significant differences between the three groups concerning ficolin-1, ficolin-3, MBL, MASP-3, MAP-1, C4c, PTX-3 (Table 1). MASP-3 levels were lower in males (mean value, 5031 (95% confidence interval, 4842–5227) ng/mL) than in females (5432 (5198–5677) ng/mL; p = 0.011); apart from MASP-3, no influence on mediator levels was exerted by age or sex.

CSF Levels in ALS Patients, NCs, and NHCs

In CSF, only ficolin-3 and the complement activation products C4c, C3bc, and sC5b-9 were measured (Table 1). ALS patients had higher ficolin-3 and lower C3bc levels compared to NCs and NHCs (Figure 3A and C); levels of C4c and C5b-9 did not differ between groups. For ALS patients, blood levels of ficolin-3 and C3bc correlated positively with CSF levels (Figure 3B and D); no correlation was found for C4c and C5b-9.

Association Between Complement Markers and Clinical Score

Collectin-11, C3 and C3bc correlated negatively with the ALSFRS-S score (Figure 4). The remaining biomarkers (ficolin-1, ficolin-2, ficolin-3, MBL, MASP-3, MAP-1, MAP-2, C4c, PTX-3, MASP-1, MASP-2, MBL, and C4c) did not correlate with the ALSFRS-S score.
PTX3, C4, C4c, and sC5b-9) were not associated with the ALSFRS-R score (Supplementary Table 2, Supplementary Figures 1 and 2).

**Association Between Complement Markers and Survival**

Eighty-five ALS patients were included for the survival analysis; the remaining seven patients were already receiving invasive mechanical ventilatory support on the test date (Supplementary Table 3). Among the included ALS patients, 70 patients reached the primary end-point (initiation of invasive mechanical ventilatory treatment or death) at censor date. Of those, the median survival time from onset of symptoms until the primary end point was 34 months. The median survival time from the time of diagnosis until the primary end point was 18 months. According to the Cox proportional hazard model, none of the complement markers was associated with survival time.

**Discussion**

In the present study, we analysed several markers and activation products of the lectin complement system and found that several of these were markedly changed...
as assessed by variance analysis among the groups of ALS patients, neurological controls and neurologically healthy controls. Even though correction for multiple testing rendered some differences non-significant when these changes were further compared between groups, the point estimates for the circulating levels of...
collectin-11, C4, and sC5b-9 were increased in the ALS group compared to both control groups, whereas ficolin-2 was decreased both in the ALS and neurological controls.

Even though sC5-9 complex was increased in the ALS patients compared to the two control groups, as previously reported,^{20} we observed no difference between the three groups regarding C4c, an activation marker which is supposedly increased by either classical or lectin pathway activation. However, the activation marker C3bc was increased only in neurological patients compared to ALS and neurologically healthy controls. These differences between the three different complement activation markers are challenging to explain. Still, it could indicate that the terminal complement route is activated by a non-canonical mechanism not involving the traditional activation routes. Apart from the activation by the classical, lectin, and alternative pathways direct activation of C5 has been shown, resulting in the generation of sC5b-9 without the involvement of C3 complement convertases. This has been documented by the action of thrombin on C5, which is not very likely in ALS, but other proteases might be involved. C3 activation without downstream generation of an efficient C5 convertase and generation of sC5b-9 has been demonstrated and might explain why we only observed increased levels of C3bc in neurological controls. In these patients, complement activation might be related to the alternative route and not the classical and lectin pathway, explaining the absence of C4c activation. However, to further complicate the picture, C3 activation via the lectin pathway enzymes without the involvement of C4 and C2 has recently been reported.^{46}

In CSF, we were able to detect ficolin-3, C4c, and C3bc. Trace amounts of sC5-9 were only detected in some individuals, while the other components were undetectable by the methods used. In contrast to the results found in plasma, the ficolin-3 level was significantly increased in the CSF from ALS patients compared with the two control groups. As for plasma, CSF C4c did not differ between ALS and controls, while the same difference as was observed in plasma was also observed for C3bc in CSF. The level of sC5b-9 in CSF was too low to allow a meaningful comparison between the groups.

Previously, increased levels of ficolin-3 have been detected by mass spectrometry in ALS patients and confirmed by ELISA in a small group of patients.^{26,47} We could not confirm these observations in our study. There might be several reasons for this. ALS is a heterogeneous disease, and protein levels might differ over time. The strength of our study was that we carefully selected the control groups to be able to detect differences specific for ALS and not just a random difference from the diseased individuals compared with healthy controls. Thus, the former difference might have been due to a type 1 error.

Nevertheless, we did observe an increased concentration of ficolin-3 in CSF compared with the controls. Whether this is because ficolin-3 might be differentially regulated in the different compartments at different stages of the disease cannot be answered from our study, but is indeed an indication of the involvement of ficolin-3 at least as a biological marker in ALS. The decreased level of ficolin-2 in both ALS and neurological controls is relevant since it is consistent with findings that ficolin-2 is either consumed or down-regulated during certain inflammatory processes.^{48}

Two novel findings of this study were the selective increase of collectin-11 and C4 in plasma. The knowledge of the biological role of collectin-11 is not well established, but there is an emerging acceptance that it plays an important role in organ development during embryogenesis and in tissue homeostatic mechanisms. C4 has been suggested to be important for the development of the brain, as increased C4 gene copy numbers are associated with the development of neuropsychiatric diseases like schizophrenia. Thus, it would be interesting to investigate if this phenomenon is also associated with ALS.

This study has several limitations. First, we obtained the samples when it was appropriate according to the patients’ clinical workup after they had consented to the study. For the majority of patients, this was during the diagnostic workup, whereas a few patients with confirmed ALS were also included. This potentially increases the heterogeneity of the population. Second, the controls were not matched with regard to age and sex of the patients, mainly for practical purposes; since the characteristics of the lectin-complement system change both with age and sex,^{51} this makes any comparison between patients and controls harder. Along the same line of reasoning, it would have been interesting to compare the ALS patients to other patients with chronic neurological disease. As controls with neurological disease, we included only patients suspected of an acute neurological disease (subarachnoid hemorrhage), which provided an acute indication for lumbar puncture and as such was logistically attractive. By contrast, including patients with chronic neurological disease was not clinically feasible. Third, the neurologically healthy patients’ did not undergo specific testing
to exclude neurological disease, but were simply assumed to be healthy based on the available medical information on the day of surgery. Fourth, we included patients only with definite or probable ALS; this may have excluded patients during the onset or early stages of ALS and thus obviated our chances of studying inflammatory processes as part of the pathogenesis of ALS. On the other hand, any association between inflammation and progression of ALS should be more marked in patients with established ALS. The present findings do not support the presence of a persistent inflammatory process to drive the progression of ALS, at least not one that includes the lectin-complement system. They do, however, not exclude temporary or ‘pulsatile’ inflammatory phenomena that were not present at the time of sampling. Fifth, we did not obtain data regarding general systemic or neuroinflammation, such as blood or CSF white blood cell and differential counts or plasma C-reactive protein. Finally, ALS may be associated with cognitive impairment; we did not attempt to distinguish between patients with and without cognitive difficulties. Neither did we attempt to characterize the functional level of patients beyond that which was comprised in the ALSFRS-R scale; a more granular clinical staging could potentially have been obtained using tools such as the Kings ALS clinical staging.52

Conclusion
This explorative study substantiates that the terminal part of the complement system is activated in ALS patients in a non-canonical manner. Collectin-11, C3 and C3bc correlated negatively with the ALSFRS-S score. We also observed that collectin-11 and C4 were significantly increased in plasma compared to controls, but not after correction for multiple comparisons. Thus, further analysis in independent cohorts is necessary to validate these observations. Nevertheless, this study supports the notion that the complement system might be involved in the pathophysiology of ALS.

Abbreviations
ALS, amyotrophic lateral sclerosis; ANOVA, analysis of variance; AU, arbitrary unit; AUROC, area under the ROC curve; CNS, central nervous system; CSF, cerebrospinal fluid; EDTA, ethylenediamine tetraacetic acid; ELISA, enzyme-linked immunosorbert assay; fALS, familial amyotrophic lateral sclerosis; g, the relative centrifugal force; Log, logarithmic; MAP-1, MBL/ficolin/collectin associated protein 1 (MAP-1); MASP-3, MBL/ficolin/collectin-associated serine protease 3; MBL, Mannose-binding lectin; NMJ, neuromuscular junction; NCs, neurological controls; NHCs, neurologically healthy controls; NPV, negative predictive value; PPV, positive predictive value; PTX-3, pentraxin-3; ROC, receiver operating characteristic; SAH: aneurysmal subarachnoid haemorrhage; sALS, sporadic amyotrophic lateral sclerosis; Sqrt, Square root.

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Author Contributions
All authors made a significant contribution to the work reported, whether that is in the conception, study design, execution, acquisition of data, analysis and interpretation, or in all these areas; took part in drafting, revising or critically reviewing the article; gave final approval of the version to be published; have agreed on the journal to which the article has been submitted; and agreed to be accountable for all aspects of the work.

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Disclosure
The authors report no conflicts of interest for this work.

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