

ORIGINAL WORK



Longitudinal Assessment of Blood-Based Inflammatory, Neuromuscular, and Neurovascular Biomarker Profiles in Intensive Care Unit–Acquired Weakness: A Prospective Single-Center Cohort Study

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Abstract

Background: The diagnosis of intensive care unit (ICU)-acquired weakness (ICUAW) and critical illness neuromyopathy (CINM) is frequently hampered in the clinical routine. We evaluated a novel panel of blood-based inflammatory, neuromuscular, and neurovascular biomarkers as an alternative diagnostic approach for ICUAW and CINM.

Methods: Patients admitted to the ICU with a Sequential Organ Failure Assessment score of ≥ 8 on 3 consecutive days within the first 5 days as well as healthy controls were enrolled. The Medical Research Council Sum Score (MRCSS) was calculated, and motor and sensory electroneurography (ENG) for assessment of peripheral nerve function were performed at days 3 and 10. ICUAW was defined by an MRCSS < 48 and CINM by pathological ENG alterations, both at day 10. Blood samples were taken at days 3, 10, and 17 for quantitative analysis of 18 different biomarkers (white blood cell count, C-reactive protein, procalcitonin, C-terminal agrin filament, fatty-acid-binding protein 3, growth and differentiation factor 15, syndecan 1, troponin I, interferon- γ , tumor necrosis factor- α , interleukin-1 α [IL-1 α], IL-1 β , IL-4, IL-6, IL-8, IL-10, IL-13, and monocyte chemoattractant protein 1). Results of the biomarker analysis were categorized according to the ICUAW and CINM status. Clinical outcome was assessed after 3 months.

Results: Between October 2016 and December 2018, 38 critically ill patients, grouped into ICUAW (18 with and 20 without) and CINM (18 with and 17 without), as well as ten healthy volunteers were included. Biomarkers were significantly elevated in critically ill patients compared to healthy controls and correlated with disease severity and 3-month outcome parameters. However, none of the biomarkers enabled discrimination of patients with and without neuromuscular impairment, irrespective of applied classification.

Conclusions: Blood-based biomarkers are generally elevated in ICU patients but do not identify patients with ICUAW or CINM.

Trial registration: ClinicalTrials.gov identifier: NCT02706314.

Keywords: Muscle weakness, Critical illness, Biomarkers, Neuromuscular diseases, Prospective study

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Introduction

In critically ill patients, intensive care unit (ICU)-acquired weakness (ICUAW) is one of the most common causes of neuromuscular impairment. This is associated with increased morbidity, mortality, and long-term impairment in quality of life up to 5 years after ICU discharge [1]. In daily routine practice, clinical assessment and manual muscle strength testing using established scores such as the Medical Research Council Sum Score (MRCSS) represent the most frequently performed and recommended diagnostic tests for ICUAW [2–4]. However, prolonged sedation, mechanical ventilation, and unconsciousness due to neurological impairment frequently hampers clinical examination during the acute phase of critical illness [5]. Muscle biopsies and electroneurography (ENG) are established procedures for the diagnosis of critical illness neuromyopathy (CINM) but are rarely performed in daily clinical routine because of their invasiveness and required expertise [6]. Although ultrasonography can be considered as a valuable tool for the detection of neuromuscular pathologies, evidence with focus on ICUAW and CINM is still limited because of the absence of well-powered studies [7]. In contrast, the diagnostic and prognostic value of blood-based biomarkers has been widely evaluated in several disease entities within critically ill patients, including delirium [8, 9] and sepsis-associated encephalopathy [10]. Advantages of blood-based biomarkers include their easy assessment through simple blood sampling and the opportunity of serial measurements to monitor disease progression. In the field of ICUAW and CINM, the specific value of blood-based biomarkers for the detection and monitoring of ICUAW as well as the prediction of patient outcomes remains unclear. Because sepsis is considered one of the main risk factors for ICUAW, multiple cytokines, including tumor necrosis factor- α (TNF α), interleukin-1 (IL-1), IL-6, IL-10, and interferon- γ (IFN γ), have been suggested to promote muscle proteolysis, activate the ubiquitin proteasome system, induce contractile dysfunction, and inhibit muscle protein synthesis, but in contrast, they have also been suggested to enhance muscle regeneration following injury [11–16]. Cleavage products of vascular endothelium glycocalyx shredding in sepsis might therefore also be detectable in ICUAW [17, 18]. In case of CINM, direct markers of neuromuscular damage might include proteins released from the nerval and muscular compartments as well as from the

interlinking neuromuscular junction following inflammation, immobilization, and mechanical silencing [19].

To overcome this gap, we conducted a prospective observational study using a broad panel of cytokines and inflammatory (white blood cell count, C-reactive protein, procalcitonin, IFN γ , TNF α , IL-1 α , IL-1 β , IL-4, IL-6, IL-8, IL-10, IL-13, monocyte chemoattractant protein 1, and growth and differentiation factor 15 [GDF15]), neurovascular (syndecan 1) and neuromuscular (C-terminal agrin filament [CAF], troponin I, and fatty-acid-binding protein 3 [FABP3]) biomarkers. We hypothesized, that distinct longitudinal biomarker profiles help to differentiate between patients with and without ICUAW and correlate with clinical outcome parameters.

Methods

Study design and inclusion and exclusion criteria

We conducted a prospective single-center observational cohort study. The study was performed in accordance with the Declaration of Helsinki and approved by the local ethics committee of the University of Rostock (ethics identifier: AS 2016–0016). Written informed consent for participation was given by healthy volunteers, patients, or a legal representative prior to enrollment. The present investigation represents the laboratory analysis part of a comprehensive and combined clinical–experimental study. Registration was made prior to the trial beginning and before the first patient enrollment (registered at ClinicalTrials.gov: NCT02706314). Results of the clinical and ex vivo experimental parts have already been published [20, 21]. Briefly, patients being at least 17 years of age and presenting with a Sequential Organ Failure Assessment (SOFA) score ≥ 8 on 3 consecutive days within the first 5 days after ICU admission were defined as critically ill and were eligible for enrollment. Main exclusion criteria comprised preexisting neuromuscular disorders and high-dosage corticosteroid treatment with ≥ 300 mg of hydrocortisone or equivalent per day. Further details are published elsewhere [18]. Clinical and neurophysiologic examinations were performed on days 3 and 10 after enrollment, and blood was sampled at study days 3, 10, and 17. The Barthel Index (BI) and the modified Rankin scale (mRS) 3 months after enrollment were considered as markers of the long-term patient outcome. Additionally, healthy volunteers without any preexisting neuromuscular disorders from the Institute of Physiology of the Rostock University Medical Center

were recruited and received the same clinical and neurophysiological tests as the patients in one session. This trial was conceptualized as a pilot study.

Clinical assessment and definition of ICUAW

Clinical evaluation of muscle strength was performed by the MRCSS assessment at study days 3 and 10. Intubated and ventilated patients received a sedation holiday prior to clinical examination. Patients were considered eligible for muscle strength testing using the MRCSS based on the score of five questions [22, 23]. ICUAW was considered by an MRCSS < 48 at day 10 after enrollment, and patients were subsequently defined as ICUAW positive (ICUAW+); otherwise, they were defined as ICUAW negative (ICUAW−) [24].

Electrophysiology and definition of CINM

ENG was performed as described in detail before [21]. Briefly, at study days 3 and 10, compound motor action potentials were assessed over the musculus abductor digiti minimi and the musculus extensor digitorum brevis. Stimuli were applied via the stimulator module of an Epoch XP EMG machine (Axon Systems), and recordings of evoked responses were recorded at 10 kHz with 1-Hz high-pass and mains filtering using a PowerLab28T system (AD instruments, New Zealand) via surface EMG electrodes (Kendall, Covidien, Ireland). A nonpathological response was defined by a minimum amplitude of 4 mV. Sensory nerve action potentials were recorded using the antidromic technique over the superficial radial and the sural nerves with a threshold of 7.5 μ V for a normal response in the averaged trace of ten repeated recordings. CINM was diagnosed as an absence of muscle reflexes, reduced muscle tone, and movements induced by painful stimuli as well as reduced compound motor action potentials and sensory nerve action potentials in four or more recording sites, and patients were defined as CINM positive (CINM+); otherwise, patients were defined as CINM negative (CINM−).

Biomarker measurements in plasma and serum samples

Analyses were performed using commercially available enzyme-linked immunosorbent assay (ELISA) kits. The biomarker panel comprised skeletal muscle troponin I (catalog number: MBS765801, Human TNNI1 ELISA Kit, Mybiosource, San Diego, CA), FABP3 (catalog number: BMS 2263, Human FABP-3 ELISA, Thermo Fisher Scientific, Waltham, MA), and CAF (catalog number: MBS7606926, Human CAF ELISA Kit, Mybiosource, San Diego, CA). Syndecan 1 was used to assess vascular endothelium (catalog number: ab46506, Human Syndecan-1 ELISA Kit, Abcam, Cambridge, UK). Cytokine profiling comprised the stress-related GDF15 (catalog

number: ab155432, GDF-15 Human ELISA Kit, abcam, Cambridge, UK) and inflammatory cytokines (catalog number: ab197449, Human inflammation Antibody Array A [IL-1 α , IL-1 β , IL-4, IL-6, IL-8, IL-10, IL-13, MCP1, IFN γ , TNF α] – Quantitative, abcam, Cambridge, UK). Plasma and serum samples were centrifuged at 4 °C with 2500 rpm for 15 min and aliquoted and stored at −80 °C until analysis. All ELISA measurements were performed according to manufacturer-specific protocols, and all samples were assessed in duplicate.

Statistics

Microsoft Excel 2010 (Microsoft, Redmond, WA) was used for data curation. Statistical analysis was done with IBM SPSS Statistics (Version 25, IBM Corp., Armonk, NY). The Shapiro–Wilk test was used for assessment of normal distribution of continuous data. Results are presented as frequency (percentage), median (interquartile range [IQR]), or mean (standard deviation) as appropriate. Student's *t*-test, the Mann–Whitney *U*-test, Fisher's exact test, and the χ^2 test were employed. For correlation analysis between normally and nonnormally distributed data, the Pearson correlation coefficient and the Spearman rank correlation coefficient were computed, respectively. Statistical significance was indicated by *p* < 0.05. All statistical tests were two-sided.

Results

Study population characteristics and outcome data

In total, 51 critically ill patients and ten healthy controls were recruited between October 2016 and December 2018 (Fig. 1). Three patients died within the first days after enrollment, and these data were therefore excluded from the analysis. Furthermore, the MRCSS at study day 10 was not assessable in ten patients because of prolonged sedation, delirium, or patient noncompliance. Thus, data from 38 patients were available for subsequent analysis. Baseline characteristics of these patients are given in Table 1. Subsequently, patients were categorized according to the presence of ICUAW and CINM. All patients with ICUAW, except one with an MRCSS of 52 at study day 10, were also classified as CINM+. All patients, regardless of their classification, were comparable in terms of their activities of daily living and neuromuscular function prior to hospital admission. Because of inconclusive ENG data, we had to exclude three patients from the CINM subgroup analysis. We also collected data from ten healthy controls (six men, mean age 53.1 ± 8 years) without any preexisting neuromuscular disorder (Supplementary Table 1). Patients with and without ICUAW as well as with and without CINM were comparable for age, sex distribution, type of surgery, sepsis, and renal

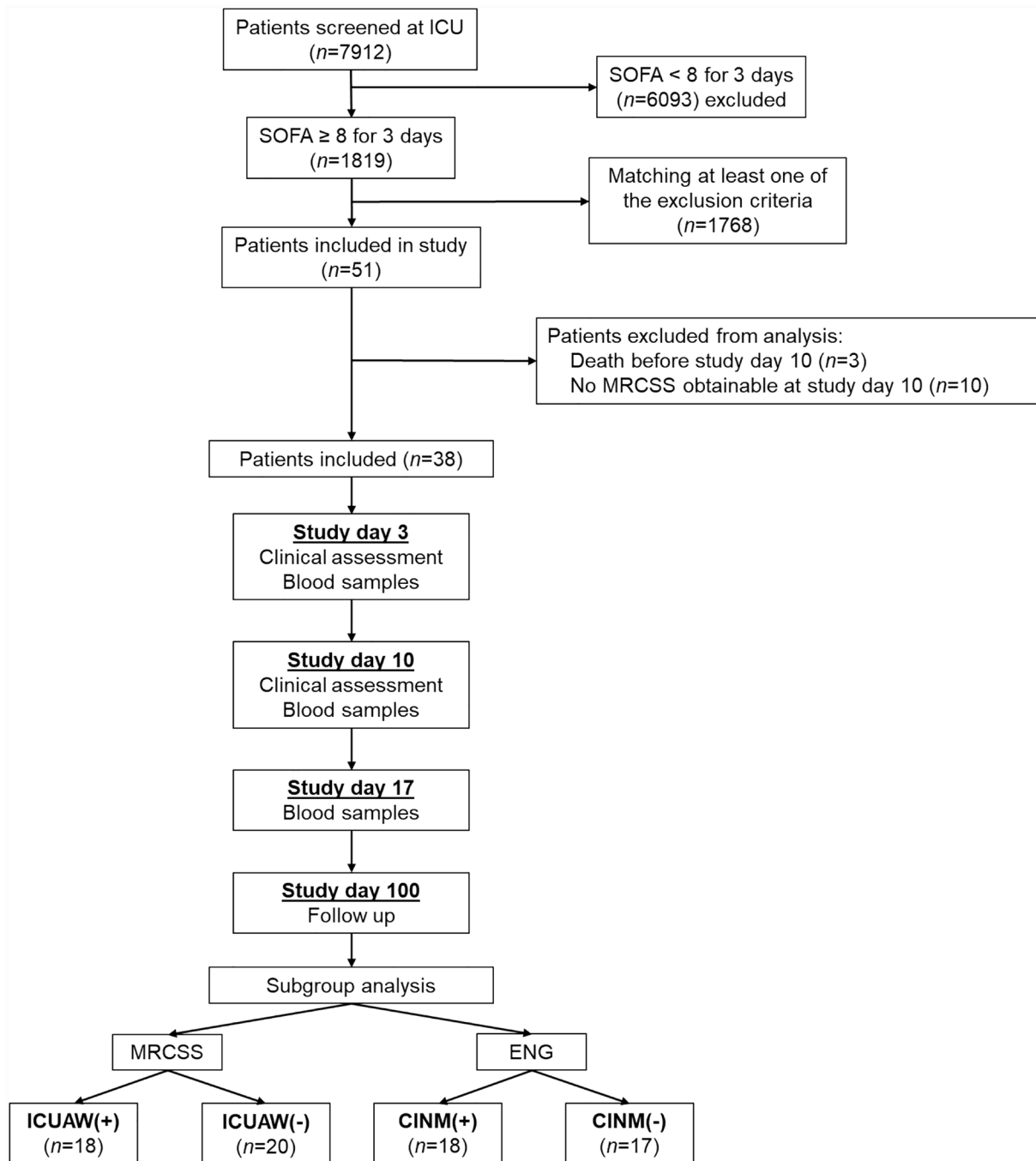


Fig. 1 Study flowchart. CINM critical illness neuromyopathy, ENG electroneurography, ICU intensive care unit, ICUAW(+) patients with intensive-care-unit-acquired weakness, ICUAW(–) patients without intensive-care-unit-acquired weakness, MRCSS Medical Research Council Sum Score, SOFA Sequential Organ Failure Assessment score

dysfunction. The SOFA scores and the MRCSS (Fig. 2) at study days 3 and 10 were significantly higher in patients with ICUAW and CINM compared with the corresponding negative groups. No differences in the

degree of neurological impairment represented by the mRS and overall survival were observed between the groups. Only the BI 3 months after enrollment was significantly lower in the ICUAW + group ($p=0.03$).

Table 1 Study population characteristics and outcome parameters

	ICUAW –	ICUAW +	<i>p</i> value	CINM –	CINM +	<i>p</i> value
Basic demographic data						
Total, <i>n</i> (%)	20 (42)	18 (38)	N/A	17 (49)	18 (51)	N/A
Male, <i>n</i> (%)	15 (75)	10 (56)	0.307	13 (77)	11 (61)	0.328
Age in years, mean (SD)	68.1 (14.2)	70.8 (11.5)	0.52	67.9 (15.2)	70.4 (11.6)	0.577
Cardiac and vascular surgery, <i>n</i> (%)	11 (55.0)	10 (55.6)	0.998	10 (59)	9 (50)	0.600
Thoracic surgery (noncardiac), <i>n</i> (%)	2 (10.0)	0 (0)	0.488	1 (6)	1 (6)	0.966
Visceral surgery, <i>n</i> (%)	3 (15.0)	5 (27.8)	0.438	2 (12)	4 (22)	0.411
Trauma surgery, <i>n</i> (%)	2 (10.0)	1 (5.5)	0.989	2 (12)	1 (6)	0.512
General surgery, <i>n</i> (%)	1 (5.0)	0 (0)	0.999	1 (6)	1 (6)	0.966
Urology, <i>n</i> (%)	1 (5.0)	0 (0)	0.999	1 (6)	0	0.485
Medical, <i>n</i> (%)	0 (0)	2 (11.0)	0.218	0	2 (11)	0.486
Sepsis, <i>n</i> (%)	3 (15)	4 (22)	0.687	3 (17)	0	0.104
APACHE II, mean (SD)	23.7 (6.2)	26.2 (3.7)	0.14	25.1 (4.9)	25.3 (5.7)	0.716
SOFA day 3, mean (SD)	10.1 (2.3)	12.9 (2.8)	0.001	10.1 (2.4)	12.6 (2.7)	0.006
SOFA day 10, mean (SD)	3.6 (3.0)	6.6 (2.8)	0.001	3.3 (2.5)	6.3 (2.9)	0.001
MRCSS day 3, mean (SD)	51.2 (12.4)	30.5 (8.7)	0.01	50.3 (13.1)	30.5 (8.7)	0.017
MRCSS day 10, mean (SD)	55.4 (4.2)	29.5 (12.9)	<0.0001	56.0 (4.2)	30.7 (13.9)	<0.001
Dialysis needed, <i>n</i> (%)	7 (35.0)	5 (27.8)	0.734	5 (29)	4 (22.2)	0.551
Creatinine (μmol/L) day 3, median (IQR)	126.0 (104.0–297.0)	114.5 (83.0–218.0)	0.362	125.5 (89.3–302.0)	114.5 (89.9–221.3)	0.617
Creatinine (μmol/L) day 10, median (IQR)	129.0 (74.1–275.0)	106.5 (76.7–182.3)	0.693	112.5 (64.3–297.0)	101.2 (70.8–180.8)	0.756
Creatinine (μmol/L) day 17, median (IQR)	129.5 (73.1–175.5)	86.5 (60.3–157.0)	0.315	131.0 (68.1–253.0)	86.4 (59.2–135.0)	0.169
Outcome parameters						
mRS after 3 months, mean (SD)	2.1 (2.6)	3.1 (2.2)	0.32	1.7 (2.5)	2.9 (2.2)	0.164
Barthel Index at admission, mean (SD)	97.5 (4.4)	93.9 (16.4)	0.68	98.2 (3.5)	93.4 (16.4)	1.0
Barthel Index after 3 months, mean (SD)	87.9 (25.5)	63.7 (36.0)	0.03	93.8 (13.2)	70.0 (32.2)	0.02
28-day survival, <i>n</i> (%)	18 (90)	18 (100)	0.49	15 (88)	18 (100)	0.134
3-month survival, <i>n</i> (%)	16 (80)	16 (88.9)	0.66	13 (77)	15 (83)	0.576

Significant *p* values are marked in bold

APACHE II, Acute Physiology and Chronic Health Evaluation II score, CINM, critical illness neuromyopathy, ICUAW, intensive-care-unit-acquired weakness, IQR, interquartile range, MRCSS, Medical Research Council Sum Score, mRS, modified Rankin Scale, N/A, not applicable, SOFA, Sequential Organ Failure Assessment score

Skeletal muscle biomarker profiles

Blood levels of skeletal muscle biomarkers were similar between the subgroups at study days 3, 10, and 17 (Table 2). Only FABP3 levels tended to be higher in ICUAW – compared with ICUAW + patients but sharply missed significance (median 14,785 [IQR 5836–24,000] pg/mL vs. 4125 [IQR 1089–15,536] pg/mL, $p=0.059$). Except for CAFE, concentrations of FABP3, GDF15, syndecan 1, and skeletal muscle troponin I were significantly higher in ICUAW + and ICUAW – patients compared with healthy controls (Supplementary Table 2).

Cytokine and inflammatory biomarker profiles

Biomarker levels of both ICUAW + and ICUAW – patients were elevated in similar patterns compared with healthy controls (Supplementary Table 3). Except for IL-1 α at study day 17 (median 16.7 [IQR 5.9–27.3] pg/mL vs. 7.9 [IQR 2.7–12.7] pg/mL, $p=0.023$) in the ICUAW subgroup comparisons and IL-10 at study day 17 (median 4.7 [IQR 2.2–5.2] pg/mL

vs. 1.8 [IQR 1.4–3.3] pg/mL, $p=0.038$) in patients with and without CINM, we found no statistically significant differences in biomarker levels between the subgroups (Table 3).

Longitudinal assessment of biomarker profiles

Results from longitudinal monitoring of skeletal muscle biomarker profiles are presented in Fig. 3. FABP3 levels decreased significantly in both ICUAW + and CINM + subgroups until study day 17. In contrast, biomarker concentrations tended to increase between study days 10 and 17 in ICUAW – and CINM – patients. Similar patterns were observed for GDF15 but without statistical significance over time.

Correlations of biomarker levels in patients with and without ICUAW

Skeletal muscle biomarkers correlated positively with measures of disease severity in both study groups (Figs. 4 and 5). In ICUAW + patients, GDF15, CAFE, and syndecan

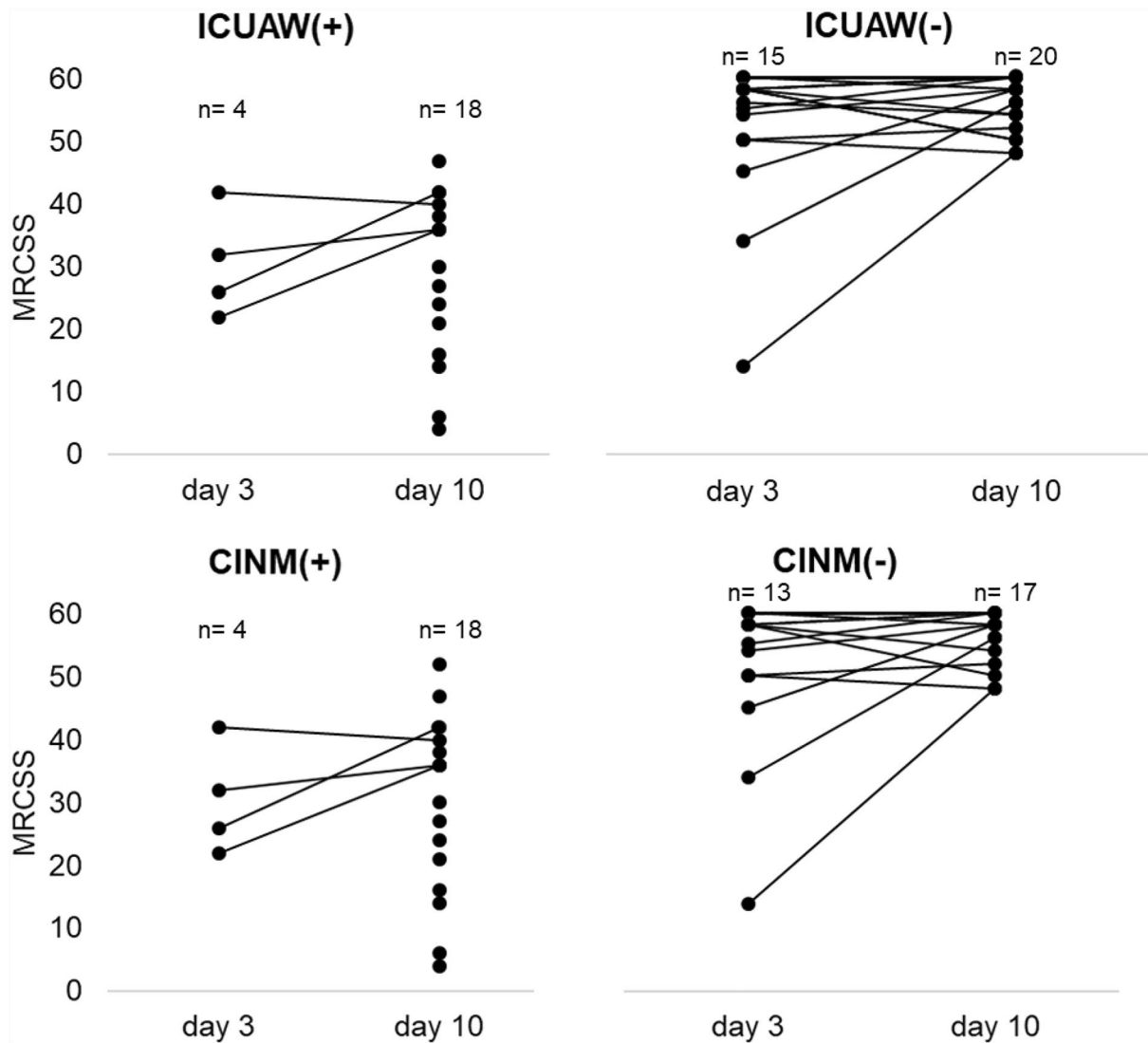


Fig. 2 Muscle strength assessment. CINM critical illness neuromyopathy, ICUAW intensive-care-unit-acquired weakness, MRCSS Medical Research Council Sum Score

1 levels correlated with the early SOFA score at day 3. Similarly, multiple positive correlations with overall disease severity assessments and outcome parameters in ICUAW – patients were observed. FABP3 at day 3 correlated with the Acute Physiology and Chronic Health Evaluation (APACHE II) score and the SOFA score. Elevated syndecan 1 serum concentrations were associated with a higher APACHE II score. Conversely, skeletal muscle troponin I was negatively correlated with the APACHE II score. Neither in ICUAW + nor in ICUAW – patients were significant correlations found between skeletal muscle biomarkers and muscle strength.

A correlation analysis revealed multiple moderate to strong correlations of inflammatory biomarkers with

distinct measures of disease severity. IL-4 levels at study day 3 as well as IL-1 β levels at study day 10 correlated with the SOFA score at days 3 and 10 (IL-4: $r=0.52$, $p=0.037$; IL-1 β : $r=-0.64$, $p=0.014$) in ICUAW + patients. In contrast, in ICUAW – patients, IL-13 levels were negatively correlated with the APACHE II score at all study days (day 3: $r=-0.40$, $p=0.09$; day 10: $r=-0.60$, $p=0.007$; day 17: $r=-0.71$, $p=0.033$).

Correlations of biomarker levels in patients with and without CINM

Similar to the ICUAW subgroups, skeletal muscle biomarker levels of CINM + patients correlated with the SOFA score at day 3 (CAF day 3: $r=0.58$, $p=0.03$;

Table 2 Skeletal muscle biomarker levels

Muscle and endothelial biomarkers	ICUAW −, median (IQR)	ICUAW +, median (IQR)	<i>p</i> value	CINM −, median (IQR)	CINM +, median (IQR)	<i>p</i> value
CAF (pg/mL) day 3	106.5 (65.9–127.8)	152.5 (72.1–230.5)	0.120	97.8 (52.3–128.4)	129.5 (62.5–222.5)	0.143
CAF (pg/mL) day 10	79.1 (54.8–105.1)	72.4 (44.6–226.9)	0.937	78.9 (36.5–103.6)	72.4 (41.5–215.4)	0.853
CAF (pg/mL) day 17	76.1 (25.9–105.4)	90.5 (15.6–311.0)	0.452	76.1 (22.9–105.4)	70.2 (14.2–273.8)	0.730
FABP3 (pg/mL) day 3	34,525 (14,850–102,650)	25,900 (17,103–87,050)	0.843	67,100 (10,650–117,500)	30,050 (15,077–10,0025)	0.925
FABP3 (pg/mL) day 10	7047 (5055–31,125)	13,125 (5763–22,288)	0.640	5805 (3225–34,750)	13,125 (4594–26,413)	0.766
FABP3 (pg/mL) day 17	14,785 (5836–24,000)	4125 (1089–15,537)	0.059	16,910 (5836–33,300)	4125 (1195–16,350)	0.081
GDF15 (pg/mL) day 3	14,281 (4569–23,738)	13,448 (6213–27,813)	0.654	12,210 (4050–17,250)	8633 (5114–25,738)	0.752
GDF15 (pg/mL) day 10	7665 (3914–20,550)	10,610 (5168–22,556)	0.579	6630 (3,153–23,175)	8355 (4221–19,269)	0.644
GDF15 (pg/mL) day 17	11,799 (8545–16,288)	9790 (2939–13,475)	0.346	11,799 (8545–16,288)	7900 (2505–14,600)	0.245
Syndecan 1 (pg/mL) day 3	278.2 (206.0–350.6)	310.0 (289.9–365.1)	0.384	294.7 (140.9–378.3)	306.2 (247.0–359.7)	0.857
Syndecan 1 (pg/mL) day 10	317.5 (224.3–379.2)	328.1 (279.7–378.8)	0.739	318.2 (206.9–379.4)	321.1 (243.1–377.8)	0.914
Syndecan 1 (pg/mL) day 17	369.5 (331.1–378.5)	374.1 (281.1–375.9)	0.764	369.5 (205.5–378.5)	374.1 (306.1–376.3)	0.905
Troponin I (pg/mL) day 3	4.1 (3.5–4.7)	3.9 (3.6–4.2)	0.532	4.1 (3.3–4.7)	3.9 (3.4–4.4)	0.517
Troponin I (pg/mL) day 10	4.1 (3.6–4.6)	4.0 (3.5–4.2)	0.498	4.0 (3.2–4.8)	4.0 (3.5–4.4)	0.509
Troponin I (pg/mL) day 17	4.1 (3.8–4.4)	3.7 (3.5–4.4)	0.502	4.0 (3.3–4.4)	3.7 (3.4–4.5)	0.872

CAF, C-terminal agrin filament; CINM, critical illness neuromyopathy; FABP, fatty acid binding protein; GDF, growth and differentiation factor; ICUAW, intensive-care-unit-acquired weakness; IQR, interquartile range

FABP3 day 17: $r=0.57$, $p=0.025$; GDF15 day 3: $r=0.67$, $p=0.005$ and day 10 (CAF day 17: $r=0.59$, $p=0.034$) as well as with the BI at 3 months (FABP3 day 10: $r=-0.53$, $p=0.041$). In CINM− patients, we observed multiple correlations of biomarker levels with outcome data, including the APACHE II score (FABP3 day 3: $r=0.58$, $p=0.024$; GDF15 day 10: $r=0.51$, $p=0.036$; syndecan 1 day 3: $r=0.59$, $p=0.017$; syndecan 1 day 10: $r=0.59$, $p=0.016$), the mRS at day 100 (GDF15 day 3: $r=0.69$, $p=0.008$; GDF15 day 10: $r=0.62$, $p=0.015$), and the SOFA score at day 10 (FABP3 day 10: $r=0.74$, $p<0.001$; FABP3 day 17: $r=0.80$, $p=0.015$; GDF15 day 3: $r=0.77$, $p=0.003$; GDF15 day 10: $r=0.86$, $p<0.001$; GDF15 day 17: $r=0.743$, $p=0.035$; syndecan 1 day 3: $r=0.55$, $p=0.026$; syndecan 1 day 10: $r=0.56$, $p=0.025$). Regarding the inflammatory biomarkers, no relevant correlations between biomarker levels and outcome parameters were observed.

Discussion

We investigated the diagnostic value of a novel blood-based biomarker panel to differentiate between critically ill patients with and without acquired neuromuscular weakness. Skeletal muscle and inflammatory biomarkers are elevated in critically ill patients compared with healthy controls but do not discriminate between patients with and without ICUAW or CINM. Comparison of

absolute biomarker concentrations between groups and longitudinal changes of blood biomarker levels over the course of 17 days revealed similar patterns in patients with and without ICUAW and CINM, despite clinically significant differences of overall limb muscle strength. Importantly, both subgroups were comparable for age, sex distribution, and renal dysfunction, which represent typical confounding factors in biomarker analysis. Nevertheless, other factors might have contributed to these results. First, although the MRCSS has been proven to be a valid and reliable clinical tool for the assessment of overall muscle strength in critically ill patients [25], a detailed differentiation between predominant neuropathy and/or myopathy is not possible and requires more elaborate methods [2]. Therefore, the reduction in muscle strength within the ICUAW+ group might be due to a higher proportion of patients with predominant CIP rather than direct muscle damage. Therefore, we additionally performed ENG examinations, which showed similar observations. Unfortunately, serum biomarkers of neuroaxonal damage, such as neurofilament light and heavy chains, have not been assessed within the present trial but also showed no clear benefit in the detection of patients with ICUAW in former studies [26]. Second, except for troponin I, which is an integral part of the sarcomere troponin complex, no other structural components of the actual skeletal muscle contractile apparatus

Table 3 Cytokine and inflammatory biomarker levels

Cytokines and inflammatory biomarkers	ICUAW −, median (IQR)	ICUAW +, median (IQR)	p value	CINM −, median (IQR)	CINM +, median (IQR)	p value
CRP (mg/L) day 3	202.0 (105.0–272.5)	213.0 (129.0–243.3)	0.788	201.0 (75.1–316.5)	216.5 (119.1–259.8)	0.558
CRP (mg/L) day 10	121.0 (75.2–193.0)	68.9 (45.1–118.5)	0.068	110.3 (74.–05.8)	73.5 (39.0–135.3)	0.120
WBC (10 ⁹ /L) day 3	11.9 (9.1–18.2)	14.0 (9.1–16.2)	0.989	11.3 (8.6–18.1)	12.9 (8.8–16.4)	0.741
WBC (10 ⁹ /L) day 10	12.3 (9.4–17.9)	13.4 (11.1–16.4)	0.726	12.1 (8.6–18.5)	13.4 (10.8–16.8)	0.448
PCT (ng/mL) day 3	1.0 (0.4–4.9)	2.5 (1.0–9.0)	0.193	0.9 (0.4–6.2)	2.5 (0.9–10.6)	0.181
PCT (ng/mL) day 10	0.2 (0.1–0.8)	0.3 (0.2–0.7)	0.334	0.2 (0.1–0.8)	0.3 (0.2–0.8)	0.235
IFN γ (pg/mL) day 3	1.9 (1.1–2.3)	1.6 (1.1–1.8)	0.579	1.8 (0.8–2.3)	1.5 (1.0–2.1)	0.564
IFN γ (pg/mL) day 10	1.7 (1.2–2.5)	1.5 (1.0–2.7)	0.812	1.7 (1.0–2.5)	1.7 (1.0–2.8)	0.801
IFN γ (pg/mL) day 17	1.7 (0.6–2.4)	1.2 (0.6–2.0)	0.460	1.4 (0.4–2.5)	1.5 (0.5–2.2)	0.616
IL-10 (pg/mL) day 3	3.3 (3.0–3.7)	2.6 (2.0–3.3)	0.099	3.4 (2.9–4.0)	2.6 (1.8–4.1)	0.122
IL-10 (pg/mL) day 10	2.9 (1.8–4.3)	2.2 (1.6–3.2)	0.366	2.9 (1.7–4.4)	2.2 (1.4–4.4)	0.517
IL-10 (pg/mL) day 17	4.0 (2.2–4.8)	1.7 (1.4–2.6)	0.079	4.7 (2.2–5.2)	1.8 (1.4–3.3)	0.038
IL-13 (pg/mL) day 3	2.1 (1.4–3.1)	1.5 (1.4–1.8)	0.188	2.0 (1.4–3.3)	1.6 (1.4–1.8)	0.387
IL-13 (pg/mL) day 10	1.9 (1.6–3.5)	1.6 (1.3–2.8)	0.358	1.8 (1.5–3.5)	1.8 (1.4–3.6)	0.759
IL-13 (pg/mL) day 17	2.4 (1.4–2.7)	1.5 (1.2–1.7)	0.089	2.4 (1.4–2.7)	1.5 (1.1–2.3)	0.192
IL-1 α (pg/mL) day 3	5.4 (4.4–9.1)	5.7 (4.3–7.1)	0.558	5.4 (3.5–8.6)	5.8 (4.5–8.3)	0.857
IL-1 α (pg/mL) day 10	6.7 (4.8–9.6)	5.5 (4.3–8.8)	0.334	6.4 (4.7–10.3)	5.5 (3.5–12.3)	0.494
IL-1 α (pg/mL) day 17	16.7 (5.9–27.3)	7.9 (2.7–12.7)	0.023	16.8 (1.5–38.2)	8.1 (2.7–14.0)	0.307
IL-1 β (pg/mL) day 3	2.9 (0.5–7.4)	1.4 (0.9–4.5)	0.716	2.3 (0.4–7.4)	1.5 (0.9–6.3)	0.792
IL-1 β (pg/mL) day 10	2.0 (0.5–8.5)	2.1 (0.6–6.7)	0.899	1.7 (0.5–12.7)	3.0 (0.6–9.3)	0.603
IL-1 β (pg/mL) day 17	7.9 (2.0–32.6)	5.0 (0.9–20.4)	0.571	7.9 (2.0–32.6)	9.9 (1.3–23.6)	0.941
IL-4 (pg/mL) day 3	2.0 (0.6–2.7)	1.6 (0.9–3.3)	0.974	1.2 (0.5–2.8)	1.7 (0.8–3.6)	0.624
IL-4 (pg/mL) day 10	2.5 (0.6–4.4)	1.0 (0.5–4.3)	0.486	2.2 (0.4–4.7)	1.3 (0.6–5.4)	0.943
IL-4 (pg/mL) day 17	2.3 (0.4–13.3)	1.5 (0.6–3.5)	0.378	2.3 (0.2–13.3)	2.4 (0.9–4.2)	0.941
IL-6 (pg/mL) day 3	12.0 (6.9–23.8)	10.4 (7.4–30.1)	0.643	14.2 (6.8–27.7)	10.4 (7.4–31.6)	0.880
IL-6 (pg/mL) day 10	9.3 (8.0–16.1)	8.4 (5.1–12.7)	0.231	9.3 (7.0–16.6)	8.5 (5.3–12.9)	0.553
IL-6 (pg/mL) day 17	13.6 (11.4–21.7)	9.4 (7.8–15.7)	0.144	13.6 (11.4–21.7)	11.6 (7.8–17.2)	0.217
IL-8 (pg/mL) day 3	9.4 (7.7–11.3)	10.2 (6.0–11.7)	0.526	10.0 (7.7–12.5)	7.7 (5.5–12.1)	0.214
IL-8 (pg/mL) day 10	10.1 (7.9–12.3)	8.3 (5.3–10.0)	0.178	10.1 (7.6–12.5)	8.6 (5.2–13.5)	0.296
IL-8 (pg/mL) day 17	16.6 (13.3–28.9)	11.3 (9.0–20.0)	0.245	16.6 (13.3–29.6)	11.3 (8.2–22.2)	0.192
MCP1 (pg/mL) day 3	138.7 (76.0–218.9)	82.8 (61.8–262.4)	0.579	156.4 (89.1–234.5)	77.3 (58.1–266.6)	0.280
MCP1 (pg/mL) day 10	109.0 (67.5–158.6)	60.8 (43.2–119.4)	0.261	138.0 (56.9–192.6)	67.5 (43.2–152.0)	0.249
MCP1 (pg/mL) day 17	217.9 (197.4–275.0)	229.3 (168.7–290.0)	0.913	217.9 (197.4–275.0)	229.3 (168.5–309.3)	0.916
TNF α (pg/mL) day 3	4.3 (3.7–4.8)	4.7 (3.1–5.6)	0.669	4.1 (3.4–4.9)	4.7 (3.1–5.7)	0.589
TNF α (pg/mL) day 10	4.4 (3.3–5.5)	4.5 (3.2–6.5)	0.837	4.2 (3.3–5.4)	5.0 (3.2–8.2)	0.428
TNF α (pg/mL) day 17	5.3 (4.1–12.3)	4.3 (1.6–7.2)	0.250	5.3 (4.1–12.6)	5.8 (1.2–7.6)	0.597

Significant *p* values are marked in bold

CINM, critical illness neuromyopathy, CRP, C-reactive protein, ICUAW, intensive-care-unit-acquired weakness, IQR, interquartile range, IFN, interferon, IL, interleukin, MCP1, monocyte chemoattractant protein 1, PCT, procalcitonin, TNF, tumor necrosis factor

have been assessed as potential serum biomarkers here. Myosin filaments are known to be severely affected in ICUAW and CINM by insufficient protein synthesis and enhanced protein degradation, even within the first days after the onset of critical illness [27]. However, proteins within other structural and functional compartments of the skeletal muscle might be more resilient to these early

alterations. FABP3, also known as heart-type fatty-acid-binding protein, is abundantly expressed in cardiomyocytes and slow-type skeletal muscle fibers [28]. Elevated serum levels of FABP3 have been observed in polymyositis/dermatomyositis [29] and generalized sarcopenia [30]. Hereby, recent animal data point toward a critical role of FABP3 for muscle atrophy and endothelial dysfunction

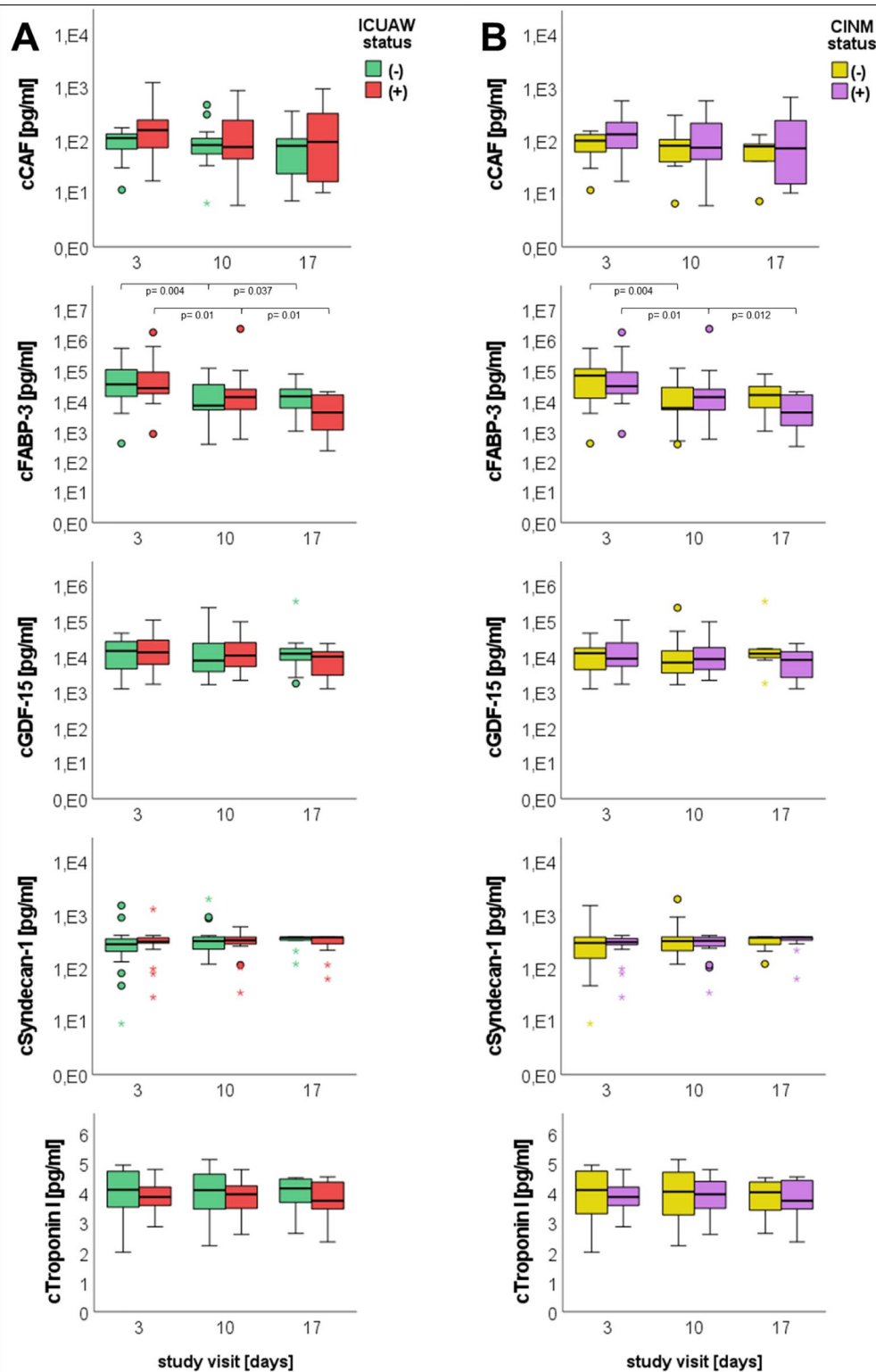
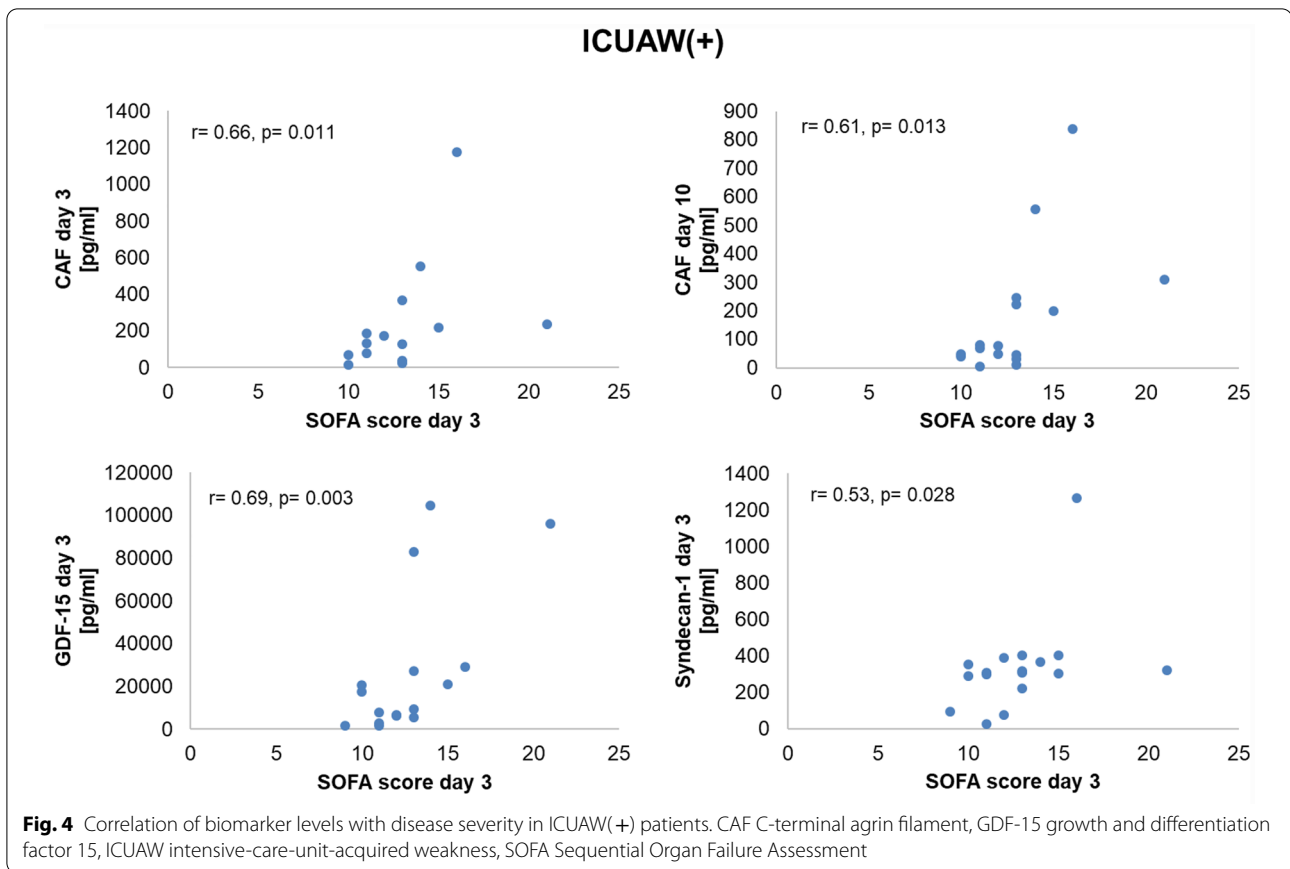


Fig. 3 Longitudinal assessment of blood-based biomarker levels. **a**, Comparison between patients with (red box plots) and without (green box plots) ICUAW. **b**, Comparison between patients with (purple box plots) and without (yellow box plots) CINM. Box plots depict the median and the first and third quartiles, whiskers show values within the 1.5×interquartile range. Dots represent values between 1.5×and 3.0×and stars values beyond the 3.0×of the interquartile range. CAF C-terminal agrin filament, CINM critical illness neuromyopathy, FABP-3 fatty-acid-binding protein 3, GDF-15 growth and differentiation factor 15, ICUAW intensive-care-unit-acquired weakness (Color figure online)



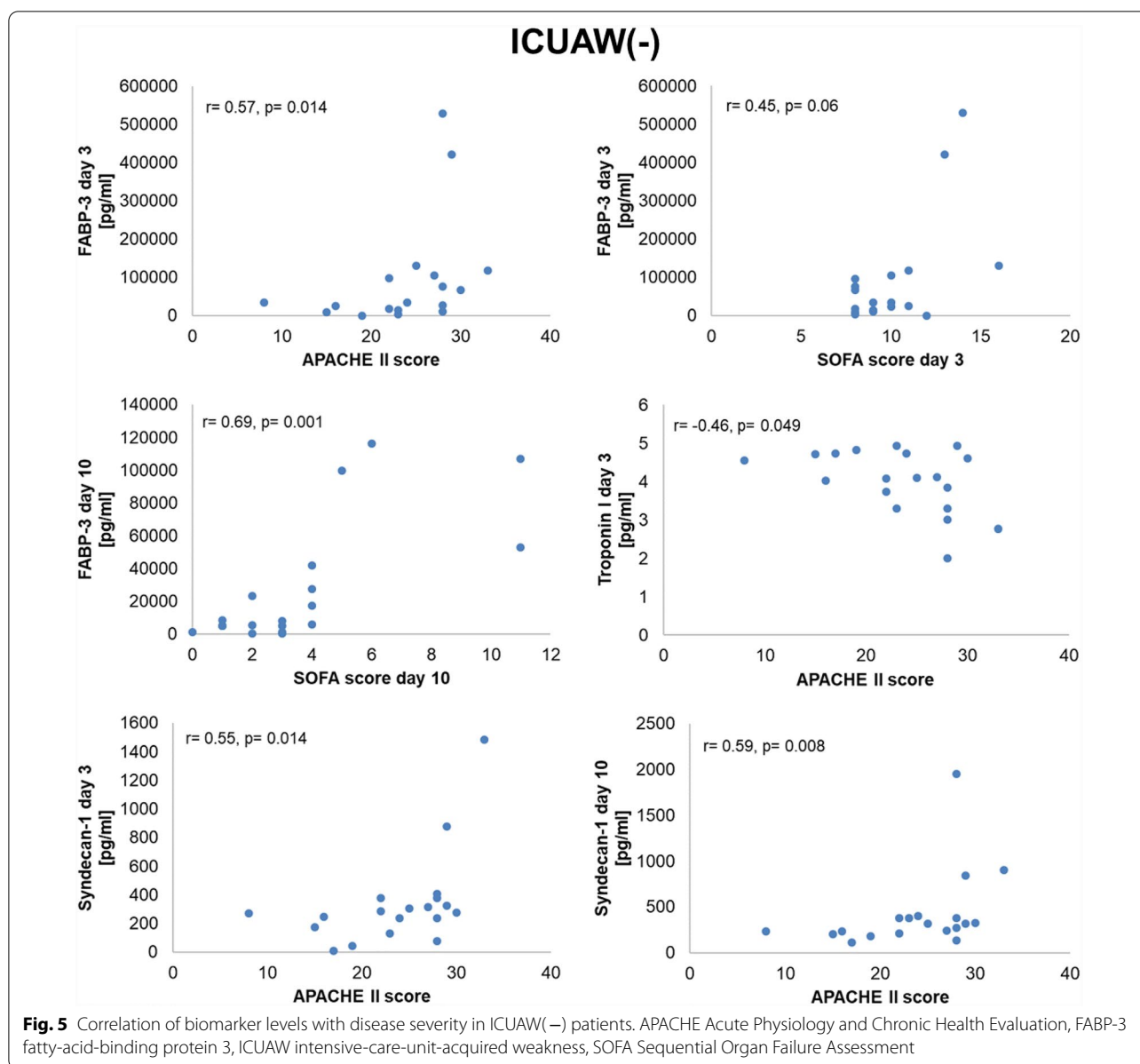
[31, 32]. These results might point toward a possible role of FABP3 in the development of sepsis-induced skeletal muscle microvascular dysfunction, potentially contributing to CIM [33]. In our study, serum levels of FABP3 were markedly elevated in both subgroups at the first assessment but progressively declined in ICUAW+ and CINM+ patients over time. In contrast, FABP3 serum levels elevated again significantly in ICUAW- and CINM- patients at study day 17. However, to date, evidence regarding the significance of FABP3 in critical-illness-induced muscle weakness and muscle atrophy is lacking.

CAF is an integral component of the neuromuscular junction and has been targeted as a potential serum biomarker in age-related sarcopenia, muscle disuse, and chronic muscle wasting and after stroke [34, 35]. Within the present study, serum CAF levels tended to be higher in patients with ICUAW compared to the ICUAW- group early during critical illness but without statistical significance, indicating similar neuromuscular impairment.

In a study by Bloch et al. [36], serum levels of GDF15 protein and GDF15 muscle mRNA were elevated in 20 critically ill patients with ICUAW compared with healthy

controls, and the GDF15 serum levels of about 7000 pg/mL are comparable to the results in the present study. In contrast, Xie et al. [37] compared serum GDF15 in 50 patients with ICUAW and 45 patients without ICUAW and found significantly higher levels of serum GDF15 in patients with clinically relevant neuromuscular weakness after 7 days of critical illness. In our study, GDF15 levels were not different within both subgroups, but serum levels were up to sixfold higher compared to the results by Xie and coworkers [37]. This might be explained by the overall high disease severity of our study cohort, with markedly elevated APACHE II and SOFA scores compared to former studies.

Cytokines and biomarkers of inflammation also revealed no differences between patients with and without ICUAW or CINM. A possible explanation might be the similar proportion of patients with sepsis in both groups, indicating a comparable degree of systemic inflammation, regardless of underlying neuromuscular dysfunction. As for the skeletal muscle biomarker panel, we found no relevant correlations between serum cytokine levels and muscle strength, but we did find relevant correlations with disease severity scores. These results are comparable with a study by Winkelmann et al.



[38], who investigated IL-8, IL-15, and TNF α serum levels in critically ill patients and observed a positive association between IL-8 concentrations after mobilization and physical activity scores after ICU discharge but not with muscle strength.

A major strength of the present study is the investigation of a broad panel of inflammatory, neurovascular, and neuromuscular biomarkers. In total, we measured 18 different blood-based biomarkers, which is, to our knowledge, the largest biomarker panel ever assessed in the context of ICUAW and CINM. Furthermore, biomarker levels were longitudinally measured over 17 days after ICU admission, which allowed us to compare acute and

subacute changes of biomarker concentrations related to the clinical outcome. Another advantage of the present study is the combined clinical and electrophysiological assessment of patients, providing two subgroup comparisons of biomarker data. However, some limitations have to be mentioned. Because of the relatively small study cohort, the present results may not have reached sufficient statistical power to show possible significant differences in biomarker levels between the individual groups. However, a power analysis was not performed, as the study was conceptualized as a pilot trial. Although multiple group comparisons at different study visits were calculated, we did not apply a statistical correction

of the p value for multiple testing (e.g., a Bonferroni correction) because all comparisons were already statistically insignificant. Furthermore, despite good validation in ICU patients, the MRCSS remains a subjective measure, assessing skeletal muscle strength only in part and depending on the operator's experience as well as patients' compliance. Other tests such as dynamometry might have provided a more detailed assessment of overall muscle strength, but this assessment would also be dependent on patients' cooperation, which supports the main idea of the present study to identify blood-based biomarkers independently indicating neuromuscular impairment. Furthermore, baseline assessment of patients before the onset of critical illness was not possible because of the study design.

Conclusions

Blood-based inflammatory, neurovascular, and neuromuscular biomarkers are frequently elevated in critically ill patients but cannot distinguish between patients with and without acquired neuromuscular weakness. As far as we can conclude from our analysis, biomarker levels are not helpful to identify and monitor patients with ICUAW, although a certain correlation with overall disease severity was observed. Other diagnostic tools such as neuromuscular ultrasound might offer better opportunities in the future and should be analyzed in large-scale studies [7].

Supplementary Information

The online version contains supplementary material available at <https://doi.org/10.1007/s12028-024-02050-x>.

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Author contributions

Conceptualization: JE, RP, and UW; methodology: JE, RP, and D-CF; software: FK and RP; validation: JE and RP; formal analysis: FK, RP, and FL; investigation: FK, JE, RP, RB, LD, FL, AR, and KP; resources: RP, D-CF, and AR; data curation: FK, JE, and RP; writing (original draft preparation): FK and JE; writing (review and editing): all authors; visualization: FK; supervision: JE and RP; project administration: FK, JE, and RP; funding acquisition: FK and JE. All authors read and approved the final version of the manuscript.

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Data Availability

Data are available on reasonable request.

Conflicts of Interest

UW has received speaker honoraria and travel grants from Bristol-Myers Squibb, Boehringer Ingelheim Pharma, Daiichi-Sankyo, Ipsen Pharma, Merz Pharmaceuticals, and Pfizer Pharma and an unrestricted research grant from Merz Pharmaceuticals independent from this study. He serves as joint editor-in-chief of the *European Journal of Ultrasound* (Thieme, Stuttgart, Germany). JE received financial support for attending study meetings from B. Braun Melsungen AG independent from this study. FK, FL, D-CF, AR, KP, LD, RP, and RB have nothing to declare.

Ethical Approval/Informed Consent

The study was performed in accordance with the Declaration of Helsinki and was approved by the local ethics committee of the University of Rostock (ethics identifier AS 2016–0016). Written informed consent for participation was given by volunteers, patients, or a legal representative before enrollment.

Clinical Trial Registration

ClinicalTrials.gov identifier: NCT02706314.

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