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Toro, Camilo Jain, Sonia Sun, Shelly <u>et al.</u>

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Association of Brain Injury Biomarkers and Circulatory Shock Following Moderate-Severe Traumatic Brain Injury: A TRACK-TBI Study

Camilo Toro, BA¹, Sonia Jain, PhD², Shelly Sun, MS², Nancy Temkin, PhD^{3,4}, Jason Barber, MS⁴, Geoffrey Manley, MD, PhD⁵, Jordan M. Komisarow, MD⁶, Tetsu Ohnuma, MD, MPH, PhD^{7,8}, Brandon Foreman, MD⁹, Frederick Korley, MD, PhD¹⁰, Michael L. James, MD^{7,11}, Daniel Laskowitz, MD, MHS^{7,11}, Monica S. Vavilala, MD¹², Adrian Hernandez, MD, MHS¹³, Joseph P. Mathew, MD, MHSc, MBA⁷, Amy J. Markowitz, JD⁵, Vijay Krishnamoorthy, MD, MPH, PhD^{7,8,14},

TRACK-TBI Investigators

⁷Department of Anesthesiology, Duke University. Durham, NC.

⁶Department of Neurosurgery, Duke University. Durham, NC.

¹¹Department of Neurology, Duke University. Durham, NC.

¹³Department of Medicine, Duke University. Durham, NC.

¹⁴Department of Population Health Sciences, Duke University. Durham, NC.

⁸Critical Care and Perioperative Population Health Research (CAPER) Unit, Department of Anesthesiology, Duke University. Durham, NC.

³Department of Biostatistics, Anesthesiology and Pain Medicine, University of Washington. Seattle, WA.

⁴Department of Neurosurgery, Anesthesiology and Pain Medicine, University of Washington. Seattle, WA.

⁹Department of Neurology and Rehabilitation Medicine, University of Cincinnati. Cincinnati, OH.

¹⁰Department of Emergency Medicine, University of Michigan. Ann Arbor, MI.

¹Duke University School of Medicine. Durham, NC.

²Biostatistics Research Center, Herbert Wertheim School of Public Health and Human Longevity Science, University of California, San Diego. San Diego, CA.

⁵Brain and Spinal Injury Center, University of California, San Francisco. San Francisco, CA.

SUPPLEMENTARY MATERIAL

Supplemental digital content 1

Supplementary Tables 1–5 showing temporal changes in biomarker levels Supplemental Digital Content 1.pdf

Corresponding Author: Vijay Krishnamoorthy, MD, MPH, PhD, Duke University Hospital, 2301 Erwin Road, Durham, NC 27710, 312-403-0718, Vijay.krishnamoorthy@duke.edu.

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¹²Department of Anesthesiology and Pain Medicine, and Harborview Injury Prevention and Research Center, University of Washington, Seattle, WA

Abstract

Introduction—Early circulatory shock following traumatic brain injury (TBI) is a multifactorial process; however, the impact of brain injury biomarkers on the risk of shock has not been evaluated. We examined the association between neuronal injury biomarker levels and the development of circulatory shock following moderate-severe TBI.

Methods—In this retrospective cohort study, we examined adults with moderate-severe TBI (Glasgow Coma Scale score <13) enrolled in the TRACK-TBI study, an 18-center prospective TBI cohort study. The exposures were day-1 levels of neuronal injury biomarkers (glial fibrillary acidic protein, ubiquitin C-terminal hydrolase-L1 [UCH-L1], S100 calcium binding protein B [S100B], neuron specific enolase), and of an inflammatory biomarker (high-sensitivity C-reactive protein). The primary outcome was development of circulatory shock, defined as cardiovascular Sequential Organ Failure Assessment score 2 within 72 hours of admission. Association between day-1 biomarker levels and the development of circulatory shock was assessed with regression analysis.

Results—The study included 392 subjects, with a mean age of 40 years; 314 (80%) were male and 165 (42%) developed circulatory shock. Median [interquartile range] day-1 levels of UCH-L1 (994.8 [518.7–1988.2] pg/ml vs. 548.1 [280.2–1151.9] pg/ml; p <.0001) and S100B (0.47 ug/ml [0.25–0.88] vs. 0.27 [0.16–0.46] ug/ml; p <.0001) were elevated in those who developed early circulatory shock compared to those who did not. In multivariable regression, there were associations between levels of both UCH-L1 (odds ratio, 1.63 [95% confidence interval, 1.25–2.12]; p <.0005) and S100B (odds ratio, 1.73 [95% confidence interval 1.27–2.36]; p <.0005) with the development of circulatory shock.

Conclusion—Neuronal injury biomarkers may provide improved mechanistic understanding and possibly early identification of patients at risk for early circulatory shock following moderate-severe TBI.

Keywords

traumatic brain injury; circulatory shock; biomarkers

INTRODUCTION

Traumatic brain injury (TBI) is a leading cause of death and disability in the United States¹ and worldwide. Compared to mild TBI, moderate-severe TBI has been associated with a greater incidence of hypotension, increased mortality, and worse functional outcomes². The critical care management of moderate-severe TBI is aimed at limiting damage from the primary brain injury and reducing secondary brain injury, including by maintenance of hemodynamic stability and prevention of multi-organ dysfunction^{3 4}. Early circulatory shock following moderate-severe TBI is a multifactorial process driven by the underlying brain injury, cardiac dysfunction, and impaired vascular autoregulation. Circulatory shock has long been recognized for its association with poor clinical outcomes following TBI, including cerebral ischemia, disrupted cerebral hemodynamics, and increased mortality^{5–10}.

Therefore, improving early recognition, diagnosis and treatment of circulatory shock could represent an important therapeutic target to improve moderate-severe TBI outcomes.

Blood-based biomarkers have the potential to identify patients who may be at risk for clinical deterioration and would benefit from novel therapies aimed at specific pathophysiologic mechanisms. Current research on neural injury biomarkers, including glial fibrillary acidic protein (GFAP), ubiquitin C-terminal hydrolase-L1 (UCH-L1), S100 calcium binding protein B (S100B), and neuron specific enolase (NSE), and the inflammatory biomarker high-sensitivity C-reactive protein (hsCRP), demonstrate promise to assist in the management of TBI^{11-19} . Given that the development of circulatory shock is associated with primary brain injury severity, direct neuronal and glial injury may contribute to the dysfunction in circulatory autoregulation commonly seen following TBI²⁰. As such, examining the association of these biomarkers with hemodynamic failure may elucidate mechanistic pathways contributing to the development of circulatory shock following brain injury, and may be useful for identifying patients at risk of circulatory shock. No studies to date have been conducted to explore an association of these biomarkers with the development of circulatory shock following TBI. To address this gap, the aim of our study was to determine the association of neuronal injury and inflammatory biomarker levels with the development of early circulatory shock following moderate-severe TBI.

METHODS

We conducted a retrospective cohort study of adult patients enrolled in the Transforming Clinical Research and Knowledge in TBI (TRACK-TBI) study. TRACK-TBI was a prospective 18-center cohort study of patients evaluated in a level 1 trauma center Emergency Department within 24 hours of suffering blunt TBI, conducted between February 2014 and July 2018. This was a secondary retrospective analysis, as the current research question was not pre-specified when designing the original cohort. In addition to the collection of a multi-dimensional outcome battery over the year following injury, detailed hospital encounter data, including time-stamped diagnostic, pharmacy, and laboratory information, were collected in TRACK-TBI²¹. Subjects were excluded from the TRACK-TBI cohort if they met the following criteria: significant history of pre-existing conditions that would interfere with follow-up and outcome assessment; prisoners or patients in custody; pregnancy; on psychiatric hold; major debilitating baseline mental health disorders or major debilitating neurologic disease; participants in an interventional trial; or penetrating head or spinal cord injury with American Spinal Injury Association impairment scale score of C or worse²². Data were collected by trained research coordinators, using structured data collection tools. The present study was approved by the Institutional Review Board at Duke University (Reference 00100061).

Study population

We examined adults (age 17 years) in the TRACK-TBI cohort with moderate-severe TBI, defined as Glasgow Coma Scale (GCS) score <13 after resuscitation, who were admitted to an Intensive Care Unit (ICU) from the Emergency Department. To remove the confounding effects of extracranial injuries on the development of circulatory shock, we excluded patients

with significant extracranial injury, defined as non-head/neck Abbreviated Injury Scale score >3. Among these patients, we included only those who had biomarker data available on the day of hospital admission (day-1).

Exposure, outcomes, and covariates

The exposures were the levels of four day-1 neuronal injury biomarkers and an inflammatory biomarker; biomarker levels at days 3, 5, and 14 were also collected for patients with available information. Biomarker data collected were GFAP (pg/ml), UCH-L1 (pg/ml), S100B (ug/l), NSE (ng/ml), and hsCRP (mg/l). The first batch of GFAP and UCH-L1 concentrations (n=170) were measured using the prototype point-of-care i-STATTM Alinity[™] System. The second batch of GFAP and UCHL1 concentrations (n=222) were measured on the prototype core lab Abbott ARCHITECT® platform for faster throughput. For the i-STATTM AlinityTM assays, the test limit of detection (LoD) and limit of quantitation (LoQ) for GFAP were <15 pg/mL and <25 pg/mL respectively, and, for UCHL-1 <10 pg/mL and <20 pg/mL, respectively. For the ARCHITECT® assays, the GFAP LoD and LoQ were 2 pg/mL and 5 pg/mL, respectively, and the LoD and LoQ for UCHL-1 were10 pg/mL and 20 pg/mL, respectively. All samples were tested neat, without dilution, and in duplicate. Samples reading greater than the calibration range were reported as greater than the reportable range and were not diluted. ARCHITECT® values were converted to iSTAT equivalents using two previously derived equations: iSTAT= -12.36+1.02*ARCHITECT for GFAP (Spearman's correlation coefficient, 0.985) and iSTAT=-3.29+0.72*ARCHITECT for UCHL1 (Spearman's correlation coefficient, 0.933).²³ Biomarker measurements were blinded to outcome assessments. Biomarker levels were log-transformed for the statistical models because of the skewness of the distributions.

The primary outcome of interest was the development of circulatory shock, defined by the cardiovascular component of the Sequential Organ Failure Assessment (SOFA) score 2 within 72 hours of admission. The SOFA score was developed to assess acute morbidity in a range of organ systems (respiratory, cardiovascular, hepatic, coagulation, renal, and neurological), for which both the sub-scores and the total score are associated with prognosis²⁴. Covariates were ascertained by examination of time-stamped demographic, clinical, diagnostic, pharmacy, and laboratory information in the TRACK-TBI cohort; these included age, gender, race, ethnicity, injury cause, Injury Severity Score (ISS), computed tomography (CT) confirmed intracranial injury, CT Marshall Score, CT Rotterdam Score, and GCS upon arrival to the ED (3–8 vs. 9–12).

Statistical analysis

Descriptive statistics were used to examine demographics, clinical characteristics, and day-1 biomarker levels, stratified by early circulatory shock status. All biomarker levels were reported using median and interquartile range (IQR). Group comparisons used Wilcoxon Rank Sum test for continuous variables and Fisher's exact test for categorical variables. Categorical variables are reported as number (percentage) and continuous variables as mean and standard deviation (SD). Descriptive statistics were also used to examine biomarker levels at days-3, 5, and 14, stratified by early circulatory shock status. Multivariable logistic regression models were performed to assess the association between each of the day-1

biomarker levels and development of early circulatory shock. Models were fit with and without the following covariates: age, gender, race, non-head ISS, CT Rotterdam Score, and GCS. Odds ratios (OR) per unit increase in each biomarker level (in log scale) were presented with 95% confidence intervals (CI). A p-value < 0.05 was considered statistically significant. All analyses were performed using Statistical software R version 3.6 (http://www.r-project.org).

Results

The study population included 392 subjects, of which 165 (42%) developed circulatory shock (Figure 1). The mean age of participants was 39.9 ± 16.8 years; 313 (79.9%) participants were male, 307 (80.6%) participants were white, and 76 participants (19.9%) identified as Hispanic. The median ISS non-head/neck of the study cohort was 4 [IQR,1–10] and mean CT Rotterdam Score was 3.4 ± 1.2 . Mean GCS upon arrival was 5.8 ± 3.1 ; 74.2% of patient cohort had severe TBI (GCS 3–8) and 25.8% had moderate TBI (GCS 9–12). Details of the demographic and clinical characteristics of the study population stratified by development of circulatory shock within 72 hours of admission are shown in Table 1.

Biomarker levels

Figure 2 shows the median (25th/75th) percentiles of day 1 biomarkers, stratified by early circulatory status. Median day-1 UCH-L1 levels were significantly elevated in those who developed early circulatory shock compared to those who did not (994.8 [IQR, 518.7-1988.2] pg/ml vs. 548.1 [IQR, 280.2–1151.9] pg/ml, respectively; p <.0001) Similarly, S100b levels were significantly elevated in those who developed early circulatory shock (0.47 ug/ml [0.25–0.88] vs. 0.27 [0.16–0.46] ug/ml; p <.0001). Temporal changes in biomarker levels on days 1, 3, 5 and 14 are shown in Figure 3 and in the supplementary material (Supplemental digital content 1: Supplementary tables 1–5). There were no significant differences in day-1 GFAP levels between those who did and did not develop circulatory shock. GFAP levels were significantly elevated on days-3, 5, and 14 amongst those who developed circulatory shock, with levels in both groups gradually declining by day-14. UCH-L1 levels were significantly elevated on day-1 and day-3 in those who developed early circulatory shock, and levels gradually declined in both groups by day-14. S100b levels were significantly elevated on days-1, 3, and 5 in those who developed circulatory shock, and also gradually declined in both groups over 14 days. There were no significant between-group differences in NSE levels on days-1, 3, 5 and 14; in both groups, NSE levels declined between day-1 and day-5 and then increased by day-14. There were no significant between-group differences in day-1 hsCRP levels; however, day-3, 5, and 14 hsCRP levels were significantly increased in those who developed early shock. Compared to day-1, hsCRP levels were higher on day-3 and day-5 in both groups.

Association of day-1 biomarker levels with the development of early shock

Table 2 shows the association between day-1 biomarker levels and the development of early circulatory shock, before and after adjustment with covariates. Overall, 42% of the study population developed early circulatory shock. In multivariable logistic regression models, there were significant associations between day-1 levels of UCH-L1 (OR, 1.63 [95% CI

1.25-2.12]; p<.0005) and day-1 levels of S100B (OR, 1.73 [95% CI 1.27-2.36]; p<.0005) and the development of early circulatory shock.

DISCUSSION

We conducted a retrospective study to examine the association of day-1 brain biomarker levels with the development of early circulatory shock following moderate-severe TBI. We found significant differences in UCH-L1, S100B, GFAP and hsCRP in the 14 days following injury between those who did and did not develop early circulatory shock, and that day-1 levels of UCH-L1 and S100B were associated with the development of early circulatory shock.

This association between neuronal and glial injury and circulatory shock helps further elucidate the multifactorial etiology of circulatory shock following moderate-severe TBI⁴ ²⁵. Current research demonstrates that TBI-associated circulatory shock occurs secondary to the initial primary brain injury, during which mechanical forces lead to axonal shearing and necrosis, and the subsequent secondary injury which is driven by inflammation, blood–brain barrier disruption, apoptosis, metabolic disturbances, and oxidative stress²⁶. The release of a cascade of autonomic and inflammatory mediators into the circulation leads to changes in central and peripheral autonomic tone and widespread catecholamine release through activation leads to a direct effect on the function of a range of organ systems, including cardiopulmonary dysfunction and subsequent shock⁴ ³⁰. Markers of neurological injury severity, such as low initial GCS (3–5), increase the risk of developing ICU hypotension 3.37 fold²⁰. Taken together, this evidence suggests that circulatory shock following TBI is caused by a complex cascade of dysfunction driven by injury to the brain and its cellular components.

In this study, we observed a significant association between day-1 levels of UCH-L1 and S100B and the development of circulatory shock within 3 days of admission. UCH-L1 is a protein specifically expressed in neurons involved in either the addition or removal of ubiquitin from abnormal proteins (i.e., misfolded proteins, proteins damaged by oxidation, or denatured by other means) that are destined for proteasomal degradation³¹. In the first 24 hours after severe TBI, there is a significant increase in serum and cerebrospinal fluid UCH-L1, which is associated with increased hospital and 6-month mortality^{13 32}. UCHL1 has also been shown to predict abnormal head CT findings following brain injury with a sensitivity of 100% and a specificity of 39% at a cut off value of >40pg/ml³³. S100B is a low affinity calcium-binding protein expressed in astroglial and Schwann cells that regulates intracellular calcium levels; it is released during astroglial injury and becomes elevated in the cerebrospinal fluid and serum following injury³⁴. S100B has been shown to correlate with GCS score and neuroradiological findings at the time of hospital admission of TBI patients^{35–38}. A high level of S100B early after TBI can predict poor outcome, especially if it is accompanied by a second increase in serum S100B levels during the subacute phase^{16 39}. Our data demonstrated that day-1 and day-3 levels of both UCH-L1 and S100B were significantly increased in the group that developed circulatory shock. Median levels of UCH-L1 and S100B subsequently declined over the 14-day interval for the study population.

These findings suggest that UCH-L1 and S100B may be useful biomarkers for detecting the development of circulatory shock after moderate-severe TBI, and further support the hypothesis that direct neuronal and astroglial injury contributes to the development of circulatory shock in patients with moderate-severe TBI.

We also observed unique patterns of elevation of GFAP, NSE and hsCRP levels in the first 14-days following moderate-severe TBI. GFAP, which was acutely elevated in both groups in our study cohort and declined over 14 days, is a monomeric intermediate filament protein of astrocytes and considered specific for central nervous system disease^{40 41}. Previous studies demonstrate GFAP is elevated early after TBI and functions as a good predictor for TBI severity and abnormal head CT findings in patients with severe TBI^{12 42-44}. In one study, GFAP outperformed S100B in its ability to predict intracranial abnormalities on CT scan in patients with TBI across the full injury spectrum of GCS 3-15 through 24 hours post-injury⁴⁴. We did not observe significant differences in day-1 GFAP levels between those who developed circulatory shock and those who did not. Serum hsCRP, a non-specific but sensitive biomarker for systemic inflammation, has previously been shown to predict disability at 6-months after injury when measured within 2 weeks of TBI^{17 45}. Though we did not observe an association with day-1 levels in those who developed circulatory shock, we did observe a slow increase in hsCRP at days-3, 5 and 14 that was significantly increased in those who developed early shock. As such, hsCRP and therefore systemic inflammation levels following injury may not correlate with neuronal or glial injury incident to the index event, but rather secondary to downstream inflammatory events such as ventilator associated pneumonia and secondary organ injury. NSE is a glycolytic enzyme which is released following acute neuronal damage in the brain; it has high specificity to the brain but limited accuracy to predict brain injury due to its release in the serum during hemolysis³⁴. NSE was initially identified in the serum and cerebrospinal fluid of patients with head trauma and those in a state of coma, and its levels in cerebrospinal fluid were proportional to the severity of TBI and were associated with increased mortality in cases of moderate or severe TBI¹⁸. We found that serum NSE was not predictive of circulatory shock, and between group differences were not observed at any point during the 14-day interval of the study. As such, the utility of NSE in predicting or correlating with TBI severity and its complications may be limited.

This study has several limitations. First, vasopressor utilization was used a proxy for circulatory shock as part of cardiovascular SOFA score because detailed information on exact shock state and episodes of hypotension could not be gathered. As such, we could not control for different mechanisms of shock (i.e., distributive, hemorrhagic, cardiogenic, obstructive) nor the dosing of different vasopressors over time. Furthermore, vasopressors may have been used to augment cerebral perfusion pressure despite absence of circulatory shock, contributing to bias in our analysis. Second, the retrospective nature of the study limits the conclusions that can be drawn, and places it at risk for residual confounding. Third, the levels of biomarkers may have been affected by a number of other conditions, including acute and chronic inflammatory conditions, surgical interventions, and complications such as ventilator associated pneumonia or sepsis, which we did not account for in this study. We addressed this by limiting our exposure to day-1 biomarkers and our outcome to early circulatory shock to increase the likelihood that vasopressor treatments

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were used for injury-induced hypotension rather than for subsequent complications such as septic shock or pulmonary embolism. Fourth, brain injury biomarkers could be associated with the development of other extracranial organ dysfunction in addition to circulatory shock, but this was not included in our analysis. As well as examination of the development of these additional multiple outcomes not being the focus of this study, the development of additional extracranial organ dysfunction would potentially occur on the causal pathway between brain injury and the development of circulatory shock (making this variable a mediator); our analysis was not designed to examine mediation. Nevertheless, we believe that future studies should consider the relationship between brain injury and the development of gan injuries, above and beyond circulatory shock. Lastly, despite the large sample size of TRACK-TBI overall, a relatively small number of cases met study inclusion criteria, potentially leading to decreased precision in the risk estimates from a lack of statistical power.

CONCLUSION

In conclusion, we observed an association between elevated day-1 levels of UCH-L1 and S100B biomarkers and the development of early shock, as well as significant differences in UCH-L1, S100B, GFAP and hsCRP in the 14 days following injury between those who did and did not develop early circulatory shock. Our data support the role of neuronal injury and inflammatory biomarkers for improved mechanistic understanding and possibly early identification of patients at risk for circulatory shock following moderate-severe TBI.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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The TRACK-TBI Investigators:

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Figure 1. Study flowchart

CV SOFA, cardiovascular component of the Sequential Organ Failure Assessment score; ED, Emergency Department; GCS, Glasgow coma score; ICU, intensive care unit; TBI, traumatic brain injury



Figure 2. Boxplot of day 1 biomarker level stratified by early circulatory shock status (Median and $25^{th}/75^{th}$ percentiles)

GFAP, glial fibrillary acidic protein; hsCRP, high-sensitivity C-reactive protein; NSE, neuron specific enolase; S100B, S100 calcium binding protein B; UCH-L1, ubiquitin C-terminal hydrolase-L1

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Figure 3. Biomarker levels (log transformed) at days 1, 3, 5 and 14 stratified by early circulatory shock status

GFAP, glial fibrillary acidic protein; hsCRP, high-sensitivity C-reactive protein; NSE, neuron specific enolase; S100B, S100 calcium binding protein B; UCH-L1, ubiquitin C-terminal hydrolase-L1

Table 1

Demographic and clinical characteristics of the study population

| Variable | Total | Circulatory Shock | | |
|--------------------------------|---------------|-------------------|-------------|---------|
| | | No | Yes | p value |
| Subjects | 392 | 227 | 165 | |
| Age [mean (SD)] | 39.9.6 (16.6) | 38.5 (16.6) | 41.7 (16.6) | 0.036 |
| Unknown [no.] | 0 | 0 | 0 | |
| Age [no. (%)] | | | | |
| 17-39 years | 211 (53%) | 129 (57%) | 82 (50%) | 0.369 |
| 41-60 years | 133 (34%) | 72 (32%) | 61 (37%) | |
| >60 years | 48 (13%) | 26 (11%) | 22 (13%) | |
| Unknown | 0 | 0 | 0 | |
| Sex [no. (%)] | | | | |
| Male | 313 (80%) | 182 (80%) | 131 (79%) | 0.899 |
| Female | 79 (20%) | 45 (20%) | 33 (21%) | • |
| Unknown | 0 | 0 | 0 | |
| Race [no. (%)] | | | | |
| White | 307 (80%) | 171 (77%) | 138 (85%) | 0.001 |
| Black | 47 (12%) | 39 (18%) | 8 (5%) | • |
| Other | 27 (7%) | 13 (6%) | 16 (10%) | • |
| Unknown | 11 | 5 | 6 | |
| Hispanic [no. (%)] | | | | |
| No | 306 (80%) | 171 (77%) | 135 (84%) | 0.122 |
| Yes | 76 (20%) | 50 (23%) | 26 (16%) | • |
| Unknown | 10 | 6 | 4 | |
| Injury Cause [no. (%)] | | | | |
| Road traffic incident | 222 (57%) | 127 (57%) | 95 (58%) | 0.038 |
| Incidental fall | 96 (25%) | 57 (26%) | 39 (24%) | • |
| Violence / assault | 34 (9%) | 25 (11%) | 9 (5%) | • |
| Other | 36 (9%) | 14 (6%) | 22 (13%) | • |
| Unknown | 4 | 4 | 0 | |
| ISS non-head/neck [mean (SD)] | 5.8 (6.3) | 5.4 (6.2) | 6.5 (6.4) | 0.134 |
| Unknown [no.] | 5 | 3 | 2 | |
| CT Rotterdam Score [mean (SD)] | 3.4 (1.2) | 3.1 (1.2) | 3.8 (1.2) | < 0.000 |
| Unknown [no.] | 29 | 13 | 16 | |
| GCS ED Arrival [mean (SD)] | 5.8 (3.1) | 6.3 (3.2) | 5.1 (2.9) | < 0.000 |
| Unknown [no.] | 0 | 0 | 0 | |
| GCS ED Arrival [no. (%)] | | | | |
| Severe (3–8) | 291 (74%) | 157 (69%) | 134 (81%) | 0.007 |
| Moderate (9-12) | 101 (26%) | 70 (31%) | 31 (19%) | • |

| Variable | Total | Circulatory Shock | | |
|------------------------|-----------|-------------------|-----------|---------|
| | | No | Yes | p value |
| Unknown | 0 | 0 | 0 | |
| CT Intracranial Injury | | | | |
| CT negative | 22 (6%) | 21 (10%) | 1 (1%) | 0.0002 |
| CT positive | 346 (94%) | 196 (90%) | 150 (99%) | |
| Unknown | 24 | 10 | 14 | |
| CT Marshall Score | | | | |
| I-II | 183 (50%) | 133 (62%) | 45 (34%) | 0.0005 |
| III-IV | 36 (10%) | 22 (10%) | 14 (9%) | |
| V-VI | 144 (40) | 59 (28%) | 85 (57%) | |
| Unknown | 29 | 13 | 16 | |

CT, computed tomography; ED, Emergency Department; ISS, injury severity score; SD, standard deviation

Table 2

Association of day-1 biomarkers with early circulatory shock

| Unadjusted | | | Adjusted ¹ | | | |
|------------|------|-----------|-----------------------|------|-----------|-------|
| Biomarker | OR | 95% CI | р | OR | 95% CI | р |
| log GFAP | 1.10 | 0.96-1.26 | .175 | 1.07 | 0.91-1.27 | .417 |
| log UCH-L1 | 1.82 | 1.47-2.25 | <.001 | 1.63 | 1.25-2.12 | <.001 |
| log S100B | 1.94 | 1.51-2.48 | <.001 | 1.73 | 1.27-2.36 | <.001 |
| log NSE | 1.24 | 0.92-1.66 | .160 | 1.04 | 0.73-1.48 | .840 |
| log hsCRP | 1.10 | 0.94-1.30 | .243 | 1.05 | 0.87-1.27 | .582 |

¹Adjusted models include age, gender, race (black, other vs. white), GCS (3–8 vs. 9–12), non-head ISS, and CT Rotterdam score

CI, confidence interval; GFAP, glial fibrillary acidic protein; hsCRP, high-sensitivity C-reactive protein; NSE, neuron specific enolase; OR, odds ratio; S100B, calcium binding protein B; UCH-L1, ubiquitin C-terminal hydrolase-L1