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Serum NGAL predicts survival after resuscitation from cardiac arrest

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Abstract

Objective—In the first days after cardiac arrest, accurate prognostication is challenging. Serum biomarkers are a potentially attractive adjunct for prognostication and risk stratification. Our primary objective in this exploratory study was to identify novel early serum biomarkers that predict survival after cardiac arrest earlier than currently possible.

Design—Prospective, observational study

Setting—A single academic medical center

Subjects—Adult subjects who sustained cardiac arrest with return of spontaneous circulation

Intervention—None

Measurements and Main Results—We obtained blood samples from each subject at enrollment, 6, 12, 24, 48 and 72 hours after return of spontaneous circulation. We measured serum levels of novel biomarkers including neutrophil gelatinase-associated lipocalin (NGAL), high-mobility group protein B1 (HMGB), intracellular cell adhesion molecule 1 (ICAM-1) and leptin, as well as previously characterized biomarkers including neuron specific enolase (NSE) and S100B protein. Our primary outcome of interest was survival to hospital discharge. We compared biomarker concentrations at each time point between survivors and non-survivors and used logistic regression to test the unadjusted associations of baseline clinical characteristics and enrollment biomarker levels with survival. Finally, we constructed a series of adjusted models to explore the independent association of each enrollment biomarker level with survival.

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A total of 86 subjects were enrolled. Enrollment levels of HMGB, NGAL and S100B were higher in non-survivors than survivors. Enrollment leptin, NSE and ICAM-1 levels did not differ between non-survivors and survivors. The discriminatory power of enrollment NGAL level was greatest (*c*-statistic 0.78 (95%CI 0.66 – 0.90), and remained stable across all time points. In our adjusted models, enrollment NGAL level was independently associated with survival even after controlling for development of acute kidney injury and its addition to clinical models improved overall predictive accuracy.

Conclusion—Serum NGAL levels are strongly predictive of survival to hospital discharge after cardiac arrest.

MeSH Headings

Biomarker; cardiac arrest; outcome; acute kidney injury; prognosis

Introduction

Over 500,000 Americans suffer a cardiac arrest (CA) each year and more than 100,000 achieve return of spontaneous circulation (ROSC) and are cared for in the hospital (1, 2). Of these, 50-89% die in the hospital (3, 4). In the first 72 hours after ROSC, no known clinical finding or diagnostic test has adequate specificity for poor outcome or anticipated clinical course to inform clinical care or accurate neurological prognostication (5-7). Thus, it is difficult to prospectively identify and intervene upon high-risk patients prior to deterioration, and in most cases at least 3 days of critical care are necessary before accurate prognostication is possible, creating a burden to families and the healthcare system.

Serum biomarkers are a potentially attractive adjunct for prognostication and risk stratification. The best-described serum biomarkers after CA are neuron specific enolase (NSE) and S100B protein, which are released into the blood following anoxic brain injury and may reflect the severity of injury (8, 9). Unfortunately, NSE levels peak between 48-72 hours after ROSC, and so do not predict outcomes early after ROSC, and recent consensus guidelines have concluded that S100B lacks sufficient precision and discrimination to be clinically useful (8, 9).

Our primary objective in this exploratory study was to identify novel early serum biomarkers that predict survival after cardiac arrest earlier than currently possible. The post-arrest syndrome is characterized by multiple organ dysfunction secondary to global ischemia-reperfusion injury and a systemic inflammatory response (10-12). As such, rather than testing markers of brain injury as have been previously evaluated, we selected candidate biomarkers from molecular pathways related to systemic inflammation (T cell activation, leukocyte adhesion and activation, and toll-like receptor-mediated innate immunity) that might reflect the extent of total body injury.

Materials and Methods

Study Design

We performed a prospective, observational study enrolling consecutive subjects at a single academic medical center. The University of Pittsburgh Institutional Review Board approved all aspects of this study, and either the subject or a legally authorized proxy provided written informed consent prior to enrollment.

Patients and Setting

The UPMC Presbyterian Hospital is a 795-bed tertiary care referral center. Our Post-Cardiac Arrest Service (PCAS) provides standardized care to over 250 patients admitted after cardiac arrest annually. During the study period, this included routine use of therapeutic hypothermia for comatose patients, defined as not following commands within 6 hours of ROSC, regardless of initial arrest rhythm or arrest location (13). Our standard practice was to induce hypothermia as rapidly as possible with intravenous boluses of 4°C crystalloid, application of ice to the neck, axillae and groins, and surface or endovascular cooling systems, targeting a core temperature of 33°C. We maintained hypothermia for 24 hours, then rewarmed patients to normothermia over 16 hours at a rate of 0.25°C/hr to a goal temperature of 37°C which we maintained until 5 days after ROSC or awakening. We included subjects in the present study who suffered an in-hospital cardiac arrest (IHCA) or out-of-hospital cardiac arrest (OHCA) with return of spontaneous circulation (ROSC) and were admitted to the emergency department or intensive care unit between January 1, 2007 and December 31, 2009. We excluded subjects for age <18 years, presentation >6 hours after ROSC, pregnant, prisoner, withdrawal of life-sustaining therapy within 6 hours, and surgical or traumatic etiology of arrest. We defined “cardiac arrest” as a subject having received chest compressions or defibrillation from a health care provider, and considered arrests in the ED to be OHCA.

Measurements

We obtained blood samples from each subject at enrollment, 6, 12, 24, 48 and 72 hours after ROSC. We collected samples in 2.7mL plastic tubes with 3.2% buffered sodium citrate additive and 10mL plastic serum tubes without additives (Becton, Dickinson and Co, Franklin Lakes, NJ) that we rested for 20 minutes prior to centrifugation at 3000rpm for 10 minutes. We removed the supernatant and stored it in Eppendorf tubes at -80°C until analysis. We measured serum levels of novel biomarkers in the post-arrest population including neutrophil gelatinase-associated lipocalin (NGAL), high-mobility group protein B1 (HMGB), intracellular cell adhesion molecule 1 (ICAM-1) and leptin, as well as previously characterized biomarkers including neuron specific enolase (NSE) and S100B protein. We used Luminex immunofluorescence assays (Millipore, Billerica, MA) and analyzed samples according to manufacturer instructions (available at www.millipore.com).

We prospectively recorded baseline characteristics including subject age, sex, presenting arrest rhythm [categorized as shockable (ventricular tachycardia/fibrillation (VT/VF)], or non-shockable (pulseless electrical activity, asystole, or unknown), arrest location (IHCA or OHCA), Sequential Organ Failure Assessment (SOFA) score, use of therapeutic

hypothermia, and Pittsburgh Cardiac Arrest Category (PCAC). The PCAC is a clinical prediction tool that stratifies CA survivors by their risk of subsequent death or neurological deterioration based on clinical characteristics during the first 6h after ROSC (14). This tool divides survivors of CA into four categories that are predictive of survival and functional outcome. Because it could potentially affect biomarker clearance, we recorded each subject's temperature hourly for the first 36 hours, and then at 48 and 72 hours. As a measure of ongoing cardiovascular dysfunction and the severity of the post-cardiac arrest inflammatory syndrome, we recorded cumulative vasopressor index (CVI) hourly for the first 24 hours. The CVI is a method of standardizing vasopressor dosing (15, 16).

We defined acute kidney injury (AKI) according to the AKI Network (AKIN) criteria(17) as a rise in SCr of 0.3mg/dL within 48 hours of ROSC or 1.5-fold from baseline, or the need for renal replacement therapy, and treated AKI as a dichotomous outcome. We recorded serum creatinine (SCr) level at presentation and the peak value within 72 hours (+/- 12 hours) of ROSC, as well as each subject's pre-arrest baseline SCr. When a pre-arrest baseline was not available within the prior 6 months, we treated the lowest SCr after arrest as that subject's baseline. SCr was checked at the discretion of the treating clinician, and measurements were performed by the hospital's clinical laboratory. For patients who were discharged from the hospital less than 5 days after ROSC, we assumed that renal function did not deteriorate after discharge and so if they had not developed AKI prior to discharge they were classified as having no AKI.

We recorded each subject's outcomes including, survival to hospital discharge, neurological function at hospital discharge measured using the Pittsburgh Cerebral Performance Category (CPC), development of AKI and intensive care unit (ICU) length of stay. Finally, for non-survivors, we recorded each subject's mode of death, which we categorized as withdrawal of life-sustaining therapy based on anticipated neurological prognosis; brain death; withdrawal of life-sustaining therapy based on prior advanced directives or a surrogate's representation of the subject's wishes; or multiple system organ failure, hemodynamic instability, or re-arrest without successful resuscitation.

Statistical methods

Serum biomarker levels were not normally distributed and no transformation consistently minimized skewness. Therefore, we used nonparametric tests for all analyses. We summarized baseline clinical characteristics and report medians with interquartile ranges. Our primary outcome of interest was survival to hospital discharge. To determine whether each biomarker profile differed between those who survived to discharge and those who did not, we compared the level between survivors and non-survivors using Wilcoxon rank-sum tests at each time point, and report medians, interquartile ranges, and P values for these comparisons. Since the accuracy of early predictions of outcome is of greatest clinical interest, the remainder of our analyses focused on enrollment biomarker levels. First, we used unadjusted logistic regression to test the association between each baseline clinical characteristics and biomarker level with survival. We quantified the discriminatory power of each biomarker that was associated with survival at enrollment by generating receiver operating characteristic curves (ROC) and report area under the curve (AUC) with

associated 95% confidence intervals (CI) and P values. Next, we sequentially explored the independent association of each enrollment biomarker with after survival adjusting for baseline clinical characteristics associated with survival at a level of $P < 0.1$. We excluded SOFA score and therapeutic hypothermia from these models, as they are collinear with PCAC.

Based on these models, enrollment NGAL level demonstrated the most favorable performance characteristics, so we used three additional approaches to further explore its utility. Since biomarker levels could be influenced by the presence of AKI, (18, 19) we forced “development of AKI within 72 hours” as a binary predictor into the adjusted model constructed above. Next, since a combination of multiple biomarkers and clinical characteristics might perform better than a model including only a single biomarker, we used forward selection to determine the most predictive combination of baseline clinical characteristics and biomarkers. We selected a probability of entry and removal of 0.1 and 0.2, respectively, and included all baseline clinical characteristics and enrollment biomarker levels as potential covariates. We evaluated the predictive performance of each of these models by computing *c*-statistics. Finally, we assessed the contribution of enrollment NGAL to each model by comparing *c*-statistics and integrated discrimination improvement (IDI) for each adjusted model with and without NGAL. We performed all analyses using Stata Version 13.1 (StataCorp, College Station, TX).

Results

Baseline characteristics and outcomes

A total of 86 subjects were enrolled. Median age was 56 (IQR 45-70) years, and 58% were male (Table 1). Most patients (69%) were enrolled after an OHCA, and multiple organ system dysfunction was common at presentation (median SOFA score 5 (IQR 7-10)). AKI was present at presentation in 53 (62%) subjects, and a majority of subjects met criteria for AKI within the first 5 hospital days (Table 2). Overall, 69 (80%) of subjects survived at least 72 hours after ROSC, 39 (45%) survived to hospital discharge and 7 (18%) of survivors were discharged with a CPC of 1-2.

Unadjusted analysis

In unadjusted analysis, an initial shockable rhythm was associated with improved survival (OR 3.41 (95%CI 1.40-8.31)) and higher baseline SOFA score (OR 0.76 (95%CI 0.62 – 0.93) per unit increase) and PCAC (OR 0.30 (95% CI 0.18 – 0.51)) were associated with reduced survival (Table 3). Presence of AKI at baseline, within 72 hours and within 5 days of ROSC was not associated with outcome. Enrollment levels of HMGB, NGAL and S100B were higher in non-survivors than survivors (Figure 1). The AUC of the ROC curve for enrollment NGAL level was 0.78 (95%CI 0.66 – 0.90, and remained similar at all time points (Figure 2, Table 5). Enrollment NGAL below 58,000 ng/mL was perfectly specific for survival to hospital discharge (Figure 2). Conversely, enrollment NGAL above 188,000 ng/mL was 96% specific for non-survival. Consistent with previous work, (9) NSE levels in non-survivors began to rise after 12-24 hours and were maximally separated at 48 hours, while S100b levels began to fall after 12 hours.

Adjusted analysis

In our adjusted models, enrollment NGAL level was independently associated with survival, regardless of whether we forced development of AKI into the model (Table 4). When we removed enrollment NGAL from each of these models, the overall model fit and discriminatory power declined (Table 4). Addition of enrollment NGAL to each clinical model resulted in a significant increase in IDI and predictive accuracy (Table 4). No other enrollment biomarker was independently associated with outcome after controlling for PCAC and shockable rhythm (all $P > 0.05$, data not shown). Using forward selection, enrollment NGAL and PCAC remained significant and had stable point estimates and confidence intervals when compared with the other methods of model building. Consistent with previous reports, NGAL levels rose early in subjects who went on to develop clinically apparent AKI within 72 hours of ROSC, but was not elevated in those who already exhibited AKI at presentation (data not shown).

Discussion

The main finding of our study is that serum NGAL level is strongly predictive of outcome after cardiac arrest. At enrollment, NGAL performed better than the other candidate biomarkers, including NSE and S100b, which have been better investigated after cardiac arrest. Moreover, addition of enrollment NGAL to clinical models resulted in a significant increase in the discriminatory power of these models. Notably, the association between NGAL and outcome remained stable at all time points up to 72 hours after ROSC. Thus, even in the case of inter-facility transfer, where time from arrest to presentation at the final treating hospital may be considerably delayed, the prognostic value of NGAL is preserved.

NGAL is expressed in a variety of tissues including kidney, liver and lung (20). Plasma levels may be elevated in a variety of pathological inflammatory states including heart failure and malignancy (20). In AKI, NGAL is secreted both by neutrophils that contribute to tubular injury and directly by tubular epithelial cells themselves (21). Compared to serum creatinine, NGAL rises earlier and may be a more sensitive marker of AKI; moreover, in risk-adjusted analyses, development of AKI independently predicts mortality (18, 19). Thus, it is possible that the association between NGAL and outcomes in this cohort is related to underlying, subclinical AKI. Overt AKI detected by change in serum creatinine was not associated with mortality after cardiac arrest in past studies (22), nor did we find an association between AKI and outcome in this cohort. However, serum creatinine is relatively insensitive for AKI and it is possible that the NGAL is a more sensitive marker for subclinical but important AKI. Alternatively, as a nonspecific inflammatory marker (20), NGAL may also reflect the overall severity of systemic inflammation in the post-cardiac arrest syndrome (10-12). In this regard, NGAL may be an easily measurable surrogate measure of covariates such as duration of pulselessness, quality of CPR and total burden of anoxic and ischemia-reperfusion injury, metrics which are at present are unreliably available or difficult to quantify for clinical or research purposes (23-25).

Both NSE and S100B have been proposed to be specific markers of brain injury and evaluated in the CA population. NSE is an intracellular enzyme involved in glucose metabolism found in neurons and neuroendocrine cells, but also platelets and erythrocytes

(26). To our knowledge, the largest study NSE as a prognostic biomarker after CA was the PROPAC study, which was conducted before the widespread implementation of targeted temperature management (27). This study suggested that an NSE value above 33mcg/L uniformly predicted poor outcome. However, hypothermia attenuates the rise in NSE level, which limits the applicability of the PROPAC results to current clinical care. Furthermore, as re-demonstrated by our results in the present study, NSE rises slowly after CA and is not predictive of outcome in the first days after ROSC, limiting its utility during the first 48-72 hours after ROSC when accurate prognostication is most challenging (28). S100B is an intracellular calcium binding protein found in glial cells, and may be more specific to brain injury than NSE (26). Our results are consistent with previous studies that have shown that S100B rises early in non-survivors, although the variability of levels among survivors and non-survivors is large. (27, 29). Previous studies have suggested that S100B may not have sufficient discriminatory power to be clinically useful (27, 29).

With the exception of NGAL, the remaining novel biomarkers we investigated in the present study were inconsistently or never associated with clinical outcome. HMGB is a ubiquitous nuclear protein involved in chromatin structure, and also found in cytosol, where it is involved in sterile inflammation and autophagy (30). This offers some face validity as a marker of the severity of the post-arrest syndrome. Indeed, HMGB concentration was significantly higher in non-survivors than survivors at enrollment. However, the discriminatory power of enrollment HMGB level was lower than that of enrollment NGAL, and in contrast to NGAL, subsequent HMGB levels did not predict outcome. Leptin is a peptide expressed in neuronal tissue and extracranial adipose, and has been evaluated as a prognostic marker after traumatic brain injury (31), but did not correlate with outcome in this cohort. ICAM-1 is a marker of endothelial dysfunction and pathological central nervous system inflammatory states where it facilitates leukocyte adhesion and recruitment (32). It also did not correlate with outcome in this cohort.

Our study has several important limitations. As a tertiary care referral center, many of our patients are received in transfer from other facilities. As such, “enrollment” biomarker levels were not obtained immediately after cardiac arrest. However, all “enrollment” biomarker levels were obtained within 6 hours of ROSC. For all subsequent time points, samples were obtained at fixed intervals from return of spontaneous circulation, but there was some variability in timing of the initial sample that reflects real-world clinical practice. In the case of NGAL, this variability does not alter our conclusions since the relationship with outcome was stable over time. Also, although we used several methods of model building to generate multivariable models, we could not adjust for several potentially important covariates such as time intervals from collapse-to-CPR and CPR-to-ROSC, which we have previously shown are unreliably documented in our care system (23). It is possible that unmeasured confounders biased our findings, a limitation inherent to all observational studies.

Like many biomarker studies, we followed subjects' clinical outcomes to hospital discharge. While this is an important marker of overall recovery, in patients with acquired brain injury, survival to hospital discharge is not necessarily a patient-centered outcome as compared to a favorable neurological outcome. Future investigations may focus on subjects' neurological recovery and long-term outcomes. As a single center study, it is possible that some of our

the correlations we observed were influenced by our local standard of care. For example, the rise in many biomarkers after cardiac arrest is influenced by therapeutic hypothermia. While we systematically treated all comatose subjects in this study with therapeutic hypothermia, other local care practices may represent unmeasured confounders. Since comatose patients in this cohort were treated with hypothermia and also had a greater severity of illness, we are not able to differentiate the effects of hypothermia on biomarker levels from the effect of illness severity. The results of the recent Target Temperature Management trial, which support hypothermia to a goal of either 33 or 36°C, offer a future opportunity to disentangle this confounding in these most severely injured patients. Finally, while our local standard of care has not changed significantly since the recruitment period for the present study, it is possible that recent changes in resuscitation and general critical care management and guidelines limit the generalizability of our results to other centers' current clinical practice. Contemporary, external validation of our findings by other groups is a necessary next step in exploring the utility of NGAL as a prognostic marker in this population.

Conclusion

In this exploratory study, it appears that serum NGAL levels are strongly predictive of survival to hospital discharge after cardiac arrest. This relationship is stable over time, and addition of enrollment NGAL to clinical models improve overall model fit and discriminatory power. We believe NGAL is an appealing candidate that merits further investigation in this patient population.

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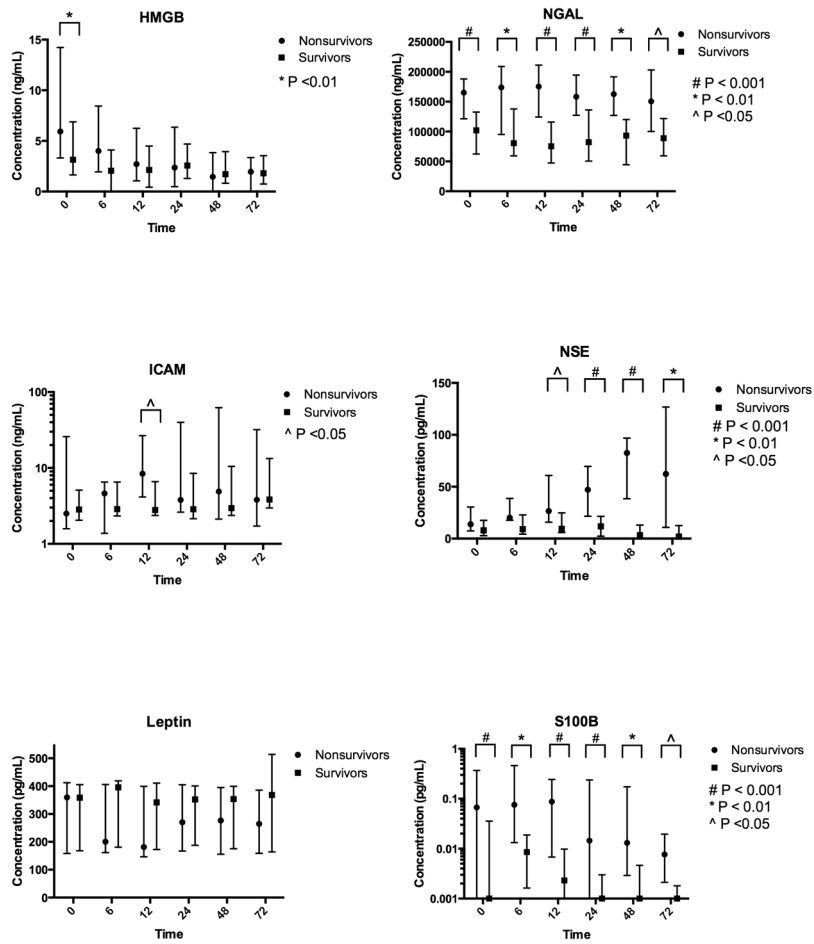


Figure 1. Biomarker profiles after cardiac arrest versus time versus time from return of spontaneous circulation, stratified by survival to hospital discharge.

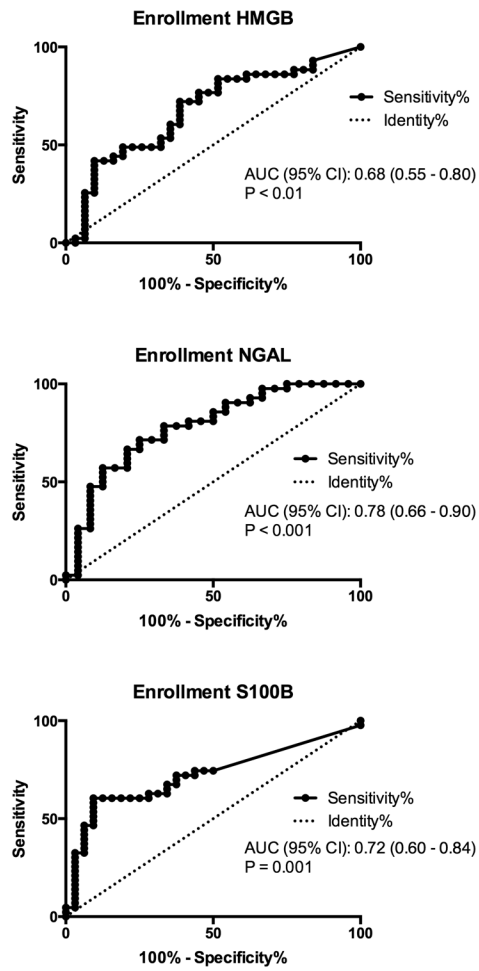


Figure 2. Receiver operating characteristic curves for enrollment biomarker levels associated with survival to hospital discharge after cardiac arrest.

Table 1
Baseline clinical characteristics after cardiac arrest in the biomarker cohort

Characteristic	Survivors (n = 39)	Non-survivors(n = 47)	Overall cohort (n = 86)
Age, years	56 [43 – 56]	56 [45 – 56]	56 [45 – 70]
Male sex	21 (54)	29 (62)	50 (58)
Presenting rhythm			
VT/VF	24 (62)	15 (32)	39 (45)
PEA	8 (21)	12 (26)	20 (23)
Asystole	4 (10)	12 (26)	16 (19)
Unknown	3 (8)	8 (17)	11 (13)
Out-of-hospital arrest	28 (72)	31 (66)	59 (69)
Pittsburgh Cardiac Arrest Category			
I	9 (23)	0 (0)	9 (10)
II	18 (46)	12 (26)	30 (35)
III	7 (18)	8 (17)	15 (17)
IV	5 (13)	27 (57)	32 (27)
Baseline SOFA score	6 [4 – 6]	9 [7 – 9]	7 [5 – 10]
Central nervous system	3 [2 – 3]	4 [4 – 4]	4 [3 – 4]
Respiratory	2 [1 – 2]	3 [2 – 3]	3 [1 – 3]
Cardiovascular	0 [0 – 0]	0.5 [0 – 0.5]	0 [0 – 1]
Liver	0 [0 – 0]	1 [0 – 1]	0 [0 – 0]
Renal	0 [0 – 0]	0 [0 – 0]	0 [0 – 2]
Coagulation	0 [0 – 0]	0 [0 – 0]	0 [0 – 0]
Cumulative vasopressor index	0 [0 – 4]	2 [0 – 38]	0 [0 – 30]
Baseline SCr, mg/dL	1 [0.9 – 1]	1.4 [0.8 – 1.4]	0.7 [0.6 – 1.0]
Therapeutic hypothermia	26 (67)	45 (98)	71 (84)

Data are presented as median [interquartile range] or raw number with corresponding percentages.

Abbreviations: VT/VF – Ventricular tachycardia/fibrillation; PEA – Pulseless electrical activity; SOFA – Sequential Organ Failure Assessment; SCr – Serum creatinine.

Table 2
In-hospital outcomes after cardiac arrest within the overall cohort

Outcome	Overall cohort (n = 86)
Survival to hospital discharge	39 (45)
Discharge CPC (% of survivors)	
1-2	7 (18)
3-4	32 (37)
ICU length of stay, days	5 [3 – 11]
Non-survivors	4 [3 – 9]
Survivors	7 [4 – 12]
Hospital length of stay, days	9 [4 – 14]
Non-survivors	4 [3 – 9]
Survivors	12 [9 – 20]

Data are presented as median [interquartile range] or raw number with corresponding percentages.

Abbreviations: CPC – Cerebral Performance Category; ICU – Intensive Care Unit

Table 3
Unadjusted predictors of survival to hospital discharge after cardiac arrest

Characteristic	Odds ratio (95% CI)	P value
Baseline clinical characteristic		
Age	0.99 (0.97 – 1.02)	0.54
Male sex	0.72 (0.31 – 1.71)	0.46
Shockable rhythm	3.41 (1.40 – 8.31)	0.007
Out-of-hospital arrest	1.31 (0.53 – 3.30)	0.56
Pittsburgh Cardiac Arrest Category	0.30 (0.18 – 0.51)	<0.001
Baseline SOFA score	0.76 (0.62 – 0.93)	0.008
Therapeutic hypothermia	0.04 (0.01 – 0.34)	0.004
Acute kidney injury		
Immediate post-arrest	1.21 (0.50 – 2.91)	0.67
Within 72 hours	0.51 (0.17– 1.49)	0.22
Within 5 days	0.58 (0.20 – 1.74)	0.34
Biomarkers at enrollment		
NGAL, per 25,000 ng/mL	0.58 (0.42 – 0.79)	<0.001
ICAM-1, per 10 ng/mL	0.73 (0.52 – 1.02)	0.07
Leptin, per 100 pg/mL	1.10 (0.81 – 1.50)	0.53
HMGB, per 10 ng/mL	0.95 (0.77 – 1.17)	0.61
NSE, per 10 pg/mL	0.85 (0.67 – 1.07)	0.17
S100B, per 0.1 pg/mL	0.65 (0.45 – 0.92)	0.01

Abbreviations: NGAL - Neutrophil gelatinase-associated lipocalin; ICAM-1 – Intracellular cell adhesion molecule 1; HMGB – High-mobility group protein B1; NSE – Neuron specific enolase.

Enrollment NGAL level improves the discriminatory value of adjusted models predicting survival after cardiac arrest, regardless of the methods used to build the model or adjustment for acute kidney injury.

Table 4

Model	Covariate	Adjusted OR (95% CI)	P value	Model performance	Full model	Reduced model (NGAL removed)	Integrated discrimination improvement with NGAL Estimate	P value
1	Enrollment NGAL	0.65 (0.44 – 0.96)	0.03	C statistic	0.90	0.79		
	Shockable rhythm	1.28 (0.28 – 5.89)	0.76	Efron's R ²	0.49	0.30	0.13	<0.001
	PCAC	0.20 (0.08 – 0.48)	<0.001	BIC	-210	-280		
2	Enrollment NGAL	0.64 (0.43 – 0.95)	0.03	C statistic	0.90	0.83		
	AKI within 72 hours	1.76 (0.41 – 7.55)	0.45	Efron's R ²	0.50	0.33		
	Shockable rhythm	1.36 (0.29 – 6.45)	0.70	BIC	-206	-279	0.12	0.002
	PCAC	0.20 (0.09 – 0.49)	<0.001					
3	Enrollment NGAL	0.52 (0.30 – 0.89)	0.02	C statistic	0.94	0.81		
	Enrollment HMGB	1.37 (1.02 – 1.86)	0.04	Efron's R ²	0.59	0.35		
	Enrollment ICAM-1	0.53 (0.28 – 1.02)	0.06	BIC	-214	-230	0.18	<0.001
	PCAC	0.12 (0.04 – 0.38)	<0.001					

Abbreviations: NGAL – Neutrophil gelatinase-associated lipocalin; PCAC – Pittsburgh Cardiac Arrest Category; AKI – Acute kidney injury; HMGB – High-mobility group protein B1; ICAM-1 – intracellular cell adhesion molecule 1; BIC – Bayesian information criterion.

Adjusted Model 1: Built including significant clinical predictors with unadjusted P < 0.1; Adjusted Model 2: Development of AKI is forced into Model 1; Adjusted Model 3: Built with forward selection. All models are presented with (full) and without (reduced) inclusion of NGAL. Odds ratios for NGAL are expressed per 25,000ng/mL change; odds ratios for HMGB and ICAM-1 are expressed per 10ng/mL change.

Table 5
Receiver operating curve characteristics for the association of serum NGAL level with survival to hospital discharge

Time point	ROC AUC (95%CI)	P value
Enrollment	0.78 (0.66 – 0.90)	0.0002
6 hours	0.80 (0.66 – 0.95)	0.003
12 hours	0.82 (0.68 – 0.96)	0.0008
24 hours	0.79 (0.67 – 0.90)	<0.0001
48 hours	0.79 (0.62 – 0.95)	0.005
72 hours	0.79 (0.61 – 0.97)	0.02

Abbreviation: ROC AUC – Receiver operating curve area under the curve.

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