



Research article

Prediction model of death risk in patients with sepsis and screening of biomarkers for prognosis of patients with myocardial injury

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ABSTRACT

This study aimed to create a robust prediction model for sepsis patient mortality and identify key biomarkers in those with myocardial injury. A retrospective analysis of 261 sepsis inpatients was conducted, with 44 deaths and 217 recoveries. Key factors were assessed via univariate and multivariate analyses, revealing myocardial injury, shock, and pulmonary infection as independent mortality risk factors. Using LASSO regression, a reliable prediction model was developed and internally validated. Additionally, procalcitonin (PCT) emerged as a sensitive biomarker for myocardial injury prediction in sepsis patients. In summary, this study highlights myocardial injury, shock, and pulmonary infection as independent risk factors for sepsis-related deaths. The LASSO-based prediction model effectively forecasts the prognosis of septic patients with myocardial injury, with PCT showing promise as a predictive biomarker.

1. Introduction

Sepsis, a condition characterized by life-threatening organ dysfunction resulting from the body's dysfunctional response to infection, poses a growing global health challenge. Recent data indicate an annual increase of 19 million sepsis cases worldwide, with a mortality rate exceeding that of stroke [1,2]. Sepsis-related mortality in hospitals and intensive care units (ICUs) ranges from 33.5% to 48.7%, particularly among patients experiencing sepsis accompanied by multiple organ dysfunction [3]. Thus, the accurate prediction of mortality risk and identification of associated risk factors in sepsis patients are crucial for informed clinical decision-making [4].

Clinical observations reveal that approximately 50% of individuals with severe sepsis also exhibit left ventricular systolic dysfunction [5]. A critical aspect of myocardial injury in sepsis is the apoptosis of cardiomyocytes [6,7]. Numerous endotoxins produced during sepsis trigger a cascade of apoptotic processes, leading to cardiomyocyte apoptosis and necrosis [8]. Earlier studies have additionally underscored that myocardial injury serves as a significant prognostic factor in sepsis patients [9,10]. One pressing clinical challenge is the prediction of outcomes in sepsis patients with myocardial injury, enabling timely interventions.

This retrospective analysis seeks to develop a predictive model for assessing the risk of death in sepsis patients, particularly focusing on those with myocardial injury. Through an in-depth examination of biomarker trends, we aim to identify markers that can help

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predict sepsis in combination with myocardial injury.

2. Methods

2.1. Patients collection

This study is a retrospective cohort analysis. Patients with sepsis who underwent standardized diagnosis and treatment in the hospital from January 1, 2018, to January 1, 2020, were collected retrospectively. All experiments involving human beings were conducted by conforming to the ethical standards of the national research committee and the 1964 Declaration of Helsinki and its later amendments or comparable ethical standards. This study was approved by the Ethics Committee of the West China Hospital of Sichuan University (2021YFS0189). All specimens were collected with the written consent of patients and ethical approval.

Inclusion criteria:

- 1) Patients with sepsis who met the Sepsis3.0 diagnostic criteria [11].
- 2) The patient was ≥ 18 years old.

Exclusion criteria:

- 1) Incomplete clinical data during hospitalization.
- 2) Automatic discharge affects the judgment of the outcome.

The criteria for diagnosing myocardial injury.

Clinical Symptoms and Signs: Chest pain or discomfort (angina) that may radiate to the arms, jaw, or back. Shortness of breath. Profound fatigue. Cold sweats. Nausea or vomiting.

Electrocardiogram (ECG or EKG):

Cardiac Biomarkers: Cardiac-specific biomarkers released into the bloodstream due to myocardial injury include: Troponin: Elevated troponin levels are a key indicator of myocardial injury. Creatine Kinase-MB (CK-MB): An increase in CK-MB levels can also suggest myocardial injury. Myoglobin: Elevated myoglobin levels may be an early sign of myocardial injury.

Imaging Studies: Echocardiogram (Echo): This ultrasound imaging technique can reveal abnormalities in the structure and function of the heart, such as reduced ejection fraction or wall motion abnormalities.

Cardiac MRI: Magnetic resonance imaging can provide detailed images of the heart and identify areas of damage.

2.2. Patient grouping

- 1) Patients with sepsis were divided into dead and alive groups according to the clinical outcome after treatment.
- 2) Patients with sepsis complicated with myocardial injury were divided into the dead group and alive group according to whether they survived after treatment.

2.3. Clinical information collection

- 1) Basic clinical data of patients: age, sex, and complications (myocardial injury and shock).
- 2) Patients' personal history: chronic obstructive pulmonary disease, hypertension, diabetes, renal insufficiency, cardiac insufficiency, tumor history, trauma history, and immunodeficiency. The study was conducted in accordance with the Declaration of the Helsinki and approved by the Institutional Review Board of the Sichuan Academy of Medical Science-Sichuan Provincial People's Hospital (Chengdu, China). Written informed consents were signed by all the enrolled individuals.
- 3) Infection sites: pulmonary infection, abdominal infection, and urinary system infection.
- 4) Pathogens: MRSA, fungi, *Escherichia coli*, *Streptococcus pneumoniae*, *Klebsiella pneumoniae*, *Pseudomonas aeruginosa*, *Acinetobacter baumannii*, *Pneumocystis carinii*, and *Staphylococcus aureus*.
- 5) Laboratory test indexes (BNP, high sensitivity troponin I, CK-MB, myoglobin, creatinine, and PCT) before treatment, 5, 10, and 15 days after treatment in sepsis patients complicated with myocardial injury.

2.4. Statistical analysis

Statistical analysis was conducted using SPSS version 26.0, R version 4.0.1, and GraphPad version 8.0.1 software. To assess the normality of continuous variables, the Shapiro-Wilk (SmurW) test was employed. For normally distributed data, descriptive statistics are presented as mean $\pm$ standard deviation, and group comparisons were conducted using independent sample t-tests. Non-normally distributed data were represented as median with interquartile range (25% and 75% quartile), and group comparisons were performed using the Mann-Whitney *U* test. In R version 4.0.3, we performed the establishment and validation of multivariate analysis and the development of a clinical prediction model. Logistic proportional hazard regression analysis was applied for both univariate and multivariate analyses, leading to the creation of the corresponding clinical prediction model. To visually illustrate the impact of each risk factor, a line chart was generated to display the scores. The robustness of the model was assessed through internal validation,

including the calculation of the C-index and the construction of a calibration curve. It's worth noting that our study yielded statistically significant test results, including the multivariate Cox proportional hazard regression outcomes.

3. Results

3.1. Comparative analysis of clinical characteristics of survival/death sepsis patients

Through inclusion, exclusion criteria, and follow-up integrity, 261 sepsis patients were included in this study, of which 44 died and 217 improved after treatment. The basic clinical information of the two groups is illustrated in Table 1. The patients in the survival group were 65.41 ± 17.44 years old, and that in the death group were 66.52 ± 20.06 years old. There were 130 male patients in the survival group and 28 in the death group. There was no significant difference in age and sex between the two groups ($P > 0.05$). The proportion of myocardial injury (41 vs. 118) and shock (32 vs. 37) in the death group was significantly higher than that in the survival group ($P < 0.05$). The results of the personal history comparison demonstrated that there was no significant difference in the proportion of patients with chronic obstructive pulmonary disease (3 vs. 15), hypertension (8 vs. 54), diabetes mellitus (9 vs. 76), renal insufficiency (4 vs. 46), history of trauma (1 vs. 4), and immunodeficiency (3 vs. 7) between the death group and the survival group. However, the proportion of cardiac insufficiency (0 vs. 22) in the death group was significantly lower than in the survival group. The proportion of patients with personal tumor history in the death group was significantly higher than that in the survival group (9 vs. 16 min, $P < 0.05$). The results indicated that the proportion of pulmonary infection in the death group was significantly higher than that in the survival group (33 vs. 113, $P < 0.05$), and the proportion of urinary infection was significantly lower than that in the survival group (1 vs. 36, $P < 0.05$). Comparative analysis of pathogens demonstrated no significant difference in the proportion of patients infected with MRSA, fungi, *Escherichia coli*, *Streptococcus pneumoniae*, *Klebsiella pneumoniae*, *Pseudomonas aeruginosa*, *Acinetobacter baumannii*, *Pneumocystis carinii*, and *Staphylococcus aureus* between the two groups.

3.2. Univariate and multivariate logistic analysis of survival/death in patients with sepsis

Univariate Logistic regression analysis revealed that age, myocardial injury, shock, pulmonary infection, *Acinetobacter baumannii* infection, and *Staphylococcus aureus* infection were the different factors between the survival group and the death group (Table 2, $P < 0.05$). The results of multivariate Logistic regression analysis revealed that myocardial injury, shock, and pulmonary infection were independent risk factors for death in sepsis patients (Table 2, $P < 0.05$).

Table 1
Baseline characteristics of sepsis patients.

		Alive(n = 217)	Dead(n = 44)	t/x ² value	P value
age		65.41 ± 17.44	66.52 ± 20.06	−0.37	0.45
Sex	Male	130	28		
	Female	87	16	0.21	0.65
Myocardial Injury		118	41	23.14	<0.01
Shock		37	32	58.31	<0.01
COPD		15	3	0.01	0.98
Hypertension		54	8	0.91	0.34
Diabetic		76	9	3.54	0.05
Renal Insufficiency		46	4	3.47	0.05
Cardiac Insufficiency		22	0	4.87	0.02
Tumor		16	9	7.23	<0.01
Trauma		4	1	0.03	0.85
Immune Insufficiency		7	3	1.28	0.26
Infection					
	Pulmonary Infection	113	33	7.64	<0.01
	Abdominal Infection	43	4	2.85	0.07
	Urinary Infection	36	1	6.16	0.01
Pathogene					
	MRSA	7	2	0.19	0.66
	Escherichia Coli	53	7	1.53	0.21
	Fungus	12	4	0.81	0.37
	streptococcus	2	0	0.41	0.52
	Pseudomonas Aeruginosa	12	2	0.07	0.79
	Acinetobacter Baumannii	15	5	1.02	0.31
	Staphylococcus Aureus	12	2	0.07	0.79
	Klebsiella Pneumoniae	26	6	0.09	0.76
	Pneumocystis Carinii	1	0	0.20	0.65

Table 2
Univariate and multivariate Logistic analysis of survival/death in patients with sepsis.

Coef	S.E.	Wald	Z	Pr(> Z)	S.E.	Wald	Z	Pr(> Z)
	Univariate				multivariate			
sex	1.7641	1.1171	1.58	0.1143				
age	0.1562	0.0757	2.06	0.039				
Myocardial Injury	4.8993	1.2329	2.54	0.0034	4.34	1.23	2.32	0.02
Shock	5.9697	1.8845	3.17	0.0015	4.54	1.56	3.12	0.01
COPD	1.2911	1.6315	0.79	0.4287				
hypertension	−0.9656	1.2602	−0.77	0.4435				
diabetic	−1.7732	1.5933	−1.11	0.2657				
Renal Insufficiency	2.0621	1.3802	1.49	0.1352				
Cardiac Insufficiency	−7.9225	30.7767	−0.26	0.7969				
tumor	1.6077	1.372	1.17	0.2413				
Trauma	−5.0811	67.0713	−0.08	0.9396				
ImmuneInsufficiency	3.6657	2.1855	1.68	0.0935				
PulmonaryInfection	3.0486	1.5099	2.02	0.0435	2.12	1.67	1.42	0.04
AbdominalInfection	−0.3744	1.7862	−0.21	0.834				
UrinaryInfection	1.2183	1.8403	0.66	0.508				
MRSA	5.6588	4.1478	1.36	0.1725				
EscherichiaColi	1.1982	1.3013	0.92	0.3572				
Fungus	4.2326	2.6802	1.58	0.1143				
streptococcus	−0.8773	108.7187	−0.01	0.9936				
PseudomonasAeruginosa	2.5048	2.1904	1.14	0.2528				
AcinetobacterBaumannii	5.8981	2.3207	2.54	0.011	1.23	0.67	1.12	0.23
StaphylococcusAureus	6.1793	2.7553	2.24	0.0249	1.36	0.88	1.94	0.45
KlebsiellaPneumoniae	2.3774	29.7421	−0.08	0.9363				

3.3. Establishment and verification of survival/death prediction model for sepsis patients

Because many factors are included in this study’s initial stage, the prediction model is further established, and the dimension reduction screening variables are selected using LASSO elastic regression. The LASSO regression results are displayed in Fig. 1a, and the appropriate variables (Table S1, Fig. 1b) were chosen when they landed. The results demonstrated that nine factors, including myocardial injury, shock, hypertension, MRSA, cardiac insufficiency, personal history of tumor, pulmonary infection, abdominal infection, and urinary system infection, were screened by LASSO regression. Furthermore, multivariate regression analysis demonstrated that myocardial injury, shock and pulmonary infection were relatively independent risk factors (Table 2, $P < 0.05$). Therefore, the nomogram is drawn based on the above nine factors, and the results can be seen in Fig. 1c. Furthermore, to evaluate the reliability of the line chart, internal verification draws calibration curve results to depict that the predicted results are in good agreement with the actual results (Fig. 1d). Internal verification draws ROC curve and calculates the curve area as 0.87 (95% CI: 0.84–0.89) (Fig. 1e–f).

3.4. Screening of biomarkers in patients with sepsis complicated with myocardial injury

Previous studies showed that the prognosis of sepsis patients with myocardial injury worsened significantly. On this basis, this study retrospectively collected the expression of patients with sepsis at different stages during treatment to clarify the value of biomarkers in judging the prognosis of patients with sepsis complicated with myocardial injury. The changing trend of each biomarker is shown in Fig. 2 a–f. The results showed that the expression levels of BNP, Troponin, CK-MB, and myoglobin in the death group were significantly higher than those in the survival group during treatment. In addition, the creatinine and PCT ex After treatment, there was no significant difference in the expression of creatinine and PCT between the two groups. Based on the biomarkers before treatment, the clinical outcome events of sepsis complicated with myocardial injury were predicted, the ROC curve was drawn, and the area under the curve was calculated. Fig. 3 a–f depicts the ROC curve for each biomarker, and Table 3 depicts the area under the curve. Table 4 results showed that the AUC of PCT was significantly higher than that of the other five biomarkers, suggesting that PCT may be an effective biomarker for predicting the prognosis of sepsis patients with myocardial injury.

4. Discussion

In this study, sepsis patients were selected retrospectively. The results showed that sepsis patients with myocardial injury, shock, and pulmonary infection were independent risk factors for death. Furthermore, based on myocardial injury, shock, hypertension, MRSA, cardiac insufficiency, personal tumor history, pulmonary infection, abdominal infection, and urinary system infection, a predictive model and internal verification results show that the model is highly reliable.

The changing trend of biomarker expression in sepsis patients with myocardial injury was observed. ROC curve showed that PCT was a sensitive biomarker for predicting sepsis with myocardial injury [12,13]. Sepsis is a life-threatening organ dysfunction caused by the imbalance of infection response, and its morbidity and mortality are very high [14]. According to the latest report in the Lancet, 48.9 million sepsis cases were recorded worldwide in 2017, and 11 million sepsis-related deaths accounted for about 20% of global

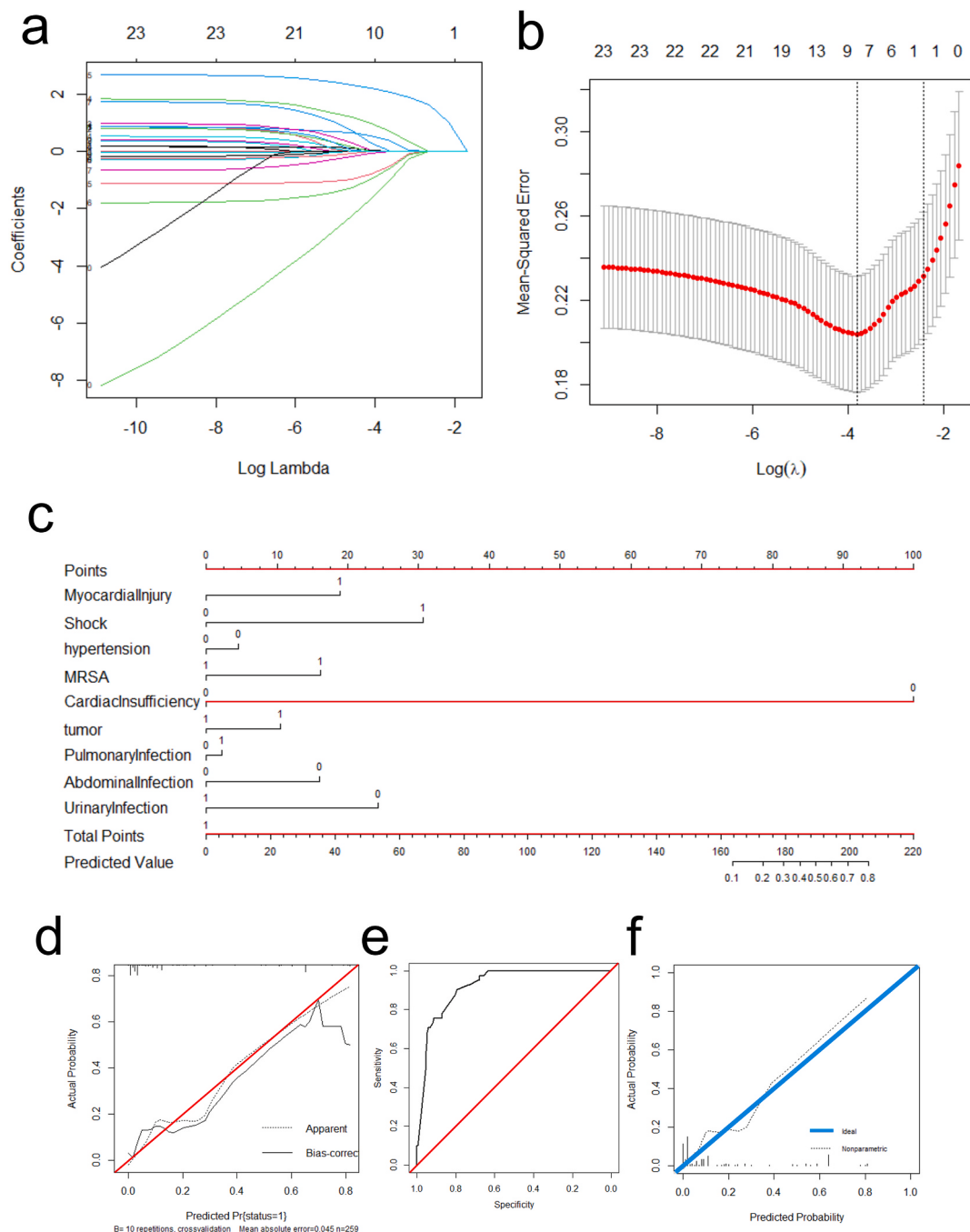


Fig. 1. Establishment and verification of survival/death prediction model for sepsis patients. (A–B) LASSO regression analysis based on each variable. (C) Nomogram of the prediction model for sepsis patients. (D) Calibration curve by internal validation. (E) ROC curve by internal validation. (F) DCA curve by internal validation.

deaths [15]. In developed countries, such as the United States, there are about 1.7 million cases of sepsis each year and about 270000 sepsis-related deaths [16]. Furthermore, sepsis is the leading cause of death among patients in intensive care units (ICU) in low-and middle-income countries, with a mortality rate of 80% [17]. The World Health Organization has recognized the significant threat of sepsis to global health and has strengthened the prevention, diagnosis, and treatment of sepsis [18]. This study included 261 patients, 44 of whom died, yielding a death rate of 16.8%. Therefore, identifying the risk factors that increase the risk of death in sepsis patients helps promote targeted clinical decision-making. Through retrospective analysis, it was finally shown that sepsis patients with myocardial injury, shock, and pulmonary infection were independent risk factors for poor prognosis.

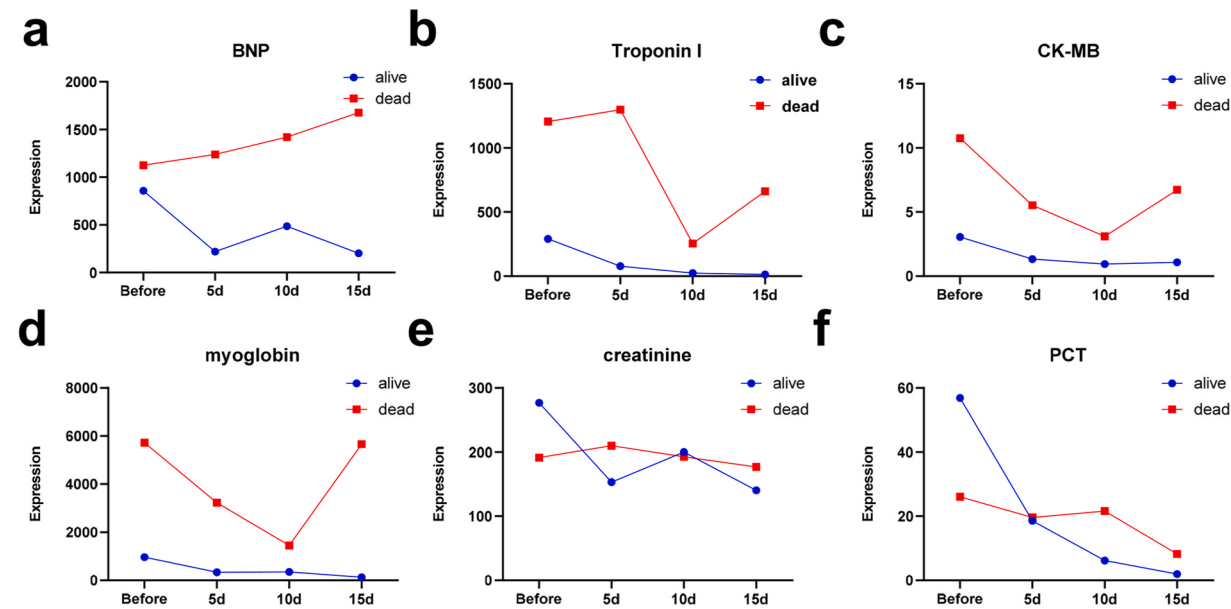


Fig. 2. Biomarkers expressions during treatment in septic patients with myocardial injury. Note: (A–F) The expression of biomarkers. BNP, Troponin I, CK-MB, myoglobin, creatinine, and PCT in different groups of septic patients with myocardial injury before treatment and 5, 10, and 15 days after treatment.

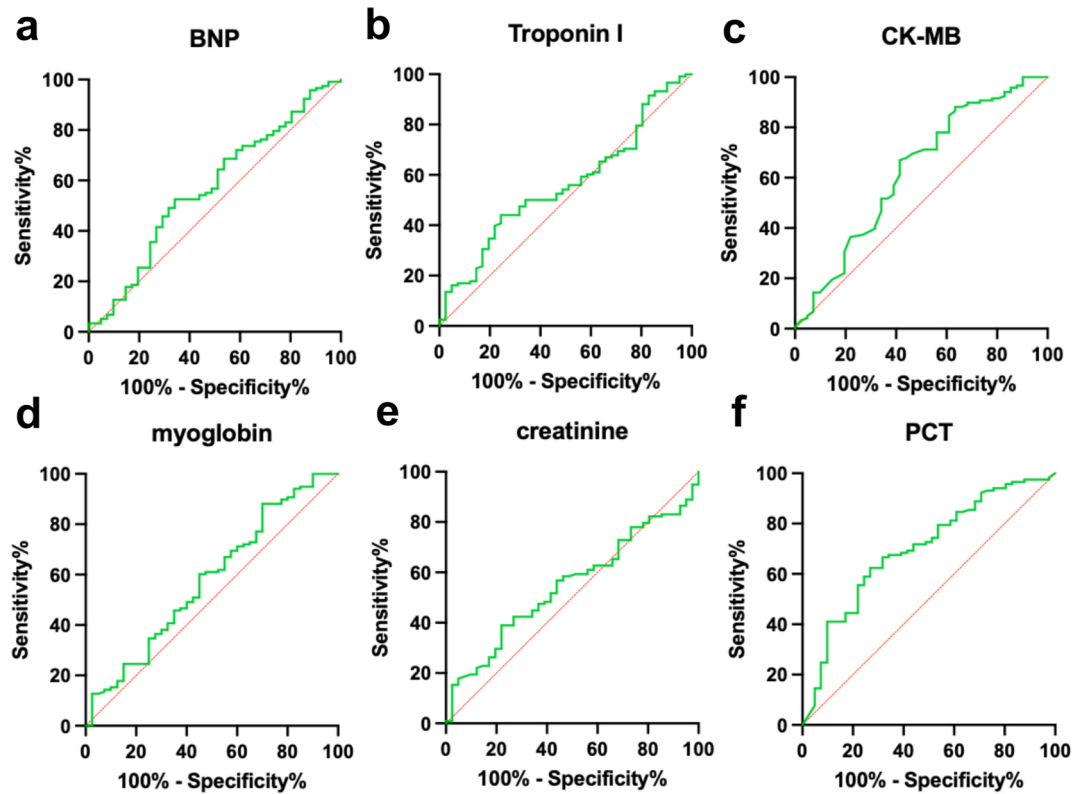


Fig. 3. ROC curves of each biomarker in septic patients with myocardial injury. (A–F) ROC curves of each biomarker (BNP, Troponin I, CK-MB, myoglobin, creatinine, and PCT) in septic patients with myocardial injury predict the survival of each patient.

Table 3
Multivariate Logistic analysis in patients with sepsis based on LASSO and univariate analysis.

Coef	S.E.	Wald	Z	Pr(> Z)
Intercept	−3.8123	0.7784	−4.9	<0.0001
MyocardialInjury	1.607	0.6752	2.38	0.0173
Shock	2.61	0.4867	5.36	<0.0001
hypertension	−0.3895	0.5263	−0.74	0.4592
MRSA	1.3773	1.1166	1.23	0.2174
CardiacInsufficiency	−8.514	22.4798	−0.38	0.7049
tumor	0.8856	0.6125	1.45	0.1482
PulmonaryInfection	0.1852	0.5661	0.33	0.7436
AbdominalInfection	−1.3566	0.7779	−1.74	0.0812
UrinaryInfection	−2.0634	1.1174	−1.85	0.0648

Table 4
The AUC of each factor under the regression analysis.

Biomarkers	AUC	95%CI
BNP	0.5682	0.46–0.67
Troponin I	0.55	0.46–0.65
CK-MB	0.62	0.51–0.73
myoglobin	0.57	0.47–0.68
creatinine	0.54	0.45–0.64
PCT	0.69	0.61–0.79

The pathophysiological changes of sepsis are based on systemic inflammatory responses induced by severe infection [19]. Inflammatory mediators, cytokines, reactive oxygen free radicals, mitochondrial dysfunction, and gene expression can directly damage cardiomyocytes or lead to the abnormal energy metabolism of cardiomyocytes, resulting in myocardial injury [20]. Sepsis3.0 defines septic shock as a subtype of sepsis. If circulation cannot be maintained after initial positive goal-directed fluid resuscitation, vasoactive drugs should be used to ensure tissue perfusion pressure [21]. However, up to now, there remains a relative lack of effective indicators that can help identify sepsis-related cardiac diastolic dysfunction in the early stage [22]. The secretion and release of natriuretic peptides increase rapidly after myocardial injury, left ventricular filling pressure, and wall tension, which has a specific diagnostic value for diastolic cardiac insufficiency and myocardial injury [23]. However, many factors such as age, renal insufficiency, and pulmonary hypertension may affect the secretion of natriuretic peptides, especially in sepsis patients, which may be related to the pathophysiological changes and volume resuscitation of sepsis [24,25]. Therefore, finding more reliable and accurate biomarkers in clinical work is necessary to promote the implementation of clinical decision-making.

By drawing ROC curves based on various biomarkers, this study showed that PCT has a high predictive value for the prognosis of sepsis patients complicated with myocardial injury. Therefore, this study suggests that PCT may be a sensitive biomarker. PCT, composed of 116AA, is a kind of procalcitonin peptide with no hormone activity, which is transcribed and synthesized by the Calc-1 gene of thyroid medulla cells, which is no more than 0.05 µg/L in healthy people. PCT can rapidly increase when there is a severe bacterial infection. When it exceeds 0.05 µg/L, we should pay attention to the possibility of sepsis. Therefore, PCT is considered an important sign of bacterial infection and a reliable index to judge the severity and prognosis of the disease. Concurrently, PCT can guide the use of early antibiotics and provide a basis for selecting antibiotics, thus reducing the mortality of infected patients. The results of the univariate analysis showed that PCT in the death group was significantly higher than that in the survival group, and unconditional Logistic regression analysis also showed that PCT was a risk factor for the prognosis of sepsis. According to some studies, the level of PCT is positively correlated with the degree of inflammation and can be reduced to normal by improving the condition. Through this study, we also confirmed the importance of dynamic monitoring of PCT levels in judging the disease and guiding the treatment of sepsis.

This study still has some limitations. As it is a retrospective study, the level of evidence is not high. Therefore, this work intends to conduct a prospective cohort study to verify the conclusions further.

Sepsis patients with myocardial injury, shock, and pulmonary infection are independent risk factors for death. Based on myocardial injury, shock, hypertension, MRSA, cardiac insufficiency, personal tumor history, pulmonary infection, abdominal infection, and urinary system infection, a predictive model and internal verification results show that the model is highly reliable. Therefore, PCT is a sensitive biomarker for predicting sepsis with myocardial injury.

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Additional information

Supplementary content related to this article has been published online at XXXX.

Data availability statement

All data generated or analyzed during this study are included in this published article.

CRediT authorship contribution statement

Weiwei Qian: Writing – review & editing, Writing – original draft, Software, Methodology, Funding acquisition, Formal analysis, Data curation, Conceptualization. **Cunqiao Han:** Writing – review & editing, Writing – original draft, Formal analysis, Data curation. **Shenglong Xie:** Writing – original draft, Visualization, Project administration, Formal analysis, Conceptualization. **Shuyun Xu:** Writing – review & editing, Writing – original draft, Visualization, Validation, Supervision, Software, Resources, Project administration, Methodology, Investigation, Funding acquisition, Formal analysis, Data curation, Conceptualization.

Declaration of competing interest

The authors declare the following financial interests/personal relationships which may be considered as potential competing interests: Shuyun Xu reports financial support was provided by Sichuan Provincial Science and Technology Department Project. If there are other authors, they declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.heliyon.2024.e27209>.

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