Serum N-terminal pro-B-type natriuretic peptide and cystatin C for acute kidney injury detection in critically ill adults in China: a prospective, observational study

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ABSTRACT

Objective Serum N-terminal pro-B-type natriuretic peptide (NT-proBNP) and cystatin C (sCysC) are available clinically and beneficial in diagnosing acute kidney injury (AKI). Our purpose is to identify the performance of their combined diagnosis for AKI in critically ill patients.

Design A prospectively recruited, observational study was performed.

Setting Adults admitted to the intensive care unit of a tertiary hospital in China.

Participants A total of 1222 critically ill patients were enrolled in the study.

Main outcome measures To identify the performance of the combined diagnosis of serum NT-proBNP and sCysC for critically ill patients. The area under the receiver operating characteristic curve (AUC-ROC), category-free net reclassification index (NRI) and incremental discrimination improvement (IDI) were utilised for comparing the discriminative powers of a combined and single biomarker adjusted model of clinical variables enriched with NT-proBNP and sCysC for AKI.

Results AKI was detected in 256 out of 1222 included patients (20.9%). AUC-ROC for NT-proBNP and sCysC to detect AKI had a significantly higher accuracy than any individual biomarker (p<0.05). After multivariate adjustment, a level of serum NT-proBNP >204 pg/mL was associated with 3.5-fold higher odds for AKI compared with those below the cut-off value. Similar results were obtained for sCysC levels (p<0.001). To detect AKI, adding NT-proBNP and sCysC to a clinical model further increased the AUC-ROC to 0.859 beyond that of the clinical model with or without sCysC (p<0.05). Moreover, the addition of these two to the clinical model significantly improved risk reclassification of AKI beyond that of the clinical model alone or with single biomarker (p<0.05), as measured by NRI and IDI.

Conclusions In critically ill individuals, serum NT-proBNP, sCysC and clinical risk factors combination improve the discriminative power for diagnosing AKI.

INTRODUCTION

Acute kidney injury (AKI) is a predominant clinical syndrome affecting more than 50% of patients who underwent treatment at intensive care unit (ICU).1–3 Increased morbidity, mortality, hospitalisation length and cost are extremely related to AKI,4–6 so early recognition of AKI is critical to guiding management. Urine production and serum creatinine were employed as diagnostic criteria for AKI in accordance with the recommendations of Kidney Disease Improving Global Outcomes (KDIGO) guidelines.7 Changes in sCr or urine production cannot recognise early renal tubular injury prior to a reduction in glomerular filtration rate.8,9 Therefore, early and reliable AKI biomarkers are necessary to promote timely intervention and minimise complications. However, it is impossible for an individual biomarker to adequately evaluate the risk of AKI as a complex multifactorial syndrome.3,10 Combining diverse biomarkers in a clinical model evaluation could enhance early detection of AKI in critically ill patients.11,12

A haemodynamic marker stress, N-terminal pro-B-type natriuretic peptide (NT-proBNP), has recently received attention as a potential
predictor of AKI in a wide diversity of clinical settings. High serum NT-proBNP level reflects haemodynamic instability, myocardial wall stress, myocardial ischaemia, volume overload, sympathetic nervous system and renin-angiotensin-aldosterone system activation, all of which may contribute to AKI incidence. However, there are finite data on clinical use of NT-proBNP for detecting AKI among critically ill patients.

Serum cystatin C (sCysC), a 122-amino acid low-molecular-weight protein (13 kDa), is a marker of glomerular filtration. CysC has a half-life of about one-third that of sCr, causing CysC to reach homeostasis three times faster. These properties promote sCysC as an alternative marker of renal function. To date, sCysC has been displayed to be conductive for the early identification of all-cause AKI.

Although serum NT-proBNP and sCysC are mainly used for the prognosis assessment of heart disease and the prediction of acute cardiac events with AKI, the diagnostic accuracy of their combination for AKI in ICU remains unknown. We performed a prospective study in the present research to evaluate the performance of serum NT-proBNP and sCysC at ICU admission, both independently and in combination, for AKI determination among critically ill adults.

METHODS

Study design and participants

At Guangdong Provincial People’s Hospital in China, a tertiary care hospital, a prospectively recruited observational research was performed in a mixed medical–surgical ICU. From December 2016 to December 2017, we consecutively enrolled patients aged 18 and up in a row. Pregnancy, renal replacement therapy prior to ICU admittance, nephrectomy, end-stage renal disease, renal transplant, rejection of consent or missing admission information were all exclusion criteria. The study protocol strengthened the reporting requirements of observational studies in epidemiology and standards for the reporting of diagnostic accuracy. This study protocol was authorised by ethics committee of Guangdong Provincial People’s Hospital. Additionally, all procedures were carried out consistent with applicable regulations and guidelines. All patients provided written informed consent.

Data collection

Clinical baseline data were collected prospectively. Within 1 hour following ICU admission, blood samples were collected contemporaneously to determine sCr, serum NT-proBNP and sCysC. Within 24 hours after collection, all samples were analysed in the Guangdong Provincial People’s Hospital central laboratory utilising standard protocol. When a patient is admitted to the ICU, sCr was measured and subsequently done at least once a day until discharge as part of routine clinical care. Throughout admission to ICU, urine production was also been recorded. We evaluated the following clinical variables: age, gender, body mass index, pre-existing clinical conditions, sepsis, admission type, baseline sCr, baseline estimated glomerular filtration rate (eGFR), Acute Physiology and Chronic Health Evaluation (APACHE) II score, ICU mortality, in-hospital mortality, duration of ICU admittance, length of hospitalisation, ICU costs and total costs. The outcome was the incidence of AKI following ICU enrolment within 1 week. To calculate eGFR, we employed the Chronic Kidney Disease (CKD) Epidemiology Collaboration creatinine equation.

Definitions

The KDIGO classification criteria were utilised to define AKI: as a rise in sCr by ≥0.3 mg/dL (26.5 µmol/L) within 48 hours or a rise in sCr to ≥1.5 times the baseline within 1 week, or urine output <0.5 mL/kg/hour for 6 hours after ICU admission.

Relying on following principles, ranked in descending order of preference, a baseline sCr was affirmed: (1) prior to ICU admittance, the most recent pre-ICU value between 30 and 365 days; (2) a stable pre-ICU value >365 days prior to ICU admittance for patients <40 years of age (stable definition is being within 15% of the lowest ICU measurement); (3) pre-ICU value >365 days prior to ICU admittance and lower than the initial sCr at ICU admission; (4) a pre-ICU value (within 3 and 39 days prior to ICU admittance) lower than or equal to the initial sCr on ICU admittance not obviously during AKI and (5) the least sCr value obtained at initial ICU admittance, the most recent ICU value or the lowest value achieved to a 365-day follow-up.

Biomarker measurement

The levels of sCysC and sCr were quantified through the UniCel DxC 800 Synchron system usage in compliance with the manufacturer’s instructions (Beckman Coulter, Brea, CA, USA). For sCysC, the intra-assay and interassay variation coefficients were 10% and 5%, respectively. Levels of serum NT-proBNP were quantified through an electrochemiluminescence immunoassay employing a Cobas e602 system usage (Roche Diagnostics, Germany). For NT-proBNP, the uppermost limit of normal for those who seem to be healthy (95th percentage) has been 125 pg/mL. The coefficient of interassay variation for NT-proBNP was <5%. Each patient’s clinical features were blinded to the personnel measuring the biomarkers.

Patient and public involvement

Patients and/or the public were not involved in the design, or conduct, or reporting or dissemination plans of this research.

Statistical analysis

For statistical analyses, we utilised SPSS V.21.0 (SPSS, Chicago, IL, USA), MedCalc V.18.2.1 (MedCalc Software, Ostend, Belgium) and R V.4.1.1 (R Foundation for Statistical Computing, Vienna, Austria). The mean±SD or median (25th to 75th percentage, IQR) had been utilised
to express continuous variables. Numbers (per cent) were utilised to represent categorical variables. For non-normally distributed continuous variables, the Wilcoxon rank-sum test was employed for intergroup difference measurement, and for categorical variables, the $\chi^2$ or Fisher’s exact test was deployed.

NT-proBNP concentrations were extremely skewed and therefore were log10 transformed before inclusion in the models. Areas under the receiver operating characteristic curves (AUC-ROCs) were computed. The method exploited by DeLong et al was used to compare AUC-ROCs between groups. The biomarkers’ sensitivity, specificity, positive and negative predictive values, positive and negative likelihood ratios of the biomarkers were calculated. Youden’s index for AKI detection identified the optimal cut-off values for individual biomarkers and their combination.

Logistic analysis was utilised to compute the ORs and 95% CIs for each factor: to identify the independent risk factors of AKI, with a forward stepwise method, in which the clinical variables with $p<0.10$ in univariate analysis were incorporated into the multivariate logistic model. We categorised NT-proBNP and sCysC levels according to their cut-off values and then performed logistic regression on the created variables. We conducted multiple logistic regression analyses to calculate the adjusted ORs of AKI, which was based on the clinical risk factors for AKI.

The performance of AKI detection after adding NT-proBNP and sCysC, or any of them, into the clinical model as categorical variables was assessed by AUC-ROC, category-free net reclassification improvement (NRI) index and integrated discrimination improvement (IDI) index, as described previously. To better quantify how accurately the reference and reclassification model would perform with independent data, we adopted a 10-fold cross-validation. All the tests were two-tailed, and $p<0.05$ was regarded statistically significant.

RESULTS

Clinical data and outcomes

A total of 150 (10.9%) of the 1372 adult patients enrolled in the study were excluded (figure 1). Therefore, 1222 patients were enrolled and AKI occurred in 256 patients (20.9%). Patient baseline variables and outcomes are exhibited in table 1. In comparison to patients without AKI, patients with AKI were elderly and were observed more frequently in those with comorbidities, including CKD, hypertension, coronary artery disease (CAD), diabetes mellitus (DM), heart failure (HF), cerebrovascular disease and chronic obstructive pulmonary disease (COPD). Increased sCr, serum NT-proBNP and sCysC levels at admission, as well as increased APACHE II scores, were more prevalent in patients with AKI. The baseline sCr and eGFR did not show significant differences between the two groups.

Indeed, patients with AKI had a higher risk of adverse outcomes, a higher percentage of ICU and in-hospital mortality, higher expenses, a longer hospitalisation duration and a longer stay in ICU (p<0.001) compared with those without AKI.

Detective abilities of the two biomarkers for AKI

To demonstrate the ability of these biomarkers for AKI detection, we used AUC-ROCs to calculate the two biomarkers, respectively, and in combination. AUC-ROCs for NT-proBNP and sCysC were computed for AKI detection (0.821 and 0.766, respectively). For AKI detection, NT-proBNP had a sensitivity of 78% and a specificity of 75%, while sCysC had high specificity but limited sensitivity. The cut-off values for NT-proBNP and sCysC were 204 pg/mL and 1.02 mg/L, respectively, yielding good sensitivity and specificity. We included NT-proBNP and sCysC in a multivariate logistic regression model to derive their combined AUC for comparison with the single biomarkers. The AUC-ROCs for AKI presented a better
performance by NT-proBNP and sCysC (0.832) than any individual biomarker (p<0.05, table 2 and figure 2).

Multivariate logistic regression analyses of the two biomarkers for AKI detection
We stratified patients based on the two cut-off values of serum NT-proBNP and sCysC levels into two categories, respectively. Compared with those with serum NT-proBNP <204 pg/mL, patients with serum NT-proBNP ≥204 pg/mL on admission exhibited a higher incidence of AKI (7.2% vs 45.1%, p<0.001). AKI was found to have higher incidence (46.5%) in patients with sCysC ≥1.02 mg/L compared with those with sCysC <1.02 mg/L (10.2%) (p<0.001). Following clinical variables adjustment (including CKD, HF, sepsis, admission type, sCr at admission and APACHE II scores, table 3), a level of serum NT-proBNP ≥204 pg/mL was associated with 3.5-fold
higher odds for AKI compared with NT-proBNP level below the cut-off value (p<0.001). Similarly, patients with sCysC ≥1.02 mg/L were linked to 2.6-fold greater odds for AKI compared with those with sCysC <1.02 mg/L (p<0.001) (table 4).

**Discrimination and reclassification of the combination of biomarkers and clinical models for AKI**

Considering the effect of adding NT-proBNP and sCysC, or any of them, to a clinical model as categorical variables for AKI detection, logistic regression analysis was employed. On ICU admittance for AKI diagnosis, potential available variables including age, sepsis, admission type, sCr and APACHE II scores as well as comorbidities, including hypertension, DM, CKD, cerebrovascular disease, COPD, CAD and HF were considered. The clinical model for detecting AKI involved CKD, HF, sepsis, admission type, sCr at admission and APACHE II scores (table 3). The cross-validated baseline performance characterised by accuracy and Kappa for the clinical model was 0.844 and 0.448.

To evaluate the enhancement of discriminative capacity, a panel of NT-proBNP and sCysC was introduced to the above-mentioned model. As shown in table 5, compared with the clinical model, the addition of NT-proBNP to the clinical model had a higher AUC-ROC (p<0.05), and no statistically significant variation existed when sCyC was added to the clinical model. However, the risk reclassification was markedly improved through the addition of NT-proBNP or sCyC to the clinical model, as measured by category-free NRI and IDI (p<0.05). Adding NT-proBNP and sCyC to a clinical model for AKI detection further increased the AUC-ROC to 0.859 beyond that of the clinical model with or without any individual biomarkers (p<0.05), with maximum NRI (0.531) and IDI (0.038).

**DISCUSSION**

The study’s key finding was that the combination of NT-proBNP and sCysC yields greater discriminative ability for AKI detection at ICU admission with or without a clinical model in critically ill adults. The finding indicates that assessing both serum NT-proBNP and sCysC levels on admission may assist with the early diagnosis and risk stratification of AKI in critically ill adults. One of the most prevalent complications occurring in a variety of clinical settings is AKI, especially for critically ill patients.4 6 36 37 The development of AKI, as the
same as its severity, is strongly associated with increased mortality. However, early identification is challenging when sCr or urine production changes are used to detect AKI, and precise clinical predictors are not widely known. Numerous studies have found and confirmed the accuracy and additional clinical benefits of these renal biomarkers for early AKI diagnosis, such as insulin-like growth factor-binding protein, matrix metalloproteinase-7, tissue inhibitor metalloproteinase-2, angiotensinogen, sCysC and neutrophil gelatinase-associated lipocalin. However, some novel biomarkers are not being used in clinical practice, due to insufficient evidence or previously unavailable commercially. Thus, the rational application of clinically available biomarkers is more practical and economical.

NT-proBNP, a widely used marker of haemodynamic stress, is a polypeptide secreted by the ventricles and its role is to facilitate natriuresis. In patients with elevated central venous pressure (CVP), the raised pressure may be transmitted to the renal veins, resulting in renal congestion and reduced glomerular filtration. It makes sense that NT-proBNP is associated with the development of AKI. The clinical application of serum NT-proBNP in cardiac disease has been extensively confirmed and finds its predictive value for AKI development in patients with HF, coronary angiography or percutaneous coronary intervention. Several studies have elucidated the link between NT-proBNP and AKI incidence after cardiac or non-cardiac surgery. However, its utility for AKI detection has not been fully evaluated in general ICU, and it is necessary to verify the reliability and universality of NT-proBNP in heterogeneous populations. The data demonstrated here both maintain NT-proBNP as a risk factor for AKI and also indicate that it can enhance the risk reclassification and discrimination for AKI in ICU.

### Table 3 Logistic analyses of clinical risk factors for AKI detection

<table>
<thead>
<tr>
<th>Variables</th>
<th>Univariate analysis</th>
<th>Multivariate model</th>
<th>P value</th>
<th>P value</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>OR (95% CI)</td>
<td>P value</td>
<td>OR (95% CI)</td>
<td>P value</td>
</tr>
<tr>
<td>Age</td>
<td>1.029 (1.019 to 1.038)</td>
<td>&lt;0.001</td>
<td>4.182 (1.250 to 13.998)</td>
<td>0.020</td>
</tr>
<tr>
<td>Hypertension</td>
<td>2.152 (1.577 to 2.937)</td>
<td>&lt;0.001</td>
<td>3.493 (2.023 to 6.032)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Diabetes mellitus</td>
<td>3.314 (2.242 to 4.898)</td>
<td>&lt;0.001</td>
<td>3.237 (2.157 to 4.858)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Chronic kidney disease</td>
<td>17.175 (6.410 to 46.023)</td>
<td>&lt;0.001</td>
<td>1.538 (1.149 to 2.058)</td>
<td>0.004</td>
</tr>
<tr>
<td>Cerebrovascular disease</td>
<td>2.802 (1.903 to 4.126)</td>
<td>&lt;0.001</td>
<td>1.538 (1.149 to 2.058)</td>
<td>0.004</td>
</tr>
<tr>
<td>Chronic obstructive pulmonary disease</td>
<td>4.222 (2.143 to 8.316)</td>
<td>&lt;0.001</td>
<td>1.538 (1.149 to 2.058)</td>
<td>0.004</td>
</tr>
<tr>
<td>Coronary artery disease</td>
<td>3.342 (1.910 to 5.849)</td>
<td>&lt;0.001</td>
<td>1.538 (1.149 to 2.058)</td>
<td>0.004</td>
</tr>
<tr>
<td>Heart failure</td>
<td>9.997 (4.544 to 21.997)</td>
<td>&lt;0.001</td>
<td>3.487 (1.365 to 8.911)</td>
<td>0.009</td>
</tr>
<tr>
<td>Sepsis</td>
<td>12.608 (7.914 to 20.084)</td>
<td>&lt;0.001</td>
<td>5.033 (2.914 to 8.692)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Admission type</td>
<td></td>
<td></td>
<td>1.0 (referent)</td>
<td>1.0 (referent)</td>
</tr>
<tr>
<td>Elective surgical</td>
<td>1.0 (referent)</td>
<td></td>
<td>1.0 (referent)</td>
<td>1.0 (referent)</td>
</tr>
<tr>
<td>Emergency surgical</td>
<td>6.121 (3.735 to 10.030)</td>
<td>&lt;0.001</td>
<td>3.493 (2.023 to 6.032)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Medical</td>
<td>9.409 (6.791 to 13.037)</td>
<td>&lt;0.001</td>
<td>3.237 (2.157 to 4.858)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>sCr</td>
<td>3.135 (2.385 to 4.120)</td>
<td>&lt;0.001</td>
<td>1.538 (1.149 to 2.058)</td>
<td>0.004</td>
</tr>
<tr>
<td>APACHE II score</td>
<td>1.215 (1.182 to 1.249)</td>
<td>&lt;0.001</td>
<td>1.538 (1.149 to 2.058)</td>
<td>0.004</td>
</tr>
</tbody>
</table>

AKI, acute kidney injury; APACHEII score, Acute Physiology and Chronic Health Evaluation II score; sCr, serum creatinine.

### Table 4 Multivariate logistic regression analyses of the two biomarkers for AKI detection

<table>
<thead>
<tr>
<th>Variables</th>
<th>AKI, %</th>
<th>Unadjusted OR (95% CI)</th>
<th>P value</th>
<th>Adjusted OR* (95% CI)</th>
<th>P value</th>
</tr>
</thead>
<tbody>
<tr>
<td>NT-proBNP (pg/mL, n)</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>&lt;204, 779</td>
<td>7.2</td>
<td>1.0 (referent)</td>
<td></td>
<td>1.0 (referent)</td>
<td></td>
</tr>
<tr>
<td>≥204, 443</td>
<td>45.1</td>
<td>10.626 (7.639 to 14.781)</td>
<td>&lt;0.001</td>
<td>3.460 (2.307 to 5.189)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>sCysC layered by the cut-off value (mg/L, n)</td>
<td></td>
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</tr>
<tr>
<td>&lt;1.02, 861</td>
<td>10.2</td>
<td>1.0 (referent)</td>
<td></td>
<td>1.0 (referent)</td>
<td></td>
</tr>
<tr>
<td>≥1.02, 361</td>
<td>46.5</td>
<td>7.646 (5.651 to 10.345)</td>
<td>&lt;0.001</td>
<td>2.649 (1.842 to 3.810)</td>
<td>&lt;0.001</td>
</tr>
</tbody>
</table>

*Adjusted for chronic kidney disease, heart failure, sepsis, admission type, serum creatinine at admission and Acute Physiology and Chronic Health Evaluation II score.

AKI, acute kidney injury; NT-proBNP, N-terminal pro-B-type natriuretic peptide; sCysC, serum cystatin C.
et al showed that combining urinary liver-detect AKI can improve predictive abilities.5 49 50 Naruse revealed that combining different biomarkers utilised to be combined to detect AKI. Numerous studies have shown that AUC-ROC analysis when two biomarkers were added to the clinical model significantly enhanced risk stratification for AKI.25 In the present cohort, sCysC had a slightly higher specificity than NT-proBNP in the detection of AKI, but its sensitivity is limited. The ability of sCysC for detecting AKI was fine, but its AUC-ROC was not as great as NT-proBNP. Moreover, the combination of NT-proBNP and sCysC at ICU admission had the highest AUC-ROC. These data indicate that the simultaneous measurement of serum NT-proBNP and sCysC at ICU admission could improve the early identification of AKI.

Biomarkers for AKI, which indicates the various underlying pathophysiological mechanisms involved in AKI incidence, might be superior to individual biomarkers alone.3 6 It is also important that these biomarkers profit from being easily measurable, readily accessible, comparatively cheap and with an elevated level of sensitivity and specificity. In the present study, for AKI detection, NT-proBNP acted as a haemodynamic stress biomarker with high sensitivity and specificity, and sCysC had high specificity as a functional biomarker. Even after clinical risk variables adjustment, elevated NT-proBNP is an independent risk factor for AKI. The NT-proBNP addition to the clinical model significantly enhanced risk binding protein and serum NT-proBNP can enhance early prediction of AKI in patients in medical cardiac ICUs. Similar to such studies, we identified that the combination of two markers, serum NT-proBNP and sCysC, improved the diagnostic performance of AKI.

CysC is a glomerular filtration biomarker that can be utilised to anticipate the development of AKI and undesirable outcomes.8 9 The application and performance of CysC for AKI prediction have been demonstrated in various clinical settings.5 11 Similar to such studies, we identified that the combination of two markers, serum NT-proBNP and sCysC, improved the diagnostic performance of AKI.

Table 5 Discrimination and reclassification of the combination of biomarkers and clinical model for AKI

<table>
<thead>
<tr>
<th>Model</th>
<th>AUC-ROC (95% CI)</th>
<th>P value*</th>
<th>Category-free NRI (95% CI)</th>
<th>P value*</th>
<th>IDI (95% CI)</th>
<th>P value*</th>
</tr>
</thead>
<tbody>
<tr>
<td>Clinical model†</td>
<td>0.840 (0.812 to 0.868)</td>
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</tr>
<tr>
<td>Clinical model+sCysC</td>
<td>0.847 (0.819 to 0.874)</td>
<td>0.163</td>
<td>0.193 (0.052 to 0.405)</td>
<td>0.036</td>
<td>0.017 (0.009 to 0.026)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Clinical model+NT-proBNP</td>
<td>0.855 (0.828 to 0.882)</td>
<td>0.013</td>
<td>0.462 (0.196 to 0.747)</td>
<td>0.001</td>
<td>0.028 (0.016 to 0.039)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Clinical model+NT-proBNP+sCysC</td>
<td>0.859 (0.832 to 0.885)</td>
<td>0.006</td>
<td>0.531 (0.238 to 0.741)</td>
<td>&lt;0.001</td>
<td>0.038 (0.025 to 0.051)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Clinical model+NT-proBNP+sCysC clinical model+sCysC</td>
<td></td>
<td></td>
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<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Clinical model+NT-proBNP+sCysC clinical model+NT-proBNP</td>
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</table>

Hosmer–Lemeshow goodness-of-fit test: for clinical model, χ² value=14.249 (p=0.075); for clinical model+sCysC, χ² value=6.971 (p=0.54); for clinical model+NT-proBNP, χ² value=9.362 (p=0.313) or clinical model+NT-proBNP+sCysC, χ² value=4.245 (p=0.834).

*Biomarker+clinical model vs clinical model.
†The clinical model for detecting AKI is composed of chronic kidney disease, heart failure, sepsis, admission type, serum creatinine at admission and Acute Physiology and Chronic Health Evaluation II score.

AKI, acute kidney injury; AUC-ROC, area under the receiver operating characteristic curve; IDI, incremental discrimination improvement; NRI, net reclassification index; NT-proBNP, N-terminal pro-B-type natriuretic peptide; sCysC, serum cystatin C.

A single biomarker is insufficient to express the multiple pathophysiological mechanisms of AKI, especially for critically ill patients, as AKI is a heterogeneous syndrome.9 10 There is no consensus on which specific markers should be combined to detect AKI. Numerous studies have revealed that combining different biomarkers utilised to detect AKI can improve predictive abilities.5 49 50 Naruse et al showed that combining urinary liver-type fatty-acid binding protein and serum NT-proBNP can enhance early prediction of AKI in patients in medical cardiac ICUs. Similar to such studies, we identified that the combination of two markers, serum NT-proBNP and sCysC, improved the diagnostic performance of AKI.

CysC is a glomerular filtration biomarker that can be utilised to anticipate the development of AKI and undesirable outcomes.8 9 The application and performance of CysC for AKI prediction have been demonstrated in various clinical settings.5 11 Similar to such studies, we identified that the combination of two markers, serum NT-proBNP and sCysC, improved the diagnostic performance of AKI.
reclassification, as demonstrated by category-free NRI and IDI. Moreover, our results identified that the addition of NT-proBNP to sCysC markedly improved their detectable abilities as biomarkers. Adding biomarkers to the clinical model further improved the diagnostic accuracy of AKI, as measured by AUC-ROCs. These data suggest that a single biomarker is insufficient for early diagnosis of AKI. Therefore, the method of combining different biomarkers may be of greater use.

There are some limitations for this study. First, it was a single-centre study, with an unproven external validity. Second, there were only 26 patients with CKD and 31 patients with HF enrolled, and hence we were unable to stratify our group relying on eGFR or cardiac function at baseline. Accordingly, future studies should be administered in these subgroups. Last, we did not accomplish routine echocardiography in all patients to associate NT-proBNP levels with ventricular dilatation or other pathways that may promote NT-proBNP release. Despite these limitations, we believe that our findings have clinical implications and should facilitate further research to confirm our results.

CONCLUSION
In the present cohort, simultaneous measurement of NT-proBNP and sCysC at ICU admission increases the early identification of AKI beyond that of biomarker in isolation, and that the combination of the two biomarkers and clinical risk factors improves the discriminative ability for AKI detection in critically ill adults.

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