



# Hepatocyte growth factor and soluble cMet levels in plasma are prognostic biomarkers of mortality in patients with severe acute kidney injury

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**Background:** Hepatocyte growth factor (HGF)/cMet pathway is necessary for repair and regeneration following acute kidney injury (AKI). We evaluated the clinical potential of plasma HGF and soluble cMet as prognostic biomarkers for severe AKI requiring continuous renal replacement therapy (CRRT).

**Methods:** One hundred thirty-six patients with severe AKI who participated in the VENUS (volume management under body composition monitoring in critically ill patients on CRRT) trial between 2017 and 2019 were enrolled in this study. We investigated associations between plasma HGF and cMet concentrations and all-cause mortality.

**Results:** Plasma HGF and soluble cMet levels were positively correlated. Patients were divided into three groups based on their HGF and soluble cMet concentrations. The day D 0, D2, and D7 highest concentration HGF groups had significantly higher in-hospital mortality after adjusting for sex, body mass index, Acute Physiology and Chronic Health Evaluation II, and age-adjusted Charlson comorbidity index score, especially on D7 (hazard ratio, 4.26; 95% confidence interval, 1.71–10.62;  $p = 0.002$ ). D7 soluble cMet level was also associated with mortality. Receiver operating characteristic curve analysis indicated that D7 HGF and soluble cMet levels were best at predicting mortality. Addition of plasma HGF and soluble cMet to conventional prognostic indices significantly improved the predictive value for mortality on D7. However, plasma HGF and soluble cMet were not associated with fluid status.

**Conclusion:** Plasma HGF and soluble cMet levels were significant predictors of the outcomes of severe AKI patients undergoing CRRT. There was no correlation between plasma HGF and soluble cMet levels and fluid balance.

**Keywords:** Acute kidney injury, Biomarkers, Continuous renal replacement therapy, Hepatocyte growth factor, Soluble c-Met

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## Introduction

Acute kidney injury (AKI) occurs in 10% of hospitalized patients and 30% of critically ill patients each year, and its incidence is increasing [1]. Severe AKI is one of the leading causes of death in critically ill patients. Despite recent technical advances in AKI management, mortality rates associated with severe AKI are approximately 40% to 50% [2,3].

Reliable biomarkers are important for treating AKI patients and predicting outcomes. Several studies have reported various novel biomarkers for AKI that can detect kidney injury before serum creatinine (sCr), such as tissue inhibitor of metalloproteinase (TIMP)-2, insulin-like growth factor-binding protein 7 (IGFBP7) [4], and urinary matrix metalloproteinase 7 [5]. However, few studies have investigated prognostic predictors of the outcomes of patients with severe AKI undergoing continuous renal replacement therapy (CRRT).

Hepatocyte growth factor (HGF) and its tyrosine kinase receptor, cMet, have important roles in regulating wound healing, cell proliferation, and tissue fibrosis [6-8]. Several experimental studies have indicated that HGF has multiple protective effects on physiological and pathophysiological processes in the kidney, including accelerated DNA synthesis and improved kidney cell regeneration, apoptosis, and necrosis (including that of proximal tubular cells, mesangial cells, podocytes, and endothelial cells) [7,8]. Activation of the HGF/cMet axis can improve acute and chronic kidney disease (CKD) by inhibiting oxidative stress, apoptosis, fibrosis, and inflammation [9,10]. We have also confirmed that cMet activation by agonistic antibodies can improve renal fibrosis [11].

A recent study showed that plasma HGF level predicted all-cause mortality and cardiovascular mortality in the general population [12]. Additionally, HGF and cMet can predict clinical outcomes of patients with various cancers [13,14]. Recent studies have reported that HGF and cMet are biomarkers for CKD [15,16]. However, the clinical predictive abilities of HGF and soluble cMet in AKI patients have not been well studied.

In this study, we investigated whether plasma HGF and soluble cMet levels could predict clinical outcomes of AKI patients who participated in the VENUS (volume management under body composition monitoring in critically ill patients on CRRT) trial. Furthermore, we investigated whether

plasma HGF and soluble cMet levels were associated with fluid balance.

## Methods

### Study design and population

A total of 136 severe AKI patients admitted to seven hospitals in Korea who participated in the VENUS trial between 2017 and 2019 were enrolled in this study [3]. The VENUS trial was designed to determine whether bioelectrical impedance analysis-guided fluid management could help achieve euvolemic status in patients treated with CRRT more efficiently than fluid management guided by a generally used quantification method. All participants were selected from patients who were scheduled to undergo CRRT for at least 72 hours. There was no mandatory standardized CRRT protocol. CRRT initiation was decided by each institution's physicians. CRRT settings, effluent dose, and target I/O balance were freely applied to maintain stable blood pressure to meet metabolic demands. In general, blood flow was set to 100–130 mL/min, and target effluent dose was set to 35 mL/kg/hr.

Exclusion criteria were the following: age younger than 18 years; imminent death (<24 hours); maintenance dialysis used before current hospitalization; any other major illness that, according to the investigators' judgment, would substantially increase the risk associated with the subject's participation in this study; and withdrawal of patient consent.

### Clinical data and sample collection

Clinical baseline characteristics data at the time of study enrollment, including age, sex, mean arterial pressure (MAP), body mass index (BMI), laboratory findings, contributing factors for AKI, all-cause mortality, outcome events include sepsis, cardiogenic diseases, pulmonary diseases, and cancer, among others, and Acute Physiology and Chronic Health Evaluation II (APACHE II) scores, and age-adjusted Charlson comorbidity index (Age-CCI) scores, were collected. Age-CCI is the combination of the age equivalence index and Charlson comorbidity index (CCI) [17]. For patients over 40 years old, the cumulative score was 1 point for each additional 10 years of age, and the score for age was added to the CCI. A completely resolved condition or current inactive surgery history was not considered a comorbid disease [18].

Laboratory evaluations included measurements of complete blood cell counts, electrolytes, sCr, total protein, albumin, calcium, and high-sensitivity C-reactive protein. Blood samples were centrifuged to extract plasma. The plasma was frozen, stored at  $-70^{\circ}\text{C}$ , and thawed before analysis.

All procedures were performed in accordance with the ethical standards of the institutional and/or national research committee and the 1964 Declaration of Helsinki and its later amendments or comparable ethical standards. All patients provided written informed consent at the time of their enrollment. The APACHE II score was designed to measure the disease severity of adult patients admitted to the intensive care unit (ICU) during the first 24 hours after admission. It is calculated using the patient's age and 12 routine physiological measurements (alveolar-arterial oxygen gradient [AaDO<sub>2</sub>] or partial pressure of oxygen [PaO<sub>2</sub>] depending on the fraction of inspired oxygen [FiO<sub>2</sub>], temperature, MAP, pH, heart rate, respiratory rate, sodium level, potassium level, creatinine level, hematocrit, white blood cell count, and Glasgow Coma Scale score). Approval to perform the study was obtained from the Institutional Review Boards of all participating centers (No. 20-2019-79). The trial protocol has been registered at <http://www.clinicaltrials.gov> (NCT03330626).

### Measurement of plasma soluble cMet and hepatocyte growth factor

Enzyme-linked immunosorbent assay kits were used to measure the plasma concentrations of HGF (DY294; Becton Dickinson, Minneapolis, MN, USA) and soluble cMet (KHO 2031; Invitrogen, Vienna, Austria) on day (D) 0, D2, and D7. Plasma HGF concentrations were measured after a 10-fold dilution and plasma soluble cMet concentration was measured after a two-fold dilution. Enzyme-linked immunosorbent assays were performed according to the manufacturers' instructions. All laboratory investigators were blinded to the sample source, and all measurements were performed in duplicate.

### Assessment of fluid status

Fluid status was measured by bioimpedance analysis using InBody S10 (InBody, Seoul, Korea) on D0, D2, and D7. Measurements of total body water (TBW), intracellular water

(ICW), and extracellular water (ECW) were obtained with InBody S10. Fluid state was based on ECW/TBW and TBW/height squared ( $\text{H}^2$ ).

### Clinical outcomes

Primary outcome was all-cause mortality after CRRT. We investigated the associations between HGF concentration and soluble cMet concentration and all-cause mortality. In this study, survival duration was calculated from the day of enrollment in the study to the day of discharge from the hospital or death. The relationship between all-cause mortality after 2 years of follow-up was also assessed, starting from the first enrolled patient. Secondary outcome was fluid status according to HGF and soluble cMet concentrations.

### Statistical analysis

Patients were classified into three groups based on their D0 HGF and soluble cMet concentrations. [Table 1](#) shows the HGF and soluble cMet concentrations of each group. Categorical variables, which were expressed as frequencies and proportions, were compared using the chi-square tests. After testing for normality, normally distributed continuous variables were expressed as mean  $\pm$  standard deviations and compared using Student t test or one-way analysis of variance. Nonnormally distributed variables were expressed as medians (interquartile ranges) and compared using the Mann-Whitney U or Kruskal-Wallis tests. To investigate the impact of D0, D2, and D7 plasma soluble cMet and HGF levels on mortality, Kaplan-Meier survival curves were constructed. Cox proportional hazard models with plasma soluble cMet or HGF levels were used for multivariate survival analyses. To examine the prognostic value of plasma HGF and soluble cMet levels at multiple time points, we calculated the statistical significance of differences between the areas under the curve (AUC) at three-time points (D0, D2, and D7). Survival duration was calculated from day 0 to evaluate the predictive effects of D2 and D7 soluble cMet and HGF. Receiver operating characteristic (ROC) curves and AUC were generated to evaluate the accuracy and predictive capability of each indicator of survival. To examine the incremental prognostic value before and after inclusion of plasma HGF and soluble cMet levels with traditional indices, including APACHE II and Age-CCI, we calculated the statis-

**Table 1. Baseline characteristics at day 0**

Clinical parameter	DO HGF			DO cMet			p-value	p-value
	Total (n = 136)	Group 1 (n = 45)	Group 2 (n = 46)	Group 3 (n = 45)	Group 1 (n = 45)	Group 2 (n = 46)		
Plasma HGF (pg/mL)	7,675.6 (5,231.7- 10,119.5)	1,881.2 (1,668.9-2,093.4)	4,270.6 (3,972.1-4,569.1)	16,821.9 (10,137.9- 23,506.0)	9,547.7 (2,596.7- 16,498.8)	6,250.8(4,184.6- 8,317.1)	7,228.2 (5,345.6-9,110.7)	0.54
Plasma cMet (pg/mL)	801.1 (715.2-887.0)	627.4 (562.3-692.4)	889.5 (704.6-1,074.4)	884.3 (720.4-1,048.2)	423.4 (389.4-457.4)	684.7 (666.8-702.7)	1,297.6 (1,117.7-1,477.5)	<0.001
Male sex	103 (75.7)	33 (73.3)	31 (67.4)	39 (86.7)	39 (86.7)	34 (73.9)	30 (66.7)	0.08
Age (yr)	66.2 (63.7-68.7)	66.4 (62.2-70.6)	69.2 (64.9-73.4)	63.0 (58.5-67.6)	62.8 (58.0-67.6)	68.3 (64.9-71.7)	67.5 (62.8-72.1)	0.16
Comorbidity								
Diabetes mellitus	57 (41.9)	21 (46.7)	14 (30.4)	22 (48.9)	14 (31.1)	23 (50.0)	20 (44.4)	0.17
Hypertension	60 (44.1)	20 (44.4)	20 (43.5)	20 (44.4)	17 (37.8)	22 (47.8)	21 (46.7)	0.57
Congestive heart disease	17 (12.5)	8 (17.8)	6 (13.0)	3 (6.7)	6 (13.3)	6 (13.0)	5 (11.1)	0.94
Cerebrovascular attack	12 (8.8)	2 (4.4)	6 (13.0)	4 (8.9)	5 (11.1)	4 (8.7)	3 (6.7)	0.76
Malignancy	34 (25.0)	7 (15.9)	15 (32.6)	12 (26.7)	9 (20.0)	12 (26.1)	13 (28.9)	0.61
Body mass index (kg/m <sup>2</sup> )	25.0 (24.3-25.6)	25.1 (24.0-26.3)	24.9 (23.7-26.1)	24.9 (23.9-25.8)	24.9 (23.8-26.0)	24.9 (23.8-26.0)	25.1 (24.0-26.2)	0.95
MAP (mmHg)	83.2 (80.5-85.9)	86.8 (82.7-90.9)	85.4 (80.6-90.1)	77.3 (72.3-82.4)	81.3 (76.0-86.5)	84.2 (79.5-88.8)	84.1 (79.7-88.6)	0.62
Baseline sCr (mg/dL)	1.9 (1.5-2.3)	2.1 (1.4-2.8)	1.9 (1.4-2.5)	1.5 (0.9-2.1)	1.7 (1.1-2.2)	2.0 (1.2-2.8)	2.0 (1.4-2.5)	0.69
Factor contributing to AKI								
Septic	73 (53.7)	21 (46.7)	23 (50.0)	29 (64.4)	26 (57.8)	23 (50.0)	24 (53.3)	0.76
Nephrotoxic	4 (2.9)	0 (0)	3 (6.5)	1 (2.2)	2 (4.4)	1 (2.2)	1 (2.2)	0.77
Major surgery	6 (4.4)	3 (6.7)	3 (6.5)	0 (0)	4 (8.9)	1 (2.2)	1 (2.2)	0.20
Ischemic	25 (18.4)	10 (22.2)	10 (21.7)	5 (11.1)	5 (11.1)	10 (21.7)	10 (22.2)	0.31
Others	38 (27.9)	15 (33.3)	13 (28.3)	10 (22.2)	12 (26.7)	14 (30.4)	12 (26.7)	0.90
APACHE II	30.3 (28.6-32.0)	28.0 (25.3-30.7)	29.8 (26.9-32.7)	33.0 (29.7-36.4)	27.2 (24.2-30.2)	30.7 (27.9-33.5)	32.9 (29.9-35.9)	0.03
SOFA	11.5 (10.5-12.4)	12.4 (10.0-14.8)	10.1 (9.0-11.2)	11.9 (10.7-13.2)	12.0 (10.3-13.7)	12.2 (10.1-14.2)	10.1 (9.0-11.3)	0.17
Age-CCI	2.9 (2.5-3.2)	2.6 (2.0-3.3)	3.2 (2.6-3.8)	2.8 (2.1-3.4)	2.4 (1.7-3.1)	2.8 (2.3-3.3)	3.4 (2.6-4.1)	0.12
Vasopressor	97 (71.3)	27 (60.0)	31 (67.4)	39 (86.7)	32 (71.1)	33 (71.7)	32 (71.1)	0.997
Duration from ICU to CRRT (day)	1.6 (1.0-2.6)	1.8 (0.6-3.0)	1.7 (0.5-2.9)	1.4 (0.6-2.3)	2.0 (0.7-3.2)	1.4 (0.6-2.2)	1.6 (0.4-2.8)	0.73
Laboratory findings at CRRT initiation								
WBC (10 <sup>3</sup> /μL)	14.5 (12.4-16.7)	12.0 (8.4-15.7)	16.3 (11.8-20.9)	15.2 (12.1-18.2)	14.3 (10.5-18.2)	14.0 (11.6-16.4)	15.2 (10.3-20.2)	0.89
Hemoglobin (g/dL)	10.0 (9.6-10.5)	9.6 (9.0-10.3)	9.7 (8.9-10.5)	10.7 (9.8-11.6)	10.1 (9.3-10.9)	10.0 (9.3-10.6)	10.0 (9.1-10.8)	0.94
Platelet (10 <sup>3</sup> /μL)	111.0 (98.0-124.0)	113.0 (92.3-133.8)	117.4 (90.2-144.6)	102.4 (82.3-122.6)	117.1 (95.1-139.1)	103.8 (81.1-126.5)	112.3 (88.0-136.6)	0.71
sCr (mg/dL)	3.0 (2.7-3.3)	2.7 (2.2-3.2)	2.8 (2.4-3.3)	3.4 (2.8-4.0)	2.8 (2.4-3.3)	2.9 (2.5-3.4)	3.2 (2.5-3.8)	0.69
Albumin (mg/dL)	2.7 (2.6-2.8)	2.7 (2.5-2.9)	2.7 (2.5-2.8)	2.7 (2.6-2.9)	2.7 (2.5-2.9)	2.8 (2.6-2.9)	2.6 (2.5-2.8)	0.36
CRP (mg/L)	14.2 (12.2-16.2)	14.4 (10.2-18.6)	13.5 (10.4-16.6)	14.7 (11.1-18.3)	17.1 (12.7-21.6)	12.9 (10.1-15.6)	12.7 (9.3-16.0)	0.14
Lactate (mmol/L)	5.7 (4.6-6.7)	2.8 (1.9-3.7)	6.1 (4.5-7.6)	8.1 (5.6-10.5)	5.5 (3.4-7.5)	5.7 (4.0-7.4)	5.8 (3.9-7.6)	0.98
pH	7.3 (7.3-7.4)	7.4 (7.3-7.4)	7.3 (7.3-7.4)	7.3 (7.3-7.3)	7.3 (7.3-7.4)	7.3 (7.3-7.4)	7.3 (7.3-7.4)	0.63

(Continued to the next page)

**Table 1. Continued**

Clinical parameter	D0 HGF			D0 cMet				
	Total (n = 136)	Group 1 (n = 45)	Group 2 (n = 46)	Group 3 (n = 45)	Group 1 (n = 45)	Group 2 (n = 46)	Group 3 (n = 45)	p-value
Outcome event	66 (48.5)	16 (35.6)	24 (52.2)	26 (57.8)	21 (46.7)	19 (41.3)	26 (57.8)	0.28
Sepsis	20 (14.7)	4 (8.9)	9 (19.6)	7 (15.6)	9 (20)	4 (8.7)	7 (15.6)	0.31
Pulmonary disease	11 (8.1)	4 (8.9)	3 (6.5)	4 (8.9)	4 (8.9)	4 (8.7)	3 (6.7)	0.91
Cardiologic disease	12 (8.8)	4 (8.9)	2 (4.3)	6 (13.3)	5 (11.1)	3 (6.5)	4 (8.9)	0.74
Cancer	7 (5.1)	2 (4.4)	4 (8.7)	1 (2.2)	1 (2.2)	2 (4.3)	4 (8.9)	0.34
Others	16 (11.8)	2 (4.4)	6 (13.0)	8 (17.8)	2 (4.4)	6 (13.0)	8 (17.8)	0.14

Data are expressed as median (interquartile range) or number (%).

Age-CrI, age-adjusted Charlson comorbidity index; AKI, acute kidney injury; APACHE II, Acute Physiology and Chronic Health Evaluation II; CRP, C-reactive protein; CRRT, continuous renal replacement therapy; D, day; HGF, hepatocyte growth factor; ICU, intensive care unit; MAP, mean arterial pressure; sCr, serum creatinine; SOFA, sequential organ failure assessment.

tical significance of differences between the AUC, integrated discrimination improvement (IDI) index, and category-free net reclassification improvement (cfNRI) index. We divided patients into four groups based on median HGF and soluble cMet levels to determine the relationship between the two indicators and mortality rate. Patients with low concentrations of HGF and soluble cMet were classified into group 1; individuals with high concentrations of both HGF and soluble cMet were classified into group 4. Those patients with a low HGF concentration but high soluble cMet concentration were assigned to group 2; those with a high concentration of HGF combined with a low concentration of soluble cMet were assigned to group 3. We evaluated fluid status on D0, D2, and D7 according to HGF and soluble cMet concentrations using Student t test. All statistical analyses were performed using IBM SPSS version 23 (IBM Corp., Armonk, NY, USA) and R program version 3.2.5 (The R Foundation for Statistical Computing, Vienna, Austria). The p-value of <0.05 indicated statistical significance.

**Results**

**Baseline characteristics**

The study cohort consisted of 136 patients divided into three groups based on their HGF and soluble cMet concentrations on D0. Table 1 provides clinical baseline characteristics based on D0 HGF and soluble cMet concentrations. There were no significant differences in diabetes, hypertension, BMI, sCr, or blood urea nitrogen among the three groups based on D0 HGF and soluble cMet concentrations. In the group with the highest D0 HGF concentration, the soluble cMet concentration was significantly higher (p = 0.02), MAP was significantly lower (p = 0.009), and more patients were on vasopressors (p = 0.02). The laboratory findings of patients in the highest D0 HGF group indicated higher lactate levels in this group (p < 0.001) and a lower pH (p = 0.004). The group with the highest D0 soluble cMet concentration had a higher APACHE II score than the other two groups (p = 0.03).

**Variations and correlations between hepatocyte growth factor and soluble cMet concentrations after continuous renal replacement therapy**

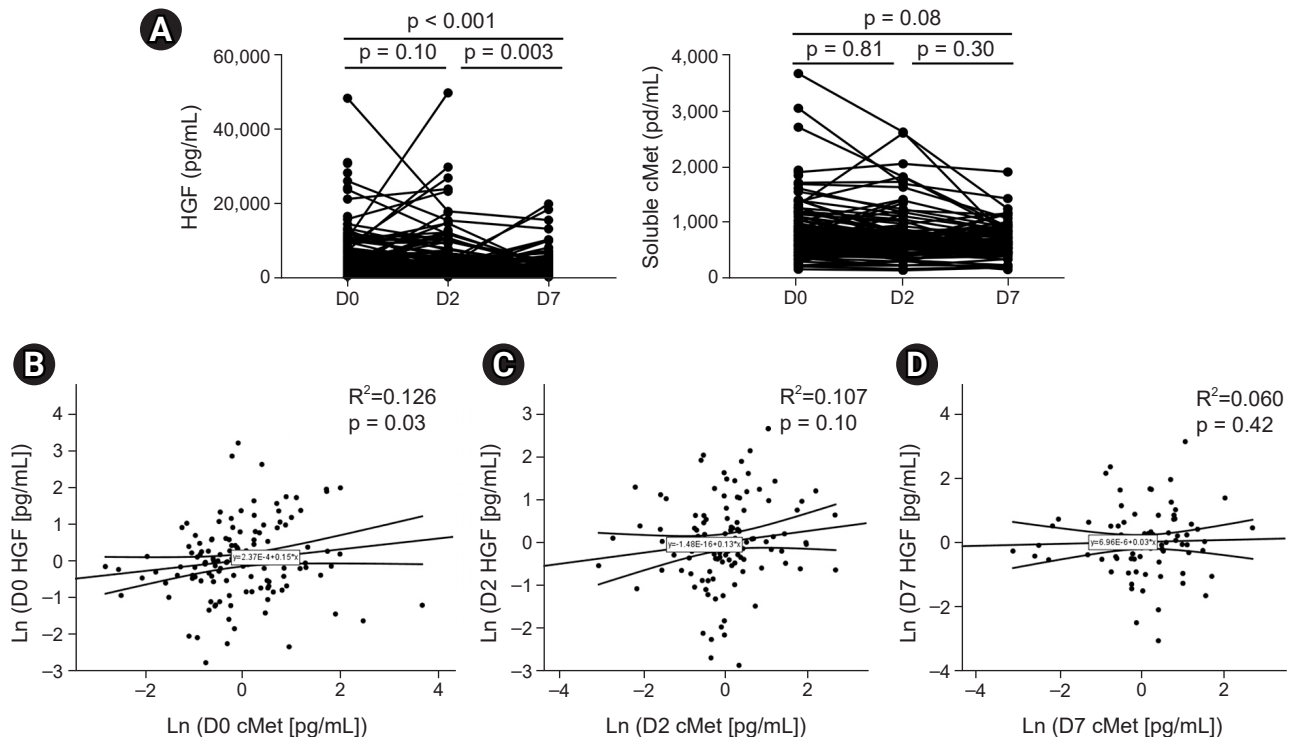
Both HGF and soluble cMet concentrations tended to

decrease after CRRT. HGF concentrations on D7 were significantly reduced compared to those on D0 (Fig. 1A). Concentrations of HGF and soluble cMet on D0 were positively correlated with each other ( $R^2 = 0.126$ ;  $p = 0.03$ ) (Fig. 1B). HGF and soluble cMet concentrations were not significantly correlated on D2 ( $R^2 = 0.107$  and  $p = 0.09$ ) or D7 ( $R^2 = 0.06$  and  $p = 0.42$ ) (Fig. 1C, D).

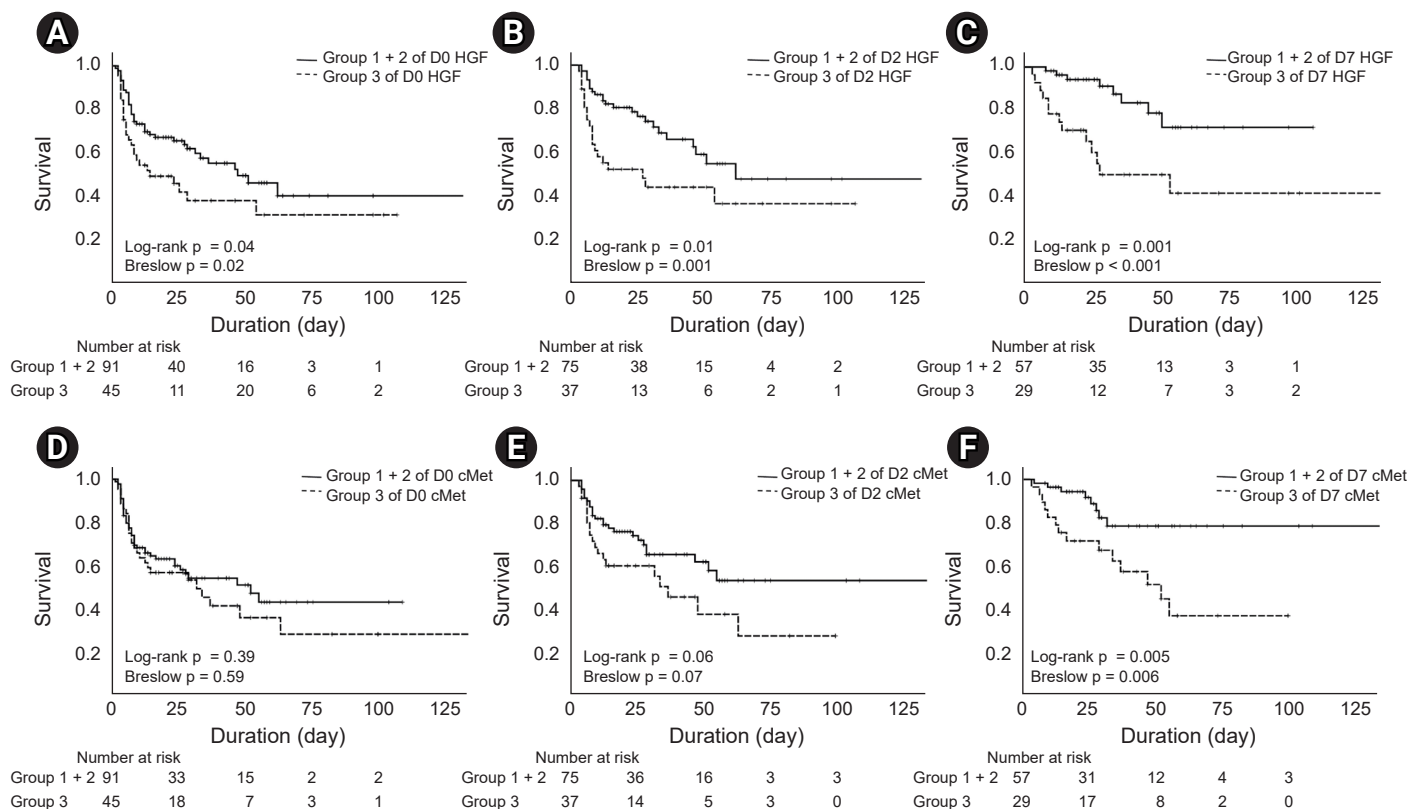
### Effects of plasma hepatocyte growth factor levels on clinical outcomes

Kaplan-Meier survival curves and log-rank tests were used to investigate the associations between plasma HGF concentrations on D0, D2, and D7 and mortality. We divided patients into three groups and analyzed survival rates. Survival rate distribution of the first and second groups based on HGF and soluble cMet levels were similar, but different from those of the third group that had the highest concentration

of both HGF and soluble cMet. Therefore, we combined the first group and the second group for analysis (Supplementary Fig. 1, available online). The risk of all-cause mortality was increased in patients with the highest D0 HGF levels ( $p = 0.04$ ) (Fig. 2A). D2 and D7 HGF levels displayed similar associations with in-hospital mortality, especially D7 HGF level ( $p = 0.01$  and  $p = 0.001$ , respectively) (Fig. 2B, C). In the 2-year follow-up data, plasma D0, D2, and D7 HGF levels were still closely related to all-cause mortality (Supplementary Fig. 2A-C, available online). Next, we performed multivariate Cox proportional analysis to investigate the independent effects of D0, D2, and D7 HGF levels on patient outcomes (Table 2). Elevated HGF values remained an independent variable associated with clinical outcomes after adjusting for confounding variables, including sex, BMI, APACHE II score, and Age-CCI (D0 HGF: hazard ratio [HR] 1.71, 95% confidence intervals [CI] 1.02–2.86,  $p = 0.04$ ; D2 HGF: HR 2.57, 95% CI 1.37–4.83,  $p = 0.003$ ; D7 HGF: HR 4.26,



**Figure 1. Variations and correlations between HGF and soluble cMet levels.** (A) Trends of HGF and soluble cMet concentrations in plasma after CRRT. (B) Plasma HGF level converted to a natural logarithm was positively correlated with plasma soluble cMet level on D0 after CRRT was initiated. Pearson correlation coefficient was  $R^2 = 0.126$  with a  $p$ -value of 0.03. No correlation was observed between plasma HGF levels and cMet levels on D2 (C) or D7 (D) after initiating CRRT. CRRT, continuous renal replacement therapy; D, day; HGF, hepatocyte growth factor.



**Figure 2. Survival rates of AKI patients who underwent CRRT according to plasma HGF and soluble cMet concentrations.** (A) Patients in group 3 had a significantly lower survival rate than patients in group 1 and group 2 according to plasma HGF concentration on D0 (log-rank  $p = 0.04$ ). (B and C) Patients in HGF group 3 had a significantly lower survival rate than those in groups 1 and 2 on D2 (log-rank  $p = 0.01$ ) and D7 (log-rank  $p = 0.001$ ). Patients in group 3 were divided according to plasma soluble cMet concentrations on D0, D2, and D7. (D and E) No difference in mortality rates was found between cMet groups 1 and 2 and group 3 on D0 and D2. (F) The risk of all-cause mortality was significantly increased in the patient group with the highest D7 cMet levels ( $p = 0.005$ ). AKI, acute kidney injury; CRRT, continuous renal replacement therapy; D, day; HGF, hepatocyte growth factor.

95% CI 1.71-10.62,  $p = 0.002$ ).

Additionally, we evaluated the effects of changes in HGF concentrations between different time points during the week after CRRT initiation on clinical outcomes. Changes in HGF concentrations were calculated as differences between D2 and D0, between D7 and D0, and between D7 and D2. Changes in HGF concentrations were not related to mortality (Supplementary Fig. 3A-C, available online).

### Association between plasma soluble cMet level and clinical outcomes

We used the same method to investigate the associations between plasma soluble cMet concentrations on D0, D2, and D7 and in-hospital mortality. D0 and D2 soluble cMet

levels did not predict all-cause mortality (Fig. 2D-E). On D7, the risk of all-cause mortality (Fig. 2F) was significantly increased in the patient group with the highest soluble cMet level ( $p = 0.005$ ). In the 2-year follow-up data, plasma D0 soluble cMet level was still closely related to all-cause mortality (Supplementary Fig. 2D-F). Multivariate Cox regression analysis showed that the group with the highest soluble cMet level had a significantly higher mortality rate than the other groups after adjusting for other risk factors such as sex, BMI, APACHE II score, and Age-CCI (HR, 4.18; 95% CI, 1.69-10.32;  $p = 0.002$ ) (Table 3). Additionally, changes in soluble cMet concentrations between D2 and D0, between D7 and D0, and between D7 and D2 were not associated with all-cause mortality (Supplementary Fig. 3D-F).

**Table 2. Multivariate Cox proportional analysis model for HGF (D0, D2, D7)**

Variable	Unadjusted		Model 1		Model 2		Model 3	
	HR (95% CI)	p-value	HR (95% CI)	p-value	HR (95% CI)	p-value	HR (95% CI)	p-value
D0 Group 1 + 2 (n = 91)	Reference		Reference		Reference		Reference	
Group 3 (n = 45)	1.65 (1.00–2.71)	0.05	1.74 (1.05–2.88)	0.03	1.68 (1.01–2.82)	0.05	1.71 (1.02–2.86)	0.04
Continuous variable (n = 136)	1.30 (1.05–1.61)	0.02	1.41 (1.11–1.79)	0.004	1.39 (1.09–1.77)	0.007	1.38 (1.08–1.77)	0.01
D2 Group 1 + 2 (n = 75)	Reference		Reference		Reference		Reference	
Group 3 (n = 37)	2.10 (1.16–3.81)	0.01	2.46 (1.33–4.54)	0.004	2.45 (1.32–4.55)	0.005	2.57 (1.37–4.83)	0.003
Continuous variable (n = 112)	1.56 (1.14–2.12)	0.006	1.60 (1.17–2.18)	0.003	1.58 (1.16–2.16)	0.004	1.68 (1.21–2.33)	0.002
D7 Group 1 + 2 (n = 57)	Reference		Reference		Reference		Reference	
Group 3 (n = 28)	3.87 (1.60–9.35)	0.003	4.35 (1.76–10.75)	0.001	4.39 (1.77–10.86)	0.001	4.26 (1.71–10.62)	0.002
Continuous variable (n = 85)	1.42 (0.98–2.05)	0.06	1.54 (1.03–2.31)	0.04	1.54 (1.02–2.31)	0.04	1.56 (1.03–2.35)	0.04

CI, confidence interval; D, day; HGF, hepatocyte growth factor; HR, hazard ratio.

Model 1: adjusted for sex and body mass index; model 2: adjusted for Acute Physiology and Chronic Health Evaluation II score in addition to model 1 adjusted variables; model 3: adjusted for age-adjusted Charlson comorbidity index score in addition to model 1 adjusted variables.

**Table 3. Multivariate Cox proportional model for cMet (D0, D2, D7)**

Variable	Unadjusted		Model 1		Model 2		Model 3	
	HR (95% CI)	p-value	HR (95% CI)	p-value	HR (95% CI)	p-value	HR (95% CI)	p-value
D0 Group 1 + 2 (n = 91)	Reference		Reference		Reference		Reference	
Group 3 (n = 45)	1.24 (0.75–2.04)	0.40	1.22 (0.74–2.03)	0.45	1.20 (0.72–1.99)	0.49	1.10 (0.66–1.84)	0.71
Continuous variable (n = 136)	1.16 (0.92–1.46)	0.21	1.18 (0.92–1.51)	0.20	1.15 (0.88–1.77)	0.30	1.08 (0.84–1.41)	0.53
D2 Group 1 + 2 (n = 75)	Reference		Reference		Reference		Reference	
Group 3 (n = 37)	1.77 (0.97–3.21)	0.06	1.61 (0.88–2.96)	0.12	1.62 (0.89–2.97)	0.12	1.52 (0.82–2.81)	0.18
Continuous variable (n = 112)	1.30 (0.98–1.73)	0.07	1.32 (0.97–1.79)	0.08	1.34 (0.97–1.86)	0.08	1.305 (0.94–1.82)	0.11
D7 Group 1 + 2 (n = 57)	Reference		Reference		Reference		Reference	
Group 3 (n = 28)	3.25 (1.36–7.76)	0.008	3.84 (1.59–9.24)	0.003	3.82 (1.58–9.24)	0.003	4.18 (1.69–10.32)	0.002
Continuous variable (n = 85)	2.07 (1.18–3.64)	0.01	2.54 (1.37–4.72)	0.003	2.56 (1.37–4.79)	0.003	2.52 (1.38–4.60)	0.003

CI, confidence interval; D, day; HR, hazard ratio.

Model 1: adjusted for sex and body mass index; model 2: adjusted for Acute Physiology and Chronic Health Evaluation II score in addition to model 1 adjusted variables; model 3: adjusted for age-adjusted Charlson comorbidity index score in addition to model 1 adjusted variables.



### Ability of plasma hepatocyte growth factor and soluble cMet to predict outcomes at different time points

Levels of HGF and soluble cMet that best predicted mortality were investigated using ROC curves. The AUCs (95% CI) of D0, D2, and D7 HGF were 0.576 (0.440–0.711), 0.540 (0.392–0.689), and 0.716 (0.598–0.838), respectively (Fig. 3A). The AUCs (95% CI) of D0, D2, and D7 soluble cMet were 0.579 (0.440–0.719), 0.570 (0.429–0.711), and 0.678 (0.553–0.825), respectively (Fig. 3B).

### Plasma hepatocyte growth factor and soluble cMet as biomarkers for predicting mortality

Next, we evaluated whether the addition of plasma HGF and soluble cMet to conventional prognostic markers such as APACHE II and Age-CCI improved prediction of mortality (Table 4). In ROC analysis, the addition of log(D7 HGF) and log(D7 cMet) increased the AUC. The AUCs (95% CIs) of the APACHE II (model 1), APACHE II + log(D7 HGF) (model 2), and APACHE II + Age-CCI + log(D7 HGF) + log(D7 cMet) (model 3) were 0.523 (0.393–0.653), 0.710 (0.588–0.832), and 0.749 (0.635–0.863), respectively. Additionally, the

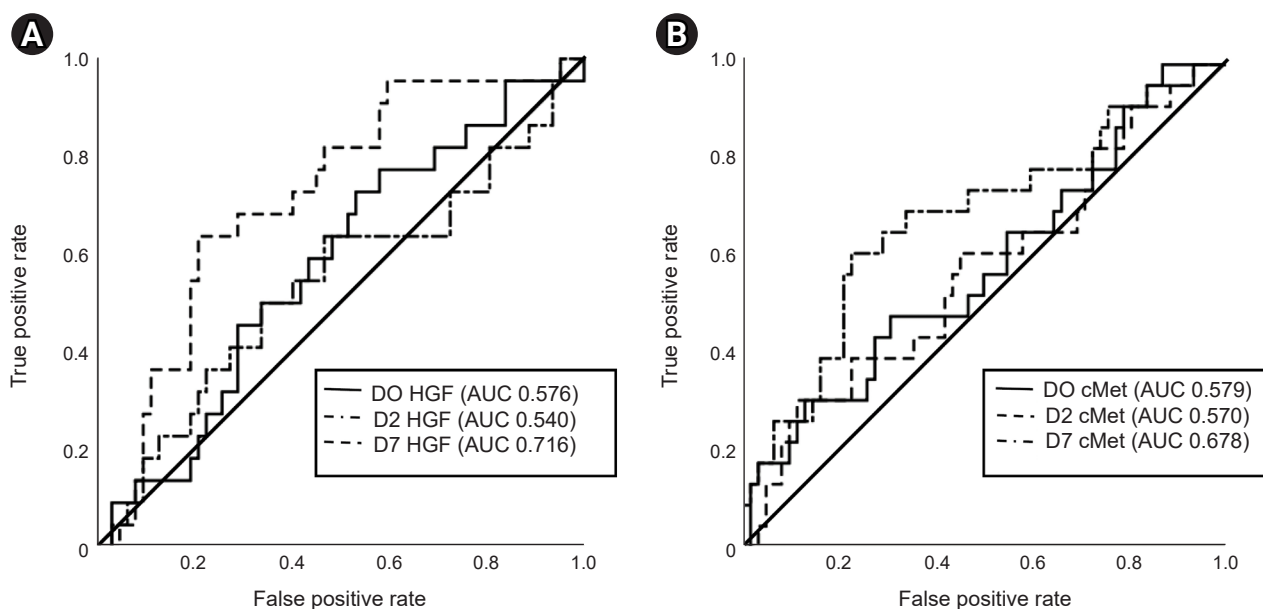
IDI and cNRI for predicting mortality (model 1 vs. model 3) were 0.166 (0.070–0.262;  $p < 0.001$ ) and 75.1% (29.9%–120.3%;  $p < 0.001$ ), suggesting that addition of plasma log(D7 HGF) and log(D7 cMet) to conventional predictors of mortality significantly increased predictive value.

### Prediction of clinical outcomes according to combined hepatocyte growth factor and soluble cMet concentrations

The ability to predict patient outcomes using the combination of HGF and soluble cMet concentrations was evaluated. Group 1 had a significantly higher mortality rate than the other groups ( $p = 0.04$ ). Furthermore, the difference in mortality between these groups was statistically significant when using D7 HGF and cMet concentrations ( $p = 0.02$ ) (Supplementary Fig. 4A–C, available online).

### Fluid status according to hepatocyte growth factor and soluble cMet concentrations

We investigated fluid status according to HGF and soluble cMet concentrations using parameters of fluid status measured using InBody S10. There was no significant difference



**Figure 3. Comparison of the ROC curves for plasma HGF and soluble cMet at various time points.** (A) ROC curve and AUC for plasma HGF concentrations at various time points. (B) ROC curve and AUC for plasma soluble cMet concentrations at various time points.

AUC, area under the curve; D, day; HGF, hepatocyte growth factor; ROC, receiver operating characteristic.

**Table 4. Incremental values of HGF and soluble cMet compared to traditional risk factors for predicting mortality due to severe acute kidney injury (n = 84)**

Day	Model	ROC analysis (DeLong test)		IDI analysis		Category-free NRI analysis	
		AUC (95% CI)	p-value	IDI (95% CI)	p-value	NRI (95% CI)	p-value
0	APACHE II	0.523 (0.393-0.653)	Reference		Reference		Reference
	APACHE II + Age-CCI	0.581 (0.426-0.736)	0.55	0.034 (-0.012 to 0.080)	0.15	0.123 (-0.362 to 0.608)	0.62
	APACHE II + log(HGF)	0.584 (0.448-0.720)	0.40	0.006 (-0.010 to 0.022)	0.49	0.285 (-0.198 to 0.767)	0.25
	APACHE II + Age-CCI + log(HGF)	0.595 (0.442-0.749)	0.45	0.043 (-0.009 to 0.095)	0.11	0.155 (-0.329 to 0.640)	0.53
	APACHE II + Age-CCI + log(HGF) + log(cMet)	0.631 (0.482-0.781)	0.29	0.063 (0.002 to 0.124)	0.04	0.311 (-0.169 to 0.790)	0.20
2	APACHE II + log(HGF)	0.547 (0.401-0.692)	0.66	0.001 (-0.006 to 0.009)	0.75	0.246 (-0.234 to 0.727)	0.32
	APACHE II + Age-CCI + log(HGF)	0.585 (0.433-0.737)	0.51	0.035 (-0.012 to 0.082)	0.14	0.188 (-0.297 to 0.672)	0.45
	APACHE II + Age-CCI + log(HGF) + log(cMet)	0.593 (0.447-0.739)	0.46	0.047 (-0.006 to 0.100)	0.08	0.167 (-0.314 to 0.658)	0.50
7	APACHE II + log(HGF)	0.710 (0.588-0.832)	0.02	0.061 (0.012 to 0.109)	0.01	0.719 (0.265 to 1.3172)	0.002
	APACHE II + Age-CCI + log(HGF)	0.681 (0.546-0.816)	0.08	0.095 (0.023 to 0.168)	0.01	0.414 (-0.065 to 0.892)	0.09
	APACHE II + Age-CCI + log(HGF) + log(cMet)	0.749 (0.635-0.863)	0.02	0.166 (0.070 to 0.262)	0.001	0.751 (0.299 to 1.203)	0.001

Age-CCI, age-adjusted Charlson comorbidity index; APACHE II, Acute Physiology and Chronic Health Evaluation II; AUC, area under the curve; HGF, hepatocyte growth factor; IDI, integrated discrimination improvement; NRI, net reclassification improvement; ROC, receiver operating curve.

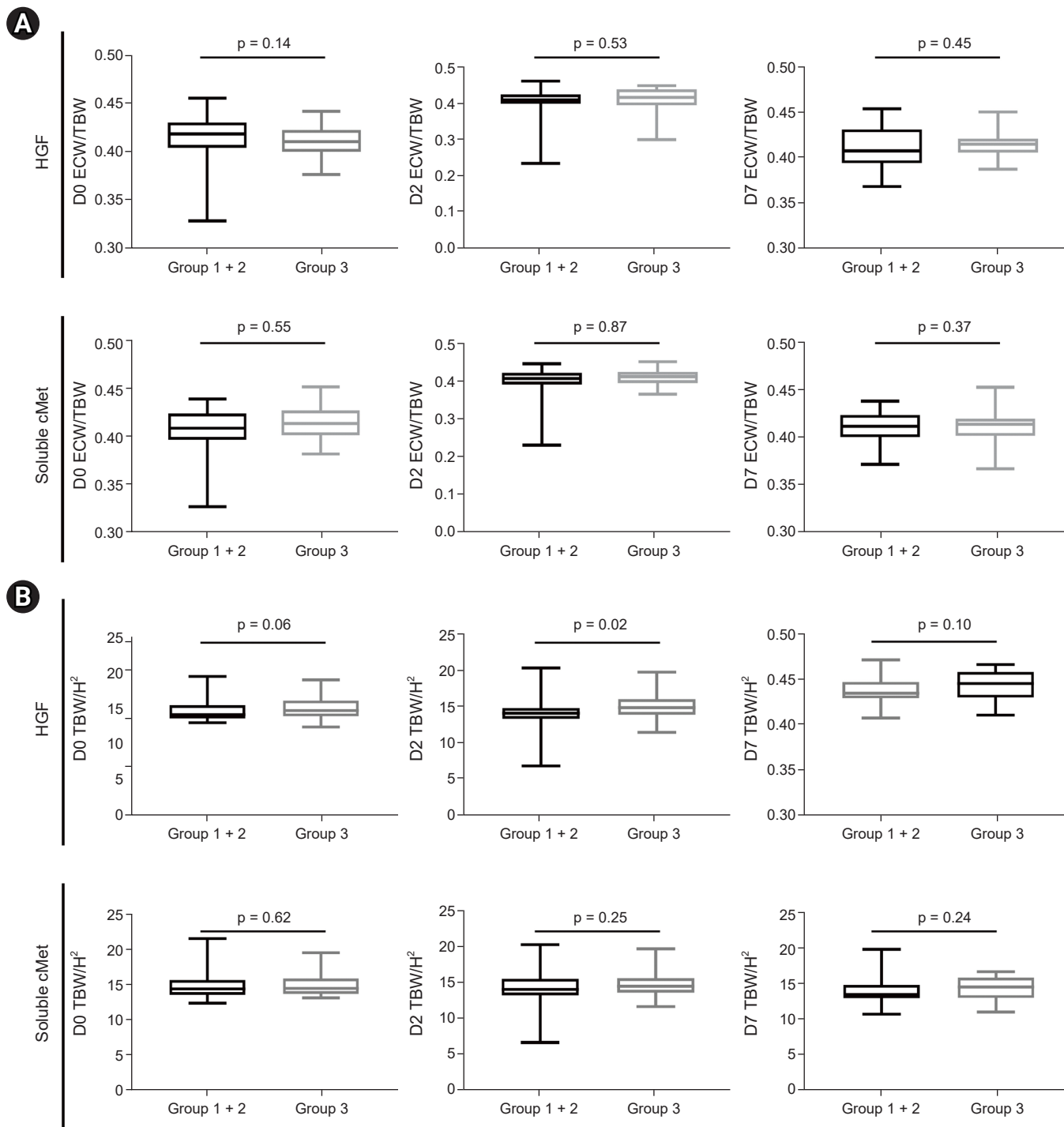
between ECW/TBW and plasma HGF and soluble cMet levels on D0, D2, or D7 (Fig. 4A). There was no significant difference between plasma HGF levels on D0 and D7 and soluble cMet levels on D0, D2, and D7. TBW/H<sup>2</sup> increased significantly (p = 0.02) only in the group of patients with the highest D2 HGF level (Fig. 4B).

## Discussion

In this study, we found that increased plasma HGF and soluble cMet levels could predict the outcomes of patients with severe AKI undergoing CRRT. Plasma HGF and soluble cMet levels on D7 were the most valuable predictors of patient outcomes. The inclusion of plasma HGF and soluble cMet levels with conventional prognostic index markers improved the predictive power of the composite indices. Finally, there was no correlation between plasma HGF and soluble cMet levels and fluid balance. To our knowledge, this is the first study to show that increased plasma HGF and soluble cMet levels of patients with severe AKI can predict clinical outcomes.

HGF and its receptor cMet are key components of a signaling pathway with critical roles in cellular regeneration, proliferation, differentiation, invasion, angiogenesis, anti-apoptosis, tissue fibrosis, and wound healing [19,20]. The HGF/ cMet signaling pathway has therefore been identified as a target to alleviate the various types of AKI and prevent progression to CKD. The expression of HGF has been shown to be increased in various acute and CKDs including experimental acute ischemic injury, toxic elements, and unilateral nephrectomy [21,22].

HGF may act as an endocrine and paracrine effector for kidney repair during kidney injury. Administration of exogenous HGF promotes tubular repair and recovery [22]. Previously, we demonstrated that administration of cMet agonistic antibodies halted the progression of CKD [11]. Our group also confirmed that cMet agonistic antibodies attenuate apoptosis in AKI [23]. Conversely, disruption of HGF signaling aggravates renal interstitial fibrosis after obstructive injury [24]. However, many studies have reported that elevated plasma HGF and soluble cMet concentrations are associated with poorer clinical prognosis. Santalahti et al. [12] reported that levels of HGF and placental growth factor in plasma could predict mortality in a general population. Elevated circulating HGF levels have been observed in many patho-



**Figure 4. Euvolemic status on D7 after CRRT according to HGF and soluble cMet concentrations.** (A) Comparison of ECW/TBW and plasma HGF or soluble cMet on D0, D2, and D7. (B) Comparison of TBW/H<sup>2</sup> and plasma HGF or soluble cMet levels. CRRT, continuous renal replacement therapy; D, day; ECW, extracellular water; HGF, hepatocyte growth factor; H<sup>2</sup>, height squared; TBW, total body water.

logic liver diseases, including hepatitis and hepatocellular carcinoma, and have been shown to be correlated with more severe liver cirrhosis [25]. Likewise, HGF levels are associated with sepsis and correlated with established markers of endothelial cell injury. Elevated HGF level in sepsis patients is a significant indicator of a poor prognosis [26]. Moreover, an increase in soluble cMet concentration has been shown to be associated with a poor clinical prognosis. As reported in various malignant tumors, soluble cMet concentration is associated with progression, metastasis, and a poor prognosis [27-29]. Our group previously demonstrated increased urinary cMet in diabetic nephropathy patients was strongly associated with progression to end-stage renal disease [15].

There are several explanations for why elevated HGF and soluble cMet concentrations are highly related to a poorer prognosis. First, the greater the extent of the disease, the greater the activation of the HGF/cMet signaling pathway for cell regeneration. Overexpression of the HGF/cMet pathway and increased plasma HGF and soluble cMet concentrations may be closely related to disease progression. In addition, an increase in soluble cMet inhibits the phosphorylation of cMet, thereby inhibiting the HGF/cMet signaling pathway [30].

One study reported a marked increase in urinary HGF levels in patients with AKI [23]. Another study showed that HGF concentration in serum was significantly increased in AKI patients relative to participants receiving hemodialysis (approximately 20-fold) [31]. Clinical trials have indicated that HGF and other biomarkers in urine (such as neutrophil gelatinase-associated lipocalin [NGAL]) have the potential to predict AKI [32]. These studies suggested that HGF was correlated with AKI severity. Despite these data, these studies only examined AKI patients with urine volumes that were maintained and measured at one-time point; furthermore, cMet concentration was not considered. Additionally, the mortality rate of patients was not assessed.

We previously reported that the expression of HGF and soluble cMet in plasma was significantly increased in AKI patients [23]. In this study, we evaluated if plasma HGF and soluble cMet concentrations could predict clinical longitudinal outcomes of severe AKI patients undergoing CRRT. We confirmed that mortality rates during hospitalization were significantly higher in patients with severe AKI and patients with the highest HGF levels on D0, D2, and D7. Soluble cMet level on D7 was a significant predictor of patient mortality.

After adjusting for sex, BMI, APACHE II score, Age-CCI, HGF and soluble cMet levels were still significant predictors of patient mortality. A large amount of HGF may be released as an injury protection mechanism to activate downstream signaling pathways. HGF is released not only by damaged kidneys but also distant organs (lung, liver, spleen) and can participate in tubular repair both as an endocrine factor and paracrine substance [33]. This suggests that plasma HGF and soluble cMet levels can be used as predictors of the prognosis of patients with severe AKI and of clinical risk and recovery after AKI.

A previous study showed that a panel of urine biomarkers measured on D1, D7, and D14 yielded significantly different results for those recovering from AKI compared with those patients who did not recover [34]. We analyzed the correlation between plasma HGF and soluble cMet concentrations and mortality at multiple time points after the initiation of CRRT (D0, D2, and D7). We found that plasma HGF as a clinical predictor was not affected by CRRT and that plasma HGF was a better predictor of mortality than soluble cMet. Furthermore, we examined whether HGF and soluble cMet levels on D0, D2, and D7 were reliable biomarkers of mortality using ROC curves and AUCs. D7 HGF and soluble cMet concentrations were prognostic biomarkers with high sensitivity and specificity. The best time to measure plasma HGF and soluble cMet levels may be on D7 after the start of CRRT for severe AKI; repeated measurements can improve prediction accuracy.

An individual biomarker is rarely sufficient for clearly defining a particular pathologic state [35,36]. Vaidya et al. [32] measured kidney injury molecule-1, HGF, NGAL, and interleukin-18 levels simultaneously in the same aliquot of urine. The specificity and sensitivity of the combination of these urinary biomarkers for the diagnosis of AKI were significantly higher than those of single biomarkers. The urinary [TIMP-2] · [IGFBP7] test can be used to identify critically ill patients at high risk for imminent AKI [5]. Therefore, multiple biomarkers measured in the same biological sample at the same time are extremely useful for predicting outcomes. Circulating HGF and cMet are attractive potential alternative biomarkers for ligand overexpression and receptor overexpression, respectively [37]. In this study, we analyzed the ability of combined plasma HGF and soluble cMet levels to predict mortality in patients with severe AKI. We found that the mortality rates of groups with high HGF and soluble

cMet concentrations on D2 and D7 were significantly higher than those of other groups; furthermore, the inclusion of soluble cMet and HGF with conventional prognostic indices such as APACHE II and Age-CCI significantly improved predictive power.

Rhee et al. [38] demonstrated that TBW/H<sup>2</sup> ( $\geq 13$  L/m<sup>2</sup>) and ICW/H<sup>2</sup> were independently associated with higher in-hospital mortality for patients with AKI undergoing CRRT. That study showed that the fluid status could be assessed using ECW/TBW in critically ill patients requiring CRRT and that ECW/TBW could predict mortality [39]. All patients who participated in this study were admitted to the ICU, and their fluid balance was maintained during CRRT. We used ECW/TBW to evaluate the correlation between fluid status and plasma HGF or soluble cMet levels; however, no significant correlation was found.

The present study had some limitations. First, although ethnicity can influence several clinical outcomes, data were acquired only from a Korean population, and this study was limited to Korean tertiary hospitals. Second, only mortality during hospitalization was assessed and long-term outcomes were not considered. Finally, plasma HGF and soluble cMet concentrations could be affected by CRRT. It is necessary to identify HGF and soluble cMet levels in CRRT waste liquid or blood before and after CRRT to confirm that the CRRT waste liquid does not contain HGF or soluble cMet in future studies.

In summary, plasma HGF measurements at multiple time points predicted the clinical outcomes of patients with severe AKI undergoing CRRT. Furthermore, plasma soluble cMet measurements at a specific time point were able to predict clinical outcomes. For accurate mortality predictions, the most valuable time to perform plasma HGF and soluble cMet measurements is D7 after CRRT, and repeated measurements can improve accuracy. We found no correlation between plasma HGF and soluble cMet levels and fluid balance in this study.

### Conflicts of interest

All authors have no conflicts of interest to declare.

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