

LETTER

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Authors' response to letter "Prediction of acute kidney injury in intensive care unit patients"

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See related letter by Guo et al., <https://ccforum.biomedcentral.com/articles/10.1186/s13054-018-2248-x>

We thank Dr. Guo and coworkers for their interest and comments [1] on our article [2]. We have provided responses to their comments. First, we agree that the patient severity of illness and level of organ failure upon admission to medical cardiac intensive care units (MCICUs) may be important predictors for the development of acute kidney injury (AKI). Hence, we evaluated the predictive ability of urinary liver-type fatty acid-binding protein (L-FABP) and serum N-terminal pro-B-type natriuretic peptide (NT-proBNP) for AKI in the analytical model that included the Sequential Organ Failure Assessment (SOFA) score. In the multivariate logistic regression analysis, L-FABP, NT-proBNP, and the SOFA score were all independent predictors of AKI (Table 1). According to these findings, we speculate that a novel panel consisting of L-FABP, NT-proBNP, and the SOFA score may improve the accuracy for predicting AKI in patients treated in MCICUs. Furthermore, the addition of both L-FABP and NT-proBNP to a baseline model that included established risk factors and the SOFA score further enhanced the net reclassification and integrated discrimination im-

provement; this difference was greater than that obtained for either of the biomarkers and the baseline model alone (Table 2). Therefore, upon admission of patients to MCICUs, combining the measurements of the two independent predictors of AKI—L-FABP and NT-proBNP—may improve the accuracy for the early prediction of AKI beyond that achieved with either predictor alone.

Second, unfortunately, the serum creatinine (SCr) concentration used for the diagnosis of AKI in our study had not been corrected according to fluid balance because of the inconsistent data recorded. Adjustment of the SCr concentration was proposed according to the assumption that the SCr concentration may be diluted by positive fluid balance [3]. However, Shen et al. have suggested that a large proportion of the infused fluid eventually leaks into the third space instead of contributing to blood volume [4]. Therefore, the use of adjusted SCr might have overestimated the AKI incidence [4]. Further studies are needed to clarify this issue.

Finally, the C-index observed following the addition of both L-FABP and NT-proBNP showed the improvement beyond that of the baseline

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Table 1 Multivariate logistic regression analyses for predictors of acute kidney injury

Variables	Multivariate model 1		Multivariate model 2	
	OR (95% CI)	P Value	OR (95% CI)	P value
Age (per 10 years increment)	1.18 (1.00–1.39)	0.05	1.21 (1.03–1.42)	0.02
IABP before admission	2.33 (1.32–4.10)	0.003	2.46 (1.42–4.27)	0.001
NT-proBNP (per 10-fold increment)	1.67 (1.22–2.29)	0.001		
Tertile of NT-proBNP (pg/mL)				
First (< 425)			1.0	
Second (425–2730)			2.10 (1.27–3.47)	0.004
Third (> 2730)			2.16 (1.20–3.88)	0.01
Urinary L-FABP (per 10-fold increment)	2.69 (2.06–3.50)	< 0.001		
Tertile of Urinary L-FABP (ng/mL)				
First (< 3.3)			1.0	
Second (3.3–11.5)			1.50 (0.92–2.44)	0.10
Third (> 11.5)			3.72 (2.34–5.93)	< 0.001
SOFA score (per 1 point increment)	1.12 (1.04–1.21)	0.004		
Tertile of SOFA score (point)				
First (< 2)			1.0	
Second (2–3)			1.17 (0.73–1.87)	0.51
Third (> 3)			2.04 (1.28–3.24)	0.003

Multivariate model adjusted for all baseline variables with $P < 0.05$ by univariate analysis. NT-proBNP, L-FABP, and the SOFA score were assessed as either continuous variables (model 1) or variables categorized into tertiles (model 2)

CI confidence interval, IABP intraaortic balloon pump, L-FABP liver-type fatty acid-binding protein, NT-proBNP N-terminal pro-B-type natriuretic peptide, OR odds ratio, SOFA Sequential Organ Failure Assessment

Table 2 Discrimination and reclassification of combination of L-FABP and NT-proBNP for acute kidney injury

	C-index	P value	NRI	P value	IDI	P value
Baseline model	0.752	Ref.		Ref.		Ref.
Baseline model + NT-proBNP	0.772	0.40	0.350	< 0.001	0.016	0.002
Baseline model + L-FABP	0.797	0.06	0.615	< 0.001	0.085	< 0.001
Baseline model + NT-proBNP + L-FABP	0.806	0.02	0.630	< 0.001	0.093	< 0.001
Baseline model + NT-proBNP + L-FABP vs. Baseline model + NT-proBNP	0.034*	0.14	0.571	< 0.001	0.077	< 0.001
Baseline model + NT-proBNP + L-FABP vs. Baseline model + L-FABP	0.008*	0.73	0.230	< 0.001	0.008	0.007

Baseline model included age, sex, hypertension, dyslipidemia, diabetes, smoking status, atrial fibrillation, acute decompensated heart failure, previous myocardial infarction, previous coronary revascularization, heart rate, emergent coronary angiography or percutaneous coronary intervention before admission, intraaortic balloon pump before admission, and the SOFA score
IDI integrated discrimination improvement, L-FABP liver-type fatty acid-binding protein, NT-proBNP N-terminal pro-B-type natriuretic peptide, NRI net reclassification improvement, Ref. reference
*Estimated differences between two groups

model alone (Table 2). On performing the calibration using the Hosmer–Lemeshow test, the model involving the addition of both L-FABP and NT-proBNP to the baseline model showed a good fit, whereas the model involving the addition of a single biomarker or the baseline model alone showed a poor fit.

Abbreviations

AKI: Acute kidney injury; L-FABP: Liver-type fatty acid-binding protein; MCICUs: Medical cardiac intensive care units; NT-proBNP: N-terminal pro-B-type natriuretic peptide; SCr: Serum creatinine; SOFA: Sequential Organ Failure Assessment

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