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Can copeptin predict the severity of coronavirus disease 2019 infection?

Erdal İn¹* , Mutlu Kuluöztürk² , Selda Telo³ , Zülal Aşçı Toraman⁴ , Ercan Karabulut⁵

SUMMARY

OBJETIVE: Coronavirus disease 2019 (COVID-19) has quickly turned into a health problem globally. Early and effective predictors of disease severity are needed to improve the management of the patients affected with COVID-19. Copeptin, a 39-amino acid glycopeptide, is known as a C-terminal unit of the precursor pre-provasopressin (pre-proAVP). Activation of AVP system stimulates copeptin secretion in equimolar amounts with AVP. This study aimed to determine serum copeptin levels in the patients with COVID-19 and to examine the relationship between serum copeptin levels and the severity of the disease.

METHODS: The study included 90 patients with COVID-19. The patients with COVID-19 were divided into two groups according to disease severity as mild/moderate disease (n=35) and severe disease (n=55). All basic demographic and clinical data of the patients were recorded and blood samples were collected.

RESULTS: Copeptin levels were significantly higher in the patients with severe COVID-19 compared with the patients with mild/moderate COVID-19 (p<0.001). Copeptin levels were correlated with ferritin and fibrinogen levels positively (r=0.32, p=0.002 and r=0.25, p=0.019, respectively), and correlated with oxygen saturation negatively (r=-0.37, p<0.001). In the multivariate logistic regression analysis, it was revealed that copeptin (OR: 2.647, 95%CI 1.272–5.510; p=0.009) was an independent predictor of severe COVID-19 disease. A cutoff value of 7.84 ng/mL for copeptin predicted severe COVID-19 with a sensitivity of 78% and a specificity of 80% (AUC: 0.869, 95%CI 0.797–0.940; p<0.001).

CONCLUSION: Copeptin could be used as a favorable prognostic biomarker while determining the disease severity in COVID-19.

KEYWORDS: COVID-19. Mild/Moderate COVID-19. Severe COVID-19. Copeptin. Biomarkers.

INTRODUCTION

In the city of Wuhan in China, a pneumonia outbreak that developed due to a novel coronavirus was detected in December 2019. The outbreak could not be taken under control and it spread all around the world, resulting in a pandemic. The novel coronavirus was defined as severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2). Accordingly, the disease that resulted from the virus was defined as coronavirus disease 2019 (COVID-19) by the World Health Organization^{1,2}. As of May

24, 2021, approximately 167 million COVID-19 cases and more than 3.4 million deaths were reported all around the world³.

Copeptin is a glycopeptide that consists of 39 amino acids. It is also related to arginine vasopressin (AVP). Copeptin is derived from the C-terminal part of pre-pro-AVP, which is an AVP precursor molecule. Together with AVP, copeptin is expressed from neurohypophysis simultaneously either with osmotic or hemodynamic signals. Therefore, it strongly correlates with plasma levels. AVP is

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¹Malatya Turgut Ozal University, Faculty of Medicine, Department of Chest Diseases – Malatya, Turkey.

²Fırat University, Faculty of Medicine, Department of Chest Diseases – Elazig, Turkey.

³Firat University, Faculty of Dentistry, Department of Biochemistry – Elazig, Turkey.

⁴Fırat University, Faculty of Medicine, Department of Medical Microbiology – Elazig, Turkey.

⁵Ankara Yildirim Beyazit Üniversity, Faculty of Medicine, Department of Medical Pharmacology – Ankara, Turkey.

^{*}Corresponding author: inerda@gmail.com

a hormone with antidiuretic and vasoconstrictive qualities. Accordingly, it exhibits endogenous stress responses and stimuli, such as hypotension, hypoxia, hyperosmolarity, acidosis, and infections increase its expression. Nevertheless, it cannot be used as a biomarker due to circadian rhythm, short half-life, and unstable molecule status of AVP. On the other hand, copeptin is a peptin that is more stable and its levels can be detected in blood easily⁴⁻⁶.

In previous studies, it was demonstrated that levels of serum copeptin were increased in diseases, such as community-acquired pneumonia (CAP), ventilator-associated pneumonia (VAP), lower respiratory tract infections, acute exacerbation of chronic obstructive pulmonary disease (COPD), and sepsis, and these increases were related to poor prognosis⁷⁻¹².

It is suggested that the assessment and management of COVID-19 should be conducted according to the disease severity. In terms of the initial data that were obtained from China, it was observed that 81% of the patients with COVID-19 experienced mild or moderate diseases, which were similar to cold and mild pneumonia, whereas 14% of the patients had severe diseases in addition to 5% of the patients, who had critical and fatal diseases¹³. At this point, early recognition of severe forms of the disease is vital for early hospitalization and appropriate treatment of the patients. The clinical status, oxygen saturation, and comorbidity mainly determine the need for hospitalization, while certain laboratory parameters can also facilitate the assessment of the severity of the disease¹⁴. In various studies, it was demonstrated that biomarkers, such as CRP (C-reactive protein), D-dimer, ferritin, cardiac troponin, interleukin-6 (IL-6), and lymphocyte counts, could be used in estimating the severity of the disease in risk classification¹⁵. In a very recent study, copeptin was found to be a useful biomarker in distinguishing COVID-19-associated pneumonia from CAP16. However, in the literature, it was observed that no study evaluated the relationship between COVID-19 severity and copeptin.

In this study, it was aimed to measure the serum copeptin levels in the patients with COVID-19, to evaluate the relationship of serum copeptin levels with disease severity, and to evaluate the relationships between copeptin and certain inflammatory parameters.

METHODS

Study subjects

In this study, 90 consecutive patients with COVID-19 diagnosis, who were admitted to the Pandemic Clinic in the Faculty of Medicine at Firat University between August and October

2020, were included in the sample according to the inclusion criteria. Patients older than 18 years were included in the study.

COVID-19 diagnosis was defined as a SARS-CoV-2 positive real-time reverse-transcriptase polymerase chain reaction (RT-PCR) from a nasal and/or throat swab together with signs, symptoms, or radiological that suggest COVID-19 infection. In the study, the clinical parameters were recorded as well as the demographic data. Following the investigation of the histories and physical examinations of the subjects, the blood samples were drawn from the patients before the treatment process.

Upon admission to the hospital, the patients with COVID-19 were categorized into three groups by considering the clinical findings, respiratory rates, oxygen saturation (SpO₂) levels, and low-dose chest CT findings¹⁷. The details of the classification are as follows:

Mild illness: mild clinical symptoms, no sign of pneumonia on low-dose CT.

Moderate illness: mild respiratory symptoms, positive signs of pneumonia on low-dose CT, and SpO₂ ≥94% on room air. Severe illness: Those who meet any of the following criteria:

- 1. Respiratory rate >30 times per min;
- 2. $SpO_3 < 94\%$ at room air;
- 3. Lung infiltrates >50% on low-dose CT.

In the study, the exclusion criteria covered the patients who had heart failure, renal failure, COPD, acute myocardial infarction, acute coronary syndrome, interstitial lung disease, any organ malignancy, or immunosuppression (HIV infection, solid organ or stem cell transplantation, or any immunosuppressive treatment). Additionally, pregnancy was an exclusion criterion in the study.

Ethics approval

The study was conducted within the framework of the Helsinki Declaration. The study was also approved by the Ethical Committee of the Medicine Faculty of Firat University (issue: 402252/20,07,2020). The subjects who participated in the study also provided their written consent to be included in the study.

Measurement of serum copeptin levels

In the measurements, the separation of the serum was conducted by centrifuging the samples at 4,000 g for 10 min. Then the samples were frozen at -80°C for further analysis. A double-antibody sandwich enzyme-linked immunosorbent assay kit (Catalog No. 201-12-5463 Human copeptin Elisa Kit: Sunred Biological Technology Co. Ltd. Shanghai) was used to measure the serum copeptin levels. The assay had a sensitivity of 0.067 ng/mL. The inter-assay and intra-assay calculation values were <12% and <10%, respectively. Furthermore, the detection range of copeptin was 0.07–20 ng/mL.

Statistical analyses

In the statistical analyses, IBM SPSS Statistics 21 (Version 21, authorization code: d91314f638c364094170, Armonk, NY, USA) software was used. The results obtained in the analyses were stated as mean±SD. The level of statistical significance was regarded as p<0.05. To compare two independent samples, the student's t-test was conducted. To compare the distribution of sex between the groups, Chi-square (χ^2) test was conducted. The parametric variables were evaluated by the Pearson correlation analysis. In the study, univariate and multivariate analyses were conducted by binary logistic regression to evaluate the variables that could predict severe COVID-19 disease. In the analysis, a 95% confidence interval (CI) was adopted while calculating the odds ratios (ORs). The receiver operating characteristic (ROC) analysis method was to determine the cutoff value for copeptin. With the ROC curve, the area under curve (AUC) value was determined.

RESULTS

Comparison of groups

In the study population, 35 patients were assigned to a mild/moderate disease and 55 patients were allocated to a severe disease. The mean serum copeptin levels of the patients with mild/moderate and severe COVID-19 were 6.4±2.2 and 11.7±4.6 ng/mL, respectively (p<0.001). Comparison of all laboratory data of the patients with COVID-19 was demonstrated in Table 1.

Correlation analysis

In the analyses, it was observed that serum copeptin levels positively correlated with serum ferritin and fibrinogen levels (r=0.32, p=0.002 and r=0.25, p=0.019, respectively), whereas negatively correlating with SaO, levels (r=-0.37, p<0.001; Figure 1).

Table 1. Comparison of the demographical and laboratory data of two groups of patients with COVID-19.

	Mild/ moderate disease (n=35)	Severe disease (n=55)	р			
Age, years	44.5±14.9	58.8±16.8	<0.001			
Sex, male, n(%)	18 (51.4)	35 (63.6)	0.25			
SaO ₂ , %	94.7±2.5	88.4±4.4	<0.001			
Complete blood count						
Leukocyte, ×10 ⁹ /L	6.4±2.8	6.0±3.2	0.53			
Hemoglobin, g/dL	13.5±1.9	13.6±2.0	0.87			
Platelet, ×10 ⁹ /L	194.5±40.7	179.8±83.1	0.33			
Biochemical markers						
Urea, mg/dL	32.7±12.7	39.8±16.2	0.03			
Creatinine, mg/dL	0.81±0.23	0.89±0.25	0.13			
ALT, U/L	22.1±10.8	32.9±20.1	0.004			
AST, U/L	27.0±15.8	40.7±20.2	0.001			
LDH, U/L	262.9±73.2	344.7±137.8	0.002			
D-dimer, mg/L	0.8±1.1	1.2±1.8	0.24			
Inflammatory markers						
CRP, mg/L	23.8±40.6	84.6±64.2	<0.001			
Procalcitonin, mg/L	0.34±0.5	0.30±0.41	0.61			
Ferritin, ng/mL	185.2±198.0	601.9±364.4	<0.001			
Fibrinogen, mg/dL	402.7±128.7	536.8±159.9	<0.001			
Cardiac markers						
CK, U/L	84.6±121.3	162.5±184.6	0.03			
CK-MB, U/L	20.9±19.5	31.2±32.4	0.09			
Troponin I, μg/L	0.02±0.1	0.13±0.8 0.46				
Copeptin, ng/mL	6.4±2.2	11.7±4.6	<0.001			

SaO₂: oxygen saturation; ALT: alanine transaminase; AST: aspartate transaminase; LDH: lactate dehydrogenase; CRP: C-reactive protein; CK: creatine kinase. Bold values are statistically significant values (p< 0.05).

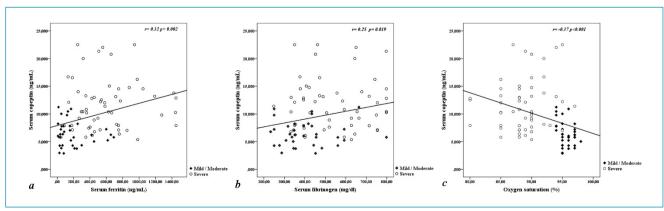


Figure 1. Correlation between serum copeptin levels and (a) ferritin, (b) fibrinogen, and (c) oxygen saturation levels in patients with COVID-19.

Logistic regression analysis

The univariable logistic regression model demonstrated the following parameters that were statistically significant, which included age, copeptin, CRP, fibrinogen, AST, ALT, LDH, and ferritin. In the multivariable logistic regression model, it was observed that that older age (OR: 1.139, 95%CI 1.036–1.251; p=0.007) and increased copeptin (OR: 2.647, 95%CI 1.272–5.510; p=0.009) and ferritin (OR: 1.005, 95%CI 1.000–1.010; p=0.034) levels were independent predictors of severe COVID-19 disease (Table 2).

ROC curve analysis

The efficacy of biochemical and inflammatory markers for the prediction of severe disease was evaluated by ROC analysis. AUC of copeptin, ferritin, CRP, fibrinogen, AST, LDH, and CK were found as 0.869 (95%CI 0.797–0.940; p<0.001), 0.846 (95%CI 0.765–0.928; p<0.001), 0.826 (95%CI 0.734–0.918; p<0.001),

Table 2. Results of binary logistic regression analysis of potential predictors of severe COVID-19 disease.

	Univariable model		Multivariable model	
	р	OR (95%CI)	р	OR (95%CI)
Age, years	<0.001	1.055 (1.024–1.086)	0.007	1.139 (1.036–1.251)
Sex, male	0.25	1.653 (0.699–3.909)		
Copeptin, ng/mL	<0.001	1.740 (1.362–2.223)	0.009	2.647 (1.272–5.510)
CRP, mg/L	<0.001	1.033 (1.017–1.050)		
Urea, mg/ dL	0.57	1.005 (0.988–1.022)		
Fibrinogen, mg/dL	<0.001	1.006 (1.003–1.010)		
AST, U/L	0.003	1.054 (1.018–1.091)		
ALT, U/L	0.008	1.048 (1.012–1.084)		
LDH, U/L	0.005	1.008 (1.002–1.013)		
Ferritin, ng/mL	<0.001	1.006 (1.003–1.009)	0.034	1.005 (1.000–1.010)
CK, U/L	0.06	1.005 (1.000–1.011)		
Troponin, μg/L	0.59	1.616 (0.288–9.075)		

CRP: C-reactive protein; AST: aspartate transaminase; ALT: alanine transaminase; LDH: lactate dehydrogenase; CK: creatine kinase.

0.743 (95%CI 0.639–0.847; p<0.001), 0.740 (95%CI 0.635–0.845; p<0.001), 0.676 (95%CI 0.563–0.789; p=0.005), and 0.698 (95%CI 0.590–0.807; p=0.002), respectively. Among these parameters, copeptin occupied the maximum area with 0.869. When the cutoff value for copeptin in predicting of severe disease was determined to be 7.84, the sensitivity was determined as 78%, whereas the specificity was 80%.

DISCUSSION

Biomarkers that predict the outcomes of diseases, should be higher in the disease groups compared with the healthy groups and should reflect the severity of the disease. The significance of copeptin in the patients with COVID-19 and its role as a biomarker in predicting disease severity are unknown. Our study is the first one to evaluate the relationship between severity of COVID-19 and copeptin and striking results were obtained in our study. It was determined that copeptin levels of the patients with severe COVID-19 were significantly higher than the patients with mild/moderate COVID-19, and this increase was correlated with disease severity.

It was demonstrated that copeptin played active roles in lung diseases, such as pneumonia, and it was superior to traditional inflammatory markers as a prognostic marker in the patients with CAP^{5,18}. The levels of copeptin in the patients with COVID-19 could be increased due to hemodynamic, osmotic, or inflammatory reasons. In the patients with pneumonia, it was discovered that the levels of copeptin were elevated as the severity of sepsis was elevated¹². Similarly, as the severity of the disease increases in the patients with COVID-19, the expression of copeptin, simultaneously with AVP, from neurohypophysis could be increased with hemodynamic and osmotic signals due to changes in blood pressure and plasma osmolality. Furthermore, it was reported that the endotoxins and inflammatory markers, which increased during respiratory infections, stimulated AVP secretion^{4,6}. In our study, it was determined that inflammatory markers, such as CRP, ferritin, and fibrinogen, were elevated in the patients with severe COVID-19, whereas positive correlations were discovered between serum copeptin levels, serum ferritin, and fibrinogen levels. As the severity of the disease was increased, it was believed that the inflammatory markers and cytokines contributed to the increase in copeptin levels. Additionally, it is known that the gas exchange in the lungs during lower respiratory tract infections results in changes in the AVP system. AVP expression could be induced by certain factors such as acidosis, pain, hypoxia, or neuroendocrine stress as well⁶. In the patients with severe COVID-19, it is known that oxygen levels are lower and the stress responses are higher. Conditions such as hypoxia and pain, which develop as the

severity of pneumonia increases, could contribute to copeptin expression. In the current study, it was discovered that the saturation levels were lower in severe patients and saturation levels and copeptin levels were strongly correlated.

Up to date, the prognostic value of copeptin in CAP patients was investigated in several studies in the literature. Masiá et al.¹⁰ evaluated 174 CAP patients and reported that copeptin was an independent marker of mortality. In the current study, it was determined that copeptin yielded high diagnostic accuracy in predicting mortality when the cutoff value was regarded as >18.9 pmol/L (sensitivity: 71.4%, specificity: 79.5%). Similarly, Kolditz et al. 19 evaluated 51 CAP patients and investigated copeptin's diagnostic value in predicting admission to the intensive care unit 7 days from hospitalization and predicting mortality by using ROC analysis. In this study, it was determined that copeptin had 78% sensitivity and 79% specificity when the cutoff value was regarded as 35 pmol/L. In our study, it was observed that copeptin had 78% sensitivity and 80% specificity in terms of distinguishing the patients with severe COVID-19 from those patients with mild/moderate when the optimal cutoff value was regarded as 7.84 ng/mL. According to the findings of our study, copeptin had a higher predictive value in predicting severe disease compared with biochemical and inflammatory markers such as CRP, ferritin, fibrinogen, and LDH used routinely. Furthermore, copeptin was found to be an independent predictor of severe COVID-19 disease in the multivariable logistic regression analysis. Considering the findings of our study, copeptin can be a simple and useful marker for predicting the severity of COVID-19

with high sensitivity and specificity while yielding rapid results. Nevertheless, our study is a first on this subject, and a series of studies are required to support these results.

In conclusion, in this study, it was determined that serum copeptin levels were higher in the patients with severe COVID-19 compared with the patients with mild/moderate COVID-19 and related to disease severity. Moreover, multivariable logistic regression analysis identified copeptin as an independent predictor for COVID-19 severity. Therefore, copeptin could be a useful and prognostic biomarker that can be used in the patients with distinguishing severe COVID-19 from the patients with mild/moderate COVID-19.

AUTHORS' CONTRIBUTIONS

İE: Conceptualization, Data curation, Formal analysis, Methodology, Investigation, Resources, Writing – original draft, Writing – review & editing. **KM:** Conceptualization, Methodology, Investigation, Methodology, Resources. **TS:** Methodology, Data Curation, Formal analysis, Investigation, Resources. **ATZ:** Methodology, Data Curation, Investigation, Supervision, Writing – review & editing. **KE:** Formal analysis, Supervision, Writing – review & editing. All authors read and approved the final manuscript.

AVAILABILITY OF DATA

The data that support the findings of this study are available from the corresponding author on reasonable request.

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