Cardiac biomarkers as prognosticators among SARS-CoV-19 patients in a tertiary hospital in **Philippines**

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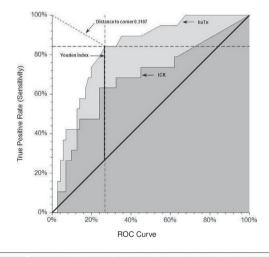
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Background: There is overwhelming volume of confirmed cases of COVID-19, despite this numerous knowledge gaps remain in the diagnosis, management, and prognostication of this novel coronavirus infection. making prevention and control a challenge.

Methods: This retrospective cohort study included patients with realtime reverse transcriptase polymerase chain reaction (rRT-PCR)-confirmed COVID-19. Binary logistic regression was used to determine the association between the cardiac biomarkers and in-hospital mortality. ROC, AUC, and cutoff analyses were used to determine optimal cutoff values for the cardiac biomarkers.

Results: A total of 90 subjects with a complete panel of cardiac biomarkers out of the 224 rRT-PCR confirmed cases were included. The median age was 57 years (IQR, 47-67 years), majority were males. Sixty-six (77.6%) subjects survived while 19 (22.4%) expired. The most common presenting symptom was fever (75.6%), and the most common comorbidity was hypertension (67.8%). Spearman rho correlation analysis showed moderate positive association of high sensitivity troponin I (hsTnI) with in-hospital mortality (R, 0.434, p = <0.001). Multivariate binary logistic regression analysis showed that creatine kinase and hsTnI were independently associated with in-hospital mortality (OR, 4.103 [95% CI, 1.241-13.563], p=0.021; and OR, 7.899 [95% CI, 2.430-25.675], p=0.001, respectively). ROC curve analysis showed that hsTnI was a good predictor for in-hospital mortality (AUC, 0.829 [95% CI, 0.735-0.923], p = <0.001) and that creatine kinase was a poor predictor (AUC, 0.677 [95% CI, 0.531-0.823], p=0.018). Optimal cutoff point derived from the ROC curve for hsTnI was 0.010 ng/ml (J, 0.574) with a sensitivity of 84% (TPR, 0.842 [95% CI, 0.604-0.966]), specificity of 73% (TNR, 0.732 [95% CI, 0.614-0.386]), and an adjusted negative predictive value of 99% (Known prevalence*adjusted NPV, 0.989), a positive likelihood ratio of 20% (LR+, 3.147 [95% CI, 2.044-4.844]) and a negative likelihood ratio of 30% (LR-, 0.216 [95% CI, 0.076-0.615]).

Conclusion: High sensitivity troponin I level was a good tool with a very high negative predictive value in significantly predicting in-hospital mortality among rRT-PCR positive COVID-19 patients.



ROC Curve, Cutoff and Area Under the Curve Analysis for Elevated Cardiac Biomarkers with n-hospital Mortality Among COVID-19 Patients

	hs Troponin I	Creatine kinase
Reference value used	0.015 ng/ml	168 U/liter
Area under the Curve (AUC)	0.829 (0.735-0.923)	0.677 (0.531-0.823)
Sig.*	< 0.001	0.018
Distance to corner (d)	0.311	0.439
Youden index (J)	0.574	0.392
Derived optimal cutoff value	0.010 ng/ml	173 U/lite
Sensitivity (TPR)	0.842 (0.604-0.966)	0.632 (0.384-0.837
Specificity (TNR)	0.732 (0.614-0.831)	0.761 (0.645-0.854
False positive rate	0.268 (0.169-0.386)	0.239 (0.146-0.355
False negative rate	0.158 (0.034-0.396)	0.368 (0.163-0.616
Positive predictive value (PPV)	0.457 (0.288-0.633)	0.414 (0.235-0.611
Known prevalence*Adjusted PPV [‡]	0.142	0.12
Negative predictive value (NPV)	0.945 (0.849-0.989)	0.885 (0.778-0.953
Known prevalence*Adjusted NPV [‡]	0.989	0.975
Prevalence	0.211 (0.132-0.310)	0.211 (0.132-0.310
Accuracy	0.756 (0.654-0.840)	0.733 (0.630-0.821
Positive likelihood ratio (LR+)	3.147 (2.044-4.844)	2.638 (1.540-4.519
Negative likelihood ratio (LR-)	0.216 (0.076-0.615)	0.484 (0.265-0.885
Diagnostics odds ratio	14.596 (3.821-55.766)	5.445 (1.850-16.032

Diagnostics odds ratio 14.596 (3.821-55.766) 5.445 (1.880-16.0. 74 value loss flow (0.65) according to the constraint of the sensitive state sta distance to the top-left corner of the ROC curve for each cutoff value is given by $d = \sqrt{(1 - sensitivity)^2 + (1 - specificity)^2}$. The accuracy reflects the total proportion of units that are correctly predicted or classified.