

The COVID-19 Lab Score: An Accurate Dynamic Tool to Predict In-Hospital Outcomes in COVID-19 Patients

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Abstract

Purpose: Deterioration is sometimes unexpected in SARS-CoV2 infection. The aim of our study is to establish laboratory predictors of mortality in COVID-19 disease which can help to identify high risk patients.

Methods: All patients admitted to hospital due to Covid-19 disease were included. Laboratory biomarkers that contributed with significant predictive value for predicting mortality to the clinical model were included. Cut-off points were established, and finally a risk score was built.

Results: 893 patients were included. Median age was 68.2 years(CI 95% 53.0-83.4). 87(9.7%) were admitted to Intensive Care Unit(ICU) and 72(8.1%) also needed mechanical ventilation support. 171(19.1%) patients died. A Covid-19 Lab score ranging from 0 to 30 points was calculated on the basis of a multivariate logistic regression model in order to predict mortality with a weighted score that included haemoglobin, erythrocytes, leukocytes, neutrophils, lymphocytes, creatinine, C-reactive protein, interleukin-6, procalcitonin, lactate dehydrogenase (LDH), and D-dimer. Three groups were established. Low mortality risk group under 12 points, 12 to 18 were included as moderate risk, and high risk group were those with 19 or more points. Low risk group as reference, moderate and high patients showed mortality OR 4.75(CI95% 2.60-8.68) and 23.86(CI 95% 13.61-41.84), respectively. C-statistic was 0-85(0.82-0.88) and Hosmer-Lemeshow p-value 0.63.

Conclusion: Covid-19 Lab score can very easily predict mortality in patients at any moment during admission secondary to SARS-CoV2 infection. It is a simple and dynamic score, and it can be very easily replicated. It could help physicians to identify high risk patients to foresee clinical deterioration.

Introduction

In December 2019, an outbreak of a new syndrome emerged in the Chinese province of Hubei. A new betacoronavirus was described, called SARS-CoV-2[1]. By april 18th, the World Health Organization reported more than 2 million of confirmed cases of COVID-19 disease (Coronavirus Disease 2019), affecting 200 different countries all around the world [2]. Deaths rose to almost 140 thousand people.

One of the most encouraging characteristic of Covid-19 disease, is the huge differences in the clinical manifestations depending of each patient [3,4,5]. It is suspected that some of them are paucisymptomatic, or even completely asymptomatic. On the other hand, in some patients SARS-CoV-2 could trigger a big inflammation response, causing a severe disease. Severe Acute Respiratory Syndrome is one of the most common and severe complications, due to the predisposition of the coronavirus for the respiratory system, requiring in some times admission in Intensive Care Units (ICU), mechanical ventilation and with an elevated death rate [6].

There are described 3 different phases [7]. First one is the early infection, consisting in the inoculation and the incubation, associated with non-specific symptoms. Secondary, there is the pulmonary involvement, because of the affinity for pulmonary ACE2 receptors, causing bilateral infiltrates, viral multiplication and lung inflammation. Finally, a stage of hyperinflammation is described, affecting extrapulmonary organs consisting in a systemic cytokine storm. Elevated levels of some biomarkers [5] like Interleukin (IL)-2, -6, -7, TNF α , IFN γ , C-reactive protein, ferritin, D-Dimer, procalcitonin, troponin are described in this severe patients. At this stage, mechanical ventilation and admission at ICU are frequently needed, because of hypoxemia, respiratory failure, and even shock due to vasoplegia.

The aim of the study was to identify high risk patients affected with COVID-19, those with a poor outcome, depending on laboratory biomarkers. Mortality was selected as the primary endpoint of interest. Our goal is to stratify the risk of death for patients with Covid-19 at any moment during the admission, using a very simple and reproducible analytic tool, which is made only with laboratory parameters.

Methods

This is a multicentre, retrospective, observational study performed at three university hospitals covering a population of more than one million of inhabitants. In our registry we included all the confirmed cases for SARS-CoV-2 infection from our health areas who were admitted to hospital. This study was approved by the Ethical Committee of Investigation of Santiago and Lugo, and it fulfils the Declaration of Helsinki of 1975.

The most deviant from normal results of the laboratory tests for each biomarker during the admission were selected for the score (they were mostly the higher ones, except for the haemoglobin and erythrocytes).

Normal ranges for the different parameters were: procalcitonin < 0.05 ng/mL; Haemoglobin 13.5-17.5 g/dL for men and 12.2-16.1 g/dL for women; erythrocytes 4.5-5.5 per 10⁶/mm³ ; leukocytes 4.09-10.8 per 10³/mm³ ; neutrophils 1.7-7.33 per 10³/mm³ ; creatinine 0.4-1.3 mg/dL for men and 0.4-1.1 mg/dL for women; C-reactive protein 0-0.5 mg/L; interleukin-6 0-5.0 pg/mL; lactate dehydrogenase (LDH) 200-446 UI/L; D-dimer 0-500 ng/mL.

OUTCOMES

This work is conducted to develop an easily replicable, objective and dynamic score to help physicians to assess the risk of adverse outcomes at any moment during the admission due to COVID-19.

All the information about the clinical evolution and the complications developed was recollected along the admission and continued also when discharged, thanks to remote monitoring. Mortality for any cause was selected as the main outcome, and was recollected from clinical records.

DATA COLLECTION

Standardized forms were used for the setting-up of the database, including demographic information, epidemiological data, previous comorbidities and chronic treatments, the clinical data available at the moment when they were admitted to hospital (symptoms, fever and peripheral O₂ saturation (SpO₂), the results of all laboratory tests done during the admission and the treatment received. Laboratory test and clinical decisions were taking according to physician's criteria. Informed consent was obtained for study participation of each patient. The data in source documents was confirmed independently by at least two physicians.

STATISTICAL ANALYSIS

Continuous variables are presented as mean (SD), whereas discrete variables are presented as percentages. Comparisons between discrete variables were performed using the χ^2 test or Fisher exact test as required, and comparisons between continuous variables using the Student t-test (Table 1).

Backward stepwise logistic regression analysis was performed to determine the predictive factors for mortality. Variables that were significantly associated with mortality in the univariate analysis (Supplementary Table 1) were included in the multivariate model (Table 2). The multivariate adjustment was performed according these variables, in addition to biomarkers: age, comorbidities (hypertension, dyslipidemia, diabetes mellitus, peripheral artery disease, heart disease, Chronic Obstructive Pulmonary Disease (COPD)/asthma), days of symptoms, respiratory insufficiency, and in-hospital drugs (antiviral, chloroquine, ceftriaxone, corticosteroids, anticoagulation, antiplatelet). The incremental value of each significant biomarker, when it was added to a clinical base model, was assessed with the change in the c-index (Table 3).

The optimal cut-point definition for different biomarkers was defined based on the receiver operating characteristic (ROC) curve, using the Youden index (Figure 1). Sensitivity and specificity were reported for cut-points.

After categorizing the biomarkers according to their cut-points, those resulting as independent predictors of mortality by the multivariate logistic regression analysis were incorporated into a risk score (Table 4). The scores assigned to each biomarker were determined according to the value of the odds ratio (OR). The predicted probability of death based on the risk score was graphically represented after modelling by fractional polynomials (Figure 2). The performance of this risk score was tested by assessing its discrimination and calibration capacity (Table 5), for all-cause death. Discrimination was evaluated by calculating the C statistic, and calibration was assessed by the Hosmer-Lemeshow test.

All p values were two-sided and values <0.05 were considered as significant. All statistical analyses were performed using STATA software, version 15.1 (Stata Corp, College Station, Texas, USA), and IBM SPSS software, version 24.0 (IBM, Armonk, New York, USA).

Results

A total of 893 patients were admitted to hospital due to COVID-19. Baseline characteristics are described in Table 1. Median age was 68.2 ± 15.2 years, and 453 were female (50.7%). 87 (9.7%) patients were admitted to Intensive Care Unit (ICU) and 72 (8.1%) also needed mechanical ventilation support and 171 (19.1%) died.

In relation with cardiovascular outcomes, heart failure was the most common complication (61 cases, 6.8%). Acute myocardial infarction, myocarditis and pericarditis were also described (3 (0.3%); 2 (0.2%); 3 (0.3%), respectively). Pulmonary embolism was diagnosed in 8 patients (0.9%).

Univariate analysis with clinical characteristics and treatment to predict mortality was made (Supplementary table 1). Those that were found to have significant difference ($p < 0.05$) (age, comorbidities (hypertension, dyslipemia, diabetes mellitus, peripheral artery disease, heart disease, COPD/asthma), days of symptoms, respiratory insufficiency, in-hospital drugs (antiviral, chloroquine, ceftriaxone, corticosteroids, anticoagulation, antiplatelet)) were included in the multivariate analysis.

After multivariate adjustment, most laboratory parameters were significantly associated with mortality (haemoglobin, leukocytes, neutrophils, neutrophils/lymphocytes ratio, lymphocytes $\times 100$ leucocyte, GOT, GGT, creatinine, CRP, IL-6, procalcitonin, LDH and D-dimer), only lymphocytes, platelets and ferritin values were not associated with high mortality (Table 2). The incremental value of each of these significant biomarkers, when it was added to a clinical base model, was assessed with the change in the c-index, achieving all of them significant difference regarding the reference (Table 3).

In order to convert these variables to categorical ones, the optimal cut-off point definition for each different biomarker was established based on ROC curves, as is represented in Figure 1.

After categorizing the biomarkers according to their cut-off points, those resulting as independent predictors of mortality by the multivariate logistic regression analysis were incorporated into a risk score (Table 4). The scores assigned to each biomarker were determined according to the value of the odds ratio (OR). The one with an individual higher score was procalcitonin < 0.2 ng/mL with 5 points (OR 5.72, CI 95% 3.35-9.76, $p < 0.001$), in relation with the severity of bacterial coinfection. Hemoglobin < 12 g/dL (OR 1.07, CI 95% 1.05-1.09, $p < 0.001$; 1 point), erythrocytes < 4.1 per $10^6/\text{mm}^3$ (OR 2.14, CI 95% 1.19-3.84, $p = 0.011$; 2 points), leukocytes > 8.3 per $10^3/\text{mm}^3$ (OR 2.51, CI 95% 1.56-4.03, $p < 0.001$; 3 points), neutrophils > 8.1 per $10^3/\text{mm}^3$ (OR 2.13, CI 95% 1.14-3.95, $p = 0.017$; 2 points), lymphocytes < 6.5 per 100 leukocytes (OR 2.85, CI 95% 1.82-4.46, $p < 0.001$; 3 points), creatinine (OR 4.10, CI 95% 2.56-6.55, $p < 0.001$; 4 points), C-reactive protein > 4.5 mg/L (OR 3.05, CI 95% 1.08-8.58, $p = 0.035$; 4 points), interleukin-6 > 24 pg/mL (OR 1.83, CI 95% 1.17-2.88, $p = 0.009$; 2 points), lactate dehydrogenase (LDH) ≥ 393 UI/L (OR 4.29, CI 95% 2.49-7.39, $p < 0.001$; 4 points), and D-dimer > 1116 ng/mL (OR 1.92, CI 95% 1.22-3.02, 2 points).

With these individual scores, the COVID-19 Lab score is performed, ranging from 0 to 30 points. The predicted probability of death based on the risk score was graphically represented after modelling by fractional polynomials (Figure 2).

Depending on the COVID-19 Lab score, mortality varies as shown in Figures 2 and 3, the higher is the score, the poorer outcome. Three groups were set, dividing patients in low (<12 points), moderate (12-18 points) and high (19 or higher points) risk of death, with significant mortality differences between them (3.9%, 16.1%, 49.1% respectively, $p > 0.001$).

The performance of this risk score was tested by assessing its discrimination and calibration capacity for all-cause death. Discrimination was evaluated by calculating the C statistic, 0.85 (0.82-0.88), and calibration was assessed by the Hosmer-Lemeshow test, p-value 0.63 (table 5)

Discussion

We constructed a simple score in a cohort of occidental population with COVID-19, which can stratify patients who was admitted to the hospital depending on the risk of death. The most remarkable characteristics of our score are the good performance for mortality risk stratification, that it can be easily obtained. This is because it includes only objective laboratory parameters and due to its dynamic nature, allowing a continuous mortality risk assessment that may help physicians to support clinical decisions in this group of patients. Above all it has to be stated that the experienced clinician and his clinical examination and estimation of the patient's condition remain indispensable.

On account of this, Covid-19 Lab score could have important implications in the future as COVID-19 disease has very different clinical manifestations depending on the patient, and also depending on the stage of the disease. Because of the inflammation and cytokine storm reported in previous studies, patient's status can rapidly change from one day to another or even in a shorter period of time [7,8]. Patients with mild or even no symptoms, can develop acute respiratory insufficiency in a few hours. That is the main reason why not only identifying those cases with higher risk of death, but also trying to foresee a clinical deterioration will be mandatory to implement a close monitoring and to use a more aggressive treatment.

Hyperinflammation and cytokine storm are previously described as the most important reasons of bad outcomes in different retrospective studies about SARS-CoV2 infection [5,9]. This uncontrolled immunity response is the main reason why acute respiratory syndrome emerges in this patients, causing clinical deterioration and the need of mechanical respiratory support. Inflammation biomarkers are known to be elevated in COVID-19 disease, such as interleukins (IL), TNF α , ferritin, D-dimer, C-reactive protein (CRP) or procalcitonin [8,10]. Although procalcitonin in COVID-19 is controversially discussed in literature, it has a high correlation with bacterial superinfection, which increases risk of poor outcomes. Akbarshakh et al, showed that some biomarkers like IL-6, ferritin, CRP and troponin were associated with higher mortality. Furthermore, also patients with elevated D-Dimer levels and lymphocytopenia were assessed to have worse outcomes, probably in relation with a procoagulability and immunodeficiency status [3,11,12].

It is not the first time these types of biomarkers are evaluated in viral infections. Inflammatory reaction and elevated plasma levels were also describe in other viral infections due to other coronavirus like SARS-CoV and MERS-CoV [13,14,15]. Other study conducted by Bautista et al [16] showed that inflammatory

biomarkers were associated with higher risk of death in patients with severe acute respiratory syndrome (SARS), secondary to Influenza A H1N1 infection,.

There are also clinical features associated with higher mortality rate in SARS-Cov2 infection. An older age, comorbidities, previous cardiovascular disease, the presence of fever or respiratory insufficiency are associated with poorer outcomes [10,11,12,17].

Others scores like CURB-65 and MuLBTSA were previously assessed for predicting mortality in bacterial and viral pneumonias respectively. Although Lingxi et al [18] proposed this MuLTBSA score predicting mortality better than CURB65 [19] for viral pneumonia, it is a static score. This characteristic makes it incapable of showing changes in the risk of death from day to another depending on the progression of the infection and it does not include inflammatory parameters. Furthermore, the main difference with our score is that ours can be replicated very easily only with blood test because all variables are laboratory biomarkers.

The clinical manifestations of COVID-19 disease change rapidly in relation with this hyperinflammation reaction, and patients develop acute respiratory insufficiency needing ventilator mechanical support. Having a score like ours, that is not static, it is very simple, and that predicts a hard outcome like mortality using only laboratory biomarkers, could help the physicians to anticipate to clinical deterioration, or even to account for changes to the positive.

Finally, a mobile app that will be available worldwide is being developed to help to calculate quickly the Covid-19 Lab score in order simplify even more the physician´s decisions.

Our study had some limitations. This is a retrospective and observational study, even when this score composed of laboratory biomarkers adjusted by other clinical characteristics emerges to predict mortality in COVID-19 disease in an excellent way, other cofounders might be underestimated. Secondly, antiviral treatment was not the same in all patients, because of the lack of evidence of them at the time of the study. Furthermore, follow up is still ongoing, and some complications or events could be missed. Despite these limitations, it is a strong study, including all patients admitted to hospital from three different centres, based on objective parameters. Finally, for a more reliable assessment of our score performance, it must be validated in other cohorts of patients from different centres and geographic areas.

Conclusion

Covid-19 Lab score is a dynamic simple score only using laboratory biomarkers that can be easily obtained. It can predict in an excellent and dynamic way mortality risk in patients admitted to hospital due to SARS-CoV2 infection in different occasions during the hospital stay.

Not only identifying those patients with high risk of death, but also anticipating the clinical deterioration will be necessary to implement a more aggressive treatment and to keep a close patient monitoring. On

the other hand, should be remarked that the experienced clinician and his clinical examination and estimation of the patient's condition remain indispensable.

Declarations

AUTHOR CONTRIBUTIONS STATEMENTS

Pablo Jose Antunez Muiños and Diego López Otero wrote the main manuscript text and prepared figures 1-3 and tables. All authors reviewed the manuscript

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Tables

Table 1. Baseline characteristics of total population (n=893)	
Baseline characteristic	Value
Age	68.2 ± 15.2
Female sex	453 (50.7%)
Obesity	110 (12.3%)
Institutionalized person	15 (1.7%)
Dementia	17 (1.9%)
Dependency	43 (4.8%)
Health worker	13 (1.5%)
Active smoking	40 (4.5%)
Hypertension	444 (49.7%)
Diabetes mellitus	174 (19.5%)
Dyslipidemia	362 (40.5%)
Peripheral artery disease	21 (2.4%)
Heart disease	149 (16.7%)
Ischemic heart disease	75 (8.4%)
Myocardopathy or depressed LVEF	49 (5.5%)
Valvular heart disease	12 (1.3%)
Atrial fibrillation	20 (2.2%)
Pulmonary disease	55 (6.2%)
COPD or Asthma	84 (9.4%)
Prior stroke	19 (2.1%)
Prior cancer	12 (1.3%)
Hypothyroidism	13 (1.5%)
Autoimmune disease	18 (2.0%)
Anticoagulation	201 (11.4%)
Antiplatelet therapy	144 (16.1%)
ACEI/ARB	333 (37.3%)
Antialdosteronic drug	21 (2.4%)

B-blockers	158 (17.7%)
Calcium channel blocker	65 (7.3%)
Diuretic drugs	113 (12.7%)
Statin	294 (32.9%)
Corticosteroid	24 (2.7%)
Immunosuppression	12 (1.3%)
Days of symptoms	7.3 ± 5.1
Fever	562 (62.9%)
Respiratory insufficiency	350 (39.2%)
Antiviral	751 (84.1%)
Chloroquine	772 (86.5%)
Interferon	33 (3.7%)
Tocilizumab	91 (10.2%)
Azithromycin	619 (69.3%)
Ceftriaxone	450 (50.4%)
Corticosteroids	223 (25.0%)
Anticoagulation	347 (38.9%)
Antiplatelets	542 (60.7%)

Table 1. Baseline characteristics of total population (n=893). Categorical variables in n (%) and quantitative variables in mean ± standard deviation. LVEF: left ventricle ejection fraction

Table2. Univariate and multivariate analysis for biomarkers to predict mortality.						
Laboratory data	Univariate			Multivariate		
	OR	95% CI	p	OR	95% CI	p
Hemoglobin, per g/dL	0.74	0.68-0.80	<0.001	0.82	0.75-0.91	<0.001
Leukocytes, per 10 ³ /mm ³	1.11	1.08-1.14	<0.001	1.08	1.05-1.11	<0.001
Neutrophils, per 10 ³ /mm ³	1.10	1.06-1.14	<0.001	1.07	1.03-1.12	0.001
Lymphocytes, per 10 ³ /mm ³	0.31	0.21-0.48	<0.001	0.72	0.50-1.04	0.083
Platelets, per 10 ⁵ /mm ³	0.80	0.68-0.95	0.010	1.00	0.99-1.01	0.903
Neutrophil-Lymphocyte ratio	1.04	1.02-1.05	<0.001	1.02	1.01-1.03	0.012
Lymphocytes per 100 Leukocytes	0.88	0.86-0.91	<0.001	0.93	0.90-0.96	<0.001
Platelet-Lymphocyte ratio	1.84	1.19-2.86	0.006	1.14	0.63-2.08	0.665
GOT, per 10 UI/L	1.05	1.02-1.08	<0.001	1.05	1.01-1.08	0.009
GGT, per 10 UI/L	1.01	1.00-1.02	0.005	1.02	1.00-1.03	0.019
Creatinine, per mg/dL	2.79	2.19-3.57	<0.001	1.97	1.58-2.47	<0.001
CRP, per 10 mg/L	1.10	1.08-1.12	<0.001	1.09	1.06-1.12	<0.001
IL-6, per 10 pg/mL	1.01	1.00-1.02	0.017	1.01	1.00-1.02	0.014
Ferritin, per 1000 ug/mL	1.25	1.09-1.44	0.002	1.19	0.98-1.43	0.079
Procalcitonin, per ng/mL	1.05	1.02-1.08	0.002	1.05	1.02-1.08	0.001
LDH, per 100 UI/L	1.19	1.12-1.26	<0.001	1.34	1.21-1.49	<0.001
D-Dimer, per 1000 ng/mL	1.05	1.03-1.07	<0.001	1.03	1.01-1.05	0.001

Table2. Univariate and multivariate analysis for biomarkers to predict mortality. Multivariate adjustment: Age, comorbidities (hypertension, dyslipemia, diabetes mellitus, peripheral artery disease, heart disease, COPD/asthma), days of symptoms, respiratory insufficiency, in-hospital drugs (antiviral, chloroquine, ceftriaxone, corticosteroids, anticoagulation, antiplatelet).

Table 3. Model performance to predict mortality basing on the addition of biomarkers.			
Model	C-statistic	95% CI	P-value
Base model (as reference)	0.891	0.866-0.916	ref
Base model + Hemoglobin	0.900	0.877-0.923	0.012
Base model + Leukocytes	0.907	0.884-0.929	0.001
Base model + Neutrophils	0.934	0.914-0.955	<0.001
Base model + Neu-Lymph ratio	0.934	0.914-0.955	<0.001
Base model + Lymph x 100 Leu	0.905	0.884-0.927	0.015
Base model + GOT	0.915	0.892-0.938	0.002
Base model + GGT	0.906	0.881-0.931	0.014
Base model + Creatinine	0.910	0.887-0.933	<0.001
Base model + CRP	0.915	0.892-0.937	0.001
Base model + IL-6	0.916	0.883-0.949	0.021
Base model + Procalcitonin	0.916	0.877-0.926	0.026
Base model + LDH	0.925	0.905-0.946	<0.001
Base model + D-Dimer	0.903	0.879-0.926	0.036

Table 3. Model performance to predict mortality basing on the addition of biomarkers. Model performance after the addition of different biomarker to the base model [age, comorbidities (hypertension, dyslipidemia, diabetes mellitus, peripheral artery disease, heart disease, COPD/asthma), days of symptoms, respiratory insufficiency, in-hospital drugs (antiviral, chloroquine, ceftriaxone, corticosteroids, anticoagulation, antiplatelet)]

Table 4. Multivariate analysis for biomarkers to predict mortality basing on cut-off points (as categorical variables).					
Laboratory data	Multivariate			Points	
	OR	95% CI	p-value		
Hemoglobin < 12 g/dL	1.07	1.05-1.09	<0.001	1	
Erythrocytes < 4.1 per 10 ⁶ /mm ³	2.14	1.19-3.84	0.011	2	
Leukocytes > 8.3 per 10 ³ /mm ³	2.51	1.56-4.03	<0.001	3	
Neutrophils > 8.1 per 10 ³ /mm ³	2.13	1.14-3.95	0.017	2	
Lymphocytes < 6.5 per 100 Leukocytes	2.85	1.82-4.46	<0.001	3	
Creatinine > 1.1 mg/dL	4.10	2.56-6.55	<0.001	4	
CRP > 4.5 mg/L	3.05	1.08-8.58	0.035	4	
IL-6 > 24 pg/mL	1.83	1.17-2.88	0.009	2	
Procalcitonin > 0.2 ng/mL	5.72	3.35-9.76	<0.001	5	
LDH ≥ 393 100 UI/L	4.29	2.49-7.39	<0.001	4	
D-Dimer > 1116 ng/mL	1.92	1.22-3.02	0.005	2	

Table 4. Multivariate analysis for biomarkers to predict mortality basing on cut-off points (as categorical variables). Multivariate adjustment: Age, comorbidities (hypertension, dyslipemia, diabetes mellitus, peripheral artery disease, heart disease, COPD/asthma), days of symptoms, respiratory insufficiency, in-hospital drugs (antiviral, chloroquine, ceftriaxone, corticosteroids, anticoagulation, antiplatelet).

Table 5. Predictive ability of COVID-19 Lab Score for mortality.			
COVID-19 Lab Score		Mortality	
Odds Ratio	Continuous		1.23 (1.19-1.26)
	Categorical	Low risk	ref
		Moderate risk	4.75 (2.60-8.68)
		High risk	23.86 (13.61-41.84)
Discrimination	C-statistics	0.85 (0.82-0.88)	
Calibration	Hosmer-Lemeshow p-value		0.63
	Chi ²		6.11

Table 5. Predictive ability of COVID-19 Lab Score for mortality.

Figures

Figure 1. Discrimination of biomarkers

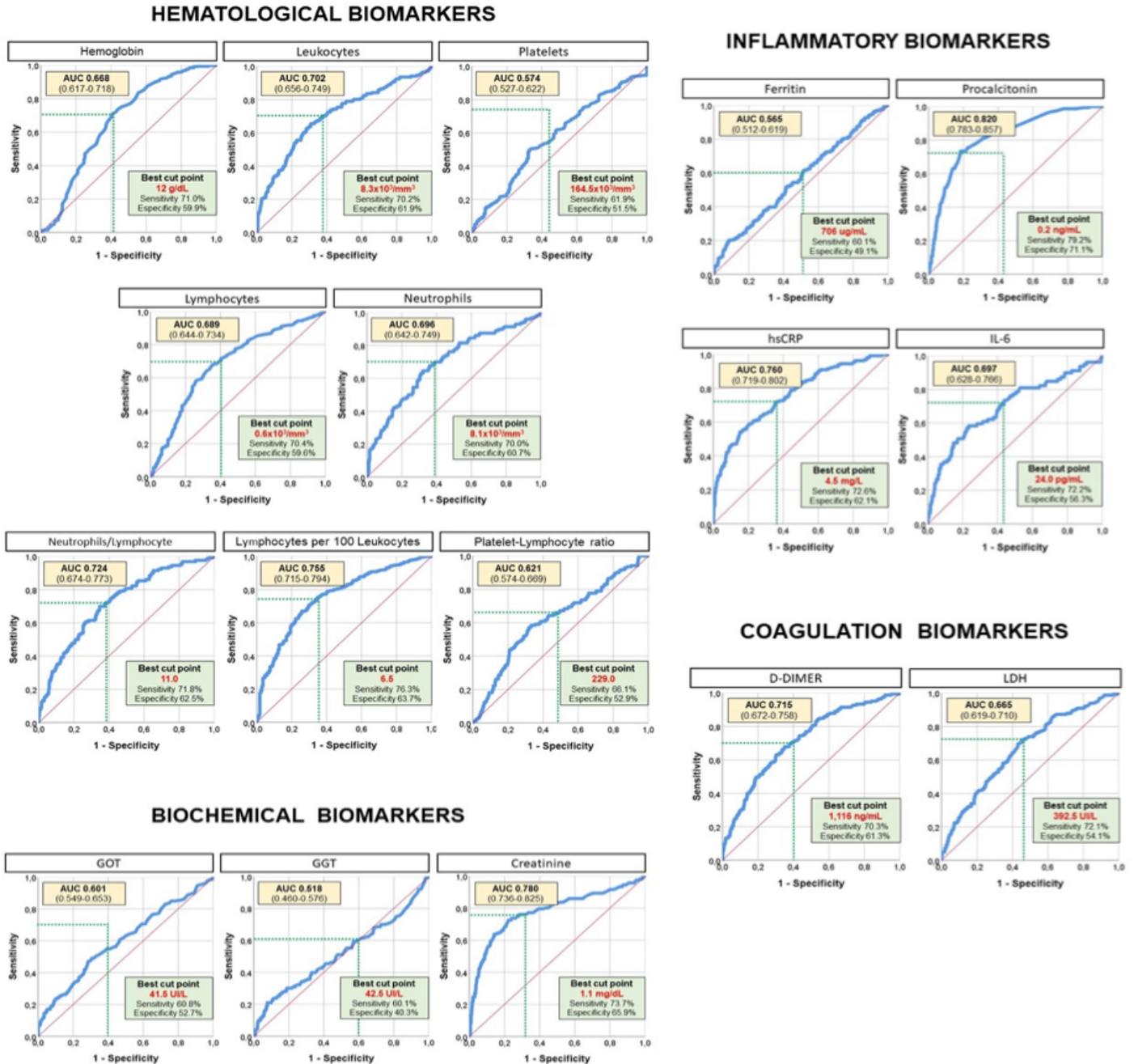


Figure 1

Discrimination of biomarkers. AUC: Area under the curve for each different biomarker

Figure 2. COVID-19 Lab Score: Histogram and predicted mortality.

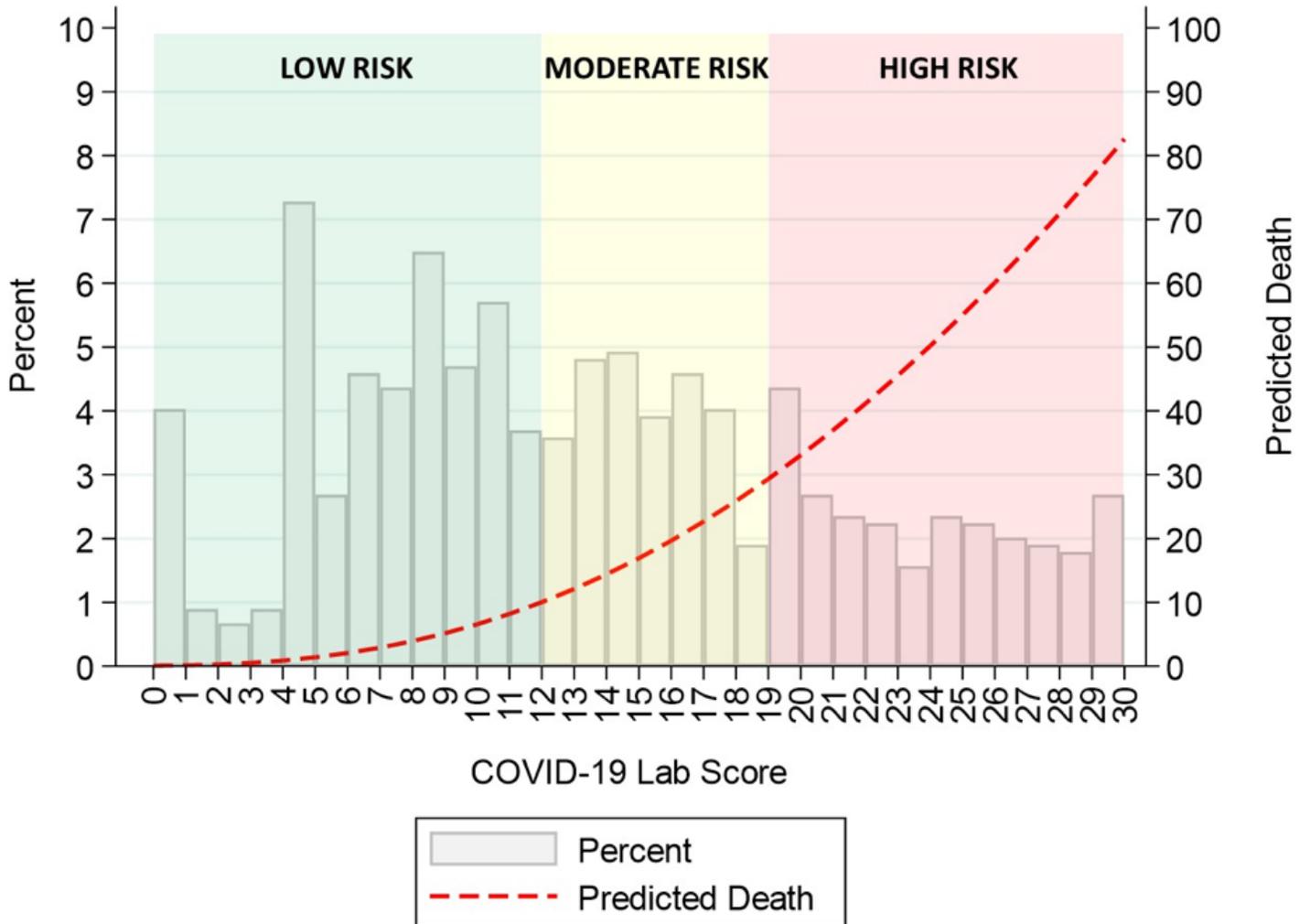


Figure 2

COVID-19 Lab Score: Histogram and predicted mortality. Represents the risk of mortality depending on the score, divided in three different groups.

Figure 3. Outcomes by risk groups of COVID-19 Lab Score

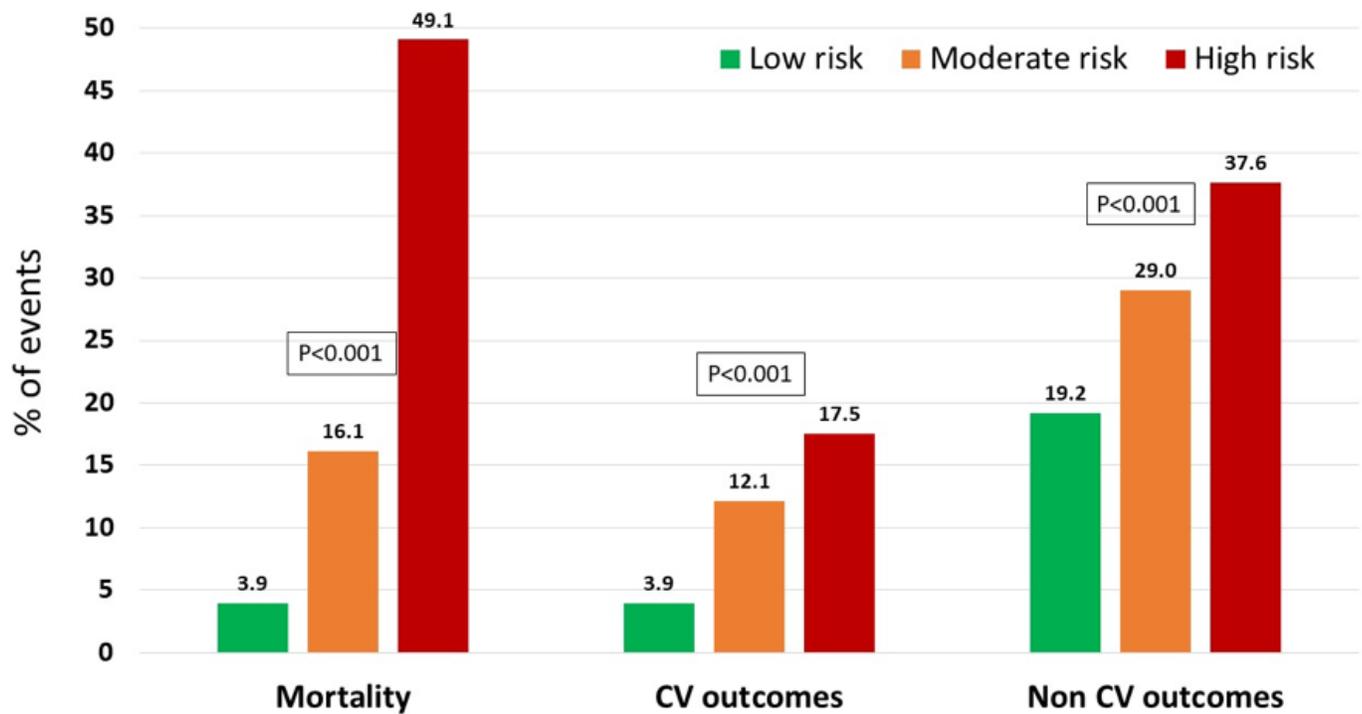


Figure 3

Outcomes by risk groups of COVID-19 Lab Score. CV: cardiovascular (myocardial infarction, hospitalizations due to heart failure, stroke)

Supplementary Files

This is a list of supplementary files associated with this preprint. Click to download.

- [Tablasuplbiomarkers.docx](#)