

An Analysis COVID-19 in Mexico: a Prediction of Severity



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BACKGROUND: Coronavirus disease 2019 (COVID-19) causes a mild illness in most cases; forecasting COVID-19-associated mortality and the demand for hospital beds and ventilators are crucial for rationing countries' resources.

OBJECTIVE: To evaluate factors associated with the severity of COVID-19 in Mexico and to develop and validate a score to predict severity in patients with COVID-19 infection in Mexico.

DESIGN: Retrospective cohort.

PARTICIPANTS: We included 1,435,316 patients with COVID-19 included before the first vaccine application in Mexico; 725,289 (50.5%) were men; patient's mean age (standard deviation (SD)) was 43.9 (16.9) years; 21.7% of patients were considered severe COVID-19 because they were hospitalized, died or both.

MAIN MEASURES: We assessed demographic variables, smoking status, pregnancy, and comorbidities. Backward selection of variables was used to derive and validate a model to predict the severity of COVID-19.

KEY RESULTS: We developed a logistic regression model with 14 main variables, splines, and interactions that may predict the probability of COVID-19 severity (area under the curve for the validation cohort=82.4%).

CONCLUSIONS: We developed a new model able to predict the severity of COVID-19 in Mexican patients. This model could be helpful in epidemiology and medical decisions.

Keywords COVID-19 · Mortality · Hospitalization · Severity

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INTRODUCTION

In December 2019, in a Wuhan animal market, clusters of patients with pneumonia were identified. The etiology of this pneumonia was a novel coronavirus: the severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) ^{1–3}. The World Health Organization announced the name of the illness as the coronavirus disease 19 (COVID-19). Up to June 7th, 2021, more than 173 million cases of COVID-19 have been confirmed worldwide, with over 3.5 million deaths recorded across the countries (Johns Hopkins University Web page) ⁴. Latin-American countries, including Mexico, are among the highest number of deaths.

Forecasting COVID-19-associated mortality and the demand for hospital beds and ventilators are crucial for rationing countries' resources ⁵. Number of people in critical conditions has compromised the healthcare capacity in some countries during the outbreak ⁶.

Even though COVID-19 causes a mild illness in most cases (nearly 50–75% of positive to SARS-CoV-2 remain without symptoms) ^{7,8}, mortality and hospital admissions for COVID-19 are a burden in Mexico; moreover, approximately 10% of symptomatic patients will develop any of dyspnea, interstitial pneumonia, acute respiratory distress syndrome, or multiorgan dysfunction ⁷. Advanced age, male gender, obesity, and the presence of non-communicable diseases, such as type 2 diabetes, hypertension, cardiovascular disease (CVD), chronic obstructive pulmonary disease (COPD), and cancer are factors associated with severe COVID-19, higher mortality, and, consequently, hospitalization ^{2,9–13}. In Mexico, almost 50% of patients with COVID-19 reported at least one comorbidity ¹⁴. Remarkably, 38.8% of patients with COVID-19 in Mexico were hospitalized, and the risk of hospitalization increases with the number of comorbidities ¹⁵.

This study aimed to identify factors associated with severe COVID-19 (death or hospitalization) in Mexico and develop a model using readily available clinical variables to predict severe COVID-19.

MATERIALS AND METHODS

Study Design

This retrospective cohort analysis uses the Mexican Epidemiological Surveillance System data for Viral Respiratory Diseases. We obtained all data from publicly available sources. Informed consent was not required as the information was de-identified.

Data Collection

We used free license data from the Mexican Epidemiological Surveillance System for Viral Respiratory Disease. In brief, all patients with COVID-19 symptoms who were tested for SARS-CoV-2 infection receive a medical questionnaire at the test time. This questionnaire includes demographic variables and comorbidities; moreover, all data for positive patients were updated daily if the patient was hospitalized or died (the patient was followed by the unit that made the test).

From the registered variables in the questionnaire, we selected the following for this study: demographic variables, smoking status, pregnancy, and comorbidities, including hypertension, type 2 diabetes, COPD, asthma, and others. In addition, we excluded variables reported after the SARS-CoV-2 confirmation or after hospitalization by COVID-19 (intubation, pneumonia, admission to the intensive care unit). Severe COVID-19 was defined as death or requirement of hospitalization. Hospitalization, death, and recovery status are updated daily in the database.

We performed a summary analysis of data included up to June 7th, 2021. However, we selected patients included in the database for the model development before applying the first dose of the vaccine in Mexico (December 23rd, 2020) because of a potential modification of the vaccine in the severity of COVID-19 (Fig. 1, Supplementary material).

Statistical Analysis

Descriptive statistics were used to summarize the characteristics of all patients from derivation and validation cohorts. We expressed categorical variables as frequency and percentages; univariate comparison for categorical variables was performed with χ^2 or Fisher's test.

We randomly selected 70% of observations to derive the logistic regression model (derivation cohort); the other 30% of the patients were assigned to validate the model (validation cohort).

For this study, we considered severe COVID-19 if the patients were hospitalized or died. We developed a logistic regression model to estimate the probability of severe COVID-19.

All variables had missingness less than 5% (supplementary material Table 1); for the model development and

validation, we eliminated the patients with missing data for the variables included in the final model (supplementary material Table S1).

After confirming the absence of collinearity, we included all possible covariates to develop the model. We explored trends of variables with non-parametric regression models that describe the relationship between a response variable and the predictors without assuming a linear relationship among the variables. We began with an initial saturated logistic model including a flexible non-linear age effect with two-way interactions. Interactions were removed from the model (one by one) when the p -value was > 0.05 (backward selection) up to maintain only the significant interactions variables; the main variables and splines were held in the model if there were significant interactions (including the non-significant primary variable). Finally, we estimated the probability of being hospitalized using the model. We conducted all statistical analysis with R version 4.0.4 (The R Foundation for Statistical Computing) and RStudio Version 1.3.1093 © 2009–2020 RStudio, Inc.

RESULTS

The initial database included 7,134,254 patients with suspected COVID-19; 4,699,692 were negative to SARS-CoV-2 or were considered without data of COVID-19 (Fig. 1 in supplementary appendix). In R, we performed mining and cleaning data; for example, the database included pregnant patients less than 5 years or older than 50 (we corrected these values as non-pregnant). Moreover, male patients had missing data in the variable of pregnancy when the real value was non-pregnant.

A total of 2,434,562 patients were considered with the diagnosis of COVID-19 up to June 7th, 2021; mean age (SD) was 43.6 (17.1) years, 1,218,425 (50%) were men, 456,909 (18.8%) required hospitalization, and 228,828 patients (9.4%) died; severe COVID-19, including hospitalized or death patients, occurred in 476,512 patients (19.6%). Table 1 shows the frequency of smokers and comorbidities registered in the database according to the severity (all patients included up to June 7th, 2021).

We excluded 1,216,137 patients admitted in the database after December 23rd, 2020; therefore, the model development and validation database included 1,435,316 patients (Fig. 1 supplementary appendix). Table 2 shows the analyzed characteristics and the comparison between severe and non-severe patients with COVID-19 and the total cohort of patients before the vaccine in the univariate analysis. The proportion of indigenous patients, smokers, and with chronic conditions was significantly higher in severe than in non-severe COVID-19 (Table 2); in contrast, pregnancy and asthma were more frequent in non-severe COVID-19 ("protective factors"). Figure 1 shows a univariable summary that describes the most substantial effect of chronic renal failure

Table 1 Comparison of Severe and Non-severe Patients, Including All Patients up to June 7th, 2021

	Overall	Non-severe	Severe	<i>p</i> -value
	<i>n</i> = 2,434,562	1,958,050	476,512	
Sex = male (%)	1,218,425 (50.0)	934,326 (47.7)	284,099 (59.6)	<0.001
Age (mean (SD))	43.61 (17.07)	40.1 (15.3)	58.2 (16.3)	<0.001
Hospitalization (%)	456,909 (18.8)	0 (0.0)	456,909 (95.9)	<0.001
Indigenous (%)	20,052 (0.9)	13,661 (0.7)	6391 (1.4)	<0.001
Pregnancy (%)	15,650 (0.6)	12,824 (0.7)	2826 (0.6)	<0.001
Diabetes (%)	319,597 (13.2)	166,102 (8.5)	153,495 (32.4)	<0.001
COPD (%)	26,565 (1.1)	10,057 (0.5)	16,508 (3.5)	<0.001
Asthma (%)	52,784 (2.2)	43,375 (2.2)	9409 (2.0)	<0.001
Immunosuppression (%)	19,857 (0.8)	10,176 (0.5)	9681 (2.0)	<0.001
Hypertension (%)	415,287 (17.1)	233,790 (12.0)	181,497 (38.3)	<0.001
Other comorbidities (%)	46,640 (1.9)	24,844 (1.3)	21,796 (4.6)	<0.001
CVD (%)	37,237 (1.5)	17,470 (0.9)	19,767 (4.2)	<0.001
Obesity (%)	344,173 (14.2)	243,962 (12.5)	100,211 (21.1)	<0.001
CRF (%)	35,898 (1.5)	11,206 (0.6)	24,692 (5.2)	<0.001
Smoker (%)	177,712 (7.3)	142,361 (7.3)	35,351 (7.5)	<0.001
Deceased (%)	228,838 (9.4)	0 (0.0)	228,838 (48.0)	<0.001

Abbreviations: *COPD* chronic obstructive pulmonary disease, *CVD* cardiovascular disease, *CRF* chronic renal failure

Table 2 Comparison of Patients with Severe and Non-severe COVID-19 (Patients Admitted in the Database Before December 23rd, 2020)

	Overall	Non-severe	Severe	<i>p</i> -value
<i>n</i>	1,435,316	1,124,126	311,190	
Sex = male (%)	725,289 (50.5)	537,214 (47.8)	188,075 (60.4)	<0.001
Age (mean (SD))	43.94 (16.88)	40.2 (15.0)	57.6 (16.3)	<0.001
Hospitalization (%)	297,964 (20.8)	0 (0.0)	297,964 (95.7)	<0.001
Indigenous (%)	13,181 (1.0)	8668 (0.8)	4513 (1.5)	<0.001
Pregnancy (%)	9304 (0.6)	7484 (0.7)	1820 (0.6)	<0.001
Diabetes (%)	201,755 (14.1)	100,053 (8.9)	101,702 (32.9)	<0.001
COPD (%)	17,820 (1.2)	6506 (0.6)	11,314 (3.7)	<0.001
Asthma (%)	34,323 (2.4)	27,732 (2.5)	6591 (2.1)	<0.001
Immunosuppression (%)	13,541 (0.9)	6676 (0.6)	6865 (2.2)	<0.001
Hypertension (%)	259,529 (18.1)	141,363 (12.6)	118,166 (38.2)	<0.001
Other comorbidities (%)	30,130 (2.1)	15,507 (1.4)	14,623 (4.7)	<0.001
CVD (%)	24,870 (1.7)	11,529 (1.0)	13,341 (4.3)	<0.001
Obesity (%)	226,154 (15.8)	157,725 (14.1)	68,429 (22.1)	<0.001
CRF (%)	23,831 (1.7)	7165 (0.6)	16,666 (5.4)	<0.001
Smoker (%)	107,398 (7.5)	83,621 (7.5)	23,777 (7.7)	<0.001
Deceased (%)	147,180 (10.3)	0 (0.0)	147,180 (47.3)	<0.001

Abbreviations: *COPD* chronic obstructive pulmonary disease, *CVD* cardiovascular disease, *CRF* chronic renal failure

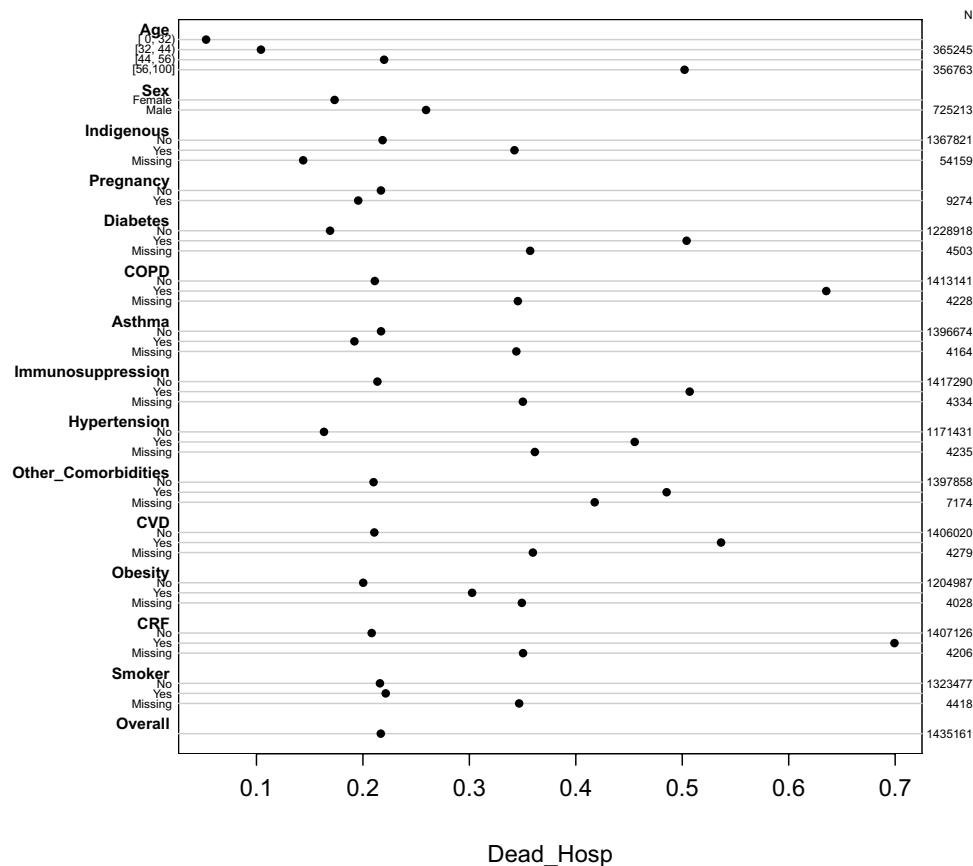
(CRF), chronic obstructive pulmonary disease (COPD), and cardiovascular disease (CVD) on the severity of COVID-19.

In the model development (derivation cohort—960,201 patients), we explored trends with non-parametric regression (with loess fit). As a result, we obtained the age distribution: Fig. 2 in the supplementary appendix shows the association between age and the probability of severe COVID-19 in different statuses of the categorical variables; these figures

showed non-linearity of age and interactions between age and categorical variables. We modeled age as a restricted cubic spline (or natural splines) with 5 degrees of freedom.

In the initial analysis, we eliminated the observations with missing data (Table 1 in supplementary material); the initial saturated logistic regression model allows for a flexible non-linear effect of age and two-way interactions. We reduced the model by removing the interactions when the *p*-value

Fig. 1 Univariable summaries of data used for development and validation. The figure shows missing data by variables. Abbreviations: COPD, chronic obstructive pulmonary disease; CVD, cardiovascular disease; CRF, chronic renal failure; Dead_Hosp, dead or hospitalization or both



was > 0.05 (backward selection). Because of the non-linearity, the log-odds is the best form to express the results shown in Fig. 2; Fig. 2 shows the effect on log odds of smoker men and women is higher in older people. Figure 2 also shows the impact of asthma according to age in men and women. These figures show the only reduction in log odds of severity for younger people. Still, there is a notorious increase in the odds of severity in people older than 25 years for smokers.

Model discrimination was good with an area under the curve of 82.5% in the derivation, and 82.4% in the validation cohort (Supplementary material 1, Fig. 3); the calibration plot showed a good agreement between predicted and observed risks (Fig. 4).

The final model—with 14 main variables, splines, and interactions—is shown in Fig. 3 supplementary material. In addition, this supplemental material shows the beta coefficients, standard errors, and *p*-values; with these values, we developed an application in shiny for R and RStudio to calculate the probability of COVID-19 severity at: <https://marcomtzmtz.shinyapps.io/HospitalizationCOVID/>.

DISCUSSION

This study focused on creating a score that helps distinguish patients with a severe COVID-19 in México using accessible clinical variables. We report developing a model that predicts severe-COVID-19 based on readily available clinical

variables asked by the Mexican Epidemiological Surveillance System for Viral Respiratory Diseases. In Mexican patients with COVID-19, 14 variables can predict severe COVID-19 with a well-calibrated model with proper discrimination.

Definition of severe COVID-19 may be different across authors: severe illness usually begins with dyspnea and hypoxemia, and these patients commonly meet criteria for ARDS requiring hospitalization or concluding in death¹⁶. We considered severe COVID-19 when patients died or were hospitalized because of the characteristics of our database (these data were daily updated for all patients).

Since the pandemic began, all countries have made substantial efforts to ascertain optimal strategies for the medical attention of the outbreak. COVID-19 pandemic represents a significant public health challenge for the world^{17,18}. Even though 80% of patients with COVID-19 have a mild illness², estimating required hospital beds is critical to scaling the health services capacity⁵. In the Latin American region, Mexico is one of the countries with the highest number of deaths cases of COVID-19⁴. The Mexican health sector has 49,083 hospital beds, 2,446 intensive care unit beds, and 5,523 mechanical ventilators¹⁹.

The probability of COVID-19 severity increases in known factors: obesity²⁰, type-2 diabetes²¹, coronary heart disease²¹, systemic hypertension²², and chronic renal failure²³. Mexico is facing the double burden of malnutrition and the prevalence of chronic diseases, including type 2 diabetes, obesity, and hypertension; these diseases caused almost half of all deaths in

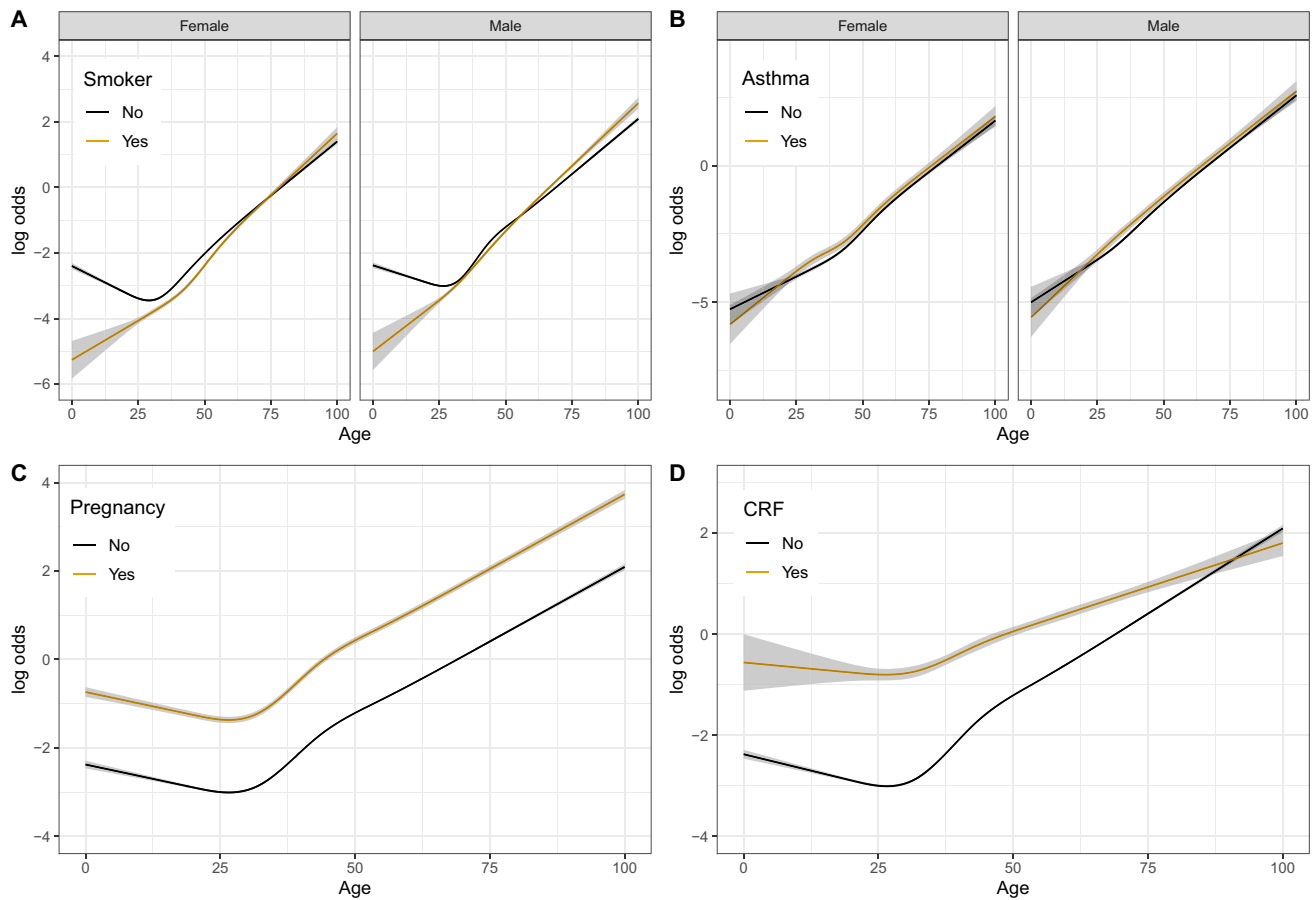


Fig. 2 Predictions based on the model for smoking, asthma, pregnancy, and chronic renal failure (CRF)

Mexico before COVID-19 and are related to COVID-19 severe upon admission in Mexican patients²⁴.

In our study, men have a higher risk of severe COVID-19 than women; this finding is consistent with the study reported by Jin et al., who describe worse outcomes and death for men²⁵.

Moreover, sex and age play a central role in the expression of membrane-bound angiotensin-converting enzyme 2, also associated with a severe COVID-19²⁶.

Our study shows an increased probability of severity for pregnant patients. The hospitalization odds of pregnancy increased in our study, probably related more to the monitoring of the pregnant patient than to the severity of the disease. Case series inform lower or no mortality of pregnant patients with COVID-19 compared to other coronavirus infections (severe acute respiratory syndrome (SARS) and the Middle East respiratory syndrome (MERS)^{27,28}. A systematic review of 77 studies showed that pregnant women with COVID-19 have an increased risk of admission to an intensive care unit, preterm birth, and reported risk factors for severe COVID-19, including advanced maternal age, high body mass index, and pre-existing comorbidities²⁹.

In our study, in the univariate analysis, asthma reduces the odds of severity. Multivariate analysis, including interactions

and non-linear relationships of age, shows that asthma increases the odds of severity, particularly in older men. Mahdavinia et al. found prolonged intubation in patients with asthma and COVID-19, but asthma was not associated with a higher rate of acute respiratory distress syndrome nor hospitalization³⁰.

Indigenous people had a higher odd for hospitalization in our study; we consider this fact an indirect marker of social factors of the Mexican indigenous population with potential poor access to health services³¹. Moreover, other authors have reported a correlation between mortality and healthcare availability^{32,33}. Mortality for COVID-19 has been reported as higher in low-income neighborhoods and rural communities in Ecuador and Mexico^{34,35}.

There are prediction models for the diagnosis and prognosis of COVID-19; some of them require studies like chest tomography, or their prediction could be unreliable when applied in daily practice³⁶. DeCaprio et al. developed a vulnerability index but with other conditions different to COVID-19³⁷; Gong et al. developed a tool to predict severe COVID-19, but the weaknesses of this study were the small sample size and requirement of laboratory tests to predict the severity³⁸. In Spain, Martín-Rodríguez et al. showed that prehospital lactate improved the National Early Warning Score 2 (NEWS2) capacity to detect the mortality risk³⁹. The Epic Deterioration Index in the United

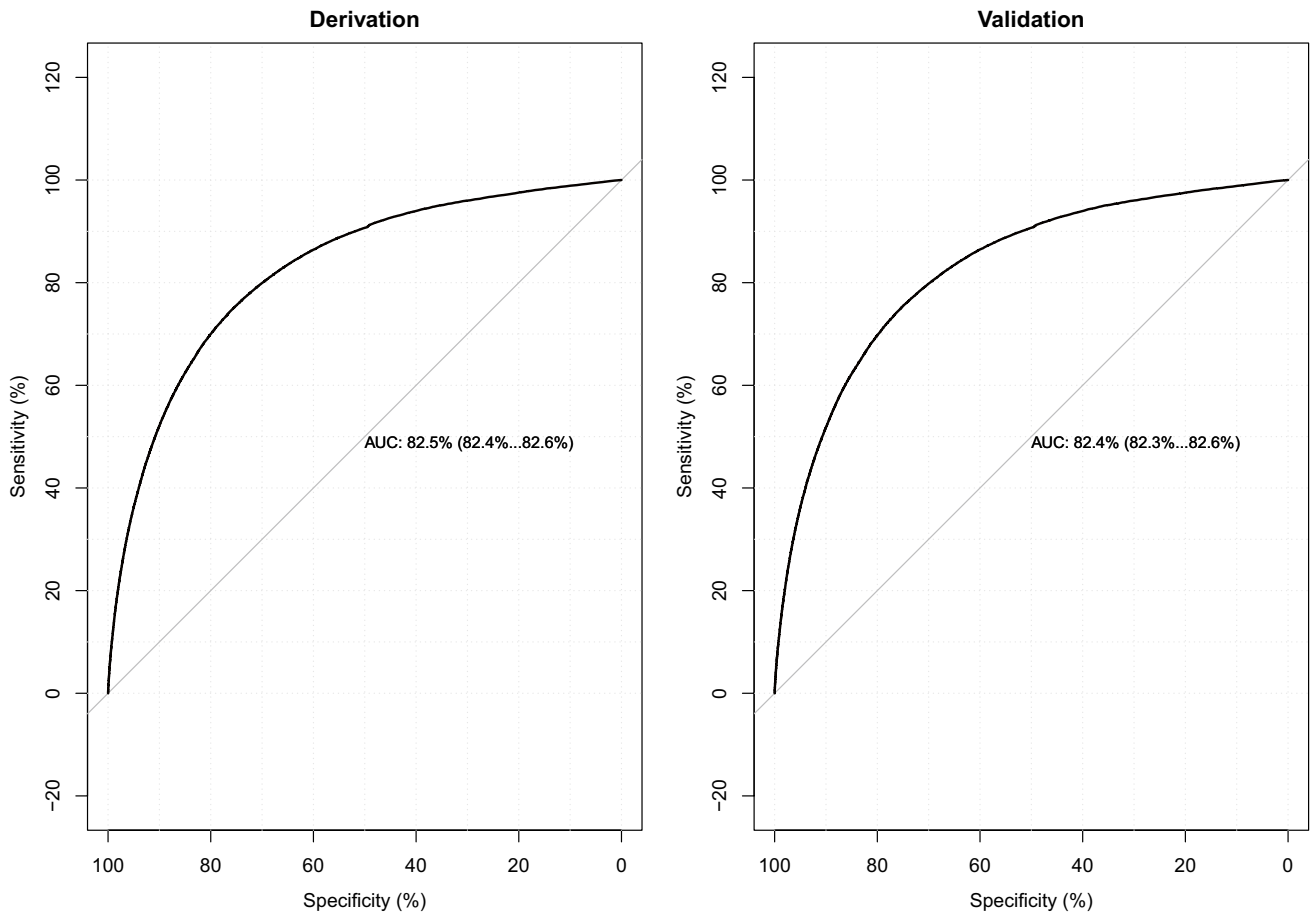
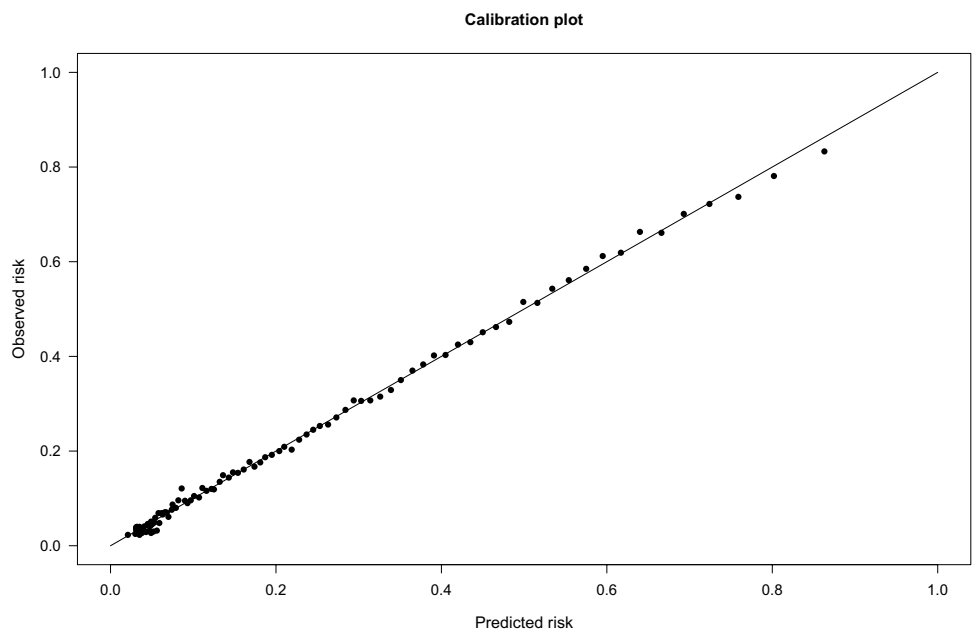


Fig. 3 Receiver operating characteristic (ROC) curves for derivation and validation cohorts

States identifies high and low risks patients; but, this index was developed in 392 hospitalized patients with low sensitivity in contrast to our study that include hospitalized and non-hospitalized large cohort⁴⁰.

We generated a reliable index that can be applied easily in clinical practice and predict the severity of COVID-19 patients. We suggest our score for a strict follow-up of patients who do not require hospitalization at the moment

Fig. 4 Calibration plot. Agreement between predicted and observed hospitalization risks. (Derivation cohort)



of diagnosis but have a high risk of severity. For example, a 51-year-old man with type 2 diabetes, systemic hypertension, chronic renal failure, and obesity scored 79.4% of the probability of being a severe COVID-19. Maybe this patient does not require hospital admission at the time of diagnosis, but, according to our score, more than 70% of patients in this group will require hospitalization or die. We suggest a strict follow-up of his symptoms and tight laboratory work. Another utility of our study is to prioritize the uses of vaccines or specific medications for COVID-19. For example, the group of 35-year-old diabetic patients and chronic renal failure have a 65.8% probability of being a severe COVID-19, compared with patients of 60 years without comorbidities having a 35.7% of being a severe COVID-19.

The strengths of our study include the number of patients and the information contained in the Mexican Epidemiological Surveillance System for Viral Respiratory Diseases. It helped us to develop a model that concluded in a helpful score to predict hospitalization. We provide a practical and inexpensive tool; our conclusions are robust because of the large sample size and validation. In Mexico, general doctors affront decisions about the care of patients with suspected COVID-19 in the community. Still, without access to a laboratory or radiological tests, this fact is typical for developing countries which our model could be helpful after validation for each country.

We consider that our score improves the prediction of Bello-Chavolla et al.⁴¹: we included all possible interactions and the non-linearity of age that must be included in every model. Furthermore, because of the robust statistics used in our model, we demonstrated the spurious “protective effect of asthma” in the severity of COVID-19^{41,42}.

A limitation of our study is the retrospective design. We did not have the information about the previous treatment and the activity of comorbidities, such as asthma or hypertension. Another limitation is that some clinical data were self-reported or reported by relatives but not confirmed as medical records were unavailable to corroborate the information.

Another limitation of our study is the no inclusion of laboratory parameters: there are some like hemogram indexes^{43,44}, hepcidin, and ferritin⁴⁵ that may have a role as valuable tools in the prediction of COVID-19 severity; however, we considered our score with a higher utility in developing countries and epidemiology because it does not require to have laboratory tests.

We have no information on which variants affect our patients; according to the date of inclusion, low or no patients with delta variant were not included, which is another limitation. Therefore, our score must be validated with the newer variants of SARS-CoV-2.

In summary, we developed and validated an inexpensive, readily available score to predict severe COVID-19 in Mexican patients with the SARS-CoV-2 infection. With this score, the clinician may suggest a stricter follow-up

or implement re-evaluation for patients who were not admitted at the first examination but have a high risk of severe COVID-19. Thus, this model could have a crucial role in epidemiology and clinician’s decision-making.

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Supplementary Information The online version contains supplementary material available at <https://doi.org/10.1007/s11606-021-07235-0>.

Declarations

Conflict of Interest All authors declare that they have no any conflict of interest.

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