

Since January 2020 Elsevier has created a COVID-19 resource centre with free information in English and Mandarin on the novel coronavirus COVID-19. The COVID-19 resource centre is hosted on Elsevier Connect, the company's public news and information website.

Elsevier hereby grants permission to make all its COVID-19-related research that is available on the COVID-19 resource centre - including this research content - immediately available in PubMed Central and other publicly funded repositories, such as the WHO COVID database with rights for unrestricted research re-use and analyses in any form or by any means with acknowledgement of the original source. These permissions are granted for free by Elsevier for as long as the COVID-19 resource centre remains active.



Contents lists available at ScienceDirect

Journal of Infection

journal homepage: www.elsevier.com/locate/jinf



Letter to the Editor

Influenza and anosmia: Important prediction factors for severity and death of COVID-19



Dear Editor,

At October 19 2020, 40 million patients had been infected with COVID-19 worldwide, and about 1.1 million had died from the disease. This virus belongs to the same coronavirus family as the MERS virus that circulated in 2015, but it is much more infectious, and the world is currently experiencing a pandemic. However, the factors affecting disease severity and mortality have not yet been clearly identified. The machine learning (ML) algorithm is a model suitable for the medical field because it has a fairly accurate prediction capability for large-scale new, never-seen-before inputs such as COVID-19 pandemic. In this paper, we have analyzed the factors affecting the severity and mortality of 8070 COVID-19 patients registered in the National Health Insurance Service (NHIS) of South Korea using ML algorithms. (NHIS-2020-1-479)

The severity of COVID-19 was defined as the end result with one of following conditions. (1) Intensive care unit (ICU) care; (2) Extracorporeal membrane oxygenation (ECMO) treatment; (3) Mechanical ventilator care; (4) Oxygen supply. The mortality of COVID-19 was also checked because the NHIS data was connected to the Korea Disease Control and Prevention Agency and Statistics Korea, which has the mortality data.

A total of 21 diseases (Hypertension (HTN), Diabetes mellitus (DM), Influenza, Cancer, Pulmonary disease, Angiotensin Converting Enzyme or Angiotensin Receptor Blocker (ARB) among hypertensive patients, Gastroesophageal reflux disease (GERD), Acute sinusitis (A_sinusitis), Chronic sinusitis (C_sinusitis), Osteoporosis, Cardiovascular disease (CVD), Angina, Peripheral vascular disease (PVD), Congestive heart failure (CHF), Depression, Rheumatologic disease (RA), Hepatitis, Myocardial infarction (MI), Inflammatory bowel disease (IBD), Non-tuberculosis mycobacterium (NTM), olfactory loss (Anosmia)) were chosen as the underlying diseases in the 8070 COVID-19 patients. NHIS-customized data for the past 5 years were selected for the patients confirmed with COVID-19, and hospital use records for the past 5 years were used to identify the following inclusion criteria.

A total of 8070 COVID-19 confirmed patients were included in this study. (Fig. 1A) Their average age was 39.9 years (SD: 19.7 years), 3236 (40.1%) males and 4834 (59.9%) females. Of the 785 patients classified as severe, 374 were men and 411 were women (p<0.001). The mean age of severely ill patients was 61.6 years (SD 16.0 years). There were a total of 248 patients who died. Among the patients who died, 136 were male and 112 were female (p = 0.0008). The average age of the patients who died was 72.1 years (10.2 years) (Fig. 1B).

Regarding the underlying diseases in COVID-19 patients, 4572 patients had a history of pulmonary disease, 674 patients with

influenza, 231 patients with ARB, and 77 patients with anosmia (Fig. 1C).

Model selection was made by comparing area under the ROC curve (AUC) values for each model. Among the various models, the model with the best prediction of severity was the neural network with an AUC value of 85.06%, followed by logistic regression elastic net (EN) (84.74%) (Fig. 1D). The most important variable for predicting severity in the neural network model was a history of influenza (relative importance: 0.083). (Fig. 1F, Table 1).

The model with the best prediction of death was the logistic regression EN model with an AUC value of 93.89%, followed by the logistic regression lasso model (93.84%), the neural network model (93.73%) (Fig. 1G). The most important variables for mortality in the EN model were age (coefficient: 2.136) and anosmia (coefficient: -1.438) (Fig. 1I, Table 1).

We analyzed 24 factors affecting severity and mortality in 8070 patients using a novel ML algorithm that has recently emerged. Foremost, influenza history was a very important variable in terms of COVID-19 severity (neural network 1st, ridge 6th) and mortality (EN 5th, lasso 3rd, ridge 5th). (Fig. 1I, Table 1) It has been reported that oseltamivir cannot prevent worsening of symptoms and disease in patients with COVID-19 as different molecular docking sites have been found in vitro and retrospective studies in COVID-19.7 Among recent papers, it has been reported that influenza vaccination can alleviate the risk of death in a pandemic situation caused by COVID-19.⁴ Since the symptoms of influenza and COVID-19 are similar, it can be confusing which disease is present, so vaccination can be important in preventing the twindemic of COVID-19 and influenza co-infection. In this paper, we studied the history of influenza and the severity of COVID-19. A history of influenza can sometimes cause pulmonary fibrosis, a common sequelae of virusinduced pneumonia, and this complication is estimated to cause increased severity and mortality of COVID-19 infection. These results are in line with the current policy recommending influenza virus vaccination, mainly considering the current COVID-19 epidemic and the prevalence of influenza during the period from autumn to spring.

Anosmia was also identified as an important variable in predicting the severity of COVID-19. The best predictive models for mortality were the EN and lasso models, and the second most important variable in both these models was anosmia. This means that the mortality rate was low in patients with olfactory loss after the COVID-19 diagnosis. There are papers which indicate that recent olfactory loss in mild to moderate COVID-19 patients is an important factor that differentiates COVID-19 from other infectious disease, and in most cases, the sense of smell recovers well.^{5,6} Another paper reports that anosmia is associated with lower inhospital mortality in COVID-19, which is in line with our research results.⁷ The novel finding in our study is that anosmia will con-

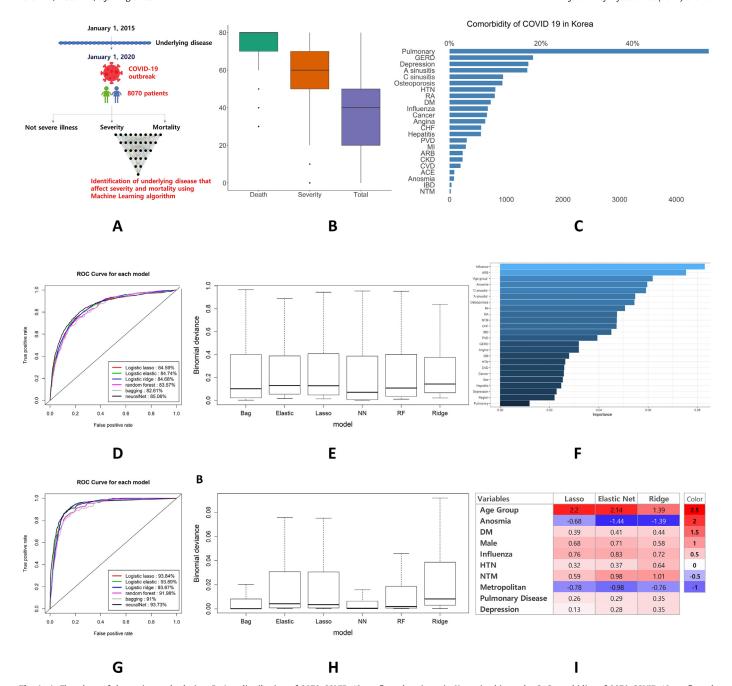


Fig. 1. A. Flowchart of the entire study design. B. Age distribution of 8070 COVID-19 confirmed patients in Korea in this study. C. Comorbidity of 8070 COVID-19 confirmed patients in Korea in this study. D. ROC curves and AUC values in the prediction of severity of COVID-19. E. BD in the prediction of severity of COVID-19. F. Variable importance of the neural network model in the prediction of severity of COVID-19. G. ROC curves and AUC values in the prediction of mortality of COVID-19. H. BD in the prediction of mortality of COVID-19. I. Coefficient Heatmap of the three logistic model in the prediction of mortality of COVID-19.

tinue to be an indicator that should be carefully examined in COVID-19 infection.⁸

Influenza was found to be a major adverse factor in COVID-19 in addition to the factors of old age and male sex, and which are already known to be related to disease severity and mortality. In addition, anosmia was found to be a major factor associated with lower severity and mortality rates. Therefore, in the current situation where there is no adequate COVID-19 treatment at present, examining the history of influenza vaccination and anosmia in addition to age and sex will be important indicators for predicting the severity and mortality of COVID-19 patients.

Abbreviations: (Receiver Operating Characteristic (ROC), Area Under the Curve (AUC), Binomial Deviances (BD), Hypertension (HTN), Diabetes mellitus (DM), Influenza, Cancer, Pulmonary disease, Angiotensin Converting Enzyme or Angiotensin Receptor Blocker (ARB) among hypertensive patients, Gastroesophageal reflux disease (GERD), Acute sinusitis (A_sinusitis), Chronic sinusitis (C_sinusitis), Osteoporosis, Cardiovascular disease (CVD), Angina, Peripheral vascular disease (PVD), Congestive heart failure (CHF), Depression, Rheumatologic disease (RA), Hepatitis, Myocardial infarction (MI), Inflammatory bowel disease (IBD), Non-tuberculosis mycobacterium (NTM), olfactory loss (Anosmia))

Outcomes	Model	Measure	Variable importance	Value	Outcomes	Model	Measure	Variable importance	Value
Severity	Lasso	Estimated	Age	1.276	Mortality	Lasso	Estimated	Age	2.203
		coefficient	DM	0.431			coefficient	Metropolitan	-0.783
			Male	0.415			which is not	Influenza	0.763
			Anosmia	-0.379			zero	Anosmia	-0.684
			HTN	0.266				Male	0.682
			ARB	0.222				NTM	0.598
			Influenza	0.211				DM	0.393
			CVD	0.209				HTN	0.322
			Pulmonary	0.135				Pulmonary	0.257
			A_Sinusitis	0.092				PVD	0.243
	Elastic	Estimated	Age	1.203		Elastic	Estimated	Age	2.136
	Liastic	coefficient	DM	0.442		Liastic	coefficient	Anosmia	-1.438
		Coefficient							
			Anosmia	-0.413			which is not	Metropolitan	-0.985
			Male	0.397			zero	NTM	0.980
			HTN	0.309				Influenza	0.830
			CVD	0.235				Male	0.710
			ARB	0.234				DM	0.405
			Influenza	0.222				HTN	0.365
			Pulmonary	0.147				Pulmonary	0.295
			A_Sinuistis	0.091				Depression	0.280
	Ridge	Estimated	Age	1.006		Ridge	Estimated	Age	1.389
		coefficient	Anosmia	-0.838			coefficient	Anosmia	-1.388
		cocincient	DM	0.480			which is not	NTM	1.002
			HTN	0.419			zero	Metropolitan	-0.761
			Male	0.419			2010	Influenza	0.722
			Influenza	0.397				HTN	0.642
			CVD	0.326				Male	0.582
			NTM	0.310				DM	0.442
			ARB	0.301				Pulmonary	0.352
			MI	-0.214				Depression	0.346
	Random	Mean	Age	174.074		Random	Mean	Age	38.970
	Forest	decrease in	HTN	51.519		Forest	decrease in	HTN	8.036
		Gini impurity	DM	36.373			Gini impurity	Male	6.669
			CVD	20.110				DM	6.427
			Osteoporosis	17.828				CVD	5.842
			Male	16.432				PVD	5.094
			Pulmonary	14.944				RA	4.963
			Cancer	14.928				Osteoporosis	4.883
			ARB	14.526				•	4.588
								Cancer	
	ъ .		A_Sinuistis	14.159		ъ .		Pulmonary	4.581
	Bagging	Mean	Age	193.724		Bagging	Mean	Age	40.825
		decrease in	HTN	60.297			decrease in	Male	9.054
		Gini impurity	DM	35.551			Gini impurity	HTN	8.948
			Male	23.011				DM	7.458
			Pulmonary	22.394				CVD	7.336
			Cancer	21.379				Pulmonary	7.193
			Osteoporosis	21.278				Cancer	7.128
			CVD	20.893				RA	7.006
			A_Sinuistis	20.869				Osteoporosis	6.700
			RA	19.921				PVD	6.366
	Neural	Relative	Influenza	0.083		Neural	Relative	CVD	0.076
	Network	importance	ARB	0.085		Network	importance		0.076
	INCLWUIK	importance				INCLWOLK	miportance	Age	
			Age	0.062				Male	0.0659
			Anosmia	0.060				RA	0.062
			C_Sinuistis	0.059				C_Sinuistis	0.053
			A_Sinuistis	0.055				Influenza	0.051
			Osteoporosis	0.054				IBD	0.048
			MI	0.051				PVD	0.045
			RA	0.047				HTN	0.045
			NTM	0.047				Pulmonary	0.044

Author Contributions

Doo Hwan Kim: Contributed to the study design, protocol and study materials, collected study data, provided data access, and helped write the first draft of the manuscript (Methods and Results sections).

Min Gul Kim: Contributed to the study design, protocol, study materials and data analysis, and helped write the first draft of the manuscript (Methods and Results sections).

Seong J. Yang: Designed the statistical plan, assisted with data analysis and interpretation of the data, and helped write the first draft of the manuscript (Methods section).

Eun Jung Lee: Contributed to the study design, protocol, and study materials, and helped write the first draft of the manuscript (Results section).

Sang Woo Yeom: Collected the study data, performed the statistical analysis, and helped write the first draft of the manuscript (Methods section).

Yeon Seok You: Contributed to the study design, protocol and study materials, collected study data.

Jong Seung Kim: Contributed to the study design, protocol and study materials, designed the statistical plan and data analysis, performed the statistical analysis, wrote the first draft of the manuscript

Supplementary material

supplementary.docx

Funding

None

Declaration of Competing Interest

None

Acknowledgments

This paper was supported by a fund of the Biomedical Research Institute at Jeonbuk National University Hospital. We specially thanks to Professor Sam Hyun Kwon for the idea of this manuscript.

Supplementary materials

Supplementary material associated with this article can be found, in the online version, at doi:10.1016/j.jinf.2021.08.024.

References

- 1. https://www.who.int/emergencies/diseases/novel-coronavirus-2019
- Goh GK, Dunker AK, Foster JA, Uversky VN. Rigidity of the outer shell predicted by a protein intrinsic disorder model sheds light on the COVID-19 (Wuhan-2019-nCoV) infectivity. Biomolecules 2020;10.
- 3. Rajkomar A, Dean J, Kohane I. Machine Learning in Medicine. N Engl J Med 2019:380:1347–58.
- Grohskopf LA, Liburd LC, Redfield RR. Addressing influenza vaccination disparities during the COVID-19 pandemic. JAMA 2020;324:1029-30.
- Lee Y, Min P, Lee S, Kim SW. Prevalence and duration of acute loss of smell or taste in COVID-19 patients. J Korean Med Sci 2020;35:e174.
- Baron-Sanchez J, Santiago C, Goizueta-San Martin G, Arca R, Fernandez R. Smell and taste disorders in Spanish patients with mild COVID-19. Neurologia 2020.
- Talavera B, García-Azorín D, Martínez-Pías E, Trigo J, Hernández-Pérez I, Valle-Peñacoba G, et al. Anosmia is associated with lower in-hospital mortality in COVID-19. J Neurol Sci 2020;419:117163.
- Calica Utku A, Budak G, Karabay O, Guclu E, Okan HD, Vatan A. Main symptoms in patients presenting in the COVID-19 period. Scott Med J 2020;65:127–32. doi:10.1177/36933020949253.

Doo Hwan Kim

Director of Big-Data Center, National Health Insurance Service (NHIS), Wonju, Republic of Korea

Min Gul Kim[†]

Department of Pharmacology, Jeonbuk National University, Jeonju, Republic of Korea

Seong J. Yang[†]

Department of Statistics (Institute of Applied Statistics), Jeonbuk National University, Jeonju, Republic of Korea

Eun Jung Lee[†]

Department of Otorhinolaryngology-Head and Neck Surgery, College of Medicine, Jeonbuk National University, Jeonju-si 54907, Republic of Korea

Research Institute of Clinical Medicine of Jeonbuk National University – Biomedical, Research Institute of Jeonbuk National University Hospital, Jeonju-si 54907, Republic of Korea

Sang Woo Yeom[†]

Department of Medical Informatics, College of Medicine, Jeonbuk National University, Jeonju-si 54907, Republic of Korea

Yeon Seok You, Jong Seung Kim*

Department of Otorhinolaryngology-Head and Neck Surgery, College of Medicine, Jeonbuk National University, Jeonju-si 54907, Republic of

Department of Medical Informatics, College of Medicine, Jeonbuk National University, Jeonju-si 54907, Republic of Korea Research Institute of Clinical Medicine of Jeonbuk National University – Biomedical, Research Institute of Jeonbuk National University Hospital, Jeonju-si 54907, Republic of Korea

*Corresponding author at: Department of Otorhinolaryngology-Head and Neck Surgery, College of Medicine, Jeonbuk, National University, Jeonju-si 54907, Republic of Korea.

E-mail address: kjsjdk@gmail.com (J.S. Kim)

† These authors contributed equally.