COVID-19 disease Severity Correlates with Smoking Status

Édouard Lansiaux

Henry Warembourg School of Medicine, Medical Student

Lille University, 59000, Lille, France

edouard.lansiaux.etu@univ-lille.fr

Jean-Laurent Picard, M.Sc

Conservatoire National des Arts et Métiers

75141 Paris, France

jl@pi.cards

Philippe P. PébaÿPh.D.
NexGen Analytics
Sheridan, WY 82801, U.S.A.
philippe.pebay@ng-analytics.com

Joachim Son-Forget, M.D.-Ph.D.

Assemblée Nationale

75355, Paris, France

joachim.son-forget@assemblee-nationale.fr

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ABSTRACT

The novel COVID-19 disease is a contagious acute respiratory infectious disease

whose causative agent has been demonstrated to be a new virus of the

coronavirus family, SARS-CoV-2. Multiple studies have already reported that risk

factors for severe disease include older age and the presence of at least one of

several underlying health conditions. However, a recent physiopathological report

and the French COVID-19 scientific council have postulated a protective effect of

tobacco smoking.

We have been able to demonstrate the statistical significance in this regard of

recent series from both China and in the US, reporting smoking status as well as

disease severity (p-values of 2.27*10⁻³ and 11.7*10⁻¹⁵, respectively).

Subsequently and using a Bayesian approach we have established that disease

severity is positively associated with smoking status. Finally, we refute claims

linking general population smoking status (N=O(10⁸) or O(10⁹)) to much smaller

disease course series (N=O(10³)). The latter point in particular is presented to

stimulate academic discussion, and must be further investigated by well-

designed studies.

Keywords: COVID-19; Coronavirus; Respiratory Distress; Tobacco; Smoking;

Correlation; Statistics; Conditional; Probability; Regression; China; U.S.A.

IMPLICATIONS

Recent preliminary estimates of the prevalence of selected underlying health conditions among patients with COVID-19, in China as well as in the United States, show that individuals with underlying health conditions appear to be at higher risk for more severe COVID-19. Thanks to a non parametric test, we have concluded that the only significant findings that can be confidently asserted from the series discussed in this article, is that COVID-19 disease severity is positively associated with smoking status.

INTRODUCTION

The novel coronavirus pneumonia (COVID-19) is a contagious acute respiratory infectious disease whose causative agent has been demonstrated to be a new virus of the coronavirus family, SARS-CoV-2. This illness was first evinced in December 2019 in the Seafood Market of Wuhan, Hubei Province, in southern China^{1,2}. Patients with the coronavirus pneumonia have a fever, and the temperature above 38 degrees Celsius with symptoms such as dry cough, fatigue, dyspnea, difficulty breathing, and diarrhea¹⁻⁵. There is a high transmission for this disease, which, as result of this and other factors such as international travel and trade, has now turned into a pandemic, with hundred thousands of individuals confirmed to be infected worldwide – and most likely millions of unreported cases⁵.

Recent preliminary estimates of the prevalence of selected underlying health conditions among patients with COVID-19, in China as well as in the United States, show that individuals with underlying health conditions appear to be at higher risk for more severe COVID-19^{3,6}. A recent Greek epidemiological review suggests that smoking is most likely associated with the negative progression and adverse outcomes of COVID-19⁷. Meanwhile, a recent physio-pathology study has formulated the hypothesis that the effectiveness of ACE inhibitors/angiotensin II receptor blockers and nicotine in patients infected with COVID-19⁸.

The smoking prevalence in China and in the U.S. have been previously and broadly documented: these are, respectively, of 26.6%, and 19.7%^{9,10}. Furthermore, the president of the COVID-19 Scientific Council of France recently postulated that nicotine may have a protective role against COVID-19 infection¹¹. Thanks to these multiple studies, we can observe that the prevalence of past/current smokers is higher in COVID-19 severe infected patients. In this context, we would like to know whether the past/current smoking status entails higher probability of a severe course of COVID-19 illness, as compared to having never smoked.

METHODS

Study and Participants

We conducted a descriptive observational cross-sectional study in order to define a hypothetical relationship between smoker status and SARS-CoV-2 severity infection. The source and targeted populations are the whole humanity in view of the ongoing COVID-19 pandemic. The eligible population is constituted by all Chinese and American already included in previous papers.

The study was conducted by a consortium of two data analysts, a MD specialized in radiology, and a medical student in clinical years. NexGen Analytics had no role in designing the study, nor in making the decision to submit manuscript to the publication, nor did it receive any fee or compensation in the context of this work. The first author vouches for the data and analyses, as well as for the fidelity of this report to the study protocol.

Enrollment

The studied and included population gathers patients from two studies: one group of patients studied in China between 2019/12/11 and 2020/01/31 (N = 1,085), and another group studied in the U.S. between 2020/02/12 and 2020/03/28 (N = 6,637)^{3,6}.

Outcome Measures

Our null hypothesis (H_0) was the orthogonality between two variables: the smoking status (X) and the COVID-19 Symptoms Severity (Y). Each variable can take two values: for the smoking status (X), it was a never smoker (X_1) or a current/past smoker (X_2) ; for the COVID-19 Symptoms Severity (Y), it was a mild (non-hospitalised) (Y_1) or a severe (hospitalised, ICU or not) (Y_2) symptoms severity.

Statistical Analysis

Each sample was analysed separately. First, we had to calculate for each sample different conditional probabilities: the probability to develop severe symptoms for non-smokers (i.e., $P(Y_2|X_1)$) and the probability to develop severe symptoms for former and current smokers combined (i.e., $P(Y_2|X_2)$). We then calculated, for each observed sample, the expected cardinalities assuming H_0 expected sample: in each case, we multiplied the two corresponding sums and subsequently divided by the actual total. In this way, we were able to apply a χ^2 test to compare the observed table to the expected table, for each sample. The p-value was subsequently computed in order to evince a statistically significant difference between the observed and the expected tables.

RESULTS

Populations

After a short treatment of the initial data from both Chinese and U.S. series (*Table 1*), in particular to eliminate incomplete data, we have obtained the observed data for each sample with their respective conditional probabilities (*Tables 2, 3*). For the Chinese sample, the probability to develop severe COVID-19 symptoms for non-smokers is of 14.5% and the probability to develop severe COVID-19 symptoms for former/current smokers is of 24.1% For the American sample, the probability to develop severe COVID-19 symptoms for non-smokers is of 21.7% and the probability to develop severe COVID-19 symptoms for former/current smokers is of 42.7%.

Outcomes

We estimated the expected data on each case by multiplying the two corresponding sums and then by dividing by the total effective, for each sample (*Tables 4, 5*). In these conditions, we were able to estimate the χ^2 parameter, for Chinese and American samples, and the p-values where the difference

between observed and expected data are significant (*Table 6*). Indeed, the difference observed between them is significant for China and US with respectively a p-value of 2.27*10⁻³ and 11.7*10⁻¹⁵.

DISCUSSION

Subsequently, by computing the empirical probability distributions conditioned on smoking status, we have established that disease severity is positively associated with smoking status, in those samples. Nevertheless, our study has some limitations:

- 1. The difference between smoking status as observed across the entire Chinese and American populations^{9,10}, and the same observed in our small clinical datasets (*Table 1*) is significant. As a result, the previously claimed relationship between former/current smoking status and a COVID-19 disease severity may not be confidently estimated from a global population.
- 2. Only adults were considered to compute smoking prevalence¹⁰.

However, regarding the first limitation above, we contend that claims linking general population smoking status, computed across populations with cardinalities N in the order of O(10⁸) or even O(10⁹) with broadly varying smoking habits (e.g. already rural vs urban divide in this regard), to much smaller disease course series (N = O(10³)), are at best imprudent and at worst marred by selection bias, especially as the latter regard a fraction of patients from the same urban area. We therefore claim that specific smoking status data, relevant to the concerned geographic areas, is required to evince a statistically significant under-representation of smokers among COVID-19 patients. An alternative argument could be made from a physio-pathological standpoint as well: indeed, previous studies and assertions describe a hypothetical protector role of nicotine against COVID-19 infection⁸, but these are largely speculative at this point and require laboratory confirmation. We therefore conclude that the only significant findings that can be confidently asserted from the series discussed in this article, is that COVID-19 disease severity is positively associated with smoking status.

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The author	's declare n	O CONTILCTS	of interest.

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TABLES

Sample	China		United States				
Sub-sample	All patients	Non severe	Severe	AII patients	Not hospitalized	Hospitalized	ICU
Cardinality	1,085	913	172	6,637	5,143	1,037	457
Never smoke	927	793	134	6,391	5,002	970	419
(%)	(85.4)	(86.9)	(77.9)	(96.3)	(97.2)	(93.5)	(91.7)
Former smoker	21	12	9	132	80	45	33
(%)	(1.9)	(1.3)	(5.2)	(2.0)	(1.6)	(4.3)	(7.2)
Current smoker	137	108	29	88	61	22	5
(%)	(12.6)	(11.8)	(16.9)	(1.7)	(1.2)	(2.2)	(1.1)

Table 1. Reported outcomes concerning smoking status on two samples.

China observed	Y 1	Y ₂	TOTAL
X 1	793	134	927
X ₂	120	38	158
TOTAL	913	172	1,085

Table 2. Reported outcomes from the China sample.

1. X₁: never smoke

2. X₂: current or former smoker

Y : severity of symptoms

1. Y₁: mild (non-hospitalized)

U.S. observed	Y 1	Y ₂	TOTAL
X 1	5,002	1,389	6,391
X ₂	141	105	246
TOTAL	5,143	1,454	6,637

Table 3. Reported outcomes from the U.S. sample.

1. X₁: never smoke

2. X₂: current or former smoker

Y : severity of symptoms

1. Y₁: mild (non-hospitalized)

China expected	Y 1	Y ₂	TOTAL
X 1	780	147	927
X ₂	133	25	158
TOTAL	913	172	1,085

Table 4. Expected outcomes from the China sample.

1. X₁: never smoke

2. X₂: current or former smoker

Y : severity of symptoms

1. Y₁: mild (non-hospitalized)

U.S. expected	Y ₁	Y ₂	TOTAL
X 1	4,952	1,459	6,391
X ₂	191	55	246
TOTAL	5,143	1,454	6,637

Table 5. Expected outcomes from the U.S. sample.

1. X₁: never smoke

2. X₂: current or former smoker

Y : severity of symptoms

1. Y₁: mild (non-hospitalized)

Sample	China	U.S.	
χ²	9.32	59.6	
<i>p</i> -value	2.27*10 ⁻³	11.7 *10 ⁻⁵	

Table 6. χ^2 test applied to both samples.