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## Letter to the Editor

### Co-infection of chlamydia pneumoniae and mycoplasma pneumoniae with SARS-CoV-2 is associated with more severe features



Dear editor,

We read with great interest the recent article published by Lansbury et al. in this Journal. They found that only 7% of hospitalized COVID-19 patients had a bacterial co-infection, increasing to 14% in studies that only included ICU patients, a percentage very low when compared to bacterial coinfections associated with previous influenza pandemics<sup>1</sup>. The Authors underline that testing for co-infecting pathogens during the course of a pandemic is important but most of the studies they screened did not report on this.

Then, few works reported coinfection with atypical pathogens such as *Mycoplasma pneumoniae*<sup>2–7</sup> and *Chlamydia pneumoniae*<sup>8</sup>, even if previous studies on severe coronavirus infections have shown a serological evidence among SARS patients that the incidence of acute or recent *Chlamydia pneumoniae* and *Mycoplasma pneumoniae* infection was about 30% and 9%, respectively<sup>9</sup>.

Here, we present a retrospective study with the aim to evaluate the prevalence of co-infections with atypical pathogens in SARS-CoV-2 infected patients compared to non-infected patients.

We included 721 hospitalized patients who underwent testing for SARS-CoV-2, *Mycoplasma pneumoniae*, *Chlamydia pneumoniae* and *Legionella pneumophyla* over a two months period between 6 March 2020 and 12 May 2020 to the Spedali Civili's Hospital, Montichiari's Hospital and Gardone Val-Trompia's Hospital, Brescia, Italy. A real time reverse polymerase chain reaction was used to detect SARS-CoV-2. Chemiluminescence immunoassays and immunoenzymatic assays were used to detect IgM antibodies for *Mycoplasma pneumoniae* and IgM/IgA/IgG for *Chlamydia pneumoniae* in the serum of patients. Legionella diagnosis was made by urinary antigen testing.

Definition of pneumonia or severe pneumonia was done according to the WHO guidelines and included clinical signs of pneumonia (fever, cough, dyspnea, fast breathing).

Categorical variables are the number (percentage), and continuous variables are the median (interquartile range [IQR]). Categorical data were compared by using the  $\chi^2$  test or the Fisher exact test, as appropriate. The data in different groups were compared with the ANOVA or independent *t*-test for normally distributed variables.

Logistic regression was used in order to perform univariate and multivariate analysis for determination of risk factors [odds ratios (ORs) and 95% CIs] associated with the presence of *Chlamydia pneumoniae* and *Mycoplasma pneumoniae* infection in SARS-CoV-2 positive patients after adjusting for confounders.

Multivariate logistic regression was used to account for differences in-hospital mortality for *Chlamydia pneumoniae* and *My-*

*coplasma pneumoniae* infection in SARS-CoV-2 positive patients. A *P* value <0.05 was considered statistically significant.

Out of 721 subjects, 443 individuals were infected with SARS-CoV-2 and 278 subjects were negative. All individuals were negative for Legionella antigen. Of the 443 patients with a SARS-CoV-2 infection, 242 individuals had an antibody positivity against *Mycoplasma* and/or *Chlamydia*, while of the 278 negative for SARS-CoV-2, only 97 showed *Mycoplasma* and/or *Chlamydia pneumoniae* antibody positivity ( $p < 0.0001$ ). Of the 242 seropositive subjects, 179 individuals were positive only for *Chlamydia pneumoniae* and 63 exhibited a positive result both for *Chlamydia* and for *Mycoplasma*. Our analyses were performed on the total of patients seropositive for both *Chlamydia* and *Mycoplasma pneumoniae* without distinction between the two pathogens.

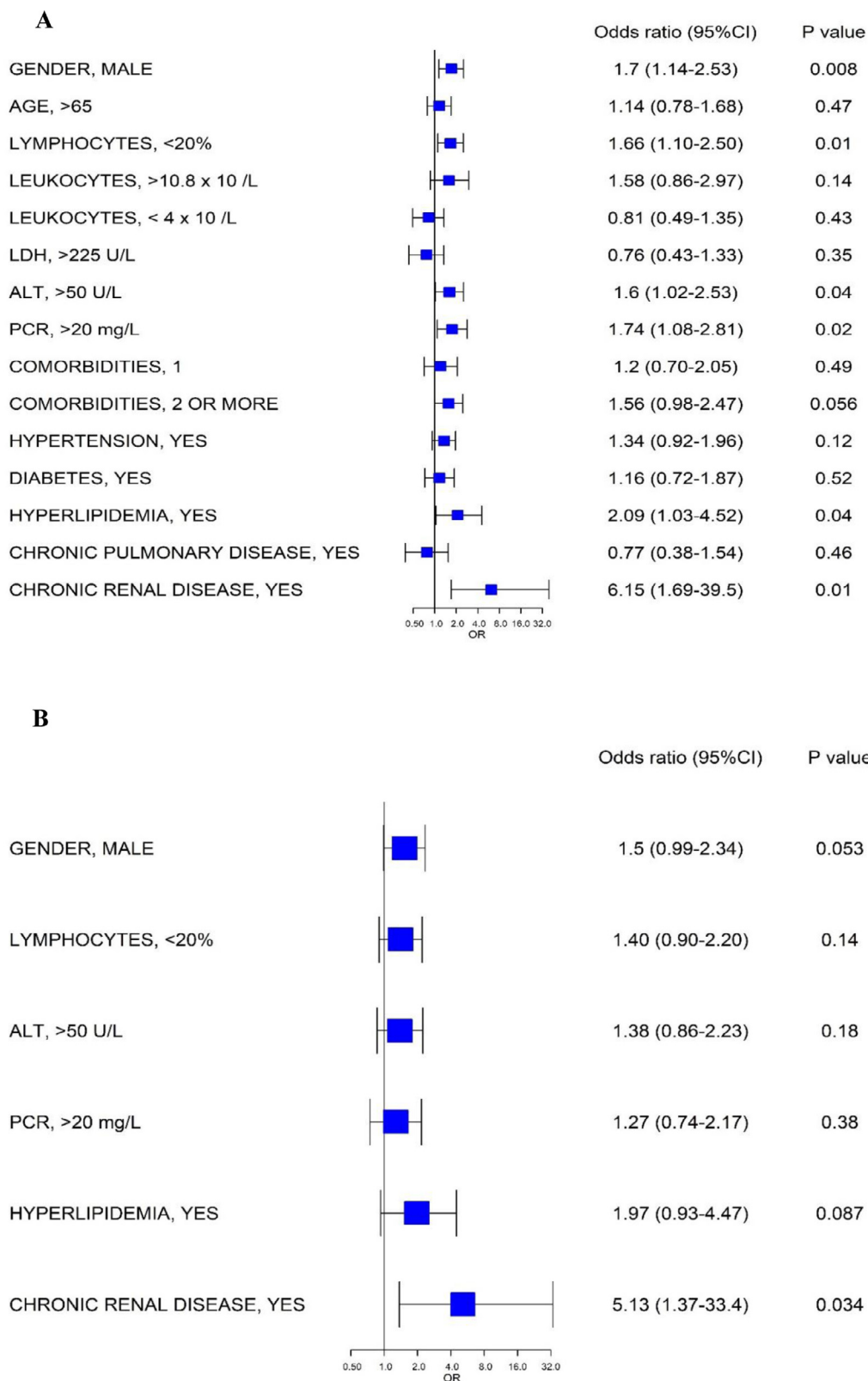
Demographic and clinical characteristics of the patients on admission are listed in Table 1. The median age was 70 (IQR, 20), 294 (66.3%) were male and 149 (34%) were female. Most of them (339, 76.5%) had one or more pre-existing comorbidities, mostly hypertension (224, 50.5%), cardiovascular disease (11, 25%), diabetes (83, 18.7%) and hyperlipidemia (32, 7.2%). The most common signs were fever (338, 76.2%), cough (223, 50.3%) and dyspnea (186, 41.9%).

Compared with those without atypical pathogen antibody positivity, patients seropositive for *Mycoplasma* and/or *Chlamydia* had a larger proportion of male, had dyspnea (47.9% vs 34.8%,  $p = 0.005$ ) and were more likely to have more than three comorbidities ( $p = 0.04$ ), in particular chronic renal disease (5.3% vs 1.4%,  $p = 0.03$ ) and hyperlipidemia (10.3% vs 3.4%,  $p = 0.005$ ) (Table 1).

Patients with SARS-CoV-2 and antibody positivity for *Mycoplasma* and/or *Chlamydia* had a higher median leukocyte count (median 6.6 [IQR, 3.6]  $\times 10^9/L$  vs median 5.7 [IQR, 3.8]  $\times 10^9/L$ ,  $p = 0.01$ ) and a lower level of lymphocytes (71.9% vs 62%,  $p = 0.003$ ) (Table S1). An increased ALT serum level (29.1% vs 19.8%,  $p = 0.02$ ) and also a higher median serum level (median 33 [IQR, 31] U/L vs 26 [IQR, 26] U/L,  $p = 0.02$ ) was found in a higher percentage of patients with SARS-CoV-2 and antibody positivity for *Mycoplasma* and/or *Chlamydia pneumoniae* (Table S1). Higher median serum level of PCR was observed in patients with SARS-CoV-2 and antibody positivity for *Mycoplasma* and/or *Chlamydia pneumoniae* (median 100.4 [IQR, 108.5] mg/L vs 64.45 [IQR, 89] mg/L,  $p = 0.006$ ).

More than three quarters of patients received antibiotic treatment and more than half patients (266, 60%) received antiviral treatment. The percentage of patients receiving azithromycin was significantly higher in those with SARS-CoV-2 and antibody positivity for *Mycoplasma* and/or *Chlamydia pneumoniae* (57% vs 46.2%,  $p = 0.02$ ) (Table S2).

The proportions of critical COVID-19 patients with atypical pathogens coinfection were higher than those of patients infected only with SARS-CoV-2 (13.2% vs 5.9%,  $p = 0.01$ ). Furthermore, requirement and use of a nasal cannula, high flow oxygen support and non-invasive ventilation was significantly higher in co-



**Fig. 1.** A) Univariate logistic regression showing the categorical variables associated with *Chlamydia* and *Mycoplasma pneumoniae* antibody positivity in SARS-CoV-2 infected patients, B) Factors showing significantly independent association with *Chlamydia* and *Mycoplasma pneumoniae* antibody positivity in SARS-CoV-2 infected patients. Odds ratio, 95%CI and P values are obtained from logistic regression modeling.

**Table 1**  
Clinical characteristics of COVID 19 patients according to their *Chlamydia* and *Mycoplasma* serology status.

	Chlamydia and Mycoplasma serology status		P value
	Positive (n = 242)	Negative (n = 201)	
Age, mean (IQR)	71 (19)	68 (20)	0.11
Age group, n°/total (%)			
0–20	0	1 (0.4)	/
21–40	5 (2)	9 (4.4)	0.2
41–60	58 (23.9)	51 (25.3)	0.8
61–80	127 (52.4)	100 (49.7)	0.6
>80	52 (21.4)	40 (19.9)	0.7
Gender, n°/total (%)			
Male	173 (71.4)	121 (60.1)	0.01
Number of comorbidities, n (%)			
None	50 (20.6)	54 (26.8)	0.15
1 or 2	99 (40.9)	89 (44.2)	0.53
≥3	93 (38.4)	58 (28.8)	0.04
Comorbidities			
Diabetes	49 (20.2)	34 (16.9)	0.4
Hypertension	130 (53.7)	94 (46.7)	0.15
Cardiovascular disease	69 (28.5)	42 (20.8)	0.08
Chronic pulmonary disease	17 (7)	18 (8.9)	0.5
Chronic kidney disease	13 (5.3)	3 (1.4)	0.03
Hyperlipidemia	25 (10.3)	7 (3.4)	0.005
Malignancy	18 (7)	16 (7.9)	0.87
Hematologic disease	8 (3.3)	8 (3.4)	0.9
Neurologic disease	11 (4.5)	5 (2.4)	0.3
Symptoms, n (%)			
Fever (temperature >37.5 °C)	174 (71.9)	164 (81.5)	0.02
Cough	127 (52.4)	96 (47.7)	0.37
Dyspnea	116 (47.9)	70 (34.8)	0.005
Chest pain	4 (1.6)	3 (1.4)	0.80
Nausea	2 (0.82)	3 (1.4)	0.83
Vomiting	13 (5.3)	11 (5.4)	0.86
Diarrhea	19 (7.8)	21 (10.4)	0.43
Myalgia	15 (6.1)	10 (4.9)	0.72
Loss of smell and taste	2 (0.82)	0	/

infected patients than in only SARS-CoV-2 positive patients (18.1% vs 3.6%,  $p < 0.0001$ ; 45% vs 23.3%,  $p < 0.0001$ ; 14.7% vs 4.6%,  $p = 0.001$ , respectively) (Table S2).

There were significantly more patients need mechanical ventilation support in SARS-CoV-2 and antibody positivity for *Mycoplasma* and/or *Chlamydia pneumoniae* than in the other group of patients (6.8% vs 2.5%,  $p = 0.04$ ) (Table S2). Furthermore, seropositive patients for *Mycoplasma* and/or *Chlamydia* were more likely to develop more complications (8.3% vs 2.5%,  $p = 0.01$ ), including acute respiratory distress (8.6% vs 3.4%,  $p = 0.03$ ), renal disease (2.8% vs 0.9%,  $p = 0.1$ ), pulmonary embolism and thrombotic events (2.8% vs 1.9%,  $p = 0.7$ ) and acute cardiac injury (2.4% vs 0,  $p = 0.03$ ).

Patients with SARS-CoV-2 and antibody positivity for *Mycoplasma* and/or *Chlamydia pneumoniae* had a slight higher fatality rate compared to the other group of patients, even if not statistically significant (24.2% vs 21.8%,  $p = 0.63$ ). However, multivariate logistic regression showed that age >65 years was an inde-

pendent risk factor for death in patients with SARS-CoV-2 infection and seropositive for *Mycoplasma* and/or *Chlamydia pneumoniae* (OR=4.02, 95%CI=1.7–11,  $p = 0.003$ ).

SARS-CoV-2 infected patients were associated with an increased probability of both *Chlamydia/Mycoplasma pneumoniae* positivity by serology (OR=2.2, 95%CI=1.6–3.0,  $p < 0.0001$ ). A logistic regression for univariate and multivariate analyses was performed in order to identify risk factors associated with *Chlamydia/Mycoplasma* infection in SARS-CoV-2 positive patients.

Six categorical variables were identified in univariate logistic regression analysis, namely: male sex, lymphopenia, ALT >50 U/L, PCR >20 mg/L, hyperlipidemia and chronic renal disease (Fig. 1A). Only chronic renal disease was demonstrated as independent risk factor based on the multivariate logistic regression model (Fig. 1B).

To the best of our knowledge, this study was the first in Europe to report a high rate of detection of antibody positivity for these atypical pathogens in SARS-CoV-2 infected patients, with 26% of the patients positive for *Mycoplasma* IgM, 18% of the patients positive for *Chlamydia* IgM and 81% of the patients positive for both *Chlamydia* IgA and IgG.

So far, in the management of COVID-19 patients, we have to keep in mind the possibility of the presence of other respiratory pathogens causing coinfections. This is important to guide the clinicians for the use of targeted therapies, aimed at treating this potential micro-organisms in order to avoid more severe outcomes of patients.

#### Declaration of Competing Interest

All the authors declare no competing interest.

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#### Ethical statement

The study was conducted in accordance with the Declaration of Helsinki and national standards, and was approved by the Brescia ethical committee (EvoSCO NP 4369).

#### Author contributions

M.A.D.F. and C. P. conceived and designed the study. M.A.D.F. analyzed the data and wrote the first draft of the manuscript. F.C., C.B., P.P. S.F., F.C. (Francesca Caccuri), V.C, and L.M. performed and analyzed microbiological tests and participated in manuscript revision. S.P. analyzed the data and performed statistical analyses. D.R., P.M., M.S., L.M., F.A., F.S., A.P. (Andrea Pilotto), A. P. (Alessandro Padovani), M.B., R.C., C.R., M.C., M.B., G.S., G.M., M.R., S.B., E.F., L.T., F.C. (Francesco Castelli), A.R., R.I., and M.M. were directly involved in the patient care, collected data and participated in manuscript revision. A.C. provided study oversight and participated in manuscript revision. All the Authors approved manuscript submission

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## Supplementary material

Supplementary material associated with this article can be found, in the online version, at doi:<https://doi.org/10.1016/j.jinf.2021.01.009>.

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