





Subphenotypes of SARS-CoV-2-Associated ARDS Overlap Each Other: A Retrospective Analysis

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Abstract

Background Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) virus-associated pneumonia and acute respiratory distress syndrome (ARDS) were often associated with hyperinflammation and elevation of several serum inflammatory markers but usually less than what is observed in non-coronavirus disease (COVID) ARDS. Elevated inflammatory markers such as C-reactive protein, interleukin (IL)-6, etc., are associated with severe infection. This study identified subphenotypes of COVID-19 ARDS patients by latent profile analysis in a cohort of Indian patients.

Methods Data of n = 233 adult Indian patients with laboratory-confirmed SARS-CoV-2 infection admitted to a tertiary care teaching hospital were analyzed in this retrospective study. Only patients with acute respiratory failure (defined by partial pressure of oxygen/fraction of inspired oxygen ratio < 200 mm Hg) and chest X-ray showing bilateral infiltrates were included.

Results The patients' mean (standard deviation) age was 53.3 (14.9) years, and 62% were male. A two subphenotypic model was formulated based on the lowest Bayesian information criterion. Neutrophil-to-lymphocyte ratio and serum IL-6 were latent variables in that model (entropy 0.91). The second phenotype (hyperinflammatory) had lower platelet count (p = 0.02), higher serum creatinine (p = 0.004), higher Greactive protein (p = 0.001), higher ferritin (p < 0.001), and serum lactate dehydrogenase (p = 0.009). Age-adjusted hospital mortality (p = 0.007), duration of hospital stay (p < 0.001), and duration of intensive care unit stay (p < 0.001) were significantly higher in the second subphenotype.

Conclusion Two distinct but overlapping subphenotypes were identified in SARS-CoV-2associated respiratory failure. Hyperinflammatory subphenotype was associated with significantly poor short-term outcomes.

Keywords

- ► ARDS
- ► COVID-19
- ► SARS-CoV-2
- ► latent class

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Introduction

Severe infection from the severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) virus can lead to pneumonia and acute respiratory distress syndrome (ARDS). Initial data from China reported that ARDS was present in nearly 42% of patients with coronavirus disease 2019 (COVID-19) pneumonia and carried mortality over 50%.² SARS-CoV-2 virusassociated pneumonia and ARDS were often associated with hyperinflammation and elevation of several serum inflammatory markers but usually less than what is observed in non-COVID ARDS.³ Elevated inflammatory markers such as C-reactive protein (CRP), interleukin (IL)-6, etc., were associated with severe infection.4

Despite an objective case definition,⁵ ARDS in intensive care units (ICUs) are often underrecognized.⁶ Significant clinical heterogeneity in ARDS is well defined as ARDS can be triggered by a myriad of pulmonary and extrapulmonary causes. The severity of ARDS is also variable and classically defined by the degree of hypoxemia, which ranges from mild to severe. Previous studies have found the existence of two distinct subphenotypes in non-COVID ARDS patients.8 The hyperinflammatory subphenotype of ARDS was associated with elevated inflammatory biomarkers, a higher requirement of vasopressors, and worse clinical outcomes.8 Identification of subtypes of ARDS is clinically significant as the majority of the successful experimental therapies were ultimately found to be ineffective in the general ARDS population. We hypothesize that severe COVID-19 infection-associated ARDS might have several latent subphenotypes and carry a different prognosis.

Methods

Nucleic acid-based laboratory test-confirmed SARS-CoV-2 infected adult (age > 18 years) patients with acute respiratory failure (defined by partial pressure of oxygen [PaO₂]/ fraction of inspired oxygen [FiO₂] ratio < 200 mm Hg) and chest X-ray showing bilateral infiltrates, admitted in an ICU of a tertiary care teaching hospital, New Delhi, India, were recruited in this study. The data collection period was between May 2020 and October 2020. Strict adherence to the Berlin definition of ARDS⁵ was not technically feasible as several patients were managed by oxygen therapy through a high-flow nasal cannula. No formal sample size estimation was performed for this study, and approximately 200 to 250 patients' data were planned to be included in this analysis.

Patients' demographic (age, sex, height, weight), clinical conditions, and preexisting comorbid illnesses were noted. In addition, patients' baseline laboratory parameters that included hematological and biochemical investigations were noted, and all investigations available within the first 24 hours of ICU admission were considered as "baseline." The respiratory support requirement at the time of ICU admission and clinical outcome (survival or death) at the time of hospital discharge was also noted for all patients.

Results

Data of n = 233 patients with laboratory-confirmed SARS-CoV-2 infection leading to acute hypoxemic respiratory failure (PaO₂/FiO₂ ratio < 200 mm Hg) were included in this study. The mean (standard deviation) age of the patients was 53.3 (15) years, and 165 of all patients (proportion [95% confidence [CI]] 70.8 [64.7-76.3]) survived hospital discharge. Sixty-two percent of all patients were male. The median (interquartile [IQR]) PaO₂/FiO₂ ratio was 150 (135-195) mm Hg. Median (IQR) duration of hospital stay and ICU stay were 11 (9-14) and 8 (6-11) days, respectively.

Initially, a two subphenotypic model was constructed with the available inflammatory markers (neutrophil-to-lymphocyte ratio [NLR], lactate dehydrogenase, IL-6, serum ferritin, and CRP). However, the model constructed by IL-6 and NLR had the lowest Akaike information criterion (AIC) (AIC 4,735, entropy 0.90, bootstrap likelihood ratio test p = 0.01, rightarrow Fig. 1). A comparison of baseline variables between subphenotype 1 and subphenotype 2 has been depicted in ►Table 1. Patients belonging to the subphenotype 2 had a lower baseline PaO₂/ FiO₂ ratio, higher CRP, ferritin, and creatinine, and lower platelet count. Male preponderance was also seen in subphenotype 2. However, baseline hemoglobin, total leukocyte count, and international normalized ratio were similar in both the subphenotypes. Unadjusted analyses revealed that the requirement of mechanical ventilation (odds ratio [OR] [95% CI] 6.4 [3.4–11.8], p < 0.001) and hospital mortality (OR [95% CI] 6.2 [3.3–11.6], p < 0.001) were higher in the subphenotype 2. In

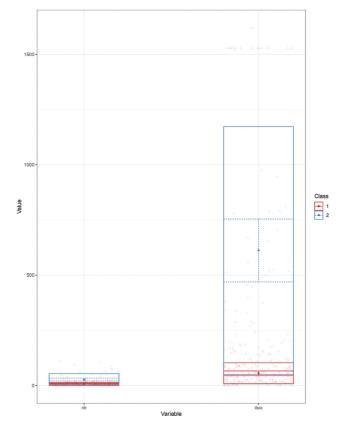


Fig. 1 Profile plot showing distribution of neutrophil-to-lymphocyte ratio and interleukin (IL)-6 in both subphenotypes.

Variable	All Patients	Subphenotype 1	Subphenotype 2	Significance
Age	55 (45–65)	55 (46–64)	56.5 (44.3–66.5)	p = 0.44
Sex (male/female)	145/88	100/67	45/21	p = 0.24
Hemoglobin	11.6 (9.1–13.5)	11.8 (9.3–13.5)	11.4 (7.9–13.1)	p = 0.14
TLC	10300 (6700–15500)	9900 (6710–15300)	10590 (6675–17775)	p = 0.44
Platelet	193 (130–270)	209 (135–289)	172 (101–224)	p = 0.005
INR	1.18 (1.08–1.3)	1.17 (1.1–1.23)	1.2 (1–1.5)	p = 0.42
Creatinine	1.2 (0.8–2.2)	1.1 (0.8–1.9)	1.6 (1.04–3.1)	p = 0.004
LDH	585 (406–808)	542 (404–739)	690 (445–937)	p = 0.009
Ferritin	716 (360–1385)	585 (329–1130)	975 (574–2000)	p < 0.001
Bilirubin	0.6 (0.4-0.95)	0.6 (0.4-0.9)	0.63 (0.5–1)	p = 0.24
PaO ₂ /FiO ₂ ratio	142 (110–174)	156 (129–177)	110 (87–155)	p < 0.001

Table 1 Comparison of baseline variables between two subphenotypes

Abbreviations: FiO₂, fraction of inspired oxygen; INR, international normalized ratio; LDH, lactate dehydrogenase; PaO₂, partial pressure of oxygen; TLC, total leukocyte count.

addition, duration of hospital stay (median [IQR] 11 [9–13] vs. 14 [11–17] days, p < 0.001) and duration of ICU stay (median [IQR] 7 [6–9] vs. 11.5 [8–14] days, p < 0.001) were significantly higher in subphenotype 2.

After adjustment of age and baseline PaO_2/FiO_2 ratio, subphenotype 2 had significantly higher mortality (OR [95% CI] 3.1 (1.4–6.9], p=0.007) and requirement of mechanical ventilation (OR [95% CI] 3 [1.1–8.1], p=0.029). In addition, duration of hospital stay (t=3.51, p>0.001) and duration of ICU stay (t=4.12, p<0.001) were also significantly higher in subphenotype 2, even after adjustment of age and baseline PaO_2/FiO_2 .

Discussion

This study identified two subphenotypes of COVID-19 infection-associated ARDS and subphenotype 2, which were associated with "hyperinflammation" and poor clinical outcomes.

Classically, ARDS is a heterogeneous disorder, and there are significant pathophysiological variations between pulmonary and extrapulmonary ARDS. Subphenotypes of ARDS were identified from secondary analyses of various randomized controlled trials. Calfee et al identified two subphenotypes of ARDS; the hyperinflammatory subphenotype was associated with the elevation of inflammatory biomarkers, acidosis, and sepsis. Clinical outcome was also worse in that subphenotype. Our findings are also quite similar to the previous data as we were also able to identify two subphenotypes, and poor clinical outcome was evident with hyperinflammation. Similar subphenotypes were identified in COVID-19 ARDS patients also. 10

Identification of subphenotypes in COVID-19 pneumonia/ARDS patients has several clinical implications. As patients with elevated inflammatory biomarkers have a worse clinical outcome and higher requirements for

mechanical ventilation, they should be managed in a wellequipped center. During the time of the pandemic, the categorization of the patients is important for proper resource utilization.

Identification of subphenotypes of COVID-19 ARDS has several therapeutic implications. First, future randomized controlled trials can be designed for a particular subphenotype to avoid heterogeneity in the clinical status of the participants. Apart from this, anti-inflammatory drugs are the mainstay of treatment in COVID-19 pneumonia patients, and they may be more useful in patients with a hyperinflammatory state.

Limitations

Our study has several limitations. Most importantly, the sample size of our study was limited and strict adherence to the Berlin definition was not practically feasible. Furthermore, we conducted this study during the so-called "first wave" of the COVID-19 pandemic in New Delhi. As the COVID-19 pandemic is evolving, variants of the SARS-COV-2 virus also evolve, and disease severity also changes. Hence, there is some probability that the study findings may not apply to other geographical locations or when other virus variants are causative agents.

Conclusion

Two distinct but overlapping subphenotypes were identified in SARS-CoV-2-associated respiratory failure. In addition, the hyperinflammatory subphenotype was associated with a significantly poor short-term outcome even after adjustment of baseline disease severity.

Note

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Conflict of Interest None declared.

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