The mNCP-SPI Score Predicting Risk of Severe COVID-19 among Mild-Pneumonia Patients on Admission

Purpose: To predict the risk of developing severe pneumonia among mild novel coronavirus pneumonia (mNCP) patients on admission.

Methods: A retrospective cohort study was conducted at three hospitals in Shanghai and Wuhan from January 2020 to February 2020. Real-time polymerase-chain-reaction assays were used to detect COVID-19. A total of 529 patients diagnosed with NCP were recruited from three hospitals and classified by four severity types during hospitalization following the standards of the Chinese Diagnosis and Treatment of Pneumonia Caused by New Coronavirus Infection (eighth version). Patients were excluded if admitted by ICU on admission (n=92), on a general ward while meeting the condition of severe or critical type on admission (n=25), or there was insufficient clinical information (n=64). In sum, 348 patients with mNCP were finally included, and 68 developed severe pneumonia.

Results: mNCP severity prognostic index values were calculated based on multivariate logistic regression: history of diabetes (OR 2.064, 95% CI 1.010–4.683; p=0.043), time from symptom onset to admission ≥7 days (OR 1.945, 95% CI 1.054–3.587; p=0.033), lymphocyte count ≤0.8 (OR 1.816, 95% CI 1.008–3.274; p=0.047), myoglobin ≥90 mg/L (OR 2.496, 95% CI 1.235–5.047; p=0.011), and D-dimer ≥0.5 mg/L (OR 2.740, 95% CI 1.395–5.380; p=0.003). This model showed a c-statistic of 0.747, with sensitivity and specificity 0.764 and 0.644, respectively, under cutoff of 165.

Conclusion: We designed a clinical predictive tool for risk of severe pneumonia among mNCP patients to provide guidance for medicines. Further studies are required for external validation.

Keywords: novel coronavirus pneumonia, predicting score, severe pneumonia

Introduction

On December 8, 2019, a cluster of cases of pneumonia emerged in Wuhan, Hubei province, China.1,2 High-throughput sequencing of isolates obtained from patients with pneumonia identified a novel coronavirus (COVID-19) as the etiology,3,4 which resembled severe acute respiratory syndrome coronavirus (SARS-CoV).3,5,6 As of February 22, 2020, a total of 76,395 patients had been confirmed to be affected with COVID-19 in China. Significantly, it is estimated that about 17% of patients have developed severe pneumonia, 5%–32% admitted to intensive care units (ICUs), and 1.36%–15% succumbed, according to different reports.1,7 A variety of studies have shown that some clinical indicators have prognostic value in novel CoV pneumonia (NCP),1,8,9 and several predictive scores, such as the
pneumonia severity index,\textsuperscript{10} CURB-65,\textsuperscript{11} and some ICU scoring systems, including Acute Physiology and Chronic Health Evaluation II score\textsuperscript{12} and Sequential Organ-Failure Assessment,\textsuperscript{13} were existed to assess the severity of pneumonia. However, none of these indices has been reported to predict the prospective severe exacerbations of NCP. Due to the seriousness of the epidemic, it is emergent to search for an early-warning model for severe NCP to assist clinicians in deciding on optimal therapeutic interventions. Herein, we unravel potential risk factors and develop a simple and practical score to predict disease-progression probability of mild NCP (mNCP) patients that can accurately stratify hospitalized patients with COVID-19 into relevant risk categories and provide guidance to make further clinical decisions.

**Methods**

**Study Design and Population**

This was a retrospective multicenter observational cohort study conducted from January 2020 to February 2020 in Ruijin Hospital (affiliated with Shanghai Jiao Tong University School of Medicine), Shanghai Public Health Clinical Center in Shanghai, and Jinyintan Hospital in Wuhan, China. The latter two are special designated local hospitals for NCP patients. The study was approved by the Shanghai Public Health Clinical Center Ethics Committee, Ruijin Hospital Ethics Committee, and Jinyintan Hospital Ethics Committee, respectively, while consent was obtained from the participants in our study.

All patients who had been diagnosed with NCP were consecutively enrolled from the three hospitals following WHO guidelines. Patients were divided into four types during overall hospitalization — mild, common, severe, and critical — according to the standards set by the Chinese Diagnosis and Treatment of Pneumonia Caused by New Coronavirus Infection (Eighth Version) as follows (assessors of the classification were blind to the final prediction scores in this study):

1. Mild: patients with mild clinical symptoms and no radiological abnormality
2. Common: patients presented with fever, cough or other respiratory symptoms, and positive radiological findings for pneumonia
3. Severe: if one of the following conditions were met:
   1. Respiratory distress \(\geq 30\) per minute
4. Finger oxygen saturation \((\text{SaO}_2)\) \(\leq 93\)% in resting state
5. Partial pressure of arterial oxygen \((\text{PaO}_2)/\text{concentration of oxygen inhaled (FiO}_2)\) \(\leq 300\) mmHg
6. Clinical symptoms had worsened progressively and pulmonary lesion had progressed significantly — \(>50\)% on thoracic imaging within 24–48 hours
7. Critical: if one of the following conditions was met:
   1. Respiratory failure and mechanical ventilation needed
   2. Shock
   3. Patients with other organ dysfunction needing ICU monitoring and treatment

Patients meeting the clinical characteristics of either the mild or common type were identified as mNCP and thus enrolled in this study. Patients were excluded if admitted to an ICU, on a general ward while meeting the conditions of severe or critical types on admission, and there was a lack of clinical information \(\geq 60\)%.

**Data Collection**

We collected epidemiological, demographic, clinical, laboratory, management, and outcome data from NCP patients’ medical records from the three hospitals. Clinical outcomes were followed up to February 21, 2020. All data were checked by two physicians. A confirmed case with COVID-19 was defined as a positive result to high-throughput sequencing or RT-PCR assay for throat swab specimens from the upper respiratory tract that had been obtained from all patients using established techniques.\textsuperscript{7} Clinical information and classification of severity on admission were available to the statisticians.

**Outcomes**

The outcome focused on in this study was patients progressing to severe NCP, defined as developing to severe or critical during hospitalization. We divided patients into two groups: severe for mNCP patients who developed severe NCP and nonsevere for other patients who remained as mNCP before discharge.

**Statistical Methods**

Normally distributed continuous variables are expressed as means ± SD and were compared using \(t\)-tests. Continuous
variables with skewed distribution are shown as medians (IQR) and compared using the Mann–Whitney test. Categorical variables are presented as numbers and percentages and compared using the χ²-test. The significance of each variable was initially assessed by univariate analysis by comparing the severe group and the nonsevere group. Collinearity diagnosis was performed by calculating the variance-inflation factor (VIF) of each predictor. Variables with VIF ≥2 were excluded from multivariate analysis. Variables with p<0.1 were regarded as potential risk factors and included in multivariate regression analysis reduced by a backward-elimination procedure (conditional likelihood-ratio test and elimination if p≥0.05). Continuous variables with statistical significance were categorized and retained for multivariate testing, and cut-off points were identified following Youden’s index of the receiver-operating characteristic (ROC) curve or a clinically relevant cutoff.

Prognostic indices were calculated based on the results of the multivariate analysis. The performance of the score was assessed by measuring the area under the ROC (AUROC), while sensitivity and specificity were also calculated. Internal validation was performed by applying the 2,000 bootstrap-resampling technique. The 2,000 samples were randomly assigned to two complementary subsets: the training set and the testing set, while the AUROC was measured for each subset. ROC curves and net reclassification improvement (NRI) analyses were used to assess improvement in risk prediction compared with CURB-65.

Validity was retested using fivefold cross-validation in the derivation cohort, which was divided randomly into five subsamples. A logistic regression model was developed using four subsamples and the score then calculated using the remaining (held out) subsample. This process was repeated for each of the five loops of cross-validation so that in the end each patient had a prediction score, for which overall (cross-validated) association and inference was tested. Statistical analysis was performed using SPSS version 22.0 and R 3.5.0. All tests were two-sided, and p<0.05 denoted statistical significance.

Results
Patient Characteristics
From January to February 2020, 529 patients diagnosed with NCP were recruited as eligible participants from the three hospitals and classified by four severity types during overall hospitalization. Among them, 92 patients had initially been admitted by an ICU, 25 were in a general ward while meeting the condition of severe or critical on admission, and 64 patients lacking clinical information ≥60% were excluded. We finally included 348 patients with mNCP, while 68 cases (19.5%) developed to severe or critical during hospitalization. Baseline characteristics of all patients and comparisons between severe and nonsevere groups are described in Supplemental Table 1. Comparison between participants with complete and incomplete data is in Supplemental Table 2.

Mean interval between onset of symptoms and admission was 7.77±5.59 days, while patients from the severe group showed significantly longer mean waiting time of 9.68 days compared with 7.31 days in the nonsevere group (p=0.002). A total of 119 patients (34.2%) had comorbidities, with hypertension being the most commonly observed (23.6%), followed by diabetes, coronary heart disease, and cancer. Overall, higher proportions of history of cardiocerebral vascular disease, diabetes, and cancer were observed in patients in the severe group than the nonsevere group.

Laboratory findings comprising leukocytes, neutrophils, total bilirubin, direct bilirubin, urea, myoglobin, D-dimer, fibrinogen, and CRP levels were obviously increased in the severe group over the nonsevere group. The severe group showed large variability in lymphocyte counts, with higher levels of lymphopenia (lymphocytes ≤0.8×10⁹/L). Hypoxemia was defined as PaO₂ ≤60 mmHg and hyperoxia PaO₂ ≥120 mmHg according to arterial blood gas, the latter appearing in 25.1% of patients in the common group. We counted lung-lobe infiltrations according to chest CT, and the mean value was 3.67 (only five lung lobes in total), indicating extensive involvement of NCP.

For coinfection, two patients were infected with Epstein–Barr virus, while no other respiratory virus infection was found. Only four hospitalized patients had bacterial coinfection. Two patients from the nonsevere group were infected with antibiotic-resistant bacteria, which were Escherichia coli and Serratia marcescens separately. One other patient was positive for Mycoplasma pneumoniae. No infection of fungi or tuberculosis was found among the patients.

As for treatment, antivirus therapy in the first 4 days after onset of symptoms was noted as early antivirus treatment, and was applied for probably half the patients in our study. Despite the low cobacterial infection rate, preventive antibiotic therapy was applied for 70% of our patients. The severe group had higher probabilities of both antibiotic and intravenous corticosteroid therapies. In our study, 23 patients died during hospital stay, indicating
a mortality rate of 6.6% for mNCP patients. All deaths occurred in the severe group. Eleven patients were admitted to an ICU before death. Immunologic tests between severe and nonsevere NCP patients are described in Table 1. Lower serum levels of CD3+, CD4+, and CD8+ T-lymphocyte subtypes were noted in the severe group.

**Table 1** Comparison of T-lymphocyte Subtypes, Humoral Immunity, and Cytokines Between Severe and Nonsevere Groups

|                | Total (n=200) | Nonsevere (n=181) | Severe (n=19) | p-value |
|----------------|--------------|-------------------|--------------|---------|
| CD3+ (%)       | 29.07±29.22  | 29.3±30.31        | 26.55±14.51  | 0.719   |
| CD4+ (%)       | 66.78±13.18  | 67.42±13.11       | 60.25±12.44  | 0.038   |
| CD8+ (%)       | 41.91±20.09  | 42.44±20.73       | 36.43±10.83  | 0.255   |
| Absolute CD3 count | 301.35±180.37 | 311.01±179.98     | 207.63±159.91| 0.017   |
| Absolute CD4 count | 809.23±432.00 | 839.30±429.62     | 522.84±349.04| 0.002   |
| Absolute CD8 count | 482.46±262.55 | 498.69±261.91     | 327.00±188.86| 0.006   |
| CD4+/CD8+ ratio | 1.75±0.73    | 1.75±0.75         | 1.75±0.44    | 0.967   |
| IgG            | 12.06±2.57   | 12.05±2.49        | 12.14±3.43   | 0.893   |
| IgA            | 2.53±0.99    | 2.50±0.97         | 2.86±1.16    | 0.187   |
| IgM            | 1.02±0.50    | 1.02±0.51         | 0.95±0.48    | 0.606   |
| IL6            | 11.07±13.10  | 10.30±8.99        | 12.44±18.29  | 0.367   |

According to the logistic regression model, five parameters were included as predictive factors (Table 2): history of diabetes (OR 2.064, 95% CI 1.010–4.683; p=0.043), time from symptom onset to admission ≥7 days (OR 1.945, 95% CI 1.054–3.587; p=0.033), lymphocyte count ≤0.8×10^9/L (OR 1.816, 95% CI 1.008–3.274; p=0.047), myoglobin ≥90 ng/mL (OR 2.496, 95% CI 1.235–5.047; p=0.011), and D-dimer ≥0.5 mg/L (OR 2.740, 95% CI 1.395–5.380; p=0.003). The VIF of each variable was <2.0. The score was named the mNCP severity prognostic index (mNCP-SPI).

**Risk Factors of Developing Severe Pneumonia**

According to the aforementioned methods and analyses, 13 categorical variables were entered in a backward stepwise logistic regression analysis (Table 2). The optimal feature subset selection showed that a model with five predictors had the minimum Bayesian information criterion, while the best subset features turned out to be diabetes, interval between onset of symptoms and admission, lymphocyte count, myoglobin, and D-dimer. The MSE value of the testing data set was 0.488.

**Table 2** Univariate and Multivariate Analysis Associated with In-Hospital Severity Progression of Patients with mNCP

| Clinical feature                  | Univariate | Multivariate |
|----------------------------------|------------|--------------|
| Male                             |            |              |
| Age ≥65 years                     | 2.000 (1.119–3.573) | 1.945 (1.054–3.587) |
| Time from symptom onset to admission ≥7days | 2.370 (1.346–4.171) | 2.064 (1.010–4.683) |
| Cardiovascular diseases           | 1.822 (1.043–3.185) | 1.816 (1.008–3.274) |
| Diabetes                         | 2.286 (1.079–4.843) | 2.740 (1.395–5.380) |
| Cancer                           | 4.365 (1.227–15.535) | 2.496 (1.235–5.047) |
| Lymphocytes ≤0.8×10^9/L           | 2.625 (1.552–4.525) | 2.740 (1.395–5.380) |
| Albumin ≤30g/L                   | 3.947 (2.138–7.286) | 2.496 (1.235–5.047) |
| Urea ≥7mmol/L                    | 2.160 (1.074–4.347) | 2.740 (1.395–5.380) |
| Myoglobin ≥90ng/mL               | 3.904 (2.027–7.519) | 2.496 (1.235–5.047) |
| D-dimer ≥0.5mg/L                 | 4.325 (2.297–8.144) | 2.740 (1.395–5.380) |
| PaO2 ≤80mmHg                     | 15.947 (6.556–38.791) | 2.740 (1.395–5.380) |
| Multilobe infiltrates ≥4         | 5.611 (1.903–16.545) | 2.740 (1.395–5.380) |

**Prognostic Index and Validation**

Relative weights were assigned according to the regression coefficient of each categorical variable (β) Prognostic score was calculated as follows:

Predictive score = 73.40221 (if history of diabetes)
The rate of development of severe NCP according to the nomogram is shown in Figure 1.

For the total study population (348 patients), this model showed a c-statistic of 0.747 (95% CI 0.682–0.814, Figure 2), and sensitivity and specificity were 0.765 and 0.644, respectively, under a score cutoff of 165. For the 2,000 bootstrap samples, the AUROC of the training set was 0.753 (95% CI 0.680–0.826), while that of the testing set was 0.738 (95% CI 0.688–0.788). Validation was also performed using fivefold cross-validation in the derivation cohort, which revealed accuracy of 0.7573 (95% CI 0.7095–0.8163, p=0.021). The mNCP-SPI was a significantly stronger predictor of severity than CURB-65 (n=348, AUROC 0.747 vs 0.538; p=0.001; Figure 2). NRI of the mNCP-SPI was also improved over CURB-65 (NRI 0.421, 95% CI 0.365–0.488; p=0.028).

Discussion
A novel predictive score was established based on 348 patients with mNCP in our study, among which 68 (19.5%) developed severe NCP during hospitalization. This score included five variables: time from symptom onset to admission ≥7 days, lymphocyte count ≤0.8×10^9/L, D-dimer ≥0.5 mg/L, myoglobin ≥90 ng/mL, and history of diabetes. This model showed sensitivity of 0.765, specificity of 0.644, and AUROC of 0.747.

During hospitalization, most patients had marked lymphopenia, and nonsurvivors developed more severe lymphopenia over time. From the 99 patients initially reported, of the nine who died, eight had lymphopenia.
Data from 1,099 patients showed that severe cases had more prominent laboratory abnormalities (ie, leukopenia, lymphopenia, thrombocytopenia) than nonsevere NCP cases.\textsuperscript{15} Consistently with these studies, our model also revealed that lymphocyte count $\leq 0.8 \times 10^9/L$ was a risk factor for developing severe pneumonia. Previous studies have shown that invasion of viral particles in severe SARS-CoV infection can damage the cytoplasmic components of lymphocytes and that lymphocyte apoptosis may lead to lymphopenia in severe Middle East Respiratory Syndrome–CoV infection. This phenomenon is also reflected in our research. Counts and absolute values of T-lymphocyte subsets (such as CD3$^+$, CD4$^+$, and CD8$^-$) in the severe group were lower than those in the nonsevere group.

Aggravation of pneumonia may trigger a cytokine storm, presenting an increase in both proinflammatory and anti-inflammatory cytokines, thereby causing depletion of immune cells and decrease in lymphocytes. It has been reported that serum IL6 levels are significantly higher in patients with COVID-19 pneumonia;\textsuperscript{16} and IL6 after tocilizumab may distinguish nonsurvivors from survivors and further support the choice of deeper targeting of IL6.\textsuperscript{17} Although there was no difference in IL6 between the mild and severe groups in our study, the indices in both groups were significantly higher than normal (normally $<3.4$ pg/mL with Immulite 2000 reagent), suggesting extensive presence of cytokine storms in NCP patients.

Time from the onset of symptoms to admission $\geq 7$ days was an independent risk factor for developing severe NCP. Data of 1,099 patients in 31 provinces/cities also showed that time from symptom onset to developing pneumonia was significantly different between the severe group and the nonsevere group and the duration of symptoms in the severe group longer.\textsuperscript{15} A clinical study of 138 hospitalized patients with NCP showed that time from onset to dyspnea was 5 days, 7 days to hospital admission, and 8 days to acute respiratory distress syndrome.\textsuperscript{18} This could be due to the inflammatory storm reaching a peak in 5–7 days in patients with delayed treatment, and which lead to multiple-organ damage. Therefore, early identification and timely treatment are important.

Among the multiple-organ damage caused by inflammatory-factor storms and depletion of T lymphocytes, myocardial damage is often a factor that is easy to ignore in patients without previous heart disease. The causes of myocardial damage by NCP are the virus infection leading directly to cardiomyocyte lysis or damage to myocardial blood vessels, immuno damage to T cells leading to viral myocarditis, acidosis or respiratory alkalosis causing myocardial damage, hypoxemia leading to myocardial ischemia and necrosis. The combination of myoglobin and other myocardial enzyme indices, such as troponin I and creatine kinase MB can sensitively and specifically reflect myocardial damage at an early stage over a wide diagnostic time,\textsuperscript{19} and is also an important parameter to evaluate the prognosis of pulmonary infection.

The storm of inflammatory factors can also cause disorders in coagulation. D-dimer is a specific molecular marker produced by the hydrolysis of plasmin after cross-linking fibrin monomer with activating factor XII. When the D-dimer concentration in plasma increases, it indicates that the body has active thrombosis and secondary hyperfibrinolysis, which is common in various thrombotic diseases, such as deep-vein thrombosis, pulmonary embolism, and disseminated intravascular coagulation.\textsuperscript{20} Studies have also pointed out that hypercoagulability is a common manifestation of severe pneumonia.\textsuperscript{21} Plasma D-dimer levels are closely related to inflammatory levels\textsuperscript{22} and severity of pneumonia, and are valuable for prognostic prediction.\textsuperscript{23} It has also been observed that SARS patients with disseminated intravascular coagulation had severer disease and higher risk of death.\textsuperscript{24} A recent study showed that prothrombin time and D-dimer level on admission were higher in ICU patients than non-ICU patients.\textsuperscript{7} The laboratory findings of 1,099 patients with COVID-19 acute respiratory distress syndrome also showed that compared with the nonsevere group, the proportion of D-dimer $\geq 0.5$ mg/L in the severe group is higher.\textsuperscript{15} D-dimer $\geq 0.5$ mg/L is also a risk factor for developing severe NCP in this model, as damage to the vascular endothelium caused by cytokine storms can lead to the up-regulation of coagulation factors and downregulation of physiological anticoagulation pathways.\textsuperscript{25}

Diabetes is a common chronic disease, and immune dysfunction in diabetic patients is at high risk of various acute and chronic infections.\textsuperscript{26} A history of diabetes and ambient hyperglycemia are independent predictors of death and morbidity in SARS patients, and metabolic control may improve the prognosis of SARS patients.\textsuperscript{27} A history of diabetes is also an independent risk factor for severe NCP in our model. Studies have reported that in patients with diabetes, anti-inflammatory macrophages can be transformed into proinflammatory macrophages, and regulatory T cells can be transformed into T$_{H1}$ and T$_{H17}$ CD4$^+$ helper T lymphocytes.\textsuperscript{28,29} This may be an explanation for why diabetic patients tend to become severer. What is more,
infection can also cause sharp fluctuations in blood glucose in diabetic patients, which will adversely affect prognosis. Therefore, great attention should be paid to the management of blood glucose in diabetic patients with NCP.

Clinical prediction rules for pneumonia, such as CURB-65, have been widely used to evaluate community-acquired pneumonia, but lack specific indicators of viral infection and are not sensitive enough to predict the severity and prognosis of viral pneumonia. In our study, it has been verified that our model has better predictive ability than CURB-65, and all variables included are easy to acquire on admission. Meanwhile, previous clinical studies have shown that risk factors, such as chronic underlying diseases, combined bacterial infections, and high-dose glucocorticoid use, are all associated with poor prognosis of viral infections. We have also established a six-parameter MuLBSTA score predicting mortality of virus infection, but it is unclear whether the model is suitable for novel CoV, considering the rapid progression and low rate of bacterial coinfection of patients with COVID-19. A prior study has also reported on the CALL scoring system with four clinical parameters proposed by Ji et al, yet the sample size was small and needs further validation. Our study provides another prediction tool with considerable ability. At the same time, our results shed light on the importance of early intervention, glycemic control, and preventive anticoagulant therapy in this long-lasting epidemic.

Our research has limitations. First of all, due to the multicenter source of patients, some data were missing during the collection of medical records, such as early anti-infection treatment, the specific usage of glucocorticoids, and control of patients’ blood glucose. At the same time, our study did not include external validation, due to the limited number of mNCP patients and lack of other independent data, which is rather important and should be further confirmed. In summary, we found that the mNCP-SPI score had significant predictive ability for development of severe pneumonia among patients with mNCP. This could assist physicians who are currently treating NCP on the front line and may help to identify potential candidates for ICU and further medical support.

Ethics Approval
This study was approved by the Coordinating Ethics Committee of Ruijin Hospital, affiliated with Shanghai Jiao Tong University School of Medicine (2017-205), and was conducted in accordance with the Declaration of Helsinki.

Author Contributions
All authors made a significant contribution to the work reported, whether in the conception, study design, execution, acquisition of data, analysis, and interpretation, or in all these areas; took part in drafting, revising, or critically reviewing the article; gave final approval to the version to be published; have agreed on the journal to which the article has been submitted; and agree to be accountable for all aspects of the work.

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Disclosure
The authors declare that they have no competing interests in this work.

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