

Low Prognostic Nutritional Index Is Closely Related To Disease Progression And Poor Prognosis In Patients With COVID-19 In Chengdu, China”.

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1. Abstract

1.1. Background:

COVID-19 patients with lower immunonutritional status have a lower prognostic nutritional index (PNI) and are more likely to have severe disease and a poor prognosis.

1.2. Objectives:

This study aims to evaluate the relationship between PNI level and disease progression and prognosis in COVID-19 patients.

1.3. Methods:

This retrospective analysis examined the PNI level of 1228 patients diagnosed with COVID-19. Patients were categorized into four groups: PNI1 group (PNI<50.65), PNI2 group (PNI 50.65-53.75), PNI3 group (PNI 53.76-57.20), and PNI4 group (PNI>57.20), based on the quartile value of PNI at admission. Comparisons were made between the four groups regarding clinical and outcome data, and assessments

were conducted to determine the association between PNI and disease progression and prognosis in patients with COVID-19.

1.4. Results:

As PNI levels decreased from the upper to the lower quartile group, HGB, ALB, TP, and TLC levels also decreased. Additionally, patient age increased, and there were longer coronavirus negative conversion times and hospitalization durations. According to the multiple regression analysis, the severity of illness was associated with PNI levels, nutritional risk, and NRS2002 score upon admission. Furthermore, the Prognostic Nutritional Index (PNI), nutritional risk, and NRS2002 score during admission are all significant prognostic factors. The PNI demonstrates a high degree of accuracy in predicting severe illness and mortality among COVID-19 patients.

1.5. Conclusions:

A low PNI is significantly correlated with disease progression and a poor prognosis in COVID-19 patients. Early intervention for malnutrition is necessary to lower the incidence of critical illness in individuals with a PNI lower than 49.12. Patients with a PNI lower than 40.45 necessitate continuous nutritional support to decrease mortality and enhance the likelihood of recovery.

2. Keywords:

2019 coronavirus disease (COVID-19), dietary risk, prognostic nutritional index (PNI), disease progression, prognosis Clinical Trial

Registry:

Chinese Clinical Trial Register Chi CTR 20000345631.

3. Introduction

The COVID-19 pandemic, caused by SARS-CoV-2, presents significant threats and challenges to global health and social development [1]. Following the World Health Organization's declaration of the pandemic on 15th March 2020, the number of confirmed cases has risen to over 762 million and resulted in 6.8 million deaths by April 2023 [2]. COVID-19 is a disease that progresses from asymptomatic or mild upper respiratory tract infection to severe pneumonia [3,4]. Most patients infected with SARS-CoV-2 are mild and have a high recovery rate. However, severe patients can rapidly progress to respiratory failure, multiple organ dysfunction, and even death, with most deaths involving elderly and sickly patients who often have poor nutritional status [5,6]. A previous study provided evidence that malnutrition is one of the main predictors of death from viral

infection [7]. Diarrhea is a common symptom in COVID-19 patients and can lead to nutrient malabsorption [8]. Decreased appetite leads to reduced nutrient intake, and the psychological pressure of the disease increases the risk of anxiety and depression in patients, further affecting food intake [9]. In addition, critically ill patients, often with severe inflammation and anorexia resulting in sharply reduced food intake, are more likely to develop severe illness. Therefore, systematic and urgent management of nutritional assessment and further scheduling of COVID-19-infected patients is essential [10,11]. Expert statements endorsed by the European Society for Clinical Nutrition and Metabolism Council and the practice guidelines for COVID-19 nutrition management both advocate and highlight the importance of nutritional status assessment [12]. However, for patients with the most severe manifestations of infection, assessing nutritional status with common tools can be quite difficult due to physical limitations and difficulties in gathering anthropometric and dietary information. Therefore, we should consider using rapid screening instruments to assess nutritional status, such as the prognostic nutritional index (PNI) and Nutrition Risk Screening 2002 (NRS2002). The European Society for Clinical Nutrition and Metabolism (ESPEN) recommends Nutrition Risk Screening 2002 (NRS2002) as a nutritional screening tool for general inpatients [13]. For hospitalized COVID-19 patients, the literature has validated the feasibility of NRS2002 for nutritional risk screening [14,15].

Our previous study found that in COVID-19 patients, nutritional risk and a high NRS 2002 score are not only closely related to disease progression but also poor prognosis. Early malnutrition intervention is needed to delay disease progression for patients with an NRS2002 score > 0.5 , and continuous nutritional support therapy is needed to reduce mortality and improve prognosis for patients with an NRS2002 score > 5.5 [16]. PNI is calculated by peripheral blood lymphocyte count and serum albumin (Alb) concentration, which can reflect the immune nutritional status of patients and was originally used for nutritional assessment of non-emergency general surgery patients [17]. There is growing evidence that PNI can predict clinical outcomes in patients with certain types of cancer, including lung, gastrointestinal, breast, and gynecological tumors [18-19]. However, the current literature on the role of PNI in assessing and predicting the clinical outcomes of COVID-19 is still limited. For hospitalized patients, some previous studies have verified that PNI has a good predictive value for both severity and prognosis and analyzed the optimal cutoff values for severity, death, and ICU admission [20-24], but the results of these articles are slightly different, and the sample size is not large. The distribution characteristics and differences in PNI among large samples of COVID-19 patients with different sexes, ages, clinical types, infection sources, infection histories and vaccination histories, disease severities and prognoses are still unclear. It is unclear whether the optimal cutoff point of PNI for predicting disease severity and death in a large COVID-19 population is consistent with those reported in the literature. Therefore, this study aimed to assess the association of immunonutritional status and PNI scores with disease progression and prognosis in patients with COVID-19. [16]

4. Methods

4.1. Subjects

A cross-sectional and short-term follow-up cohort study was conducted, retrospectively recruiting all 1228 patients with COVID-19 who presented to the Public Health Clinical Centre of Chengdu and were hospitalized in the isolation ward between January 16, 2020, and January 30, 2022. The study received approval from the Ethics Committee of the Public and Health Clinical Centre of Chengdu (ethics approval number: PJ-K 2020-26-01). Written informed consent was waived by the Ethics Commission of the designated hospital because this study concerns emerging infectious diseases.

4.2. Inclusion and exclusion criteria

The study's inclusion criteria comprised the following: no sex limit; age ≥ 18 years; COVID-19; and inpatient isolation and treatment time > 1 day. The exclusion criteria consisted of the following: age < 18 years, inpatient isolation and treatment time < 1 day.

4.3 Disease diagnosis, clinical typing and cure criteria [16,25]

The disease diagnosis, clinical typing, and cure criteria for COVID-19 patients were established according to the seventh Trial Version of the Novel coronavirus pneumonia diagnosis and treatment guidance [25]. The diagnostic criteria specified that cases must meet at least one of the following etiological markers: detection of positive nucleic acid of the novel coronavirus by real-time fluorescence RT-PCR or sequencing of viral genes [16,25]. Clinical type included asymptomatic infection, light, common, severe and critical illness. The criteria for classification were as follows: (1) Asymptomatic infection criteria can be met when there are no clinical symptoms and no pneumonia manifestations visible on imaging; (2) The light type criteria indicate mild clinical symptoms and no pneumonia manifestations on imaging; (3) The common type criteria include clinical symptoms such as fever and respiratory tract problems, and pneumonia is visible on imaging; (4) The criteria for the severe type were met by one of the following: Respiratory distress, $RR \geq 30$ times/min; In the resting state, the oxygen saturation $\leq 93\%$; Arterial blood oxygen partial pressure (PaO₂)/oxygen concentration (FiO₂) ≤ 300 mmHg (1 mmHg = 0.133 kPa), In areas with high altitude (over 1000 meters above sea level), PaO₂/FiO₂ should be corrected according to the following formula: $PaO_2/FiO_2 * [atmospheric\ pressure\ (mmHg) / 760]$; Pulmonary imaging showed that lesions progressing over 50% in 24-48 hours were classified as severe; (5) The critically ill type criteria included one of the following conditions: respiratory failure requiring mechanical ventilation, shock, or multiple organ failure warranting ICU monitoring [16,25]. The criteria for discharge upon recovery were met when the patient's body temperature had returned to normal for a minimum duration of three days. The rewas significant improvement in respiratory symptoms and lung imaging reflecting reduced exudative lesions. Additionally, the presence of negative nucleic acid results for two consecutive sputum samples, nasopharyngeal swabs or other respiratory specimens was ensured. These samples were taken at least 24 hours apart from each other. [16,25].

4.4. Grouping standards

Among the 1228 COVID-19 cases, 304, 306, 310, and 308 were classified into four groups based on PNI quartile values at admission. These groups were PNI <50.65 (PNI 1), 50.65~53.75 (PNI 2), 53.76~57.20 (PNI 3), and >57.20(PNI 4). PNI was calculated as $10 \times \text{serum albumin (g/dl)} + 0.005 \times \text{total lymphocyte count (per mm}^3\text{)}$. [26]. Among the 1228 COVID-19 cases, 1184 noncritical patients (patients with asymptomatic infection, with light and with common clinical type) were divided into the non critical group, and 44 critical patients (patients with severe and with critically illness clinical type) were divided into the critical group. Among the 1228 COVID-19 cases, 1223 surviving patients were assigned to the cured group, and 5 dead patients were divided into the death group.

4.5. Definition of the nutritional risk, viral negative conversion time, disease severity and prognosis [16, 25]

Nutritional risk was evaluated using the NRS 2002 within 48 hours of admission. The assessment comprised three categories: evaluation of nutritional status (based on weight loss, BMI, and food intake), measurement of disease severity (induced stress metabolism due to the extent of the disease), and age (whether or not it was 70 years or older). Abbreviations will be explicitly explained at first use. Scores ranged from 0 to 7. Patients with a score of 3 or more were classified as “at nutritional risk”, whereas a score below 3 indicated “not at nutritional risk”. The disease severity including critical illness (COVID-19 patients with severe or critically ill clinical type) and noncritical illness (COVID-19 patients with asymptomatic infection, light or common clinical type). Prognosis included death and survival within 4 weeks of admission. The coronavirus-negative conversion time was defined as the time from disease onset to the time when the first negative nucleic acid test met the discharge criteria.

4.6. Data collection

All clinical, laboratory, and demographic data, as well as the NRS2002 score and PNI, of 1228 cases were gathered to construct a database. The accuracy, completeness and authenticity of all data were meticulously monitored by the researchers.

4.7. Statistical analyses

Statistical analyses were conducted using SPSS 26.0 (SPSS, Chicago, IL, USA) and Graph Pad Prism 8 (GraphPad, CA, USA). Normally distributed measurements are presented as the mean and standard deviation, while non normally distributed data are presented as the median and inter quartile range (IQR). Categorical data are presented as percentages or proportions. To compare data with a normal distribution and homogeneity of variance between multiple groups, we used one-way or two-way ANOVA. Subsequently, we carried out LSD t tests to compare data between groups with different PNI levels. For data with a normal distribution and homogeneity of variance between groups with different PNI levels, we used independent sample t tests. The percentages or proportions were used to express the enumerated data, while chi-square tests were employed to compare the data between two or more groups. Spearman correlation analysis was used for two-factor correlation analysis, and binary logistic regression analysis was utilized to examine

the factors that influence disease severity and prognosis. Subject operating characteristic (ROC) analysis was conducted to evaluate lymphocytes and subpopulations' ability to distinguish between patients with non severe and severe neocoronary pneumonia and between patients who survived and those who did not. $p < 0.05$ was considered statistically significant.

4.8. Patient and Public Involvement

The research questions and study design were developed with out involvement from patients or the general public. Patients were not involved in the recruitment of subjects or the conduct of the study. In addition, investigators assessed the burden of the intervention. The eligibility of the participants was evaluated followed by data collection. General results, devoid of personally identifiable information, can be disseminated upon request. The Ethics Committee of the Public Health Clinical Centre of Chengdu approved this study (ethics approval number: PJ-K2020-26-01). Written informed consent was waived by the Ethics Commission of the designated hospital because this study is related to emerging infectious diseases.

5. Results

5.1 Characteristics of the Study Population [16]

A total of 1228 patients with COVID-19 were recruited for this investigation. The demographic and clinical characteristics of the participants are detailed in Table 1. The patients' median age was 37 years, male-dominated (78.83%), and only a small number of patients were critical illness cases (3.58%) or died cases 5 (0.41%). The median duration of hospitalization was 15.0 days, and the coronavirus negative conversion time was 11.5 days (Table 1) [16]. In addition, there were 342 (27.85%) cases with one comorbidity, 228 (18.56%) cases with two comorbidities, 318 (25.90%) cases with three or more comorbidities, and 340 (27.69%) cases without comorbidities. (Table 1) [16]. Most of them (87.21%) were imported cases, and only 12.79% of patients were domestically transmitted cases. Regarding infection history, 1180 (96.1) patients had primary infections, and only 48 (3.9) patients had reinfections. According to vaccination history, only 107 (8.7) patients were fully vaccinated, and 1121 (91.3) patients were not vaccinated (Table 1). Moreover, according to clinical type, 31.92% of patients had asymptomatic infection, 11.16% had light, 53.34% had common, 2.03% had severe, and 1.55% had critical illness. Of them, 12.79% of patients were native, and 87.21% were foreign [16]. Furthermore, the median TP level was 71.7 g/L, the ALB level was 43.9 g/L, the HGB level was 151.0 g/L [16], the total lymphocyte level was $1.9 \times 10^9/L$, and the PNI level was 53.75 g/L2 (Table 1).

Table 1: Baseline characteristics of COVID-19 patients. [16]

Variables	Total (n=1228)	Range
Male, n(%)	968 (78.83)	
Age (year), [M (IQR)]	37.0 (30.0–38.0)	18 ~ 87
Disease severity		

Noncritical illness, n (%)	1,184 (96.42)	
Critical illness, n (%)	44 (3.58)	
Number of comorbidities		
0, n (%)	340 (27.69)	
1, n (%)	342 (27.85)	
2, n (%)	228 (18.56)	
3 or more, n (%)	318 (25.90)	
Clinical type of COVID-19		
Asymptomatic, n (%)	392 (31.92)	
Light, n (%)	137 (11.16)	
Common, n (%)	655 (53.34)	
Severe, n (%)	25 (2.03)	
Critical, n (%)	19 (1.55)	
Source of cases		
Domestically transmitted cases, n (%)	157 (12.79)	
Imported cases, n (%)	1071 (87.21)	
Prognosis		
Survive, n (%)	1223 (99.59)	
Death, n (%)	5 (0.41)	
The coronavirus negative conversion time (day), [M (IQR)]	11.5 (8.0–20.0)	2 ~ 89
Duration of hospitalization (day), [M (IQR)]	15.0 (12.0–20.0)	3 ~ 91
Body mass index (kg/m ²), [M (IQR)]	24.03±3.59	15.21~38.26
ALB at admission (g/L), [M (IQR)]	43.9 (41.5–46.0)	27.3 ~ 55.3
TP at admission (g/L), [M (IQR)]	71.7 (67.9–75.6)	38.2 ~ 92.2
HGB at admission (g/L), [M (IQR)]	151.0 (138.0–159.0)	54 ~ 191
TLC at admission (g/L), [M (IQR)]	1.9 (1.45–2.46)	0.26–5.63
PNI at admission (g/L), [M (IQR)]	53.75 (50.65–57.2)	32–72

prognostic nutrition index (PNI) segments		
PNI 1, n%	304 (24.8)	32.00–50.65
PNI 2	306 (24.9)	50.65–53.75
PNI 3	310 (25.2)	53.75–57.20
PNI 4	308 (25.1)	57.20–72.00
Infection history		
reinfection	48 (3.9)	
Primary infection	1180 (96.1)	
Vaccination history		
fully vaccinated	107 (8.7)	
not vaccinated	1121 (91.3)	

Abbreviations: PNI, prognostic nutrition index; ALB, albumin; TP, total protein; HGB, hemoglobin; TLC, total lymphocytes.

5.2. Comparisons among four different PNI level groups

Compared with those in the upper quartile PNI group and the other two PNI groups, although the positivity rates of SARS-CoV-2 nucleic acid, infection history and vaccination history among the four groups were similar, in the lower quartile PNI group, the proportion of patients with three or more comorbidities and the proportion of domestically transmitted cases were larger, the critical illness rate was higher, patients with severe and critical illness clinical types were higher, and all patients who died were in this group; significant differences were all found (all $P < 0.05$) (Table 2). In the lower quartile PNI group, patient age was oldest (Figure 1A), the coronavirus negative conversion time and the duration of hospitalization were longest (Figure 1B, 1C), and the HGB level, ALB level, TP level and total lymphocyte level were the lowest (Figure 2A, 2B, 2C, 2D). From the lower quartile PNI group to the upper quartile PNI group, with the decrease in PNI level, HGB level, ALB level, TP level and total lymphocyte level gradually decreased, patient age gradually increased, and the coronavirus negative conversion time and the duration of hospitalization gradually increased (Figure 2A, 2B, 2C, 2D, 1A, 1B, 1C).

Variables	PNI group (n=1,228)	X2	P value			
	<50.65 (n=304)	50.65–53.75 (n=306)	53.76–57.20 (n=310)	>57.20 (n=308)		
Male, n (%)	199 (65.5)	235 (76.8)	264 (85.2)	270 (87.7)	55.16	<0.001
Number of comorbidities					36.65	<0.001
0, n (%)	65 (21.4)	91 (29.7)	99 (31.9)	85 (27.6)		
1, n (%)	64 (21.1)	96 (31.4)	96 (31.0)	86 (27.9)		
2, n (%)	63 (20.7)	54 (17.6)	48 (15.5)	63 (20.5)		
3 or more, n (%)	112 (36.8)	65 (21.2)	67 (21.6)	74 (24.0)		
Prognosis					15.25	0.002
Survive, n (%)	299 (98.4)	306 (100.0)	310 (100.0)	308 (100.0)		
Death, n (%)	5 (1.6)	0 (0.0)	0 (0.0)	0 (0.0)		
Disease severity					86.70	<0.001
Critical illness, n (%)	37 (12.2)	4 (1.3)	2 (0.6)	1 (0.3)		
Noncritical illness, n (%)	267 (87.8)	302 (98.7)	308 (99.4)	307 (99.7)		

Source of cases					108.74	<0.001
Domestically transmitted-cases, n (%)	90 (29.6)	34(11.1)	18(5.8)	15(4.9)		
Imported cases, n (%)	214(70.4)	272(88.9)	292(94.2)	293 (95.1)		
Clinical type of COVID-19					153.7	<0.001
Asymptomatic, n (%)	55(18.1)	87(28.4)	123(39.7)	127(41.2)		
Light, n (%)	18(5.9)	32(10.5)	50(16.1)	37(12.0)		
Common, n (%)	194 (63.8)	183(59.8)	135(43.5)	143(46.4)		
Severe, n (%)	18 (5.9)	4(1.3)	2(0.6)	1(0.3)		
Critical, n (%)	19 (6.3)	0(0.0)	0(0.0)	0 (0.0)		
Infection history					6.52	0.89
reinfection	5(1.6)	12(3.9)	17(5.5)	14(4.5)		
First infection	299(98.4)	294(96.1)	293(94.5)	293(95.5)		
Repositivity of SARS-CoV-2 nucleic acid					3.36	0.339
with	59(19.4)	49(16.0)	51(16.5)	43(14.0)		
without	245(80.6)	257(84.0)	259(83.5)	265(86.0)		
Vaccination history					6.05	0.109
fully vaccinated	24(7.9)	25(8.2)	21(6.8)	37(12.0)		
not vaccinated	280(92.1)	281(91.8)	289(93.2)	271(88.0)		

Table 2: Comparison of baseline conditions between the four groups(n=1228)

Abbreviations: NRS2002, Nutritional Risk Screening 2002.

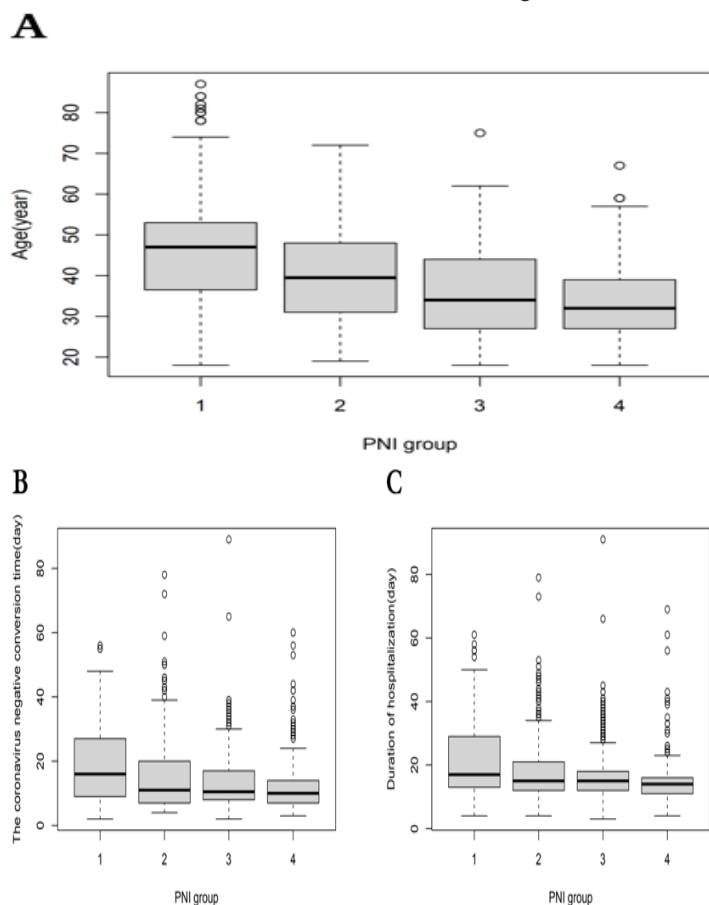


Figure 1: Comparison of age, the coronavirus negative conversion time and duration of hospitalization between four different PNI level groups. (n=1228; PNI<50.65, 50.65~53.75, 53.76~57.20 and >57.20 groups, n=304, 306, 310 and 308, respectively). (A)age. (B).the coronavirus negative conversion time. (C). duration of hospitalization. Wilcoxon rank-sum tests were used for inter group comparison.

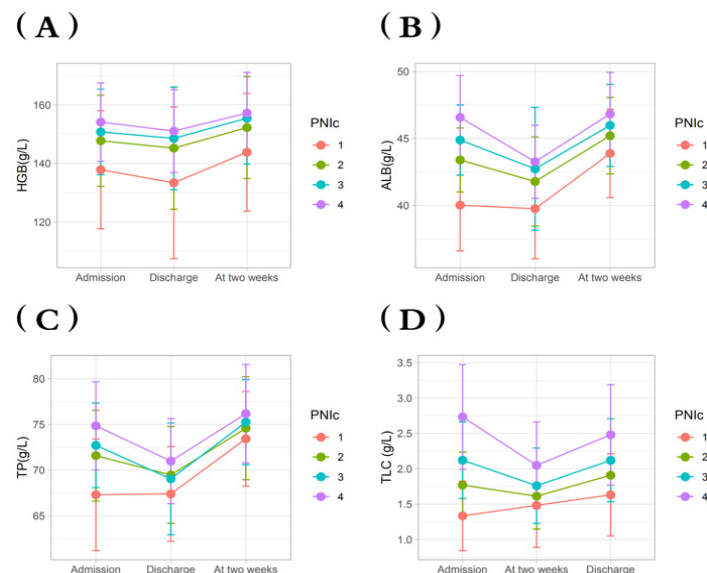


Figure 2: Comparison of HGB,ALB, TP and TLC between four different PNI level groups. (n=1228; PNI<50.65,50.65~53.75, 53.76~57.20and >57.20 groups, n=304, 306, 310 and 308, respectively). Abbreviations: HGB, hemoglobin; ALB, albumin; TP, total protein; TLC, total lymphocyte.

(A)HGB. (B)ALB. (C)TP .(D)TLC. All variables were presented as mean. Unpaired one-way ANOVA was used for intergroup comparisons. Unpaired t-tests were used for comparisons with the PNI >57.20group.

5.3. The relationship of nutritional risk, NRS 2002 score, PNI, disease severity and prognosis in patients with COVID-19

Spearman correlation analysis showed that age, sex, clinical type of COVID-19, number of comorbidities, time to negative coronavirus conversion, length of hospitalization, risk of malnutrition and NRS2002 score were positively related to disease severity [16], whereas source, ALB level, TP level, HGB level, total lymphocyte level and PNI level were negatively associated with disease severity (Table 3) [16].

Table 3: Spearman correlation analysis between nutritional parameters, NRS 2002, PNI and disease severity and prognosis (n=1228). [16]

Variables	Disease severity		Prognosis	
	(1=critical, 0=noncritical)		(1=cured, 2=death)	
	r	P	r	P
Age (year)	0.207	<0.001	0.108	<0.001
Sex	0.061	0.033	0.061	0.033
Clinical type of COVID-19	0.357	<0.001	0.121	<0.001
Number of comorbidities	0.209	<0.001	0.085	0.003
Source	-0.399	<0.001	-0.167	<0.001
The coronavirus negative conversion time (day)	0.163	<0.001	-	-
Duration of hospitalization (day)	0.068	0.017	-	-
Nutritional risk	0.496	<0.001	0.218	<0.001
NRS2002 score	0.513	<0.001	0.183	<0.001
ALB(g/L)	-0.199	<0.001	-0.107	<0.001
TP(g/L)	-0.181	<0.001	-0.101	<0.001
HGB(g/L)	-0.157	<0.001	-0.1	<0.001
TLC	-0.195	0	-0.091	0.001
PNI	-0.271	0	-0.141	<0.001

Abbreviations: NRS2002, nutritional risk screening 2002; ALB, albumin; TP, total protein; HGB, hemoglobin; PNI, prognostic nutrition index; TLC, total lymphocyte count.

When controlling for age and number of comorbidities, partial correlation analysis showed that sex, clinical type of COVID-19, coronavirus negative conversion time, duration of hospitalization, with-nutritional risk and NRS2002 score were still positively related to disease severity, while source, ALB level, TP level, HGB level, total lymphocyte level and PNI level were still negatively related to disease severity (Table 4).

Table 4: Partial correlation analysis between nutritional parameters, NRS 2002, PNI and disease severity and prognosis after controlling for age and number of comorbidities (n=1228)

Control variables	Variables	Disease severity		Prognosis	
		(1=critical, 0=noncritical)		(1=cured, 2=death)	
		r	P	r	P
Age (year)	Sex	0.07	0.014	0.058	0.044
Number of comorbidities	Clinical type of COVID-19	0.331	<0.001	0.112	<0.001
	Sour ceofcases	-0.327	<0.001	-0.112	<0.001
	The coronavirus negative conversion time (day)	0.135	<0.001	-	-
	Duration of hospitalization (day)	0.089	0.002	-	-
	Nutritional risk	0.493	<0.001	0.204	<0.001
	NRS2002 score	0.635	<0.001	0.349	<0.001
	ALB at admission (g/L)	-0.187	<0.001	-0.109	<0.001
	TP at admission (g/L)	-0.181	<0.001	-0.110	<0.001
	HGB at admission (g/L)	-0.196	<0.001	-0.171	<0.001
	TLC at admission (g/L)	-0.161	0	-0.066	0.021
	PNI at admission (g/L)	-0.208	0	-0.095	0.142

Abbreviations: NRS2002, nutritional risk screening 2002; ALB, albumin; TP, total protein; HGB, hemoglobin; PNI, prognostic nutrition index; TLC, total lymphocytes.

The risk factors for disease severity by multiple linear regression analysis were PNI level, body mass index (BMI) level, with-nutritional risk, NRS 2002 score, source, clinical type of COVID-19 and sex (Table 5). Moreover, Spearman correlation analysis showed that factors positively related to prognosis were age, sex, clinical type of COVID-19, number of comorbidities, with-nutritional risk and NRS 2002 score, while source, ALB level, TP level, HGB level, total lymphocyte level and PNI level were negatively related to prognosis (Table 3). When controlling for age and number of comorbidities, partial correlation analysis showed that sex, clinical type of COVID-19, with-nutritional risk and NRS 2002 score were still positively related to prognosis, while source, ALB level, TP

level, HGB level and total lymphocyte level were still negatively related to prognosis (Table 4). The risk factors for prognosis by multiple linear regression analysis were HGB level, nutritional risk, NRS2002 score, sex, coronavirus negative conversion time and PNI level (Table 5).

Table 5: Multiple stepwise regression analysis of influencing factors of disease severity and prognosis (n=1228)

Independent variables	B	Std. Error	Beta	t	P
Disease severity	PNI at admission (g/L)	-0.006	0.002	-3.834	<0.001
	BMI	0.005	0.001	4.36	<0.001
	Nutritional risk	-0.267	0.038	-7.027	<0.001
	NRS2002 score	0.151	0.011	13.768	<0.001
	Source of cases	-0.047	0.012	-3.907	<0.001
	Clinical type of COVID-19	0.024	0.004	6.066	<0.001
Prognosis	sex	-0.024	0.009	-2.651	0.008
	HGB	-0.0004	0.0001	-4.339	<0.001
	Nutritional risk	-0.102	0.012	-8.808	<0.001
	NRS 2002 score	0.035	0.004	10.105	<0.001
	sex	-0.009	0.004	-2.157	0.012
	The coronavirus negative conversion time (day)	-0.0003	0.0001	-2.466	0.014
	PNI at admission (g/L)	0.006	0.002	2.432	0.015

Abbreviations: NRS 2002, Nutritional Risk Screening 2002; PNI, prognostic nutrition index; BMI, body mass index, ALB, albumin; TP, total protein.

5.4. Prediction of the PNI level at admission on disease severity and prognosis inpatients with COVID-19

According to the ROC analysis, the PNI level at admission showed good utility for predicting critical COVID-19 patients (Table 6) and dead COVID-19 patients (Table 7). The areas under the curve were 0.8684 (Table 6, Figure 3) and 0.9923 (Table 7, Figure 4). The thresholds were 49.125 and 40.45, respectively (Tables 6 and 7). The sensitivities were 79.31% and 100.00%, respectively (Tables 6 and 7). The specificities were 85.15% and 99.86%, respectively (Table 6,7).

Table 6: The performance of various methods for distinguishing between severe cases and nonsevere cases (n=1228)

Variable	Cutoff point	AUC (95% CI)	Sensitivity	Specificity	False positive	False negative
PNI	49.125	0.8684	79.31%	85.15%	20.69%	14.85%

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Abbreviations: PNI, prognostic nutrition index; AUC, area under the curve; CI, confidence interval

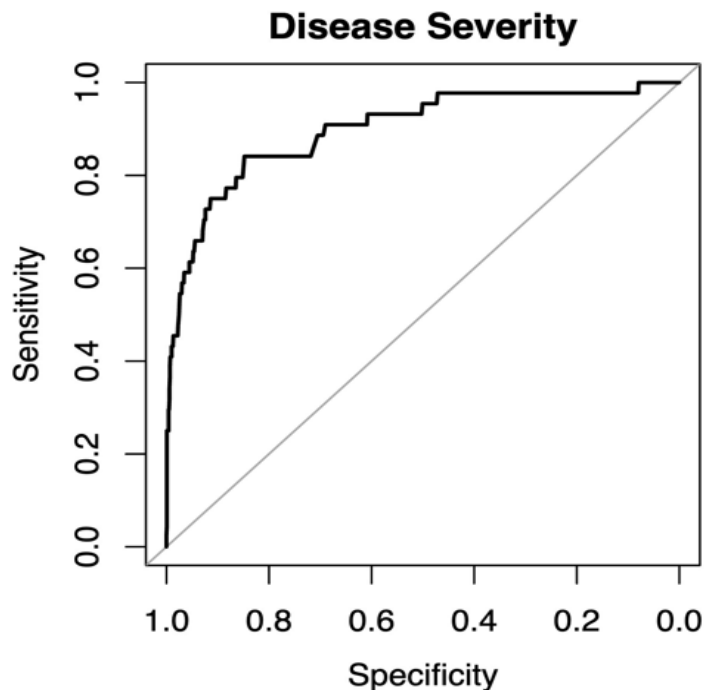


Figure 3: Using characteristics of PNI for discriminating the critical cases from the non-critical patients (n=1228; critical and non-critical groups, n=44 and 1184, respectively). ROC analysis showing the performance of PNI in distinguishing critical cases from non-critical patients. Abbreviations: ROC, receiver operating characteristic curve; AUC, area under the curve.

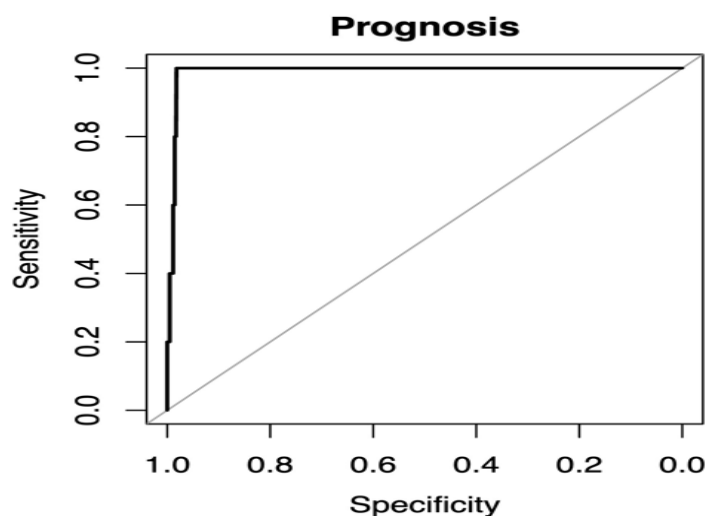


Figure 4: Using characteristics of PNI on admission for discriminating the surviving cases from the dead patients (n=1228; surviving and dead groups, n=1223 and 5, respectively). ROC analysis showing the performance of PNI in distinguishing the dead cases from the surviving

patients. Abbreviations: ROC, receiver operating characteristic curve; AUC, area under the curve.

Table 7: The performance of various methods for distinguishing between cured and dead patients (n=1228)

Variable	Cutoff point	AUC(95% CI)	Sensitivity	Specificity	False positive	False negative
PNI	40.45	0.9923	100.00%	99.86%	0.00%	0.14%

Abbreviations: PNI, prognostic nutrition index; AUC, area under the curve; CI, confidence interval.

6. Discussion

The prognostic nutritional index (PNI), calculated using peripheral blood lymphocyte counts and serum albumin (Alb) concentrations, was initially used in the nutritional assessment of nonemergency general surgical patients and is able to reflect the immunonutritional status of patients [26]. Recently, there has been growing evidence that PNI can be a useful prognostic indicator for predicting the prognosis of patients with certain types of cancer, including lung, gastrointestinal, breast, and gynecological tumors [18,19] and diseases other than cancer, such as acute heart failure, autoimmune disorders, hematologic malignancies, and chronic kidney disease in children [27-31]. In this study, PNI was applied for the first time in a large sample of COVID-19 patients in China to investigate its predictive value for disease progression and prognosis in COVID-19 patients. In this study, we found that patients in the lower PNI group were older and had longer hospitalization times and coronavirus negative conversion times, which may be related to low immunity and reduced nutritional intake and absorption in elderly individuals. This is consistent with previous studies. Nutrition plays a key role in improving immunity. In the case of viral infectious diseases, nutritional status affects the mutation of the viral genome from benign or low pathogenicity viruses to highly pathogenic viruses and their transmission within the host [32]. If nutritional risk is present, it directly affects immune defense. In addition, this study found that ALB levels, HGB levels, TP levels, and TLC levels increased with increasing PNI quartiles, whether at admission, discharge, or two weeks. ALB level, TP level, and HGB level are commonly used as indicators of malnutrition in clinical practice, with ALB levels being a significant factor in disease severity at admission, a reliable indicator of nutritional status, and a COVID-19 severity of prognosis. Therefore, we hypothesized that PNI could be used as a marker of disease progression and prognosis in COVID-19 patients.

A previous study of 140 COVID-19 patients (70 each of mild and severe cases) found that PNI was significantly lower in the severe group than in the mild group, that PNI was significantly lower in the severe death group than in the severe survival group, that PNI had a better predictive value for both the criticism and the prognosis, that those with a PNI lower than 39.08 were prone to develop a severe disease, and that those with a PNI lower than 33.05 [20] and 33.405 [21] had a higher risk of death. Overseas

studies have found that the PNI of COVID-19 patients admitted to the ICU is lower than that of non-ICU patients [22,23], and those with PNI lower than 36.7 [22] and 39.95 [23] are prone to develop severe disease, and those admitted to the ICU have a 4.4-fold higher probability of death [22]. The PNI of patients who died in the ICU was significantly lower than that of patients who survived in the ICU [20,23], and the PNI predicted that the optimal cutoff points for ICU patient death were ≤ 38.75 [11] [38] and ≤ 42 [23]. However, these studies are only small-sample studies and are not generalizable. In this study, we took a large sample of COVID-19 patients in China and found that patients with a PNI lower than 49.12 were prone to develop serious illness, and those with a PNI lower than 40.45 had a high risk of death. After ROC analysis, the PNI score showed good utility in predicting critically ill COVID-19 patients and dead COVID-19 patients. However, this study still has some limitations. It was a single-center retrospective study, and limitations inherent to retrospective studies are unavoidable and do not allow for causal inferences. In addition, we did not consider the relationship between dietary habits, psychiatric conditions, and social support and the nutritional status of patients, and we did not systematically collect data on patient-specific dietary intake, diagnosis of malnutrition, and nutritional support, information that might be useful in the management of patients with COVID-19 from a clinical point of view. Finally, the small number of severe cases, especially deaths, combined with the large number of missing data collection processes, has resulted in unclear relationships between PNI and indicators such as history of infection, vaccination history, and source of infection. As the number of prepositive positives increases, further validation of the predictive value of PNI in predicting disease progression and prognosis in patients with COVID-19 is needed.

7. Conclusions

The results of the present study indicated that not only the with-nutritional risk and NRS2002 score [16] but also the PNI level at admission are important influencing factors of COVID-19 disease severity and prognosis. COVID-19 disease severity and prognosis were positively correlated with nutritional risk and NRS2002 score [16] but negatively correlated with ALB level, TP level, HGB level [16], total lymphocyte level and PNI level. In addition, the PNI level at admission has good predictive value for disease progression and poor prognosis. For patients with PNI levels > 49.125 , early intervention should be given to malnutrition to reduce the occurrence of critical diseases. For patients with PNI levels > 40.45 , nutritional support treatment should be actively given to reduce mortality and improve prognosis.

8. Declarations

Ethics approval and consent to participate

This study was approved by the Ethics Committee of the Public and Health Clinic Centre of Chengdu (ethics approval number: PJ-K2020-26-01), and the Ethics Committee waived written informed consent because of emerging infectious diseases.

9. Consent for publication

All of the participants understand that the information will be published without their child or ward's/their relative's (circle as appropriate) name attached but that full anonymity cannot be guaranteed. All of the participants understand that the text and any pictures or videos published in the article will be freely available on the internet and may be seen by the general public. The pictures, videos and text may also appear on other websites or in print and may be translated into other languages or used for commercial purposes. All of the participants were offered the opportunity to read the manuscript.

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Authors' contributions

Concept and design: Qiqing Cui, Dafeng Liu; Data acquisition: Qiqing Cui, Fengjiao Gao, Xiaoyan Yuan, Rui Li, Yong Wang, Yanfeng Zhu, Dafeng Liu; data analysis and interpretation: Qiqing Cui, Fengjiao Gao, Xiaoyan Yuan, Rui Li, Yong Wang, Yanfeng Zhu, Dafeng Liu; Drafting the manuscript: Qiqing Cui, Fengjiao Gao, Xiaoyan Yuan, Rui Li, Yong Wang, Yanfeng Zhu, Dafeng Liu; administrative, technical, or material support: Qiqing Cui, Fengjiao Gao, Xiaoyan Yuan, Rui Li, Yong Wang, Yanfeng Zhu, Dafeng Liu; study supervision: Dafeng Liu, Yanfeng Zhu.

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