



Immune-Inflammatory Parameters in COVID-19 Cases: A Systematic Review and Meta-Analysis

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Feng X, Li S, Sun Q, Zhu J, Chen B, Xiong M and Cao G (2020) Immune-Inflammatory Parameters in COVID-19 Cases: A Systematic Review and Meta-Analysis. Front. Med. 7:301. doi: 10.3389/fmed.2020.00301 **Background:** The recent outbreak of coronavirus disease 2019 (COVID-19) has been rapidly spreading on a global scale and poses a great threat to human health. Acute respiratory distress syndrome, characterized by a rapid onset of generalized inflammation, is the leading cause of mortality in patients with COVID-19. We thus aimed to explore the effect of risk factors on the severity of the disease, focusing on immune-inflammatory parameters, which represent the immune status of patients.

Methods: A comprehensive systematic search for relevant studies published up to April 2020 was performed by using the PubMed, Web of Science, EMBASE, and China National Knowledge Internet (CNKI) databases. After extracting all available data of immune-inflammatory indicators, we statistically analyzed the risk factors of severe and non-severe COVID-19 patients with a meta-analysis.

Results: A total of 4,911 patients from 29 studies were included in the final meta-analysis. The results demonstrated that severe patients tend to present with increased white blood cell (WBC) and neutrophil counts, neutrophil-lymphocyte ratio (NLR), procalcitonin (PCT), C-reaction protein (CRP), erythrocyte sedimentation rate (ESR), and Interleukin-6 (IL-6) and a decreased number of total lymphocyte and lymphocyte subtypes, such as CD4+ T lymphocyte and CD8+ T lymphocyte, compared to the non-severe patients. In addition, the WBC count>10 × 10⁹/L, lymphocyte count<1 × 10⁹/L, PCT>0.5 ng/mL, and CRP>10 mg/L were risk factors for disease progression in patients with COVID-19 (WBC count>10 × 10⁹/L: OR = 2.92, 95% CI: 1.96–4.35; lymphocyte count<1 × 10⁹/L: OR = 4.97, 95% CI: 3.53–6.99; PCT>0.5 ng/mL: OR = 6.33, 95% CI: 3.97–10.10; CRP>10 mg/L: OR = 3.51, 95% CI: 2.38–5.16). Furthermore, we found that NLR, as a novel marker of systemic inflammatory response, can also help predict clinical severity in patients with COVID-19 (OR = 2.50, 95% CI: 2.04–3.06).

Conclusions: Immune-inflammatory parameters, such as WBC, lymphocyte, PCT, CRP, and NLR, could imply the progression of COVID-19. NLR has taken both the levels

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of neutrophil and lymphocyte into account, indicating a more complete, accurate, and reliable inspection efficiency; surveillance of NLR may help clinicians identify high-risk COVID-19 patients at an early stage.

Keywords: COVID-19, immune-inflammatory parameters, neutrophil-lymphocyte ratio, risk factor, meta-analysis

INTRODUCTION

In late 2019, a novel corona virus (SARS-CoV-2) caused an outbreak of a cluster of pneumonia cases in Wuhan, China. The new 2019 coronavirus pneumonia, COVID-19, has continued to spread rapidly around the world and has since become a global health emergency. Although control and quarantine measures have been applied to prevent a global spread, the infection has gradually increased, resulting in a pandemic (1). As of April 22, 2020, a total of 2,528,330 COVID-19 cases and 177,198 fatal cases were reported worldwide, with 84,287 cases and 4,642 deaths reported in China alone. Moreover, the number of people infected with COVID-19 in the United States accounts for about one-third of the world, with a 5.5% mortality rate.

SARS-COV-2 is a member of the coronavirus family along with SARS-CoV and MERS-CoV, but its transmission speed and infectivity are stronger than both (2, 3). SARS-CoV-2 can be transmitted through the respiratory tract, mainly causing respiratory infections and developing severe pneumonia, respiratory failure, and even death in infected patients (4, 5). Although the current situation is very grim, there is no specific medicine available for SARS-CoV-2. A recent systematic review and meta-analysis reported that the COVID-19-related mortality rate varies widely among epicenters and counties, even at the global level (6). In addition, researchers have identified several clinical characteristics associated with an increased risk of developing severe COVID-19, such as hypertension, cardiovascular disease, chronic kidney disease, and diabetes (6). Considering that infections might progress rapidly in some patients and timely clinical decisions are required, identifying patients who are at risk of developing a serious disease is particularly important for healthcare workers.

Since the pathophysiology of unusually high pathogenicity for SARS-CoV-2 has not been completely explained, information about inflammation and the immune response of patients with different severity of COVID-19 remains insufficient. Inflammation accompanied by an immune response often occurs in viral respiratory infections, and increasing evidence supports its important role in the progression of COVID-19 (7). Moreover, a previous retrospective study reported that a subgroup of patients with severe COVID-19 could have a dysregulation of the immune response that allows the development of viral hyperinflammation (8). In view of the fact that inflammation or immune indicators are very common and easily obtained, identifying risk factors in blood associated with disease severity among SARS-CoV-2-infected patients is vital for early intervention to improve mortality.

However, to date, no systematic review has been reported concerning the putative association between various inflammation indicators and the progression of COVID-19. Therefore, in the current study, a systematic review and metaanalysis was conducted to summarize the difference of several inflammation indicators between severe and non-severe COVID-19 patients and identify the relevant risk factors correlated with the progression of COVID-19.

MATERIALS AND METHODS

Literature Search Strategy

The PubMed, Web of Science, EMBASE, and China National Knowledge Internet (CNKI) databases were searched for eligible publications from December 2019 to April 2020. The search strategy was based on combination of following terms: "COVID-19" OR "Novel Coronavirus-Infected Pneumonia" OR "2019 novel coronavirus" OR "NCP" OR "2019-nCoV" OR "SARS-CoV-2." The search items in each database are also available in **Supplementary Material**. References cited in the retrieved articles were also scanned for relevant studies. Two reviewers independently screened the title and abstract of each study. Articles deemed potentially eligible were retrieved for a full-text review.

Selection and Exclusion Criteria

This systematic review and meta-analysis were conducted according to the PRISMA guidelines. Studies included in the meta-analysis had to satisfy the following criteria: (1) the study was a clinical observation in humans; (2) the study included clinical signs of a COVID-19 patient; and (3) the study included baseline information regarding inflammation indicators, such as white blood cells, neutrophils, lymphocytes, C-reactive protein (CRP), and procalcitonin (PCT). All articles of any design (randomized controlled trials, non-randomized controlled trials, case-control studies, and cross-sectional studies) were included except for narrative review, comment, opinion piece, methodological report, or conference abstract.

Data Extraction and Quality Assessment

Data were extracted separately by two reviewers, and discrepancies were resolved by consensus. The following data of each study were extracted from included articles: name of first author, publication date, study location, sample size, patients' age, gender, study design, COVID-19 severity, and inflammation indicators. The quality of the included studies was assessed using the MINORS (9) by two independent reviewers. MINORS is a valid instrument designed to assess the methodological quality of non-randomized surgical studies whether comparative or non-comparative. The global ideal score is 16 for non-comparative studies and 24 for comparative studies.



Data Analysis

Meta-analyses were conducted in order to evaluate the association between inflammation indicators and the risk of developing severe COVID-19 (10). For studies that presented continuous data as medians and inter-quartile ranges, the estimate of the means and standard deviations was performed according to the method described by Wan et al. (11). The mean difference (MD/WMD) or the standardized mean difference (SMD) and their related 95% confidence intervals (CIs) were calculated to evaluate the discrepancy of indicators between the non-severe and severe COVID-19 groups. For the pooled

analysis of the relationship between indicators and the severity of COVID-19, odds ratio (OR) and related 95% CI were pooled to calculate the effective value.

The Review Manager version 5.3 and STATA software (version 12.0) were used for data analysis. The heterogeneity between studies was assessed by the chi-squared and I-squared tests, with values of 0–25, 25.1–75, and 75.1–100% indicating a low, moderate, and high degree of heterogeneity, respectively. If $I^2 > 50\%$, a random-effect model was used to calculate the effect value. Otherwise, a fixed-effect model was performed. All *P*-values were two-tailed with a significant level at 0.05. Publication

TABLE 1 | Characteristics of included studies in this meta-analysis.

References	Location	Sample size (n, male)	Mean/median age	COVID-	19 severity	Extracted indicators	Minors score	
				Severe	Non- severe			
Bin et al. (12)	Wuhan	54 (30)	53.9	9	45	WBC; L	18	
Cai et al. (23)	Shenzhen	298 (145)	47.5	58	240	CRP; PCT; WBC; N; L; ESR; IL-6	21	
Chen et al. (33)	Wuhan	150 (84)	59	24	126	CRP	18	
Chen et al. (34)	Chongqing	139 (76)	46	31	108	WBC; N; L; NLR	18	
Chen et al. (35)	Guangzhou	296 (137)	NA	30	266	WBC; N; L; NLR	20	
Fang et al. (36)	Anhui	79 (45)	45.1	24	55	CRP; WBC; N; L	18	
Gao et al. (37)	Beijing	90 (43)	53	22	55	CRP; PCT; WBC; L; ESR; IL-6	20	
Guan et al. (38)	national	1,099 (637)	47	173	926	WBC; L	18	
Hou et al. (39)	Chengdu	56 (29)	48	11	38	L; CD3; CD4; CD8; NLR	20	
Huang et al. (13)	Wuhan	41 (30)	49	13	28	PCT; WBC; N; L	21	
Li et al. (15)	Wuhan	62 (32)	NA	22	18	CRP; PCT; WBC; N; L; CD3; CD4; CD8	19	
Li et al. (14)	Zhuzhou	80 (40)	47.8	17	63	CRP; PCT; WBC; L; ESR	20	
Li et al. (16)	Chongqing	83 (44)	45.5	25	58	CRP; PCT; WBC; N; L; M	18	
Li et al. (17)	Beijing	46 (21)	45.6	6	40	CRP; PCT; WBC; N; L	18	
Liu et al. (18)	Wuhan	78 (39)	38	11	67	CRP; PCT; WBC; N; L; ESR	18	
Long et al. (19)	Jingzhou/Xiangyang	301 (150)	51	36	245	NLR	18	
Lu et al. (20)	Wuhan	101 (34)	NA	34	67	CRP; WBC; N; L; CD3; CD4; CD8	19	
Peng et al. (21)	Wuhan	112 (53)	62	16	96	CRP; PCT; WBC; N; L; M; NLR	19	
Qin et al. (8)	Wuhan	452 (235)	58	286	166	CRP; PCT; WBC; N; L; M; CD3; CD4; CD8; ESR; IL-6; NLR	20	
Wan et al. (22)	Chongqing	153 (77)	NA	21	132	CRP; PCT; L; CD3; CD4; CD8	21	
Wan et al. (24)	Chongqing	135 (72)	47	40	95	CRP; PCT; WBC; N; L	19	
Wang et al. (25)	Wuhan	138 (75)	56	36	102	WBC; N; L; M	21	
Wu et al. (26)	Wuhan	201 (128)	51	84	117	CRP; WBC; N; L; M; CD3; CD4; CD8; ESR; IL-6	19	
Xiang et al. (32)	Jiangxi	49 (33)	42.9	9	40	CRP; PCT; WBC; N; L; CD3; CD4; CD8; ESR	18	
Yang et al. (27)	Hangzhou	93 (56)	46.4	24	69	CRP; WBC; N; L; M; NLR	19	
Yuan et al. (28)	Chongging	223 (106)	46.5	31	192	PCT; WBC; L	19	
Zhang et al. (29)	Wuhan	138 (71)	57	56	82	CRP; PCT; WBC; L	21	
Zhang et al. (30)	Beijing	74 (35)	52.7	9	56	CRP; PCT; WBC; N; L	20	
Zheng et al. (31)	Changsha	161 (80)	45	30	131	CRP; WBC; L	18	

WBC, white blood cell; N, neutrophil; L, lymphocyte; M, monocyte; CD3, T cell; CD4, helper T cell; CD8, cytotoxic T cell; ESR, erythrocyte sedimentation rate; IL-6, Interleukin-6; NLR, neutrophil-lymphocyte ratio; CRP, C-reactive protein; PCT, procalcitonin; NA, not available.

bias (number of studies >10) was evaluated using Begg's funnel plots and Begg's rank correlation test, and the significance was set to P < 0.05.

RESULTS

Literature Information

In the initial search, 10,456 potentially relevant records were found in the PubMed, Web of Science, EMBASE, and CNKI databases (**Figure 1**). All papers were screened by reading their titles, and 1,704 of them were excluded due to being duplicates found in different databases. After evaluating the abstracts, 8,441 studies were eliminated due to presenting data that were irrelevant to our aim. After carefully reading the full text of the remaining 311 studies, 282 papers that did not meet the inclusion criteria were further excluded. Finally, 29 articles met the inclusion criteria and were included in qualitative synthesis, but some indicators were not described in all articles.

Characteristics of Included Studies

A total of 29 studies (8, 11–38) published between Feb 2020 and Apr 2020 were identified, and all these studies were retrospective cohort studies of a design with 4,911 patients enrolled in this meta-analysis. The main characteristics of the included studies are summarized and presented in **Table 1**. Eleven studies were conducted in Wuhan city, the epicenter of COVID-19 outbreak. Only one study was national; it was the largest and included 1,099 COVID-19 patients. The other 17 studies were accomplished in

Immune cells	Number of studies	Participants	Mean difference (95% CI)	Р	Effects model	Heterogeneity		
						I ²	Ph	
White blood cell	25	4,278	0.83 (0.41, 1.25)	<0.001	REM	77	<0.001	
Neutrophil	18	2,446	1.50 (1.01, 1.98)	< 0.001	REM	69	< 0.001	
Lymhocyte	27	4,480	-0.36 (-0.43, -0.30)	< 0.001	REM	73	< 0.001	
T cell	7	637	-332.48 (-496.93, -168.03)	< 0.001	REM	92	< 0.001	
Helper T cell	7	637	-204.15 (-289.97, -118.33)	< 0.001	REM	88	< 0.001	
Cytotoxic T cell	7	637	-107.23 (-182.78, -31.68)	< 0.001	REM	92	< 0.001	
Monocyte	7	1,128	-0.01 (-0.03, 0.01)	0.28	FEM	0	0.28	

TABLE 2 | The association between immune cells and disease severity in patients with COVID-19.

Cl, confidence intervals; FEM, fixed-effects model; REM, random-effects model; P_h, p-value of Q-test for heterogeneity.

several cities of China outside Wuhan. All patients in selected studies were diagnosis and confirmed as COVID-19. Sample sizes of all studies ranged from 41 to 1,099. The extracted inflammation or immune indicator comprises white blood cell, neutrophil, lymphocyte, monocyte, T cell, helper T cell, cytotoxic T cell, erythrocyte sedimentation rate, Interleukin-6, neutrophillymphocyte ratio, C-reactive protein, and procalcitonin. In addition, none of the studies were considered to be seriously flawed according to the MINORS assessment. The 29 included studies scored between 18 and 21.

Immune Cells and Disease Severity in Patients With COVID-19

There is little information about the underlying mechanisms of severe COVID-19 development and further investigations are urgently needed. Qin et al. (8) previously suggested that COVID-19 might damage lymphocytes, especially T lymphocytes, and the immune system was impaired during the period of disease. Also, several studies discovered an increased level of neutrophils along with a decrease in lymphocyte numbers in patients with COVID-19 (13, 16, 29). These findings indicated that neutrophils or lymphocytes could be a potential risk factor for the progression of SARS-CoV-2-infected patients. To test this, we conducted a meta-analysis in order to investigate whether lymphocytes or other immune cells were significantly associated with increased disease severity of COVID-19 (Table 2).

Information on white blood cell (WBC) was available in 25 studies, including 4,278 patients with COVID-19. The test showed that these studies have certain heterogeneity ($I^2 = 77\%$, P < 0.001), and the random effect model was used. The estimated pooled MD for these studies revealed a significant increase in number of WBC in severe COVID-19 group (MD = 0.83, 95% CI: 0.41–1.25, P < 0.001; **Figure 2A**). In addition, our analysis revealed that, compared to COVID-19 patients with normal levels of WBC, COVID-19 patients who presented with an increased number of WBC (>10 × 10⁹/L) had an ~3-fold higher risk of developing severe disease with a combined odds ratio (OR) of 2.92 (95% CI: 1.96–4.35, P < 0.001; **Figure 2B**). Eight studies were considered to be homogeneous, and the fixed effect model

was used ($I^2 = 0\%$, P = 0.47). Therefore, we suggest that COVID-19 patients who present with an abnormal WBC count should be carefully monitored and managed according to these results.

Neutrophils are the most abundant white blood cell in the human body, numbering an average of 4,150 cells/ μ L (50–70% of the total number of WBC). Information on neutrophils was available in 18 studies, including 758 cases in the severe group and 1,688 cases in non-severe group. Since heterogeneity ($I^2 = 69\%$) exceeded 50%, a random effects model was adopted. Similarly, we observed a significant increase in number of neutrophils in severe COVID-19 group (MD = 1.50, 95% CI: 1.01–1.98, P < 0.001; **Table 2**).

Lymphocytes are an important cellular component of the body's immune response function, the main performer of almost all immune functions of the lymphatic system, and a frontline "soldier" to fight external infections and monitor cell mutations in the body. We obtained information about lymphocytes in 27 studies, including 4,480 cases. The pooled analysis revealed that the number of lymphocytes decreased in patients with severe disease compared with non-severe patients (MD = -0.36, 95% CI: from -0.43 to -0.30, P < 0.001; Figure 3A). Next, we performed a meta-analysis in order to examine the association between lymphopenia ($<1 \times 10^9$ /L) and disease severity in patients with COVID-19. We found that pronounced lymphopenia was strongly associated with increased disease severity (OR = 4.97, 95% CI: 3.53–6.99, P < 0.001; Figure 3B) with moderate heterogeneity ($I^2 = 39\%$, P = 0.12). Regarding the changes of several lymphocyte subtypes between severe and nonsevere COVID-19 patients, we found that the number of total T cells, helper T cells, and cytotoxic T cells decreased as the disease progresses (Table 2). However, our meta-analysis revealed no significant correlation between monocyte and severe COVID-19 (MD = -0.01, 95% CI: -0.03-0.01, P = 0.28; Table 2).

Neutrophil-Lymphocyte Ratio (NLR) and Disease Progression

The neutrophil-lymphocyte ratio (NLR), derived from the absolute neutrophil and absolute lymphocyte counts of a full blood count, is an potential marker of the systemic inflammatory response (40). A rising neutrophil count and a falling lymphocyte count indicate the intensity of the inflammatory response and damage to the immune system, respectively. In this study, we

	S	evere		Noi	isever			Mean Difference	Mean Difference
Study or Subgroup	Mean	SD	Total	Mean	SD	Total	Weight	IV, Random, 95	% CI IV, Random, 95% CI
Bin Y	4.62	1.77	9	4.94	1.49	45	4.1%	-0.32 [-1.56, 0	92]
Cai Q	4.8	1.9	58	4.56	1.5	240	5.8%	0.24 [-0.28, 0	76]
Chen X	5.07	2.02	31	5.13	1.65	108	5.2%	-0.06 [-0.84, 0	72]
Chen X-2	5.28	1.9	30	4.93	1.83	266	5.3%	0.35 [-0.36, 1	06]
Fang X	7.7	5.04	24	4.93	1.45	55	2.5%	2.77 [0.72, 4	82]
Gao W	4.95	1.7	22	4.13	1.45	55	5.1%	0.82 [0.01, 1	63]
Guan W	4.3	2.39	173	4.9	1.63	926	6.0%	-0.60 [-0.97, -0	23] -
Huang C	9.73	5.23	13	5.47	3.52	28	1.4%	4.26 [1.13, 7	39]
Li D	4.57	1.74	22	4.8	1.69	18	4.5%	-0.23 [-1.30, 0	84]
Li D-2	6.88	3.83	17	4.71	1.29	63	2.8%	2.17 [0.32, 4	02]
Li K	5.77	2.83	25	5.26	1.6	58	4.2%	0.51 [-0.67, 1	69]
Li L	4.93	1.96	6	4.29	1.68	40	3.2%	0.64 [-1.01, 2	29]
Liu W	6.08	2.56	11	5.18	1.63	67	3.4%	0.90 [-0.66, 2	46] +
Lu Z	5.92	2.8	34	4.97	1.75	67	4.6%	0.95 [-0.08, 1	98]
Peng Y	6.48	3.67	16	6.2	2.81	96	2.8%	0.28 [-1.60, 2	16]
Qin C	6.1	3.05	286	4.9	1.79	166	5.9%	1.20 [0.75, 1	65]
Wan S	5.67	1.54	40	5.83	3.01	95	5.2%	-0.16 [-0.93, 0	61]
Wang D	6.67	4.79	36	4.33	1.58	102	3.3%	2.34 [0.75, 3	93]
Wu C	8.2	4.62	84	5.19	2.86	117	4.4%	3.01 [1.89, 4	13]
Xiang T	9.56	6.01	9	4.61	1.88	40	1.0%	4.95 [0.98, 8	92]
Yang A	9.1	5.6	24	6.4	2.4	69	2.2%	2.70 [0.39, 5	01]
Yuan J	6.1	3.3	31	5.4	1.9	192	4.2%		
Zhang J	6.1	3.8	56	4.63	1.81	82	4.5%	1.47 [0.40, 2	54]
Zhang W	5.1	2	9	4.2	1.5	56	3.8%	0.90 [-0.46, 2	26]
Zheng F	4.23	2.77	30	4.25	1.43	131	4.6%	-0.02 [-1.04, 1	00]
Total (95% CI)			1096			3182	100.0%	0.83 [0.41, 1.	25]
Heterogeneity: Tau ² =	0.73; Ch	i ² = 1(04.48, 0	lf = 24 (P < 0.0	00001);	l ² = 77%	6	-10 -5 0 5
Test for overall effect:	Z = 3.88	(P = (0.0001)						Favours [Nonsevere] Favours [Severe]
	0.			Manaa				Odda Datia	
Chudu an Culumann		evere		Nonse		1 14/-:	alat N	Odds Ratio	Odds Ratio
Study or Subgroup		<u>nts T</u>					•	I-H, Fixed, 95% Cl	M-H, Fixed, 95% Cl
Guan W			167	39	81		4%	2.54 [1.43, 4.52]	
Huang C		7	13	5	2			5.13 [1.19, 22.11]	
Li D-2		2	17	0	6			.48 [0.94, 448.70]	
Li K		4	25	5	5	8 10	4%	2.02 [0.49, 8.26]	
Wan S		3	40	6	9	5 13	5%	1.20 [0.29, 5.07]	
Yuan J		3	31	5	19	2 5	2%	4.01 [0.91, 17.70]	
Zhang J		13	56	4	8	2 10	2%	5.90 [1.81, 19.20]	
Zheng F		0	30	3	13			0.60 [0.03, 11.96]	
Total (95% CI)			379		1459	9 100	.0%	2.92 [1.96, 4.35]	•
Total events		51		67					
Heterogeneity: Chi ² =			P = 0		0%				
	, u				0				0.002 0.1 1 10

FIGURE 2 | Correlation between WBC and disease severity of COVID-19. (A) Forest plot of mean difference in WBC count. (B) Forest plot of OR for the association of elevated WBC with disease severity.

also observed a significant increase in the number of neutrophils and a significant decrease in the lymphocyte count during the severe phase. Therefore, higher NLR could be a potential maker for predicting the disease progression. We obtained information about NLR in six studies including 1,141 cases. The estimated pooled MD for these six studies indicated that severe patients have a higher NLR than non-severe patients (MD = 0.85, 95% CI: 0.56–1.15, P < 0.001; **Figure 4A**). At the same time, elevated NLR was significantly associated with increased disease severity with the pooled OR being 2.50 (95% CI: 2.04–3.06, P < 0.001; **Figure 4B**), demonstrating that elevated NLR could be a predictor of disease progression in COVID-19 patients.

		evere			nsevere			Mean Difference		Mean Dif	
Study or Subgroup			Total				Weight			IV, Rando	m, 95% Cl
Bin Y		0.32	9		0.43	45	3.3%	-0.42 [-0.66, -0.1	-		
Cai Q		0.41	58		0.62	240	5.0%	-0.43 [-0.56, -0.3			
Chen X		0.38	31		0.49	108	4.5%	-0.31 [-0.47, -0.1			
Chen X-2		0.51	30	1.55		266	3.9%	-0.49 [-0.69, -0.2	-		
Fang X		0.55	24		0.61	55	3.0%	-0.57 [-0.84, -0.3			
Gao W	1.97	1.7	22	1.25		55	0.7%	0.72 [-0.01, 1.4		Ī	14
Guan W	0.8	0.3	173		0.45	926	6.0%	-0.27 [-0.32, -0.2		-	
Hou K		0.41	11		0.51	38	2.8%	-0.04 [-0.33, 0.2			
Huang C	0.47	0.5	13	0.93		28	2.7%	-0.46 [-0.76, -0.1	- 12 · · · ·		
Li D		0.48	22		0.56	18	2.4%	-0.24 [-0.57, 0.0	-		-
Li D-2	0.92	0.4	17		0.47	63	3.6%	-0.49 [-0.71, -0.2	-		
Li K	0.7		25		0.37	58	4.2%	-0.49 [-0.67, -0.3			
Li L		0.21	6		0.53	40	3.5%	-0.43 [-0.66, -0.2			
Liu W		0.72	11		0.54	67	1.6%	-0.36 [-0.80, 0.0	-		-
Lu Z	0.8	0.3	34		0.69	67	4.0%	-0.79 [-0.98, -0.6			
Peng Y		0.49	16		0.44	96	3.2%	-0.33 [-0.59, -0.0			
Qin C		0.37	286		0.45	166	5.7%	-0.17 [-0.25, -0.0		-	
Wan Q		0.37	21		0.61	132	4.1%	-0.54 [-0.73, -0.3			
Wan S		0.31	40	1.2		95	4.6%	-0.40 [-0.55, -0.2			
Wang D		0.31	36		0.45	102	5.0%	-0.17 [-0.30, -0.0			
Wu C		0.38	84		0.55	117	5.0%	-0.36 [-0.49, -0.2			
Xiang T		0.66	9	1.26		40	1.4%	-0.54 [-1.02, -0.0	1000		
Yang A		0.54	24		0.63	69	3.1%	-0.52 [-0.78, -0.2			
Yuan J	1		31	1.5	0.6	192	4.5%	-0.50 [-0.66, -0.3			
Zhang J		0.38	56		0.45	82	4.9%	-0.14 [-0.28, -0.0			
Zhang W	1.1	0.5	9	1.2		56	2.0%	-0.10 [-0.47, 0.2	-		
Zheng F	0.87	0.29	30	1.14	0.46	131	5.0%	-0.27 [-0.40, -0.1	4]		
Total (95% CI)			1128			3352	100.0%	-0.36 [-0.43, -0.3	01	•	
Heterogeneity: Tau ² =	0 02 [.] Cł	$ni^2 = 94$		= 26 (P				0.00 [0.40, 0.0	- H		
Test for overall effect:						, 1	1070		-2	-1 0	· ·
	L 10.0	0 (1	0.0000	, , ,						Favours [Nonsevere]	Favours [Severe]
	S	evere		Nonse	evere			Odds Ratio		Odds R	atio
Study or Subgroup		nts T			Tota	l We	eight N	1-H, Fixed, 95% C		M-H, Fixed	
Guan W	9	147	153	584			-	5.96 [2.58, 13.75]			
Huang C		11	13	15				4.77 [0.89, 25.57]		+	
Li D-2		11	17	g				1.00 [3.25, 37.24]			
Li K		22	25	22				2.00 [3.21, 44.82]			
Wan S		32	40	36				6.56 [2.72, 15.78]			
Yuan J		17	31	30				6.56 [2.92, 14.70]			
											_
Zhang J		46	56	58			4.1%	1.90 [0.83, 4.38]		_	
Zheng F		13	30	29	131	11	7.5%	2.69 [1.17, 6.18]			-
Total (95% CI)			365		1375	100	0.0%	4.97 [3.53, 6.99]			•
Total events		299		783							
Heterogeneity: Chi ² :	= 11.56	df = 7	7 (P = ().12): l ²	= 39%						<u> </u>
Test for overall effect					3070				0.01	0.1 1	10

FIGURE 3 | Correlation between lymphocyte and disease severity of COVID-19. (A) Forest plot of mean difference in lymphocyte count. (B) Forest plot of OR for the association of lymphopenia with disease severity.

Inflammation-Related Markers and Disease Severity in COVID-19

Procalcitonin (PCT), used as a marker of severe inflammation, is released during infections caused by bacteria, fungi, and parasites but is normal or only slightly elevated in viral infections (41). In this meta-analysis, we obtained information about PCT in 16 studies including 2,070 cases. The estimated pooled MD for these studies revealed a significant increase in PCT (SMD =

A Contraction of the second seco	S	evere		Nor	nsever	е	1	Std. Mean Differe	ence	Std. Mean Difference
Study or Subgroup	Mean	SD	Total	Mean	SD	Total	Weight	IV, Random, 9	5% CI	IV, Random, 95% CI
Chen X	4.61	3.14	31	3.35	1.95	108	17.8%	0.55 [0.15,	0.96]	
Chen X-2	3.91	3.45	30	1.9	1.04	266	18.1%	1.37 [0.97,	1.76]	
Hou K	6.13	6.08	11	4.01	5.62	38	11.1%	0.36 [-0.31,	1.04]	
Peng Y	8.92	4.57	16	5.33	4.7	96	14.1%	0.76 [0.22,	1.30]	
Qin C	6.27	4.99	286	3.3	2.32	166	23.9%	0.70 [0.51,	0.90]	
Yang A	20.7	24.1	24	4.8	3.5	69	15.1%	1.26 [0.76,	1.76]	
Total (95% CI)			398			743	100.0%	0.85 [0.56,	1.15]	•
Heterogeneity: Tau ² = 0	0.08; Ch	ni² = 15	.23, df	= 5 (P =	= 0.009	9); ² = 6	67%		H	-2 0 2
	7 - 5 00	(D < C	00001	۱. ۱					-4	
Test for overall effect: 2	2 = 0.69	(F < C)						Favours [Nonsevere] Favours [Severe]
В		,		,				dds Ratio		Odds Ratio
		,	ds Rat	,	SE	Weig		dds Ratio Fixed, 95% CI		
В		,	ds Rat	,		Weig 3.6	iht IV,			Odds Ratio
B Study or Subgroup		,	<u>ds Rat</u> 1.10	tio]	5404	-	<u>ht IV,</u> 3% 3.0	Fixed, 95% CI		Odds Ratio
B <u>Study or Subgrou</u> Chen X		,	<u>ds Rat</u> 1.10 0.97	, tio] 179 0.	5404 3162	3.6	<u>ht IV,</u> 5% 3.0 5% 2.6	Fixed, 95% CI		Odds Ratio
B <u>Study or Subgroup</u> Chen X Long L Yang A		,	<u>ds Rat</u> 1.10 0.97	<u>tio]</u> 79 0. 12 0.	5404 3162	3.6 10.6 85.8	int IV. 5% 3.0 5% 2.0 3% 2.4	Fixed, 95% CI 03 [1.05, 8.73] 64 [1.42, 4.91] 46 [1.98, 3.06]		Odds Ratio
B <u>Study or Subgroup</u> Chen X Long L Yang A Total (95% CI)	<u>p lo</u>	og[Ode	<u>ds Rat</u> 1.10 0.97 0.9	, 79 0. 12 0. 01 0.	5404 3162 1112	3.6 10.6 85.8 100.0	int IV. 5% 3.0 5% 2.0 3% 2.4	Fixed, 95% CI 03 [1.05, 8.73] 64 [1.42, 4.91]		Odds Ratio
B <u>Study or Subgroup</u> Chen X Long L Yang A	<u>p lo</u> = 0.17	og[Ode	<u>ds Rat</u> 1.10 0.97 0.9 2 (P =	, 79 0. 12 0. 01 0. 0.92);	5404 3162 1112 I ² = 0	3.6 10.6 85.8 100.0	int IV. 5% 3.0 5% 2.0 3% 2.4	Fixed, 95% CI 03 [1.05, 8.73] 64 [1.42, 4.91] 46 [1.98, 3.06] 60 [2.04, 3.06]	L	Odds Ratio

0.78, 95% CI: 0.34–1.22, P < 0.001; **Figure 5A**). The pooled analysis showed a high degree of heterogeneity in PCT levels ($I^2 = 93\%$, P < 0.001). Then, we performed another metaanalysis in order to examine the putative association between elevated PCT (>0.5 ng/ml) and COVID-19 severity. As shown in **Figure 5B**, our results revealed that patients who present with elevated PCT have a significantly increased risk of developing severe COVID-19 (OR = 6.33, 95% CI: 3.97–10.10, P < 0.001) with low study heterogeneity ($I^2 = 0\%$, P = 0.70).

C-reactive protein (CRP) is an acute-phase response protein synthesized by the liver and elevated in response to inflammatory diseases. It plays a vital role in protection against infection, prevention of autoimmunity, and regulation of the inflammatory response. Information on CRP was available in 20 studies, including 2,591 patients with COVID-19. The estimated pooled MD indicated that severe patients have a higher level of CRP than non-severe patients (MD = 41.02, 95% CI: 33.32-48.71, P < 0.001; Figure 6A), with moderate study heterogeneity (I^2 = 73%). Moreover, we found that elevated CRP (above normal range) was significantly correlated with increased disease severity in COVID-19 (OR = 3.51, 95% CI: 2.38-5.16, P < 0.001; **Figure 6B**) with moderate heterogeneity ($I^2 = 49\%$, P = 0.08). Regarding other major Inflammation-related markers such as erythrocyte sedimentation rate (ESR) and Interleukin-6 (IL-6), we found that ESR and IL-6 were both significantly associated with increased disease severity (Table 3).

Publication Bias

Publication bias was originally defined as the publication or nonpublication of studies depending on the direction and statistical significance of the results. Publication bias was examined by the funnel plot and Begg's rank correlation test. As shown in **Figure 7**, distribution of the funnel plot was nearly symmetric and no evidence of publication bias in lymphocyte (P = 0.967) and PCT (P = 0.964). However, it was asymmetric in the meta-analyses of WBC, neutrophil, and CRP. Therefore, the trim and fill method was adopted to adjust publication bias (42). After adjustment, distribution of the funnel plot was more symmetric than before (**Figure 7**). At the same time, we found that the adjusted pooled MD did not change significantly, which indicated that the publication bias had little impact on the analysis and the analysis results were relatively stable (**Figures S1–S3**).

DISCUSSION

The majority of COVID-19 patients have relatively mild symptoms, but a considerable number of patients progress to severe pneumonia and even eventually develop acute respiratory distress syndrome (ARDS), septic shock, and/or multiple organ failure (2). Therefore, it is of great significance to study the immunological characteristics of peripheral blood in severe patients for timely diagnosis, precise treatment, delaying or halting the progression of the disease, and reducing mortality.

As is well-known, viral infection is closely related to the human immune system; good immune function can help the body eliminate foreign microorganisms, control infection, and eventually restore health (43). Dysregulation of immune cell responses and consequently immunologic abnormality are thought to play important roles in the severity of virus-induced

		S	evere			nsever			Std. Mean Difference	Std. Mean Difference
Study	or Subgroup	Mean	SD	Total	Mean	SD	Total	Weight	IV, Random, 95% CI	IV, Random, 95% Cl
Cai Q		0.16		58	0.2	0.06	240	6.8%	-0.70 [-1.00, -0.41]	-
Gao W		0.12	0.03	22	0.12	0.03	55	6.4%	0.00 [-0.49, 0.49]	
Huang	С		0.25	13	0.1	0.001	28	6.0%	0.71 [0.03, 1.38]	
Li D			0.06	22	0.03	0.02	18	6.1%	0.63 [-0.01, 1.27]	
Li D-2			0.67	17	0.04	0.07	63	6.2%	1.37 [0.80, 1.95]	
Li K		0.12		25	0.04	0.03	58	6.4%	1.06 [0.56, 1.55]	
Li L		0.11		6	0.13	0.03	40	5.5%	-0.69 [-1.56, 0.18]	
Liu W		0.22		11	0.06	0.04	67	6.0%	1.14 [0.47, 1.80]	
Peng Y	(0.28		16	0.12	0.11	96	6.3%	1.11 [0.56, 1.66]	
Qin C			0.15	286	0.06	0.04	166	6.9%	0.33 [0.14, 0.52]	*
Wan Q		0.08	12/11/2 11	21	0.04	0.06	132	6.5%	0.65 [0.18, 1.11]	<u> </u>
Wan S			0.06	40	0.04	0.02	95	6.5%	2.17 [1.72, 2.63]	
Xiang T		0.16		9	0.04	0.03	40	5.6%	1.75 [0.94, 2.55]	
Yuan J			0.52	31	0.11	0.07	192	6.5%	2.17 [1.74, 2.60]	
Zhang			0.18	56	0.06	0.05	82	6.7%	0.74 [0.39, 1.09]	
Zhang	W	0.1	0.02	9	0.1	0.04	56	5.9%	0.00 [-0.70, 0.70]	
Total (95% CI) geneity: Tau² =	0.72; Ch	i ² = 22	642 22.59, d	lf = 15 (l	P < 0.00		100.0% ² = 93%	0.78 [0.34, 1.22]	
Total (Heterog				22.59, d		P < 0.00			0.78 [0.34, 1.22]	-4 -2 0 2 Favours [Nonsevere] Favours [Severe]
Total (Heterog	geneity: Tau² =			22.59, d		P < 0.00		² = 93%	0.78 [0.34, 1.22] ds Ratio	
Total (Heterog Test for	geneity: Tau² = r overall effect: ;	Z = 3.50) (P = 0	22.59, d 0.0005)			0001); l [:]	² = 93% Ode	ds Ratio	Favours [Nonsevere] Favours [Severe] Odds Ratio
Total (S Heterog Test for	geneity: Tau ² = r overall effect: : <u>v or Subgrou</u>	Z = 3.50) (P = 0	22.59, d 0.0005) d <u>s Rat</u>	io]	SE \	0001); l ^a <u>Neight</u>	² = 93% Ode t IV,	ds Ratio Fixed, 95% Cl	Favours [Nonsevere] Favours [Severe]
Total (S Heterog Test for <u>Study</u> Fang 2	geneity: Tau ² = r overall effect: . <u>v or Subgrou</u> X	Z = 3.50) (P = 0	22.59, d 0.0005) d <u>s Rat</u> 1.9	io] 58 1.6	<u>SE 1</u> 5514	0001); F <u>Weight</u> 2.1%	Odd t IV, 5 7.09 [ds Ratio Fixed, 95% Cl 0.28, 180.31]	Favours [Nonsevere] Favours [Severe] Odds Ratio
Total (S Heterog Test for Study Fang 2 Guan	geneity: Tau ² = r overall effect: . <u>v or Subgroup</u> X W	Z = 3.50) (P = 0	22.59, d 0.0005) d <u>s Rat</u> 1.9! 1.42 ⁻	io] 58 1.6 16 0.3	<u>SE 1</u> 5514 5564	0001); F Neight 2.1% 44.7%	Odd t IV, 5 7.09 [5 4.1	ds Ratio Fixed, 95% Cl 0.28, 180.31] 4 [2.06, 8.33]	Favours [Nonsevere] Favours [Severe] Odds Ratio
Total (S Heterog Test for Study Fang 2 Guan Huang	geneity: Tau ² = r overall effect: . <u>v or Subgroup</u> X W	Z = 3.50) (P = 0	22.59, d 0.0005) d <u>s Rat</u> 1.95 1.421 3.008	io] 58 1.6 16 0.3 88 1	<u>SE 1</u> 5514 5564 558	0001); F Neight 2.1% 44.7% 2.3%	Odd t IV, 5 7.09 [5 4.1 5 20.26 [ds Ratio Fixed, 95% Cl 0.28, 180.31] 4 [2.06, 8.33] 0.96, 429.42]	Favours [Nonsevere] Favours [Severe] Odds Ratio
Total (S Heteros Test for Study Fang 2 Guan Huang Li K	geneity: Tau ² = r overall effect: : <u>v or Subgrou</u> X W g C	Z = 3.50) (P = 0	22.59, d 0.0005) d <u>s Rat</u> 1.99 1.42 3.008 2.078	io] 58 1.6 16 0.3 88 1 81 0	SE 1 5514 5564 558 608	0001); i [*] <u>Weight</u> 2.1% 44.7% 2.3% 15.4%	Odd t IV, 5 7.09 [5 4.1 5 20.26 [5 7.99	ds Ratio Fixed, 95% Cl 0.28, 180.31] 4 [2.06, 8.33] 0.96, 429.42] [2.43, 26.30]	Favours [Nonsevere] Favours [Severe] Odds Ratio
Total (S Heterog Test for Study Fang 2 Guan Huang Li K Liu W	geneity: Tau ² = r overall effect: <u>v or Subgrou</u> X W g C	Z = 3.50) (P = 0	22.59, d 0.0005) ds Rat 1.9 1.42 3.008 2.078 1.34	io] 58 1.6 16 0.3 88 1 81 0. 31 0.9	<u>SE 1</u> 5514 5564 558 608 9894	0001); i [*] 2.1% 44.7% 2.3% 15.4% 5.8%	Odu t IV, 5 7.09 [5 4.1 5 20.26 [5 7.99 5 3.83	ds Ratio Fixed, 95% Cl 0.28, 180.31] 4 [2.06, 8.33] 0.96, 429.42] [2.43, 26.30] [0.55, 26.64]	Favours [Nonsevere] Favours [Severe] Odds Ratio
Total (S Heterog Test for Study Fang 2 Guan Huang Li K Liu W Wan S	geneity: Tau ² = r overall effect: . <u>v or Subgrou</u> X W g C	Z = 3.50) (P = 0	22.59, d 0.0005) d s Rat 1.99 1.42 3.008 2.078 1.34 1.98	io] 58 1.6 16 0.3 88 1 81 0 31 0.9 31 1.6	SE 1 5514 5564 558 608 894 6439	Neight 2.1% 44.7% 2.3% 15.4% 2.1%	Odu t IV, 5 7.09 [5 4.1 5 20.26 [5 7.99 5 3.83 5 7.25 [ds Ratio Fixed, 95% Cl 0.28, 180.31] 4 [2.06, 8.33] 0.96, 429.42] [2.43, 26.30] [0.55, 26.64] 0.29, 181.89]	Favours [Nonsevere] Favours [Severe] Odds Ratio
Total (S Heterog Test for Study Fang 2 Guan Huang Li K Liu W	geneity: Tau ² = r overall effect: . <u>v or Subgrou</u> X W g C	Z = 3.50) (P = 0	22.59, d 0.0005) ds Rat 1.9 1.42 3.008 2.078 1.34	io] 58 1.6 16 0.3 88 1 81 0 31 0.9 31 1.6	<u>SE 1</u> 5514 5564 558 608 9894	0001); i [*] 2.1% 44.7% 2.3% 15.4% 5.8%	Odu t IV, 5 7.09 [5 4.1 5 20.26 [5 7.99 5 3.83 5 7.25 [ds Ratio Fixed, 95% Cl 0.28, 180.31] 4 [2.06, 8.33] 0.96, 429.42] [2.43, 26.30] [0.55, 26.64]	Favours [Nonsevere] Favours [Severe] Odds Ratio
Total (Heteroo Test for Study Fang) Guan Huang Li K Liu W Wan S Wang	geneity: Tau ² = r overall effect: . <u>v or Subgrou</u> X W g C	Z = 3.50) (P = 0	22.59, d 0.0005) d s Rat 1.99 1.42 3.008 2.078 1.34 1.98	io] 58 1.6 16 0.3 88 1 81 0 31 0.9 31 1.6	SE V 5514 5564 558 608 9894 6439 454	Neight 2.1% 44.7% 2.3% 15.4% 2.1%	Odu t IV, 7.09 [4.1 20.26 [7.99 3.83 7.25 [10.91	ds Ratio Fixed, 95% Cl 0.28, 180.31] 4 [2.06, 8.33] 0.96, 429.42] [2.43, 26.30] [0.55, 26.64] 0.29, 181.89]	Favours [Nonsevere] Favours [Severe] Odds Ratio
Total (Heteroo Test for Study Fang) Guan Huang Li K Liu W Wan S Wang Total	geneity: Tau ² = r overall effect: : <u>v or Subgroup</u> X W g C S D (95% CI)	Z = 3.50 p log	9 (P = 0	22.59, d 0.0005) ds Rat 1.99 1.42 3.008 2.078 1.343 1.98 2.389	io] 58 1.6 16 0.3 88 1 81 0.9 31 0.9 14 1.6 96 0	SE 1 5514 5564 558 608 894 439 439 454	Meight 2.1% 44.7% 2.3% 15.4% 2.1% 2.1% 2.1% 2.1% 2.1% 100.0%	Odu t IV, 7.09 [4.1 20.26 [7.99 3.83 7.25 [10.91	ds Ratio Fixed, 95% Cl 0.28, 180.31] 4 [2.06, 8.33] 0.96, 429.42] [2.43, 26.30] [0.55, 26.64] 0.29, 181.89] [4.48, 26.56] [3.97, 10.10]	Favours [Nonsevere] Favours [Severe]
Total (S Heterog Test for Study Fang J Guan Huang Li K Liu W Wan S Wang Total Hetero	geneity: Tau ² = r overall effect: : <u>v or Subgrou</u> X W g C S D	Z = 3.50 p log	9 (P = 0 g[Odc	22.59, d 2.0005) ds Rat 1.99 1.42 3.008 2.078 1.34 1.98 2.388 6 (P =	io] 58 1.6 16 0.3 88 1 81 0.9 14 1.6 96 0 0.70);	SE 1 5514 5564 558 608 894 439 439 454	Meight 2.1% 44.7% 2.3% 15.4% 2.1% 2.1% 2.1% 2.1% 2.1% 100.0%	Odu t IV, 7.09 [4.1 20.26 [7.99 3.83 7.25 [10.91	ds Ratio Fixed, 95% Cl 0.28, 180.31] 4 [2.06, 8.33] 0.96, 429.42] [2.43, 26.30] [0.55, 26.64] 0.29, 181.89] [4.48, 26.56] [3.97, 10.10]	Favours [Nonsevere] Favours [Severe]

with disease severity.

disease (44). Indeed, previous studies of novel coronavirus infection have suggested that the severe and aberrant host immune response are responsible for the severity of COVID-19 (16, 24). At the same time, peripheral blood immuneinflammatory parameters will also change significantly with the progression of COVID-19 disease. Therefore, emerging researches were focus on these accessible laboratory data for assessing and predicting clinical severity in patients with COVID-19. One of the most prominent factors related to the severity and outcomes of the Middle East respiratory syndrome coronavirus (MERS-CoV) disease is the hematological change in the white blood cell population (45). Also, several studies have addressed the difference of baseline leukocyte counts between the clinical stages in COVID-19 patients. Mo et al. (46) reported that refractory patients had higher level of neutrophils in comparison with general patients. In patients with severe COVID-19, but

not in patients with mild disease, lymphopenia is a common feature, with drastically reduced numbers of CD4+T cells, CD8+ T cells, B cells, and natural killer (NK) cells (2, 8, 13). Qin et al. (8) investigated 452 patients with laboratory confirmed COVID-19 and found that severe cases were likely to have higher neutrophil count but lower lymphocyte count compared with non-severe patients; the NLR thus tended to be higher in the severe group. Long et al. (19) reported that neutrophillymphocyte ratio \geq 2.973 (HR = 2.64, 95% CI: 1.42–4.91, P = 0.002) was an independent risk factor for progression of COVID-19 by multivariate Cox regression analyses. Moreover, some studies showed that higher levels of inflammatory cytokines, chemokines, and NLR were correlated with the severity of the disease (13, 47). Zhu et al. (48) demonstrated that high level of IL-6 and CRP were independent risk factors for assessing the severity of COVID-19.

		S	evere		No	nsever	е		Mean Difference	Mean Difference
-	Study or Subgroup	Mean	SD	Total	Mean	SD	Total	Weight	IV, Random, 95% 0	CI IV, Random, 95% CI
	Cai Q	35.69	30.24	58	10.95	13.59	240	7.1%	24.74 [16.77, 32.71	1 -
	Chen C	96.63	72.03	24	34.97	44.85	126	3.6%	61.66 [31.80, 91.52	.]
	Fang X	50.77	60.52	24	9.27	15.68	55	4.3%	41.50 [16.94, 66.06	s]
	Gao W	42.9	32.7	22	12.3	14.62	55	6.1%	30.60 [16.40, 44.80	n – – – – – – – – – – – – – – – – – – –
	Li D	27.1	33.36	22	5.27	0.64	18	6.2%	21.83 [7.89, 35.77	n
	Li D-2	62.38	43.66	17	15.73	19.24	63	4.8%	46.65 [25.36, 67.94	j — —
	Li K	90.57	68.2	25	13.85	21.15	58	3.9%	76.72 [49.44, 104.00	n — — — — — — — — — — — — — — — — — — —
	LiL	66.35	23.82	6	19.8	17.5	40	5.1%	46.55 [26.73, 66.37	n
	Liu W	39.33	42.84	11	15.2	23.64	67	4.1%	24.13 [-1.81, 50.07	n
	Lu Z	41.12	49.94	34	18.53	28.47	67	5.4%	22.59 [4.47, 40.71	i
	Peng Y	108.1	44.05	16	40.14	50.42	96	4.4%	67.96 [44.14, 91.78	n
	Qin C	60.67	61.32	286	33.7	38.51	166	6.9%	26.97 [17.76, 36.18	ı] ———
	Wan Q	62.23	47.92	21	17.25	21.05	132	4.9%	44.98 [24.17, 65.79	
	Wan S	93.33	64.28	40	13.57	21.98	95	5.0%	79.76 [59.36, 100.16	sj
	Wu C	91.62	85.2	84	29.28	38.39	117	5.2%	62.34 [42.84, 81.84	
	Xiang T	79.86	56.87	9	16.05	20.26	40	2.7%	63.81 [26.13, 101.49	j <u> </u>
	Yang A	53.9	60.1	24	20.1	24.5	69	4.3%	33.80 [9.07, 58.53	j
	Zhang J	51.77	50.6	56	30.1	32.15	82	6.0%	21.67 [6.70, 36.64	.]
	Zhang W	51.9	43.8	9	21.2	25.7	56	3.6%	30.70 [1.30, 60.10	ı
	Zheng F	52.03	36.04	30	15.37	14.32	131	6.3%	36.66 [23.53, 49.79	
	Total (95% CI)			818			1773	100.0%	41.02 [33.32, 48.71]	•
	Heterogeneity: Tau ² =	199.63;	Chi ² = 7	0.01, d	f = 19 (P < 0.0	0001); I	² = 73%		-100 -50 0 50 100
	Test for overall effect:	Z = 10.4	5 (P < 0	.00001)					Favours [Nonsevere] Favours [Severe]
5										
								Od	ds Ratio	Odds Ratio
-	Study or Subgrou	p lo	g[Odds	s Ratio	0	SE V	Veight	IV,	Fixed, 95% CI	IV, Fixed, 95% CI
	Dong L			-0.105	4 0.64	412	9.5%	0.9	0 [0.26, 3.16]	
	Guan W			1.224			70.5%		0 [2.15, 5.40]	
	Li K			2.580			6.4%		[2.85, 61.24]	· · · · · · · · · · · · · · · · · · ·
	Liu W									
					8 0.82		5.7%		[1.18, 30.41]	
	Zhang J			1.197			6.1%		[0.69, 15.96]	-
	Zheng F			3.295	8 1.43	382	1.9%	27.00 [1.61, 452.42]	
	Total (95% CI)					1	00.0%	3.5	1 [2.38, 5.16]	•
	Heterogeneity: Chi ²	= 9.82	df = 5	(P = 0)	.08); 12	2 = 49%	6			
	Test for overall effe					,	-		0.01	
				- 0.00						Favours [Nonsevere] Favours [Severe]

TABLE 3 | The association between inflammation-related indicators and disease severity in patients with COVID-19.

Inflammation-related indicators	Number of studies	Participants	Mean difference/Std. mean difference (95% CI)	P	Effects model	Heterogeneity		
						l ²	P _h	
Procalcitonin	16	2,070	0.78 (0.34, 1.22)	<0.001	REM	93	<0.001	
C-reactive protein	20	2,591	41.02 (33.32, 48.71)	< 0.001	REM	73	< 0.001	
Erythrocyte sedimentation rate	5	706	18.37 (6.59, 30.15)	0.002	REM	84	<0.001	
Interleukin-6	3	951	0.72 (0.13, 1.30)	0.02	REM	94	< 0.001	
Neutrophil-lymphocyte ratio	6	1,141	0.85 (0.56, 1.15)	<0.001	REM	67	0.009	

Cl, confidence intervals; REM, random-effects model; Ph, p-value of Q-test for heterogeneity.



adjusted funnel plot of mean difference in neutrophil count. (E) Funnel plot of mean difference in lymphocyte count. (F) Funnel plot of Std. mean difference in PCT. (G,H) Funnel plot and adjusted funnel plot of mean difference in CRP.

This meta-analysis was conducted to assess the association between immune-inflammatory parameters and increased disease severity in COVID-19 patients. In the present study, we firstly utilized the available data from 25 included studies with a total of 4,278 patients to obtain the pooled results to evaluate the difference in WBC count between a severe and non-severe group. We found that the white blood cell counts of severe patients tended to be higher than that of less severe patients. Next, the effect of WBC count on the risk of developing severe COVID-19 was further examined. The pooled results statistically supported the conclusions that elevated WBC count was strongly associated with the deterioration of disease in COVID-19 patients (OR = 2.92, 95% CI: 1.96–4.35, *P* < 0.001). Similarly, the number of neutrophils was found to be higher in severe COVID-19 (MD = 1.50, 95% CI: 1.01-1.98, P < 0.001). However, patients with serious disease tend to have a reduced total lymphocyte count as well as virous lymphocyte subtypes count compared to non-severe COVID-19 patients. In addition, COVID-19 patients who presented with lymphopenia had an \sim 5-fold higher risk of developing severe disease (OR = 4.97, 95% CI: 3.53–6.99, P < 0.001). Consistent with previous reports, this meta-analysis also indicated that the incidence of high NLR had significant association with illness severity (MD = 0.85, 95% CI: 0.56-1.15, P < 0.001) and elevated NLR could act as a predictor for exacerbation of COVID-19 (OR = 2.50, 95% CI: 2.04–3.06, P < 0.001). Moreover, other common immune-inflammatory parameters, such as PCT and CRP, could also predict the deterioration of disease (PCT: OR = 6.33, 95%CI: 3.97–10.10, *P* < 0.001; CRP: OR = 3.51, 95% CI: 2.38–5.16, P < 0.001). In this study, ESR and IL-6 were also found to be correlated with increased disease severity in COVID-19 patients.

SARS-CoV-2 infection can activate innate and adaptive immune responses. A rapid and well-coordinated innate immune response is the first line of defense against viral infections, while uncontrolled inflammatory innate responses and impaired adaptive immune responses may lead to harmful tissue damage, both locally and systemically (49). Taking both the levels of neutrophil and lymphocyte into account, NLR may be a better biomarker for systemic inflammation and illness severity than single neutrophil or lymphocyte count. In this study, we confirmed that an increase in NLR usually indicated higher disease severity with more clinical evidence. The following reasons may explain the finding. On one hand, most patients with severe COVID-19 exhibit substantially elevated serum levels of proinflammatory cytokines. Meanwhile, neutrophils can be triggered by virus-related inflammatory factors produced by lymphocyte and endothelial cells (such as interleukin-6 and interleukin-8, tumor necrosis factor-alpha and granulocyte colony stimulating factor, and interferon-gamma) (50). The triggered neutrophils produce reactive oxygen species (ROS) and other cytotoxic mediators, which may dampen the virus infection (51). Moreover, neutrophils are able to release neutrophil extracellular traps (NETs), which are a sticky web of DNA conjugated with antimicrobial enzymes (such as myeloperoxidase and histones), resulting in the capture and the killing of different pathogens, including viruses (52-54). On the other hand, lymphocytes did not show a significant decrease in the early stage of viral infection but significantly decreased in severe and critically ill patients (34). The main reasons for the decrease or exhaustion of lymphocyte in severe cases may be the following: viruses attack target cells and directly damage cells; viral infection causes immune cells to enter an activated state and participate in the anti-viral process, resulting in severe damage and apoptosis; systematic inflammation stimulates the production of neutrophil and speed up the apoptosis of lymphocyte; severe COVID-19 patients tend to present an increase of PCT and CRP, indicating a potential bacterial co-infection; and bacterial co-infection or superinfection might affect the immune response.

Although the sample size of this study is large (4,911 COVID-19 cases from 29 clinical studies enrolled), some limitations should be noted meanwhile. Firstly, the primary research design of the studies included in this study were retrospective cohorts with insufficient demonstration ability, limiting their ability to infer definitive causality. Secondly, all included original clinical cohort studies were conducted in China, which limits the ability of this study to extrapolate other patient populations in other countries. Thirdly, the high statistic heterogeneity could be found in calculating pooled MD, which may relate to large variation among studies in the sample size.

In summary, we came to the cautious conclusion that immune-inflammatory parameters such as WBC, lymphocyte, NLR, PCT, and CRP were correlated with disease severity and could be used as potentially important risk factors for disease progression. In addition, increased NLR levels reflecting an enhanced inflammatory process may also suggest a poor prognosis. Therefore, surveillance of immune-inflammatory parameters, especially NLR, may be helpful in the diagnosis, early screening and predicting of severe illness, and treatment of COVID-19.

DATA AVAILABILITY STATEMENT

All datasets generated for this study are included in the article/**Supplementary Material**.

AUTHOR CONTRIBUTIONS

XF, BC, MX, and GC conceived and designed the study. XF, SL, and QS selected the studies, collected the data, and drafted and revised the paper. XF and JZ analyzed data. All authors interpreted the results, read, and approved the final version of the manuscript.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fmed. 2020.00301/full#supplementary-material

REFERENCES

- Chen N, Zhou M, Dong X, Qu J, Gong F, Han Y, et al. Epidemiological and clinical characteristics of 99 cases of 2019 novel coronavirus pneumonia in Wuhan, China: a descriptive study. *Lancet.* (2020) 395:507–13. doi: 10.1016/S0140-6736(20)30211-7
- Xu Z, Shi L, Wang Y, Zhang J, Huang L, Zhang C, et al. Pathological findings of COVID-19 associated with acute respiratory distress syndrome. *Lancet Respir Med.* (2020) 8:420–2. doi: 10.1016/S2213-2600(20) 30076-X
- Li Q, Guan X, Wu P, Wang X, Zhou L, Tong Y, et al. Early transmission dynamics in Wuhan, China, of novel coronavirus-infected pneumonia. *N Engl* J Med. (2020) 382:1199–207. doi: 10.1056/NEJMoa2001316
- Rodríguez-Morales AJ, MacGregor K, Kanagarajah S, Patel D, Schlagenhauf P. Going global - travel and the 2019 novel coronavirus. *Travel Med Infect Dis.* (2020) 33:101578. doi: 10.1016/j.tmaid.2020.101578
- Wang C, Horby PW, Hayden FG, Gao GF. A novel coronavirus outbreak of global health concern. *Lancet.* (2020) 395:470– 3. doi: 10.1016/S0140-6736(20)30185-9
- Wang X, Fang X, Cai Z, Wu X, Gao X, Min J, et al. Comorbid chronic diseases and acute organ injuries are strongly correlated with disease severity and mortality among COVID-19 patients: a systemic review and meta-analysis. *Research.* (2020) 2020:1–17. doi: 10.34133/2020/2402961
- Zhu N, Zhang D, Wang W, Li X, Yang B, Song J, et al. A novel coronavirus from patients with pneumonia in China, 2019. N Engl J Med. (2020) 382:727– 33. doi: 10.1056/NEJMoa2001017
- Qin C, Zhou L, Hu Z, Zhang S, Yang S, Tao Y, et al. Dysregulation of immune response in patients with COVID-19 in Wuhan, China. *Clin Infect Dis.* (2020) 58:7250–7. doi: 10.2139/ssrn.3541136
- Slim K, Nini E, Forestier D, Kwiatkowski F, Panis Y, Chipponi J. Methodological index for non-randomized studies (Minors): Development and validation of a new instrument. ANZ J Surg. (2003) 73:712– 6. doi: 10.1046/j.1445-2197.2003.02748.x
- Stroup DF, Berlin JA, Morton SC, Olkin I, Williamson GD, Rennie D, et al. Meta-analysis of observational studies in epidemiology: a proposal for reporting. Meta-analysis Of Observational Studies in Epidemiology (MOOSE) group. *JAMA*. (2000) 283:2008–12. doi: 10.1001/jama.283.1 5.2008
- Wan X, Wang W, Liu J, Tong T. Estimating the sample mean and standard deviation from the sample size, median, range and/or interquartile range. BMC Med Res Methodol. (2014) 14:1–13. doi: 10.1186/1471-2288-14-135
- Bin Y, Pan J, Liang X, Liu G, Zhang J. Clinical characteristics of 55 hospitalized patients with COVID-19 in Wuhan, China. J Guanxi Med Univ. (2020) 37:338–42. doi: 10.16190/j.cnki.45-1211/r.2020.02.034
- Huang C, Wang Y, Li X, Ren L, Zhao J, Hu Y, et al. Clinical features of patients infected with 2019 novel coronavirus in Wuhan, China. *Lancet.* (2020) 395:497–506. doi: 10.1016/S0140-6736(20)30183-5
- Li D, Long Y, Huang P, Guo W, Wu S, Zhou Q, et al. Clinical characteristics of 80 patients with COVID-19 in Zhuzhou City. *Chin J Infect Control.* (2020) 3:227–33. doi: 10.12138/j.issn.1671-9638.20206514
- Li D, Wang M, He B, Xu Y, Zhou X, Li W, et al. Laboratory test analysis of sixty-two COVID-19 patients. *Med J Wuhan Univ.* (2020). doi: 10.14188/j.1671-8852.2020.0220. [Epub ahead of print].
- Li K, Wu J, Wu F, Guo D, Chen L, Fang Z, et al. The clinical and chest CT features associated with severe and critical COVID-19 pneumonia. *Invest Radiol.* (2020) 55:327–31. doi: 10.1097/RLI.000000000000672
- Li L, Yang H, Gou C, Wang X, Luo X, Zhang J, et al. Traditional Chinese medicine characteristics of 46 patients with COVID-19. J Cap Med Univ. (2020) 41:155–60. doi: 10.3969/j.issn.1006-7795.2020.02.002
- Liu W, Tao Z-W, Lei W, Ming-Li Y, Kui L, Ling Z, et al. Analysis of factors associated with disease outcomes in hospitalized patients with 2019 novel coronavirus disease. *Chin Med J.* (2020) 133:1032–8. doi: 10.1097/CM9.0000000000 00775
- Long L, Zeng X, Zhang X, Xiao W, Guo E, Zhan W, et al. Short-term outcomes of coronavirus disease 2019 and risk factors for progression. *Eur Respir J.* (2020) 55:2000990. doi: 10.1183/13993003.00990-2020

- Lu Z, He R, Jiang W, Fan T, Geng Q. Clinical characteristics and immune function analysis of COVID-19. *Med J Wuhan Univ.* (2020) 41:529–32. doi: 10.14188/j.1671-8852.2020.0126
- Peng YD, Meng K, Guan HQ, Leng L, Zhu RR, Wang BY, et al. Clinical characteristics and outcomes of 112 cardiovascular disease patients infected by 2019-nCoV. *Zhonghua Xin Xue Guan Bing Za Zhi.* (2020) 48:E004. doi: 10.3760/cma.j.cn112148-20200220-00105
- Wan Q, Shi A, He T, Tang L. Analysis of clinical features of 153 patients with novel coronavirus pneumonia in Chongqing. *Chin J Clin Infect Dis.* (2020) 1:16–20. doi: 10.3760/cma.j.cn115673-20200212-00030
- Cai Q, Huang D, Ou P, Yu H, Zhu Z, Xia Z, et al. COVID-19 in a designated infectious diseases hospital outside Hubei Province, China. *Allergy*. (2020). doi: 10.1111/all.14309. [Epub ahead of print].
- Wan S, Xiang Y, Fang W, Zheng Y, Li B, Hu Y, et al. Clinical features and treatment of COVID-19 patients in Northeast Chongqing. *J Med Virol.* (2020). doi: 10.1002/jmv.25783. [Epub ahead of print].
- Wang D, Hu B, Hu C, Zhu F, Liu X, Zhang J, et al. Clinical characteristics of 138 hospitalized patients with 2019 novel coronavirus-infected pneumonia in Wuhan, China. JAMA. (2020) 323:1061–9. doi: 10.1001/jama.2020.1585
- 26. Wu C, Chen X, Cai Y, Xia J, Zhou X, Xu S, et al. Risk factors associated with acute respiratory distress syndrome and death in patients with coronavirus disease 2019 pneumonia in Wuhan, China. *JAMA Intern Med.* (2020). doi: 10.1001/jamainternmed.2020.0994. [Epub ahead of print].
- Yang AP, Liu J, Tao W, Li H-m. The diagnostic and predictive role of NLR, d-NLR and PLR in COVID-19 patients. *Int Immunopharmacol.* (2020) 84:106504. doi: 10.1016/j.intimp.2020.106504
- Yuan J, Sun Y, Zou Y, Chen T, Cao Q, Yuan G, et al. A retrospective analysis of the clinical characteristics of 223 NCP patients in Chongqing. J Southwest Univ Sci Ed. (2020) 42:17–24. doi: 10.13718/j.cnki.xdzk.2020.03.003
- Zhang J-J, Dong X, Cao Y-Y, Yuan Y-D, Yang Y-B, Yan Y-Q, et al. Clinical characteristics of 140 patients infected with SARS-CoV-2 in Wuhan, China. *Allergy*. (2020). doi: 10.1111/all.14238. [Epub ahead of print].
- Zhang W, Hou W, Li T, Li A, Pan W, Jin R, et al. Clinical characteristics of 74 hospitalized patients with COVID-19. J Cap Med Univ. (2020) 41:161–7. doi: 10.3969/j.issn.1006-7795.2020.02.003
- Zheng F, Tang W, Li H, Huang Y-X, Xie Y-L, Zhou Z-G. Clinical characteristics of 161 cases of corona virus disease 2019 (COVID-19) in Changsha. *Eur Rev Med Pharmacol Sci.* (2020) 24:3404–10. doi: 10.26355/eurrev_202003_20711
- Xiang T, Liu J, Xu F, Cheng N, Liu Y, Qian K, et al. Analysis of clinical characteristics of 49 patients with coronavirus disease 2019 in Jiangxi. *Chinese* J Respir Crit Care Med. (2020) 19:154–60. doi: 10.7507/1671-6205.202002070
- 33. Chen C, Chen C, Yan JT, Zhou N, Zhao JP, Wang DW. Analysis of myocardial injury in patients with COVID-19 and association between concomitant cardiovascular diseases and severity of COVID-19. *Zhonghua Xin Xue Guan Bing Za Zhi*. (2020) 48:E008. doi: 10.3760/cma.j.cn112148-20200225-00123
- 34. Chen X, Tong J, Xiang J, Hu J. Retrospective study on the epidemiological characteristics of 139 patients with novel coronavirus pneumonia on the effects of severity. *Chongqing Med.* (2020). [Epub ahead of print].
- 35. Chen X, Ou J, Huang Y, Tan M, Chen J, Lin L, et al. Diagnostic roles of several parameters in corona virus disease 2019. *Lab Med.* (2020). [Epub ahead of print].
- Fang X, Mei Q, Yang T, Zhang L, Yang Y, Wang Y, et al. Clinical characteristics and treatment strategies of 79 patients with COVID-19. *Chin Pharmacol Bull.* (2020) 36:453–9. doi: 10.3969/j.issn.1001-1978.2020. 04.002
- 37. Gao W, Yang X, Zhang XQ, Su X, Dou AH, Hui W, et al. Clinical characteristic of 90 patients with COVID-19 hospitalized in a Grade-A hospital in Beijing. *Acad J Chin PLA Med Sch.* (2020). [Epub ahead of print].
- Guan W-J, Ni Z-Y, Hu Y, Liang W-H, Ou C-Q, He J-X, et al. Clinical characteristics of coronavirus disease 2019 in China. N Engl J Med. (2020). doi: 10.1101/2020.02.06.20020974. [Epub ahead of print].
- Hou K, Zhang N, Li T, Zhou M, Shi X, Zhao G. CT features of corona virus disease 2019 (COVID-19) in different stages and its correlation with neutrophil-lymphocyte(NLR) and T lymphocyte subsets. *Radiol Pract.* (2020) 35:272–6. doi: 10.13609/j.cnki.1000-0313.2020.03.006
- 40. Guthrie GJK, Charles KA, Roxburgh CSD, Horgan PG, McMillan DC, Clarke SJ. The systemic inflammation-based neutrophil-lymphocyte ratio:

experience in patients with cancer. Crit Rev Oncol Hematol. (2013) 88:218-30. doi: 10.1016/j.critrevonc.2013.03.010

- Meisner M. Update on Procalcitonin Measurements. Ann Lab Med. (2014) 34:263. doi: 10.3343/alm.2014.34.4.263
- Duval S, Tweedie R. Trim and fill: a simple funnel-plot-based method of testing and adjusting for publication bias in meta-analysis. *Biometrics*. (2000) 56:455–63. doi: 10.1111/j.0006-341X.2000.00455.x
- Xia X, Wu J, Liu H, Xia H, Jia B, Huang W. Epidemiological and initial clinical characteristics of patients with family aggregation of COVID-19. *J Clin Virol.* (2020) 127:104360. doi: 10.1016/j.jcv.2020.104360
- Channappanavar R, Perlman S. Pathogenic human coronavirus infections: causes and consequences of cytokine storm and immunopathology. *Semin Immunopathol.* (2017) 39:529–39. doi: 10.1007/s00281-017-0629-x
- 45. Min CK, Cheon S, Ha NY, Sohn KM, Kim Y, Aigerim A, et al. Comparative and kinetic analysis of viral shedding and immunological responses in MERS patients representing a broad spectrum of disease severity. *Sci Rep.* (2016) 6:25359. doi: 10.1038/srep25359
- Mo P, Xing Y, Xiao Y, Deng L, Zhao Q, Wang H, et al. Clinical characteristics of refractory COVID-19 pneumonia in Wuhan, China. *Clin Infect Dis.* (2020). doi: 10.1093/cid/ciaa270. [Epub ahead of print].
- Liu Y, Du X, Chen J, Jin Y, Peng L, Wang HHX, et al. Neutrophil-tolymphocyte ratio as an independent risk factor for mortality in hospitalized patients with COVID-19. *J Infect.* (2020). doi: 10.1016/j.jinf.2020.04.002. [Epub ahead of print].
- Zhu Z, Cai T, Fan L, Lou K, Hua X, Huang Z, et al. Clinical value of immune-inflammatory parameters to assess the severity of coronavirus disease 2019. *Int J Infect Dis.* (2020) 592:124530. doi: 10.1016/j.ijid.2020. 04.041
- Cao X. COVID-19: immunopathology and its implications for therapy. Nat Rev Immunol. (2020) 20:269–70. doi: 10.1038/s41577-020-0308-3

- Actor JK, Actor JK. Cells and organs of the immune system. *Elsevier's Integr Rev Immunol Microbiol.* (2012). doi: 10.1016/B978-0-323-07447-6.00002-8. [Epub ahead of print].
- Agraz-Cibrian JM, Giraldo DM, Mary FM, Urcuqui-Inchima S. Understanding the molecular mechanisms of NETs and their role in antiviral innate immunity. *Virus Res.* (2017) 228:124– 33. doi: 10.1016/j.virusres.2016.11.033
- Brinkmann V, Reichard U, Goosmann C, Fauler B, Uhlemann Y, Weiss DS, et al. Neutrophil extracellular traps kill bacteria. *Science*. (2004) 303:1532– 5. doi: 10.1126/science.1092385
- Muraro SP, De Souza GF, Gallo SW, Da Silva BK, De Oliveira SD, Vinolo MAR, et al. Respiratory syncytial virus induces the classical ROS-dependent NETosis through PAD-4 and necroptosis pathways activation. *Sci Rep.* (2018) 8:1–12. doi: 10.1038/s41598-018-32576-y
- Hiroki CH, Toller-Kawahisa JE, Fumagalli MJ, Colon DF, Figueiredo LTM, Fonseca BALD, et al. Neutrophil extracellular traps effectively control acute chikungunya virus infection. *Front Immunol.* (2020) 10:3108. doi: 10.3389/fimmu.2019.03108

Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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