

CRITICAL ROLE OF INFLAMMATORY BIOCHEMICAL MARKERS DURING CORONAVIRUS DISEASE 2019:- A SYSTEMATIC REVIEW AND META-ANALYSIS

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Received : 19/03/2023
Received in revised form : 15/04/2023
Accepted : 28/04/2023

Keywords:
systematic literature review; meta-analysis; coronavirus; inflammatory biomarkers.

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DOI: 10.47009/jamp.2023.5.3.143

Source of Support: Nil,
Conflict of Interest: None declared

Int J Acad Med Pharm
2023; 5 (3); 688-696



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Abstract

Background: Biomarkers play an important role in clinical decision making in infectious diseases. The covid 19 pandemic caused by SARS-CoV-2 virus has impacted the whole world. The objective of this study is to conduct a systematic review and meta analysis to evaluate the association between select inflammatory biomarkers and outcome in covid 19 patients. This study seeks to inform subsequent research by statistically summarizing the levels of these biomarkers, in patients of covid 19 with varied severity. **Materials and Methods:** A meta-analysis and systematic literature review were carried out. The search focused on research papers that reported the laboratory results of COVID-19-positive patients and included the names of routine UK NHS laboratory tests. For each biomarker, a random effects meta-analysis of the standard deviation between the COVID-19-positive and -negative groups was carried out. Interleukin-6, ferritin, C-Reactive Protein, procalcitonin and D-dimer were taken from the studies as laboratory parameters. **Result:** Using the laboratory data, mean differences were estimated using meta-analyses with associated confidence intervals of 95 percent. Increased levels of interleukin-6, ferritin, C-Reactive Protein, procalcitonin and D-dimer were directly correlated to severe and fatal COVID-19 cases. **Conclusion:** Interleukin-6, ferritin, C-Reactive Protein, procalcitonin and D-dimer are important laboratory parameters in diagnosing coronavirus cases. Further exploration is expected to distinguish whether routine lab biomarkers can be utilized in the improvement of a clinical scoring framework to help with emergency of patients.

INTRODUCTION

The severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) has spread rapidly across the globe, resulting in significant morbidity and mortality.^[1] Without effective contact tracing, social distancing, lockdowns, or vaccination, the number of cases worldwide continues to rise and is likely to continue rising further. Fever, cough, sore throat, headache, fatigue, myalgia, breathlessness, anosmia, and ageusia are all common signs of the disease.^[2] It has been difficult for patients with COVID-19 to be identified in the laboratory. Additionally, a second or third infection peak is currently occurring in a number of nations,^[3] putting a significant strain on testing facilities in numerous locations. On nasal samples, reverse transcription polymerase chain reaction (RT-PCR),^[4] is the primary test used to support the diagnosis. Impediments of RT-PCR are

the somewhat lengthy time required to circle back as well as defective awareness.^[5]

Routine research facility biomarkers can give a general image of the wellbeing status of a patient in intense clinical settings. However, there are no routine laboratory biomarkers that can be used as a standalone diagnostic test or to assist physicians in determining which patients should receive treatment first.^[6] To aid in the diagnosis of COVID-19, some attempts have been made to combine a number of biomarkers and other parameters into a clinical scoring algorithm.^[7] However, these models are frequently poorly reported, have a high risk of bias due to poor reporting, poor methodological conduct, and lack robust validation. Involving these models by and by may bring about execution qualities that are lower than those announced in the writing.^[8]

This study sought to inform subsequent research by statistically summarizing routine laboratory biomarker measurements in COVID-19-positive and

-negative patients. Clinical scoring algorithms that are likely to be important for use in clinical settings that do not readily have access to COVID-19 point-of-care (POC) or laboratory testing could benefit from this research.

MATERIALS AND METHODS

Protocol

We adhered to the Preferred Reporting Items for Systematic Reviews and Meta-Analysis (PRISMA) and Cochrane Collaboration Handbook recommendations.^[9,10]

Search Strategy

Adapting the search to the specifics of each database, we used relevant descriptors and synonyms to search MEDLINE, EMBASE, LILACS, IBECs, and the Cochrane Central Register of Controlled Trials (CENTRAL). To find published, ongoing, and unpublished studies, we also searched the Open Grey database, the World Health Organization International Clinical Trials Registry Platform (WHO ICTRP), and ClinicalTrials.gov. Finally, we searched the references lists of the included studies using the snowballing method.

Eligibility Criteria and Study Selection

We included in the electronic search were all studies published in 2020 with no language restrictions. According to the National Health Commission of China (NHCC) Guidelines for Diagnosis and Management of COVID-19 or World Health Organization Interim Guidance for COVID-19,^[11,12] studies were included if they presented laboratory data, such as serum IL-6 levels, from mild-to-moderate, severe, or critical COVID-19 patients. Rejection standards consisted of studies that evaluated pregnant ladies, pediatric patients, people co-tainted with different microorganisms, or populaces solely connected with oncological, rheumatological, transfers, or persistent renal infection.

Data Extraction and Quality Appraisal

To collect data from the included studies, we used a form that was already set up. We specifically extracted the following characteristics from the studies and participants: age, gender, the severity of the condition, the diagnostic criteria, the number of participants who were screened, randomized, analyzed, excluded, lost to follow-up, and dropped out, the setting, the length of the study, laboratory biomarkers, outcome measures, and time points that were reported are all important factors. The modified Newcastle-Ottawa Scale (NOS) was then used to evaluate each study's inherent bias risk.^[13] When any study data or other details were missing, we attempted to get in touch with the studies' authors.

Summary Measures and Synthesis of Results

Concentrate on selection, information extraction, and evaluation of the selected studies were

performed independently by two survey creators. Discussion or, if necessary, consulting a third author was used to resolve disagreements. For the purpose of analysis, the studies were divided into two distinct groups: mortality and severity groups. IL-6, ferritin, CRP, procalcitonin, and D-dimer were analyzed in the laboratory. Between studies, the assays for detecting laboratory tests were comparable.

Statistical analysis

We performed the meta-analysis by utilizing the R software's "meta" package (version 3.4.1) following the extraction of the data.^[14] Following Hozo SP et al.'s method, we estimated the means and standard deviations for studies with continuous data presented as medians and interquartile ranges.^[15] The mean difference (MD) and 95% confidence interval (CI) for each of the laboratory parameters related to patients with confirmed COVID-19, whether they had severe disease or not, non-survivors, and survivors were calculated in a meta-analysis. The I² statistic and the chi-square (χ^2) test were used to evaluate heterogeneity among the included studies. Pooled results were calculated using a random effects model. When I² was greater than 50%, a sensitivity analysis was conducted to investigate the causes of heterogeneity. By excluding studies with unclear timepoints of collection and those that collected blood parameters seven days after hospital admission, we investigated the causes of heterogeneity. Egger's test was used to look for publication bias in meta-analyses that included at least ten studies (S2A and S2B Table).

RESULTS

Literature Retrieval

The search identified 1250 articles. After duplicate removal, title, abstract and full-text screening, 40 studies were included in this systematic review and meta-analysis.

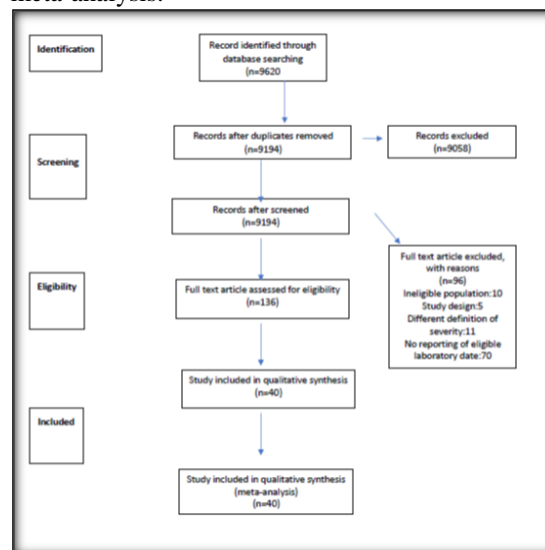


Figure 1: Study selection process flow chart based on Prisma

Characteristics of Included Studies

Most of the research (sixty percent) was carried out in China. Till January 2021, 20 % of the articles were still in pre-print, despite being published articles. Articles were either distributed or submitted between February 2020 to September 2020, and generally announced information were gathered between December 2019 to June 2020. The sample size was 340 patients on average (SD = 609.1), and

45% of those patients were found to be COVID-19 positive (SD = 22.48%). However, different definitions of COVID-19-positive patients were used in different studies. The official guidelines that were in place at the time of testing in the nation where the data were collected served as the foundation for the COVID-19 diagnoses. Instead of a composite reference standard, RT-PCR was typically used as the COVID-19 reference standard.

Table 1: Baseline and demographic characteristics of the included studies. Assessed outcome: Severity

Study ID	Severity criteria	Total enrolled patients	Severe group				Non-Severe group			
			No. (%)	Age ω	Male (%)	Comorbidity (%)	No. (%)	Age ω	Male (%)	Comorbidity (%)
Guang Chen, 2020, ^[16]	Severe vs Moderate cases according to NHCC COVID-19 Guideline (6th Edition) ^a	21	11 (52.38)	61.0 (56.5–66.0)	10 (90.9)	5 (45.5)	10 (47.62)	52.0 (42.8–56.0)	7 (70.0)	2 (20.0)
Yong Gao, 2020, ^[17]	Severe vs Mild cases according to WHO Interim Guidance for COVID-19c	43	15 (53.88)	45.20 (\pm 7.68)	9 (60)	$\omega\omega$	28 (46.12)	42.96 (\pm 14.00)	17 (60.71)	$\omega\omega$
Zhongliang Wang, 2020, ^[18]	Patients with SpO ₂ <90% (Severe) vs Patients with SpO ₂ >90% (Non-severe) according to NHCC COVID-19 Guidelines (3rd edition) ^a	69	14 (20.29)	70.5 (62.0–77.0)	7 (50)	$\omega\omega$	55 (79.71)	37.0 (32.0–51.0)	25 (45)	$\omega\omega$
Chuan Qin, 2020, ^[19]	Severe vs Moderate cases according to NHCC COVID-19 Guidelines (5th Edition) ^a	452	286 (63.27)	61 (51–69)	155 (54.2)	146 (51)	166 (36.73)	53 (41.25–62)	80 (48.2)	55 (33.1)
Chen Lei, 2020, ^[20]	Severe/Critical vs Mild cases according to NHCC COVID-19 Guidelines (4th Edition) ^a	29	14 (48.28)	NR	NR	$\omega\omega$	15 (51.72)	NR	NR	$\omega\omega$
Ruirui Wang, 2020, ^[21]	Critical (Severe or critical cases) vs Non-critical (mild or moderate) cases according to NHCC COVID-19 Guidelines (5th Edition) ^a	125	25 (20)	49.40 (\pm 13.64)	16 (64)	12 (48)	100 (80)	39.47 (\pm 14.84)	55 (55)	22 (22)
Zhe Zhu, 2020, ^[22]	Severe vs Non-severe cases according to NHCC COVID-19 Guidelines (6th Edition) ^a	127	16 (12.6)	57.50 (\pm 11.70)	9 (56.25)	12 (75)	111 (87.4)	49.95 (\pm 15.52)	73 (65.77)	40 (36.04)
Xiaohua Chen, 2020, ^[23]	Moderate vs Severe vs Critically ill cases according to NHCC COVID-19 Guidelines (6th Edition) ^a	48	Severe: 10 (20.83) Critically ill: 17 (35.42)	Severe: 63.9 (\pm 15.2) Critically ill: 79.6 (\pm 12.6)	Severe: 9 (90) Critically ill: 15 (88.2)	$\omega\omega$	21 (43.75)	52.8 (\pm 14.2)	13 (61.9)	$\omega\omega$
Ming Ding, 2020, ^[24]	Mild vs Severe vs Critical cases according to NHCC COVID-19 Guidelines (7th Edition) ^a	32	Severe: 10 (31.25) Critical: 11 (34.37)	Severe: 61.3 (\pm 17.9) Critical: 73.5 (\pm 12.3)	Severe: 5 (50) Critical: 7 (63.63)	Severe: 3 (30) Critical: 4 (36.36)	11 (34.37)	54.9 (\pm 11.3)	1 (9.09)	3 of 5 (60)
Chen LD, 2020, ^[25]	Mild (without pneumonia) vs Moderate cases with pneumonia (Non-severe) vs Severe cases with pneumonia according to NHCC COVID-19	106	25 (23.6)	60.68 (\pm 15.23)	15 (60)	11 (44.0)	Mild: 12 (11.3) Moderate: 69 (65.1)	Mild: 43.92 (\pm 13.73) Moderate: 51.41 (\pm 15.77)	Mild: 4 (33.3) Moderate: 34 (49.3)	Mild: 0 (0.0) Moderate: 14 (20.3)

	Guidelines (7th Edition)a									
Chen R, 2020, ^[26]	Mild/Moderate cases vs Severe cases according to NHCC COVID-19 Guidelines (7th Edition)a	548	Severe: 155 (28.3) Critical: 48 (8.8)	Severe: 60.9 (\pm 13.8) Critical: 61.4 (\pm 13.6)	Severe: 93 (60) Critical: 38 (79.2)	$\omega\omega$	345 (62.9)	67.3 (\pm 12.1)	182 (52.75)	$\omega\omega$
Chi Y, 2020, ^[27]	Mild vs Moderate vs Severe cases according to NHCC COVID-19 Guidelines (5th Edition)a	66	8 (12.1)	54.0 (\pm 12.38)	5 (62.5)	4 (50)	Mild: 22 (33.4) Moderate: 36 (54.5)	Mild: 43.32 (\pm 18.38) Moderate: 40.81 (\pm 11.8)	Mild: 13 (59.1) Moderate: 19 (53)	Mild: 6 (27) Moderate: 8 (22)
Hu ZI, 2020, ^[28]	Severe vs Non-severe cases according to NHCC COVID-19 Guidelines (7th Edition)a	76	13 (17.2)	61.5 (57.1–65.9)	8 (61.5)	$\omega\omega$	63 (82.8)	48.2 (46.0–50.4)	26 (41.3)	$\omega\omega$
Huang Z, 2020, ^[29]	Moderate (Non-severe) vs Severe vs Critical cases according to NHCC COVID-19 Guidelines (7th Edition)a	83	Severe: 29 (35) Critical: 33 (39.7)	Severe: 67 (60–79) Critical: 58 (49–62)	Severe: 16 (55.2) Critical: 26 (78.8)	Severe: 20 (69) Critical: 21 (63.6)	21 (25.3)	68 (57–69)	12 (57.14)	9 (42.86)
Li X, 2020, ^[30]	Severe (Severe pneumonia/ARDS) vs Non-severe cases (Mild/ Common pneumonia) according to NHCC COVID-19 Guidelines (7th Edition)a and WHO Interim Guidance for COVID-19c	215	56 (26.1)	56.5 (20–72)	36 (64.3)	$\omega\omega$	159 (73.9)	44 (32–52)	91 (57.2)	$\omega\omega$
Liu D, 2020, ^[31]	Moderate vs Severe vs Critical cases according to NHCC COVID-19 Guidelines (7th Edition)a	2044	Severe: 689 (33.7) Critical: 268 (13.1)	Severe: 64.0 (54.0–71.0) Critical: 69.0 (62.0–77.0)	Severe: 349 (50.65) Critical: 176 (65.67)	Severe: 423/687 (61.57) Critical: 212/266 (79.7)	1087 (53)	59 (46–67)	475 (43.7)	540/1086 (49.72)
Ozsurekci Y, 2020, ^[32]	Mild vs Moderate vs Severe/ Critical cases according to WHO Interim Guidance for COVID-19c	30	11 (36.7)	NR	NR	$\omega\omega$	Mild: 4 (13.4) Moderate: 15 (50)	NR	NR	$\omega\omega$
Xu X, 2020, ^[33]	Moderate (non-severe) vs Severe vs Critically ill cases according to NHCC COVID-19 Guidelines (7th Edition)a	88	Severe: 32 (36.4) Critically ill: 9 (10.2)	Severe: 59.94 (\pm 13.96) Critically ill: 74.78 (\pm 10.06)	Severe: 8 (25) Critically ill: 7 (77.78)	Severe: 17 (53.13) Critically ill: 7 (77.78)	47 (53.4)	52.49 (\pm 14.62)	21 (44.68)	17 (36.17)
Zeng YL, 2020, ^[34]	Ordinary (Non-severe) vs Severe vs Critical cases according to NHCC COVID-19 Guidelines (6th Edition)a	49	Severe: 16 (32.7) Critical: 5 (10.2)	Severe: 60 (\pm 16) Critical: 68 (\pm 20)	Severe: 8 (50) Critical: 3 (60)	$\omega\omega$	28 (57.1)	46 (\pm 19)	15 (53.6)	$\omega\omega$
Zeng Z, 2020, ^[35]	Moderate (Non-severe) vs Severe / Critical cases according to NHCC COVID-19 Guidelines (6th Edition)a	317	Severe: 167 (52.68) Critical: 57 (17.98)	Severe: 62.0 (51.0–69.0) Critical: 68.0 (57.0–77.0)	Severe: 90 (53.9) Critical: 31 (54.4)	$\omega\omega$	93 (29.34)	59.0 (46.0–68.5)	41 (44.1)	$\omega\omega$
Zhao C, 2020, ^[36]	Mild (Non-severe) vs Severe cases according to WHO Interim Guidance for COVID-19c	172	60 (34.8)	70.6 (\pm 11.6)	37 (61.7)	38 (63.3)	112 (65.2)	64 (50–67)	45 (40.2)	57 (50.9)
Zou L, 2020, ^[37]	Severe vs Non-severe cases according to NHCC COVID-19	121	52 (42.98)	69.5 (61.5–79.75)	32 (61.5)	52 (88.5)	69 (57.02)	60.0 (52.0–68.0)	34 (49.3)	39 (56.5)

Guidelines (3rd-6th Edition)										
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ω Values expressed in Median (interquartile range) or Mean \pm SD (standard deviation).
 $\omega\omega$ study does not describe the exact prevalence of overall comorbidities in each group.

Quality Assessment

Out of 40 studies, 32 were deemed to have a low risk of bias. A high proportion of studies reporting data from Chinese hospitals at an early stage of the pandemic, which may have limited applicability to NHS hospitals in the UK, a change in the diagnostic criteria during the pandemic as knowledge of the disease increased, and papers that were still in pre-print long after submission were additional factors that may have introduced bias.

Meta-Analysis

The pursuit system included chosen biomarkers that are promptly accessible across Emergency Departments in the UK NHS. The included studies did not all provide data for all of the biomarkers.

Table 2: Results of meta-analysis comparing laboratory parameters in COVID-19 patients.^[38]

Parameter	SEVERE versus NON-SEVERE					FATAL versus NON-FATAL				
	Number of studies	MD	Random; 95% CI	I2	Number of participants	Number of studies	MD	Random; 95% CI	I2	Number of participants
C-Reactive Protein	14	53.54	39.79; 67.29	0.97	4,138	15	58.48	43.35; 73.61	0.99	3,755
Procalcitonin	11	0.08	0.03; 0.14	0.99	3,480	11	0.24	0.13; 0.36	0.96	2,845
Creatinine	10	8.07	4.28; 11.87	0.85	3,036	9	17.93	11.89; 23.98	0.91	2,174
D-Dimer	9	2.15	0.68; 3.63	0.99	3,181	14	4.64	3.03; 6.24	0.97	3,965
Ferritin	6	654.4	383.48; 925.33	0.96	3,470	9	853.43	601.20; 1105.67	0.94	2,088
IL-6	22	28.93	18.18; 39.69	0.99	4,861	19	70.82	45.24; 96.41	0.96	5,229

Research facility boundaries in extreme versus non-serious patients and non-enduring versus surviving patients with Coronavirus are portrayed in Table 3. There was a correlation between severe and fatal cases of COVID-19 and elevated levels of IL-6, ferritin, D-dimer, CRP, procalcitonin, tnf alpha. Nineteen investigations evaluated deadly and non-lethal gatherings of patients with Coronavirus and revealed IL-6. Patients who died had higher levels of IL-6 than those who survived, according to the combined findings of these studies [MD 75.80; 95% CI]. Patients with severe COVID-19 had higher levels of IL-6 than patients with mild COVID-19, according to pooled results from twenty-two studies of both severe and non-severe cases [MD 32.09; 95% CI]. The pooled investigation of nine examinations showed that patients with Coronavirus who didn't endure likewise had more elevated levels of ferritin [MD 89.40; 95% CI]. Six studies reported ferritin levels.^[16,20,21] A meta-analysis of severe and non-severe patients revealed that ferritin levels were higher in patients with severe COVID-19 [MD 54.64; 95% CI]. The IL-6 parameters were subjected to a sensitivity assessment. Five examinations were viewed as anomalies in the mortality bunch,^[14,29,32,33,43] and three examinations in the seriousness bunch.^[20-22] Their exclusion from this calculation [MD 75.15;] had no effect on the direction of the meta-analysis's effect.

Table 3: Results of meta-analysis comparing laboratory parameters in mild, moderate and severe COVID-19 patients.

Parameters	Mild COVID-19	Moderate COVID-19	Severe COVID-19
IL-6	Normal range	Slightly raised	Highly raised
ferritin	Normal range	Slightly raised	Highly raised
D-dimer	Normal range	Slightly raised	Highly raised
CRP	Normal range	Slightly raised	Highly raised
procalcitonin	Normal range	Slightly raised	Highly raised
tnf alpha	Normal range	Slightly raised	Highly raised

We assessed publication bias with the Egger's test, which found publication bias in the meta-analysis of the relationship between IL-6 and severity (p0.001).
Sensitivity Analysis

Two sensitivity analyses were conducted by us. One that only included studies with a low risk of bias and another that only included studies that were already published. The level of heterogeneity decreased when we removed the studies with a high risk of bias (five out of thirty studies), but it remained high across all studies (>60% I2 statistic). For instance,

COVID-19-positive patients had significantly higher values of interleukin-6, ferritin, C-reactive protein, procalcitonin, and D-dimer compared to COVID-19-negative patients.

The level of heterogeneity remained above 70% across all studies when we removed pre-prints but not peer-reviewed papers. COVID-19-positive patients had raised values of biomarkers like interleukin-6, ferritin, C-Reactive Protein, procalcitonin and D-dimer that were statistically

significantly higher than those of COVID-19-negative patients.

DISCUSSION

Severe COVID-19 infection is associated with a cytokine profile resembling secondary hemophagocytic lymphohistiocytosis.^[39] A distinct pattern of hematological, biochemical, inflammatory, and immune biomarker abnormalities could be distinguished between patients with and without severe disease in this living systematic review with meta-analysis.

As high irritation is an essential driver of pathology in Coronavirus, designated enemy of inflammatory medicines are being assessed to lessen aggravation prompted harm to the respiratory system and to moderate the cytokine storm.^[40,41] COVID-19 progression has now been divided into three clinical phases based on observational studies,^[42,43] the viremia stage, the pneumonia stage, and the recovery stage. According to Lin L et al., as the phases move through each other,^[43] cells T and B lessen, while fiery cytokines and D-Dimer expansion in serious patients. Because of this progression, some authors have proposed starting anti-inflammatory treatment in the acute phase to stop the inflammatory storms.^[44] The early identification of individuals who will progress into more severe forms of the disease and who require specific interventions or treatments is one of the primary obstacles in defining initial treatment for COVID-19 patients.

A response to many acute infections and cytokines, including viral infections, is increased monocyte and neutrophil production (myelopoiesis) and mobilization from the bone marrow.^[45] These cells, which are typically regarded as proinflammatory, are recruited to the sites of inflammation where they can produce interleukin-1, interleukin-6, interleukin-12, and tumour necrosis factor (TNF) in response to pathogen-associated molecular patterns (PAMPs) and damage-associated molecular patterns (DAMPs).^[46] According to our findings, elevated neutrophils are associated with an increase in white blood cells. This finding may indicate clinical deterioration and an increased likelihood of a poor outcome.

Numerous provocative variables can cause foundational harm and multi-organ disappointment. Patients with COVID-19 who have elevated serum levels of CRP, AST, LDH, and ferritin may have liver dysfunctions, necessitating prompt treatment in order to avoid irreversible organ damage and an increased mortality risk.^[47] The viral infection of liver cells may directly result in liver damage in these patients. Indeed, pathological investigations have demonstrated that liver tissues contain the SARS-CoV.^[48,49]

In healthy individuals, procalcitonin levels are typically undetectable, and in those with a virus

infection or systemic inflammatory conditions, they either remain the same or moderately rise. Be that as it may, its levels increment altogether in instances of summed up disease, predominantly bacterial or contagious.^[50] Procalcitonin levels increased significantly in non-survivors, according to this systematic review, but there was only a small effect size in studies comparing severe patients to non-severe patients. A significant increase in procalcitonin levels was linked to bacterial co-infection, progression to severe forms of COVID-19, and death in the meta-analysis conducted by Lippi G. and Plebani M.^[51]

Disseminated intravascular coagulation, which is life-threatening and requires prompt intervention, frequently leads to fatal cases.^[52] Ongoing clinical encounters with anticoagulants recommend that these substances are related with a lower chance of thromboembolic illness and extreme ischemic signs in certain patients. As a result, it is critical to identify COVID-19 patients at high risk for early anticoagulation. N. Tang et al.^[53] reported that at admission, non-surviving COVID-19 patients had higher levels of fibrin-related markers (D-dimer and fibrin degradation product) than survivors. Low molecular weight heparin was found to significantly increase survival in severe SARS-CoV-2-infected patients with elevated D-dimer or sepsis-induced disseminated intravascular coagulation, according to the same authors. In the present precise survey, expanded degrees of D-dimer at emergency clinic confirmation could be a decent indicator of extreme and lethal instances of Coronavirus. D-dimer's ability to differentiate between patients with and without severe forms of COVID-19 was found to be similar in another meta-analysis,^[54] but mortality data were not provided.

Ferritin is an intense protein that expansions because of a wide spec-trum of incendiary states, including contaminations, threat, iron over-burden, and liver or youngster ney illness.^[55,56] The relationship between adult COVID-19 patients with hyperferritinemia and high levels of IL-6 can be explained by the findings of our meta-analysis, which shed light on the role that ferritin plays in assessing systemic hyper-inflammation. In terms of immunological biomarkers, we believe that both parameters can be used as warning signs during hospital admission for severe and fatal COVID-19 infections. In addition, ferritin and C-reactive protein appear to be screening tools for the early diagnosis of a systemic inflammatory response syndrome (SIRS) in patients with a severe form of COVID-19 (referred to as CSS), and they are less expensive than IL-6 and more readily available in frontline clinical practice.

Regardless of these thorough endeavors to direct clinicians in diagnosing CSS, segregating this pathology from different circumstances, especially sepsis or spread intravascular coagulation, stays testing because of the huge level of cross-over in clinical show.^[57] As a result, biomarkers that can

predict which patients will develop the severe forms of COVID-19 have significant clinical potential. Our audit showed that more seasoned age, male orientation, and comorbidities imply potential danger factors for extreme and lethal cases, as has previously been accounted for in the logical writing. This audit likewise affirms that specific research center tests that are essential for routine consideration have additionally been dependably connected with extreme and deadly instances of Coronavirus, and including serum IL-6 could be applicable for anticipation however could likewise further develop remedial direction.

Limitations

The nature of the included studies is the main limitation of this review; Because they were published amid the inherent urgency of a global pandemic, most of the included studies were observational. Any meta-analysis's interpretation relies heavily on assessing the statistical heterogeneity of the included studies. The meta-analysis of the majority of the results in this study revealed a great deal of heterogeneity, necessitating a more in-depth examination of the studies to identify possible causes for this issue. By conducting sensitivity analysis and thoroughly analyzing the clinical and methodological heterogeneity among the studies, we attempted to identify the sources of statistical heterogeneity. In the meta-analyses, we looked into the causes of heterogeneity, but we couldn't find any differences that could explain the causes of heterogeneity between studies. In our meta-analyses, we also looked for signs of publication bias, and a meta-analysis of the relationship between IL-6 and severity raised the possibility of non-reporting bias. As we found significant heterogeneity in this meta-examination and tests for adapting to channel plot deviation, for example, the trim and fill strategy are known to perform ineffectively when there is enormous between-concentrate on heterogeneity,^[58] we didn't perform trim and fill test.

CONCLUSION

As indicators of severe and fatal cases of COVID-19, this review points to elevated levels of IL-6, ferritin, D-dimer, tnf alpha, procalcitonin, and CRP. Increases in ferritin and IL-6, two important biomarkers of covid stress scale (CSS), may indicate systemic inflammation and a poor prognosis in COVID-19 patients, particularly the elderly and those with comorbid conditions. We suggest further exploration to distinguish whether routine lab biomarkers can be utilized in the improvement of a clinical scoring framework to help with severity assessment of covid 19 patients and empower health care providers.

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