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Laboratory characteristics of patients infected with the novel SARS-CoV-2 virus ☆,☆☆,★



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SUMMARY

A subgroup of COVID-19 patients develop very severe disease with requirement for ICU treatment, ventilation, and ECMO therapy. Laboratory tests indicate that the immune and clotting system show marked alterations with hyper-activation, hyper-inflammation, cytokine storm development. Furthermore, organspecific biomarkers demonstrate the involvement of cardiac muscle, kidney, and liver dysfunction in many patients. In this article the use of laboratory biomarkers is discussed with regard to their use for diagnosis, disease progression, and risk assessment.

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Introduction

Although only a minority of COVID-19 patients show critical disease progression from moderate to severe stages of the disease including requirement for ventilation and ECMO therapy, this subgroup of COVID-19 patients requires particular attention. Data collection from several regions of the world including China, Europe, and the United States clearly demonstrate that COVID-19 is not only a disease of the lung and the airways. Many other organ systems are involved and contribute to disease variety and progression. With regard to the immune system, hyper-inflammation

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together with the development of exorbitant increased cytokine production represents a hallmark of severe patients requiring ventilation. Some of these patients develop bacterial superinfections with increased levels of sepsis markers. Another important systems which recently caused increased attention is the clotting system. This is particularly highlighted by the detection of increased levels of D-dimers. Organ dysfunction has been reported in many patients including the heart (myocardial muscle damage), the kidney, and the liver. Laboratory diagnostics play not only an important role in disease diagnosis, but also in assessing progression and severity in these patients. Furthermore, laboratory diagnostics allow early detection of organ dysfunction in many cases. Moreover, biotests are used to assess an increased mortality risk in severe lethal patients. In this article we summarize the most prominent findings in COVID-19 patients and discuss the use of these markers for diagnosis, disease progression, and risk assessment. (Fig. 1 and Tables 1, 2)

Hemoglobin and white blood cells

Retrospective analyses from China demonstrated that leukocyte counts were higher among non-survivors compared to recovered patients^{1,2}; in particular, Zhou et al. reported that COVID-19 patients who did not survive had a median of 9.8×10^9 /L WBC count compared to 5.2×10^9 /L among those who survived (*p*<0.0001), although the exact time point of measurement was







Review

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Mild:	Mild symptoms, no imaging findings of pneumonia.
Moderate:	Fever <i>OR</i> respiratory symptoms.
Severe:	Respiratory distress and the respiratory rate >30/min OR saturation <93% at rest OR PaO2: FiO2 ratio
	≤300mmHg.
Critical:	Respiratory failure requiring mechanical ventilation (including ARDS) OR shock OR other organ failure
	requiring ICU.

Fig. 1. Schematic overview of key laboratory characteristics during SARS-CoV-2 infections. The latter induce an increase (depicted in red) or a reduction (depicted in green) in the concentration and/or counts of a wide range of laboratory biomarkers. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

not defined in their methods. Furthermore, another study of 140 hospitalized patients in Wuhan, demonstrated significantly higher leukocyte counts among those with severe COVID-19 disease, compared to patients with milder infection (p = 0.003).³ Finally, a series from the same center, and possibly overlapped populations with the previous study, reported significantly higher WBC counts upon hospital admission among patients requiring critical care, although median values were within normal range (WBC count median 6.6×10^9 /L for ICU vs 4.3×10^9 /L for non-ICU admission, p = 0.003).⁴

The observed leukocytosis is attributed to an elevation of neutrophils, as the other WBC populations seem to drop in severely ill and eventually fatal COVID-19 cases.⁵ Absolute lymphopenia is commonly observed in patients with COVID-19, but pronounced lymphocyte depletion is a cardinal marker of enhanced disease severity and an indicator of imminent death, that has been consistently depicted by almost all currently published reports, coming mainly from China.^{1–7} Importantly, not only the degree of lymphocyte drop, but also the persistence of low lymphocyte counts throughout the disease course have been associated with critical illness and death.^{1,2,4} In contrast to previous reports for SARS-CoV, peripheral blood smears reveal the presence of reactive lymphocytes, including some lymphoplasmacytoids, in the majority of COVID-19 patients⁷⁻⁹. Severe SARS-CoV-2 infection depletes all lymphocyte subsets, including CD4+ T cells, CD8+ T cells, B cells and natural killer (NK) cells, but CD4+/ CD8+ ratio is not inverted as seen in other viral infections.¹⁰⁻¹³ Not only the absolute numbers of T-cells are reduced, but also receptors suppressing their cytotoxic effects, like the CD94/NKG2A receptor, are up-regulated leading to diminished defense mechanisms against the virus.¹⁰

Monocyte, eosinophil and basophil counts are also decreased in COVID-19, but the magnitude of this reduction has not been associated with disease severity, in currently published data from Chinese centers.^{3,10,14} Moreover, pro-inflammatory cytokines are known to blunt erythropoiesis.¹⁵ However, aside from one study that found significantly higher frequencies of decreased hemoglobin concentrations among severe (43.6%) and critical cases (37.2%) compared to mild/moderate ones (23.1%) (p<0.001), solid evidence of significant hemoglobin reduction in severe COVID-19 has not been consistently reported as yet.^{5,11,16} In one particular study, lower hemoglobin concentration was associated with increased odds for lack of disease improvement but not death (odds ratio 1.731, p = 0.008).¹⁶

Preliminary reports imply that high neutrophil counts and persistently deep lymphocyte nadir counts during hospitalization as well as high neutrophil to lymphocyte ratios (NLR) are indicators of adverse outcomes such as ICU admission and death¹⁰. A retrospective Chinese study reported that NLR, along with the SARS-CoV-2 IgG levels, could be used as a simple discriminative tool for severity between COVID-19 patients, and further predict the clinical outcome of these patients¹⁴. However, whether these indices can actually risk stratify patients and predict poor outcomes, most importantly at an early stage of the disease, remains to be addressed and validated in large prospective trials.

 Table 1

 Laboratory parameters and associated pathophysiology in adult COVID-19 patients.
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BIOMARKER	PATHOPHYSIOLOGY	CLINICAL UTILITY IN ADULT COVID-19	REFERENCES
Hematological indices	Reduced erythropoiesis due to inflammatory	Lower levels associated with: Lack of improvement	(16 5 11)
Lymphocytes	cytokines Absolute count reduction, functional exhaustion	No clear association with disease severity and outcomes	(52, 5, 1-4, 6
Lymphocytes	of all populations (especially cytotoxic T-cells) Unknown exact mechanisms	PMN/CD8+ ratio and PMN/Lymphocyte ratio may be used as prognostic markers	(32, 3, 1–4, 6, 10–13, 7)
Monocytes/ Basophils/ Fosipophils	Absolute count reduction, Unknown exact mechanisms	No clear association with disease severity and outcomes	(14, 3, 10)
Total white blood cells/Neutrophils	Increased due to inflammation	Higher levels associated with: ↑Severity, ↑Mortality, Bacterial superinfections	(1-4)
Acute phase reactants Albumin	Reduced production due to inflammatory cytokines	Lower levels associated with: <i>†</i> Severity, <i>†</i> Mortality, Lack of improvement	(1, 2, 11, 6, 16, 19)
		Low levels on admission may be used as prognostic marker for severity	
C-reactive protein (CRP)	Increased production due to inflammatory cytokines	Higher levels associated with: ↑Severity, Lack of improvement, Bacterial superinfections	(1, 3, 16)
Erythrocyte	Increased in inflammation	Tendency for higher levels associated with: \Mortality	(1)
Ferritin	Increased production due to inflammatory	Higher levels associated with: †Severity, †Mortality	(1, 2, 6, 5)
Procalcitonin	Increased production due to inflammatory	Higher levels associated with: ↑Severity, ↑Mortality, Bacterial	(4, 3, 18, 2, 6, 5,
Serum amyloid A	cytokines Increased production due to inflammatory	superinfections High levels seen among: all COVID-19 patients	19) (3)
Biochemistry indices	cytokines		
Cholinesterase	Unknown exact mechanism	Lower levels associated with: ↑ Severity	(53)
Electrolytes (Na, K, Cl)	Multiple mechanisms (e.g. SIADH, acidosis etc.) Released by cell injury	No clear association with disease severity and outcomes	(1, 11, 18, 51) (11, 2, 1, 6, 5, 19)
(LDH)	keedsed by een injury	improvement High levels on admission may be used as prognostic marker for covority.	16)
Triglycerides	Reduced lipoprotein lipase activity due to high	Higher levels have been reported in fatal cases but not enough dat.	a (1)
TSH/FT3	Possible euthyroid sick syndrome of critical illness	Higher levels have been reported in fatal cases, not enough data	(1)
Cardiac biomarkers			(- / - / -)
Hs-troponin I	Released by myocardial injury	Higher levels associated with:↑igher levels associa, lack of improvement High levels on admission or gradual increase may be used as	(54, 31, 33, 4)
		prognostic marker for severity and mortality	
Troponin T	Released by myocardial injury	Higher levels associated with: ↑igher levels associa	(32)
CK-MB NT-proBNP	Released by myocardial injury Increased production due to heart failure	Higher levels associated with:↑igher le Higher levels associated with:↑igher le	(33, 4) (31, 33)
Creatinine	Decreased discharge due to renal injury	Higher levels associated with:	(4, 2, 55, 34, 31)
BUN	Decreased discharge due to renal injury	Higher levels associated with:	(31, 34, 55, 4)
Urinary protein Urinary erythrocyte	Possiblypositive due to renal dysfunction Possibly positive due to renal dysfunction	Tigner levels associa Proteinuria may associated with:↑roteinuri (limited data) Hematuria may associated with:↑ematuria (limited data)	(34) (34)
Liver function indices ALT	Possibly liver injury, unknown exact mechanism	Higher levels associated with:	(2, 31, 4, 18, 36)
AST	Possibly myocardial or liver injury, unknown	↑igherity, (↑Mortality, indeterminate data) Higher levels associated with:	(31, 4, 18, 36)
TBIL	exact mechanism Unknown exact mechanism	Tigher level (TMOTAIRY, Indeterminate data) Higher levels associated with:	(36, 31)
GGT	Unknown exact mechanism	Trigher levels associated with:	(36)
ALP	Increased levels in some patients, unknown	rigner ie (limited data) No clear association with disease severity and outcomes	(36, 56)
Coagulation profile	caree meetiniiiii		
D-dimer	Elevated levels possibly due to hypercoagulability and secondary fibrinolysis	Higher levels associated with: ^igher levels associa D-dimers 1 ng/mlon admission or gradual increase may be used	(4, 3, 2, 55, 31)
РТ	Prolonged PT possibly duehypercoagulability and secondary fibrinolysis	as prognostic marker for severity and mortality Higher levels associated with: ↑igher lev	(2, 4, 55, 44, 57)
		((continued on next page)

Table 1 (continued)

BIOMARKER	PATHOPHYSIOLOGY	CLINICAL UTILITY IN ADULT COVID-19	REFERENCES
INR	Elevated levels possibly duehypercoagulability and secondary fibrinolysis	Higher levels may associated with: \Severity (limited data)	(36)
APTT Fibrinogen	Unknown exact mechanism Elevated as an acute phase protein and may decreasedue to hypercoagulability	Indeterminate association with disease severity and outcomes Higher levels may associated with: ↑igher levels may ass (limited data)	(44, 5, 55) (44, 43)
Cytokines and chemokines			
IL-1 β	Increased production/Associated with CSS/sHLH	Higher levels may be associated with: ↑Mortality Indeterminate data for severity	(58, 1, 6, 11, 10)
IL-2/ soluble IL-2R IL-6	Increased production/Associated with CSS/sHLH Increased production/Associated with CSS/sHLH	 Higher levels associated with: ↑Severity, ↑Mortality Higher levels associated with: ↑Severity, ↑Mortality IL-6 levels may monitor disease progression Higher of IL-6 to IFN-γ ratio may distinguish severe from moderate cases 	(58, 1, 11, 6, 10) (5, 24, 58, 1, 11, 6, 2, 10)
IL-7	Increased production/Associated with CSS/sHLH	Higher levels associated with: ↑Severity	(58, 11)
IL-8	Increased production/Associated with CSS/SHLH	data)	(6, 10)
IL-10	Increased production by macrophages	Higher levels associated with: \uparrow Severity (also \uparrow Mortality, but not enough data)	(58, 1, 11, 6)
IL-17	Increased production/Associated with CSS/sHLH	Higher levels may be associated with: ↑Severity (Not enough data)	(10)
IP10 (CXCL10)	Increased production/Associated with CSS/sHLH	Higher levels associated with: ↑Severity	(58, 11)
G-CSF/GM-CSF	Increased production/Associated with CSS/sHLH	Higher levels associated with: ↑Severity	(58, 11, 10)
TNF-α	Increased production/Associated with CSS/sHLH	Higher levels associated with: ↑Severity(also ↑Mortality, but not enough data)	(58, 1, 11, 6, 10)
MCP1 (CCL2)	Increased production/Associated with CSS/sHLH	Higher levels associated with: ↑Severity	(58, 11, 10)
MIP-1 α (CCL3)	Increased production/Associated with CSS/sHLH	Higher levels associated with: ↑Severity	(58, 11, 6, 10)
INF-γ	Reduced production by CD4+ T cells	Lower levels may be associated with: ↑Severity Higher of IL-6 to IFN-γratio may distinguish severe from moderate cases	(24, 11, 58, 6)
Complement	Possible activation of the alternative and lectin-based complement pathways from viral proteins	Deposits of C5b-9, C4d and MASP 2 in the microvasculature of lungs (from autopsy specimens) No differences in C3/C4 levels among survivors- non survivors	(1, 27, 59)
Immunoglobulins (IgA, IgG, IgM)	In theory, increased production induced by activated B-cells	No differences in IgA/lgG/IgM levels among survivors- non survivors	(1)
Soluble urokinase plasminogen activator receptor (suPAR)	Increased due to endothelial activation	High levels may be associated with: prediction of respiratory failure	(⁵²)
parameters			
рН	Respiratory alkalosis driven by hypoxemia, metabolic acidosis due organ hypoperfusion	Conflicting data on pH and associated mortality. One study found statistically higher frequency of acidosis among fatal cases	(1, 2)
Bicarbonates	Decreased due to respiratory alkalosis and metabolic acidosis	Not enough data - possibly lower among non-survivors	(1)
PaO ₂	Decreased due to alveolar and microvasculature injury (direct and indirect)	Frequency of type I respiratory failure is significantly higher among non survivors	(1, 2)
PaCO ₂	Decreased due to high respiratory rate driven	Markediy low PaU2 (<60 mmHg) levels are seen in fatal cases Not enough data – possibly lower among non-survivors	(1)
PaO ₂ :FiO ₂ ratio	Decreased due to alveolar and microvasculature injury (direct and indirect)	PaO2:FiO2 ratio of \leq 300 associated with \uparrow Mortality	(1)

Cl: Chloride, CSS/sHLH: Cytokine storm syndrome/secondary Hemophagocytic lymphohistiocytosis, FiO₂: Fraction of inspired oxygen, FT3: Free triiodothyronine, G-CSF: Granulocyte-colony stimulating factor, GM-CSF: Granulocyte-macrophage colony-stimulating factor, IL: Interleukin, IP10:Interferon gamma-induced protein 10, K: Potassium, MASP 2: mannose binding lectin associated serine protease 2. MCP1 (CCL2): Monocyte chemoattractant protein 1, MIP-1 α (CCL3): Macrophage inflammatory protein 1-alpha), Na: Sodium, PaCO₂: Arterial carbon dioxide partial pressure, PaO₂: Arterial pressure, SIADH: Syndrome of inappropriate antidiuretic hormone secretion, soluble IL-2R: soluble Interleukin 2 receptor, TSH: Thyroid stimulating hormone, BUN, blood urea nitrogen; ALT, alanine transaminase; AST, aspartate transaminase; TBIL, total bilirubin; GGT,gamma-glutamyl transpeptidase; ALP, alkaline phosphatase; CK-MB, creatine kinase MB; PT, prothrombin time; APTT, activated partial thromboplastin time; INR, international normalized ratio; Hs-troponin I, high sensitivity troponin I.

Common inflammatory markers - Acute phase reactants

The regulation of ferritin synthesis is cytokine-controlled¹⁷; hence, the extreme immune activation in the context of the cytokine storm observed in critical, and usually fatal, cases of COVID-19, leads to an up-regulation of serum ferritin levels. Indeed, preliminary patient data demonstrate that excessive ferritin levels are observed among COVID-19 patients, ranging from 400 μ g/L to as high as >2000 μ g/L, with the highest trends being observed in severe cases and in non-survivors.^{1,2,6} Direct correlation be-tween serum ferritin concentration and poor survival, as reported by the meta-analysis conducted by Henry et al. (weighted mean difference: 408.28 μ g/L, 95%CI: 311.12–505.44 μ g/L, Cochran's Q pvalue=0.01), suggests its use as a surrogate marker of immune dysregulation and a prognostic marker of disease severity and imminent death. 5

Only scarce data have contextualized the erythrocyte sedimentation rate (ESR) kinetic in patients with COVID-19. One study reported that fatal cases had a tendency for higher ESR compared to those who recovered (median ESR 38.5 vs 28 mm/h) without reporting the statistical significance of the observed difference among the two groups.¹ A similar trend was also depicted for C-reactive protein (CRP) concentration by the same study, with median levels being 4-fold higher among non-survivors (median concentration 113 vs 26.2 mg/L).¹ Between severe and non-severe cases, reported CRP differences are not that striking (median (IQR):

Table	
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Laboratory parameters in pediatric COVID-19 patients.

BIOMARKER	PEDIATRIC COVID-19 DATA	REFERENCES			
Hematological indices					
Hemoglobin	Potentially similar to adults	(52, 53)			
Lymphocytes	Higher lymphocyte counts compared to adults	(52, 54-57)			
	Normal lymphocyte counts common				
	Lymphopenia in 0–35% of children				
	Lymphocytosis is rare				
Total white blood cells/Neutrophils	Higher levels associated with:	(52, 55-57)			
	Symptomatic disease, Younger age (<2 y.o.)				
	Lower neutrophil counts compared to adults				
	Leukocytosis is more frequent				
	Leukopenia is rare				
Acute phase reactants					
Albumin	Less frequently decreased compared to adults	(57, 53)			
C-reactive protein (CRP)	Lower CRP levels compared to adults	(52, 55, 56)			
	High CRP in 10%–83% of children				
Erythrocyte sedimentation rate	Less frequently elevated compared to adults	(57)			
Procalcitonin	Can be high in hospitalized children	(56, 57)			
	More frequently elevated compared to adults				
Biochemistry indices					
Lactate dehydrogenase (LDH)	Normal LDH levels commonly	(52, 55, 57)			
	Higher LDH levels compared to adults in one report				
Cytokines and chemokines					
IL-6	Lower IL-6 levels compared to adults	(52)			

47.6 mg/L (20.6–87.1) vs 28.7 mg/L (9.5–52.1), p<0.001), but significantly increased frequency of higher concentrations among severe and critical cases compared to mild/moderate ones are nevertheless evident (mild/moderate cases: 50.5%, severe cases: 79.2% and critical cases: 92%, p<0.001).^{3,16} Finally, one Chinese study with 663 COVID-19 patients reported that higher CRP levels are inversely associated with disease improvement (odds ratio 4.697, p<0.0001).¹⁶

Individual studies demonstrate that greater procalcitonin (PCT) concentrations (usually ≥ 0.05 ng/ml) can significantly distinguish between non-severely from severely ill and fatal cases, thus possibly acting as a prognostic marker.^{2-4,6,18,19} However, a metaanalysis found that severe from non-severe COVID-19 could be differentiated by a marginally higher PCT (by 0.2 ng/ml).⁵ Increments of both CRP and PCT may be associated, not only with the immense inflammatory response, but also with the higher frequency of bacterial superinfections among critically ill COVID-19 patients (up to 50% rate among non-survivors).⁵ The differentiation between severe SARS-CoV-2 infection and a bacterial superinfection is often difficult in clinical practice. Though markedly elevated PCT and CRP are consistent with bacterial co-infection, there is not a clear cut-off. Other markers that have been proposed as differentiators between bacterial and viral infections (such as Myxoma resistance protein (MxA1), Lipocalin 2 (Lcn2), High mobility group box one protein (HMGB1)) have not been studied in COVID-19 disease.²⁰

Albumin is a negative acute phase reactant whose synthesis is down-regulated by inflammatory cytokines.²¹ Therefore, it is not surprising that hypoalbuminemia (usually <30 g/L) has been persistently noticed among patients with severe or fatal COVID-19^{1,2,6,11}. Moreover, one study demonstrated that low albumin concentration was associated with lack of disease improvement (odds ratio 2.377, p<0.0001), while hypoalbuminemia was also introduced as a risk factor, among other parameters, in a proposed risk prediction nomogram for severe COVID-19.^{16,19}

Serum amyloid A (SAA) is another acute phase reactant inhibiting monocyte mobilization, platelet activation and various chemotactic pathways.²¹ High concentrations of SAA among all COVID-19 patients have only been reported by Zang et al., without a significant difference between severe and non-severe cases.³

Cytokines, chemokines, pathology findings and other markers

Exuberant release of pro-inflammatory cytokines is associated with multi-organ injury and acute respiratory distress syndrome (ARDS), which is inevitably fatal if left untreated.²² Fulminant hypercytokinemia has been increasingly recognized among critically ill COVID-19 patients.

Distinct pro-inflammatory cytokines (such as interleukin $(IL)-1\beta$, IL-2 and its soluble receptor, IL-6, IL-8, IL-17, Granulocyte colony-stimulating factor (G-CSF), Granulocyte-macrophage colony-stimulating factor (GM-CSF), tumor necrosis factor alpha (TNF- α)), inflammatory chemokines (such as the monocyte chemoattractant protein 1 (MCP1or CCL2) and the macrophage inflammatory protein 1-alpha (MIP-1 α or CCL3)), as well as the anti-inflammatory cytokine IL-10, have been consistently found significantly elevated in patients with severe COVID-19, those admitted to ICU or patients who died compared to milder forms of SARS-CoV-2 infection.^{1,2,6,10,11,23} Notably, monitoring of IL-6 levels has been proposed as a candidate index for disease progression.⁵ Moreover, higher of IL-6 to Interferon gamma (IFN- γ) ratios may distinguish severe from moderate COVID-19 cases (standardized mean difference of 0.739, 95% CI = 0.131-1.383). 24 All these data converge into the conclusion that major immune dysregulation occurs in severe COVID-19, leading to many clinical manifestations of the fatal form. Although measurement of these indices is not widely available, following up such markers may be an integral part of relevant prognostic and diagnostic tools.

On the other hand, complement components C3 and C4 and immunoglobulin (IgG, IgM and IgA) levels are not specific markers of the cytokine storm syndrome and the few data that are available showed no clinically significant differences between deceased and recovered patients¹. However, exuberant SARS-CoV-2-specific IgG responses were associated with increased disease severity, in a retrospective Chinese study with 222 COVID-19 patients.¹⁴

Cytokine storm can certainly, but only partially, explain the observed clinical features of COVID-19 disease. Angiotensinconverting enzyme 2 (ACE2) receptors, the mediators of SARS-CoV-2 invasion into host-cells, are expressed by numerous cells, including endothelium; therefore, direct viral endothelial injury cannot be excluded.²⁵ Indeed, preliminary histopathological data from fatal cases demonstrated lesions consistent with endotheliitis (endothelialitis) in many organs including lungs, small lung vessels' congestion, mononuclear cell infiltrates within the intima of organs' vasculature, viral inclusion bodies in peritubular spaces and viral particles in endothelial cells of the glomerular capillary loops.²⁶

Another series reported the identification of C5b-9, C4d and mannose binding lectin-associated serine protease (MASP) 2 terminal complement component deposits in pulmonary microvasculature; furthermore, co-localization of spike glycoproteins with C4d and C5b-9 in inter-alveolar septa and on skin microvasculature were evident is some cases.²⁷ This observation is consistent with a systemic complement activation leading to a catastrophic pauciinflammatory septal capillary injury and a pro-coagulant state.²⁷ Importantly, hallmarks of classic ARDS such as typical diffuse alveolar damage (DAD) were not prominent.²⁷

Moreover, the connection of viral spike protein to ACE2 receptor, down-regulates ACE2 levels in lungs; this in turn, increases the angiotensin II (AngII) levels, reduces angiotensin 1–7 (Ang-(1–7)), and imbalances the renin-angiotensin system in the lung, leading to vasoconstriction.²⁸ These data are in concordance with a notably distinct type of ARDS with highly compliant lungs, which is seen in a major subset of COVID-19 patients; this manifestation is quite possibly consistent with an underlying vasoconstriction and microvasculature injury leading to loss of lung perfusion regulation²⁹. Though neither histopathology specimens nor lung ACE2 or AngII levels are easily obtainable in daily clinical practice, they would definitely be useful in research settings in order to elucidate the disease's pathophysiology and may assist diagnosis in the future.

Cardiac biomarkers

Cardiac troponin I and T are highly sensitive and specific biomarkers of myocardial injury which can be caused by myocardial ischemia, inflammation, immune response, and toxin. ³⁰ Elevated troponin at admission was observed in more than half of dead patients and associated with increased severity and mortality in COVID-19 patients.^{1,2,31} Regardless of underlying cardiovascular disease, patients with dynamic increases of troponin during the hospitalization were more likely to have fatal outcomes. ^{2,31–33} Although some COVID-19 patients were reported with comorbidity of chronic heart disease, the underlying mechanism for troponin elevation in patients with COVID-19 is not clear. The myocardial injury in COVID-19 patients might associate with a systemic hyperinflammation^{13,31} rather than a virus attack directly. Increased levels of CK-MB and NT-proBNP can also be found in severe COVID-19 patients compared to non-severe patients.^{4,31,33}

Renal function tests

According to a cohort study ³⁴ of 701 patients with COVID-19, the proportion of proteinuria, hematuria, abnormal serum creatinine and urea nitrogen at admission and were 43.9, 26.7, 14.4 and 13.1%, respectively. In addition, there was a high prevalence (5.1%) of acute kidney injury (AKI) during the study period. The result showed proteinuria, hematuria, and elevated serum creatinine/urea nitrogen at admission and acute kidney injury (AKI) during hospitalization over stage 2 were associated with in-hospital death. However, the other largest retrospective study to date found that the prevalence of serum creatinine abnormalities and AKI was only 1.6% and 0.5%.¹⁸ This may be due to the different proportions of severe patients between the two studies and the different definitions of the normal reference range for serum creatinine. From the result of autopsy of 26 COVID-19 patients,³⁵ the histopathology of the kidney revealed significant acute tubular injury and found that the tubular epithelial cells were directly infected by SARS-CoV2. Therefore, SARS-CoV2 may cause kidney injury or exacerbate existing kidney disease. Attention should be paid to monitoring renal function and the occurrence of AKI.

Liver function tests

Abnormal liver function tests, such as increased levels of ALT, AST, TBIL, GGT and decreased level of albumin were relatively common in patients with COVID-19, and 10–33% of these patients had abnormal ALT or AST.^{2,4,18,31,36} Although patients with severe COVID-19 seem to have higher rates of liver dysfunction, it is reassuring that the levels of ALT, AST, TBIL, GGT in COVID-19 patients were not significantly different in compared with hospitalized community-acquired pneumonia patients and even the median or average transaminase level in severe COVID-19 patients was lower than twice upper reference limit.^{4,36,37} Therefore, the clinical effect of these elevated indicators may not be evident in COVID-19 patients. Liver dysfunction may be related to severe infection, inflammation induced liver injury, medication associated hepatotoxicity and hypoxia.³⁸

Coagulation profile

D-dimer is a degradation product of fibrin. Elevated D-dimer levels were consistently reported in COVID-19 patients with prevalence ranging from 43 to 68%.^{2,3,31} D-dimer>1 ng/ml at admission were associated with increased severity and odds of death with COVID-19, and the gradual increasing of D-dimer during disease course was particularly associated with disease worsening and mortality.^{2,4} Serum D-dimer can reflect fibrinolytic activities and is also an inflammatory biomarker. Furtherly, recent studies found that severe cases of COVID-19 were commonly complicated with thrombosis,^{39,40} markedly elevated D-dimer was related to thrombosis and poor prognosis of severe COVID-19 patients. Prothrombin time (PT) reflects the activity of exogenous coagulation factors. COVID-19 associated lung tissue damage may induce the release of tissue factors to circulation and promotes secondary fibrinolysis through exogenous coagulation pathways. This may explain the elevated D-dimer and prolonged PT in COVID-19 as well as CAP patients.^{2,4,18,31,36} The fibrinogen is a kind of coagulation factor, but also an acute phase protein ⁴¹. It can be induced by infection or other stress factors.⁴² Several literatures reported fibrinogen levels was elevated in severe patients or non-survivors with COVID-19.43,44 However, Du et al. found fibrinogen increased in 47.1% of fatal cases and decreased in 22.4% of fatal cases.⁴⁵ In fact, the fibrinogen would decrease when excessive consumption happened due to hypercoagulability or the worst disseminated intravascular coagulation occurred. Hence, the abnormality of the coagulation profile should be interpreted individually.

Biochemistry markers and arterial blood gases

Lactate dehydrogenase (LDH) is a cytoplasmic enzyme that is present in every tissue, and high serum concentrations indicate underlying organ damage. Thus, LDH is expected to rise in severe COVID-19 cases, where multi-organ damage occurs.²³ Current data support that critically ill patients as well as fatal cases of COVID-19 have significantly higher LDH levels (usually >320 U/L) compared to moderate infections.^{1,2,5,6,11,16} Moreover, higher LDH quadruples the odds for lack of disease improvement (odds ratio: 4.381. p<0.0001).¹⁶ Lastly, greater LDH concentrations upon admission correlate with a higher risk for serious COVID-19, and therefore it has been added in a proposed early predictive tool for severe infection.¹⁹ These data favor the utilization of LDH as a candidate prognostic marker for disease severity. Hypertriglyceridemia is commonly encountered in hyperinflammatory states, like the CSS and the secondary HLH, due to the reduced lipoprotein lipase activity driven by the high TNF- α levels.⁴⁶ Therefore, triglyceride concentration is a key component of the HScore [http://saintantoine.aphp.fr/score/] that is currently being proposed by the European Society of Intensive Care Medicine as a predictive tool for SARS-CoV-2-driven sHLH in COVID-19²²,⁴⁷ However, only scarce data on triglycerides levels in COVID-19 disease are currently available; one study reported higher concentrations in fatal cases as compared to patients who survived the disease (median 1.8 vs 1.2 mmol/L), without stating the statistical significance of this finding¹. The same study demonstrated lower thyroid stimulating hormone and free triiodothyronine concentrations in deceased patients, possibly due to critical illness-associated eythyroid sick syndrome¹.

Hyponatremia is a known sequela of lower respiratory tract infections, that is possibly induced by the inappropriate secretion of anti-diuretic hormone.^{48–50} However, not many studies currently report measurement of electrolytes, including sodium. Among these studies, none has depicted statistically or clinically significant differences of sodium or potassium concentrations between severe/fatal and less severe COVID-19 patient groups.^{1,11,18,51}

Although acid-base balance disturbances are expected among COVID-19 patients with multi-organ injury, few data are available; Zhou et al. disclosed significantly higher frequency of acidosis in non-survivors compared to survivors (30% vs 1% respectively, p<0.0001), while Chen et al. reported lower bicarbonate concentration in patients who died, without reporting the statistical significance of this finding.^{1,2} Importantly, but not surprisingly, in the latter study more than 50% of the deceased patients had arterial partial pressure of oxygen (PaO2) of <60 mmHg (compared to 0% in the survivor group), while none in the same group had a partial pressure of oxygen to fraction of inspired oxygen ratio (PaO2:FiO2) of >300¹. Hence, arterial blood gases constitute important prognostic tools for disease severity and poor outcomes, as they are directly associated with the degree of functional lung damage.

Conclusion

Recent clinical research among COVID-19 patients indicates that SARS-CoV-2 infection causes systemic disease, involving multiple organs and systems, including hyperactivation of the immune system, the nervous system and the clotting system. These in turn leading to pathologies in several organs, including the heart, liver and kidneys. In order to stratify patients at risk and to monitor high risk patients at intensive care units, tight laboratory diagnostics provide instrumental information. Laboratory tests can be used as prognostic markers for increased risk and mortality. The spectrum of currently available biomarkers is sufficient to fullfill this purpose. A major limitation of available studies is that the time point of sampling/biomarker assessment since onset of symptoms and/or presentation at health care facilities is not clearly mentioned. Furthermore, currently, there are no internationally acceptable criteria regarding disease severity, which renders evaluation of data quite subjective, depending on individual study investigations. Over the next months and years, with the use of further knowledge on the pathogenesis of SARS-CoV-2 infections, an even more comprehensive list of suitable biomarkers will be developed (Fig. 1 and Tables 1, 2).

Declaration of Competing Interest

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