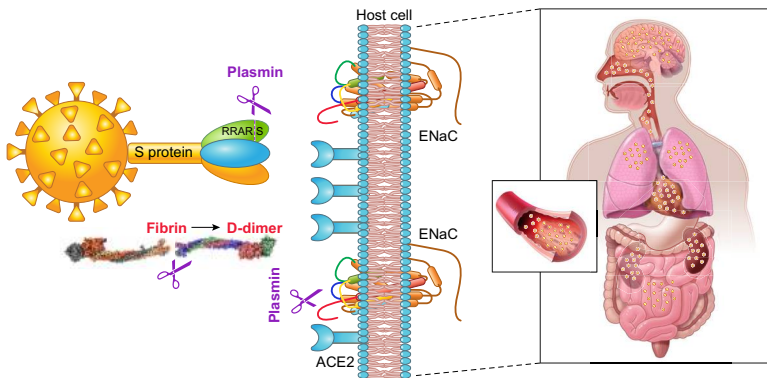


ELEVATED PLASMIN(OGEN) AS A COMMON RISK FACTOR FOR COVID-19 SUSCEPTIBILITY

GRAPHICAL ABSTRACT



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KEYWORDS

comorbidity; COVID-19; fibrinolysis; plasmin(ogen); SARS-CoV-2

CLINICAL HIGHLIGHTS

- 1) Elevated plasmin(ogen) is a common feature in people with underlying medical conditions, including hypertension, diabetes, cardiovascular disease, cerebrovascular disease, and chronic renal illness, who are susceptible to SARS-CoV-2 infection.
- 2) Plasmin enhances the virulence and infectivity of SARS-CoV-2 virus by cleaving its spike proteins.
- 3) Extremely increased D-dimer in COVID-19 patients results from plasmin-associated hyperactive fibrinolysis.
- 4) D-dimer and viral load are independent risk factors of disease severity and mortality.
- 5) Antiproteases targeting plasmin(ogen) may be a promising approach to combat COVID-19.

ELEVATED PLASMIN(OGEN) AS A COMMON RISK FACTOR FOR COVID-19 SUSCEPTIBILITY

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Ji H-L, Zhao R, Matalon S, Matthay MA. Elevated Plasmin(ogen) as a Common Risk Factor for COVID-19 Susceptibility. *Physiol Rev* 100: 1065–1075, 2020. First published March 27, 2020; doi:10.1152/physrev.00013.2020.—Patients with hypertension, diabetes, coronary heart disease, cerebrovascular illness, chronic obstructive pulmonary disease, and kidney dysfunction have worse clinical outcomes when infected with SARS-CoV-2, for unknown reasons. The purpose of this review is to summarize the evidence for the existence of elevated plasmin(ogen) in COVID-19 patients with these comorbid conditions. Plasmin, and other proteases, may cleave a newly inserted furin site in the S protein of SARS-CoV-2, extracellularly, which increases its infectivity and virulence. Hyperfibrinolysis associated with plasmin leads to elevated D-dimer in severe patients. The plasmin(ogen) system may prove a promising therapeutic target for combating COVID-19.

comorbidity; COVID-19; fibrinolysis; plasmin(ogen); SARS-CoV-2

I.	INTRODUCTION	1065
II.	CLINICAL, PATHOLOGICAL, AND...	1065
III.	CLEAVAGE OF SARS-CoV-2 S...	1066
IV.	ELEVATED PLASMIN(OGEN) LEVELS...	1068
V.	PLASMIN(OGEN) IN ARDS	1069
VI.	CLINICAL RELEVANCE AND...	1071

mortality of patients with comorbidities are unknown. The existence of significantly increased fibrin degradation products (FDPs) and reduced platelets in severe COVID-19 patients is consistent with the presence of hyperfibrinolysis. This opinion is supported by the presence of hemorrhage in multiple organs and a positive correlation between fibrinolysis and mortality. Plasmin, a key player in fibrinolysis, enhances the virulence and pathogenicity of viruses containing a furin site in their envelope proteins, as is the case with the SARS-CoV-2. The purpose of this review is to summarize the clinical and preclinical evidence for the existence of elevated plasmin(ogen) in these comorbid conditions of COVID-19 and to highlight the importance of plasmin-induced proteolytic cleavage of the SARS-COV-2 S protein and fibrin in the development of COVID-19.

II. CLINICAL, PATHOLOGICAL, AND EPIDEMIOLOGICAL FEATURES OF COVID-19

A. Epidemiology

Coronavirus disease 2019 (COVID-19) is caused by a new β -coronavirus, SARS-CoV-2. Its epicenter was in Wuhan, China, and it has been spreading globally (103). As of March 31, 2020, there have been 858,955 cases worldwide

- 1) Elevated plasmin(ogen) is a common feature in people with underlying medical conditions, including hypertension, diabetes, cardiovascular disease, cerebrovascular disease, and chronic renal illness, who are susceptible to SARS-CoV-2 infection.
- 2) Plasmin enhances the virulence and infectivity of SARS-CoV-2 virus by cleaving its spike proteins.
- 3) Extremely increased D-dimer in COVID-19 patients results from plasmin-associated hyperactive fibrinolysis.
- 4) D-dimer and viral load are independent risk factors of disease severity and mortality.
- 5) Antiproteases targeting plasmin(ogen) may be a promising approach to combat COVID-19.

I. INTRODUCTION

Patients with preexisting hypertension, diabetes, coronary heart disease, cerebrovascular disease, chronic obstructive pulmonary disease (COPD), and kidney dysfunction (comorbidities) have worse clinical outcomes when infected with SARS-CoV-2. The only treatment of COVID-19 is supportive (51), and registered clinical trials are ongoing. The mechanisms for high morbidity and

resulting in 42,119 deaths.¹ This far exceeds the total deaths caused by both severe acute respiratory syndrome (SARS) and Middle Eastern respiratory syndrome (MERS) (91). Persons older than 60 years with hypertension, diabetes, COPD, as well as cardiovascular, cerebrovascular, liver, kidney, and gastrointestinal diseases are more susceptible to the infection by SARS-CoV-2 and experience higher mortality when they develop COVID-19 (3, 80, 101). The contribution of malignant conditions is under debate due to the small number of patients (43, 92). In total, 19% of COVID-19 patients develop acute respiratory distress syndrome (ARDS) ($\text{PaO}_2/\text{FiO}_2 < 300$ mmHg) within 24–48 h after onset of symptoms.

B. Viral Load

Virus clearance is associated with the severity and survival of COVID-19. Viral load in the respiratory tract peaks at day 5–6 after the onset of symptoms (10^4 – 10^7 copies/ml) (62), and viral RNA can be found in stool and sputum samples, bronchoalveolar lavage fluid (BAL), and lung epithelial cells. Patients older than 65 years generally have higher viral load lasting up to 14 days and may develop severe acute lung injury, requiring hospitalization in the intensive care unit (ICU) with poor outcome (62). In contrast, most younger patients have a much lower viral load that is undetectable within 1 week after onset (104). Furthermore, an association between viral load and the severity of COVID-19 has been reported (46).

C. Leading Causes of Death

COVID-19 patients admitted to ICU have higher mortality (38%) than non-ICU patients (4%) (31). The mortality of patients who develop ARDS is 49% (45). Many patients with COVID-19 develop multi-organ failure (MOF). The leading causes of deaths are ARDS, septic shock with MOF, hemorrhage/coagulopathy (disseminated intravascular coagulopathy, DIC), acute heart/liver/kidney injury, and secondary bacterial infections (97, 101). Elevated FDPs and D-dimers were detected predominantly in patients with severe disease (11, 24, 31, 43, 45, 79, 85). Multivariate regression further suggests that age and D-dimer levels (>1 mg/L) are two independent risk factors for mortality (89, 101). This in addition to other factors, such as compromised immune response, may contribute to the increased morbidity and mortality of patients with COVID-19 who are older than 60 years.

D. Pathology

The pathological features of COVID-19 resemble those of SARS and MERS. In the early stages of infection, puncture

lung biopsies reveal the presence of pneumonia, edema, proteinaceous exudate with globules and focal hyperplasia of alveolar epithelial cells associated with patchy inflammatory infiltrates, and multinucleated giant cells (81). At the later stages, diffuse alveolar damage (DAD) is observed in addition to hemorrhage and some areas of interstitial fibrosis (48). Fibrotic clots and gelatinous mucus in the small airways and disseminated intravascular coagulation are also present (44, 93). Consistent with clinical observations, the lungs are the most injured organs, followed by moderate injury in the heart, liver, kidney, and brain. Systemic microthrombi in the circulatory system and hemorrhage in the affected organs result from noncoordinated responses between the coagulation and fibrinolysis systems. Although COVID-19 is characterized by hyperfibrinolysis, as evidenced by elevated levels of D-dimer, studies attempting to restore fibrinolytic function have not been reported. Elevated plasmin(ogen) in patients with preexisting conditions may be a mechanism contributing to enhanced susceptibility to SARS-CoV-2 infection and fatality.

III. CLEAVAGE OF SARS-CoV-2 S PROTEINS BY HOST FURIN AND PLASMIN

Sequencing of hundreds of SARS-CoV-2 virus isolates reveals a close relation to two bat-derived coronaviruses, bat-SL-CoVZC45 and bat-SL-CoVZXC21. These coronavirus strains have a similar receptor-binding domain structure in the Spike (S) protein for host angiotensin converting enzyme 2 (ACE2) proteins (47, 90, 102). The S protein of SARS-CoV-2 bind to human ACE2 receptors with higher affinity than that of the SARS-CoV virus (88). This may be due to a furin-like cleavage site ($^{682}\text{RRAR}/\text{S}^{686}$) inserted in the S1/S2 protease cleavage site of the SARS-CoV-2 virus (**FIGURE 1**) (13). The S1 region of the Spike protein is responsible for binding to the host cell ACE2 receptor, where the S2 region is responsible for fusion of the viral RNA and cellular membranes. Polybasic furin sites in hemagglutinin (HA) proteins are often found in highly virulent avian and human influenza viruses (10). The insertion of the furin site may augment the ability of this new SARS-CoV-2 to attach and invade human cells expressing ACE2 and CD147 receptors (12, 86).

A. Furin Proteolytically Cleaves Other Viral Proteins

The envelope proteins of numerous viruses, such as human immunodeficiency virus (HIV), human and avian influenza, herpes, Epstein-Barr, leukemia, dengue, Ebola, hepatitis B, measles, West Nile, Zika, respiratory syncytial virus (RSV), SARS, MERS, and Marburg virus, are cleaved by intracellular furin-like proteases. This increases the ability of the viruses to enter host cells (13, 34, 56). Proteolysis at a

¹These numbers are increasing daily now.

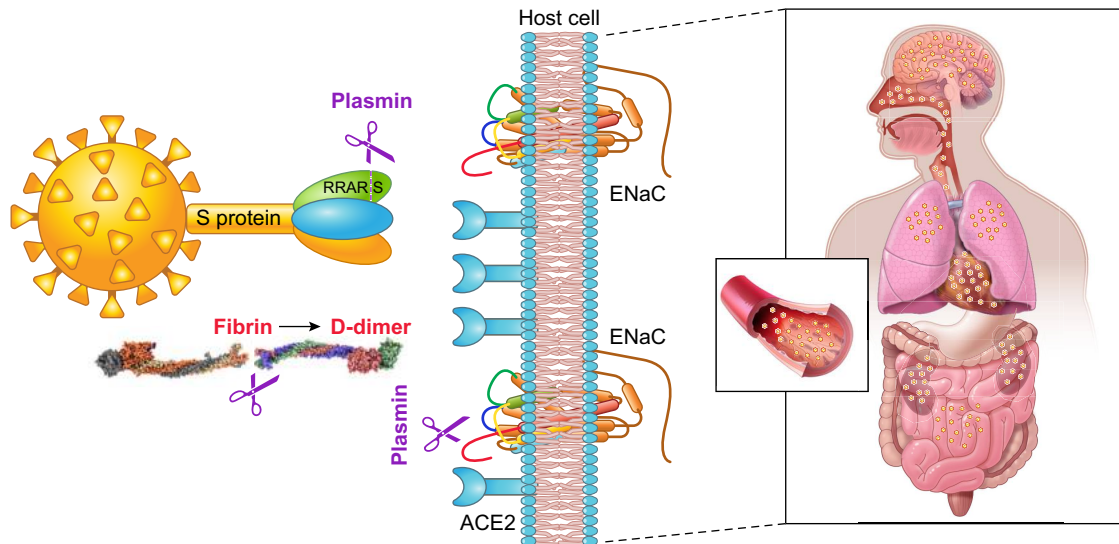


FIGURE 1. Plasmin(ogen) increases the pathogeneticity of COVID-19. Plasmin cleaves the S protein of SARS-CoV-2 extracellularly, increasing its ability to bind with angiotensin converting enzyme 2 (ACE2) receptors of host cells, and probably facilitating virus entry and fusion. Plasmin proteolytically breaks down excess fibrin to elevate D-dimer and other fibrin degradation products in both bronchoalveolar lavage fluid and plasma, which decreases platelets and results in hemorrhage. Plasmin also cleaves epithelial sodium channel (ENaC) subunits, located at the apical membranes of epithelial cells in the airway, lung, and kidney. This increases the ability of Na^+ ions to enter epithelial cells resulting in hypertension and dehydration of the fluid lining lung airways and alveolar cells.

conventional “XXXR/S” motif in the S2 region of the Spike protein may facilitate the entry of respiratory infectious viruses (such as RSV and influenza) into airway and alveolar epithelial cells (94). Both SARS and MERS have evolved an unusual two-step furin activation for fusion, suggestive of a role during the process of emergence into the human population (55). Amidst the cluster of the furin-cleaved viruses, SARS S protein is cleaved by airway proteases (trypsin, plasmin, and TMPRSS11a), expressed in human bronchial epithelial cells, subsequently enhancing pseudovirus entry via binding host ACE2 receptors (37). The protease cleavage sites are R667 in the S1 fragment and R797 in the S2 fragment of SARS virus. These two cleavage sites are preserved in other coronaviruses, HIV, human and avian influenza virus, human CMV and RSV, yellow fever virus, and Zika virus (13).

B. Cleavage of Coronavirus by Plasmin and Other Host Proteases

Single-cell profiling of human lung tissues (the LGEA portal: <https://research.cchmc.org/pbge/lunggens/mainportal.html>) reveals that furin is predominately expressed in human alveolar type II (AT2) cells in the respiratory system, while plasminogen, kallikrein, and trypsin are expressed in both airway and alveolar type I and II epithelial cells. Plasminogen is also expressed in endothelial cells. Cytosolic furin is enriched in the Golgi apparatus. The possibility for non-furin proteases to cleave viral envelope proteins is supported by the evidence that in furin-defective LoVo cells, the

cleavage-dependent process of HIV gp160 is as efficient as in normal cell lines (60). We have demonstrated that plasmin is capable of cleaving furin sites in the γ subunit of human epithelial sodium channels (ENaC) (99).

The S protein of coronaviruses may be cleaved by plasmin, trypsin, cathepsins, elastase, and TMPRSS family members; cleavage of S protein may mediate enhancement of virus entry into bronchial epithelial cells (37). Plasmin also cleaves the S proteins of SARS-CoV in vitro (37). In addition, HCoV-HKU1 S proteins are cleaved by kallikrein in the S1/S2 region and mediate the entry of HCoV-HKU1 to nonpermissive rhabdomyosarcoma cells (54). The clinical relevance of non-furin cleavage remains unknown due to the paucity of in vivo evidence for the role of plasmin cleavage of SARS-CoV. Also, it remains to be demonstrated that the envelope proteins of SARS-CoV-2 strain are cleaved by plasmin (56).

The cleavage of influenza virus by plasmin is well characterized (5, 23, 42, 57, 73, 83, 96). Proteolysis of influenza HA proteins enables fusion with the host endosome. Acidification of the endosome promotes viral membrane fusion and activates the M2 ion channel, which pumps protons (H^+) into the interior of the viral core to initiate uncoating of the M1 protein. Nuclear replication occurs, and viral gene products are transported to the plasma membrane for assembly. The fibrinolytic zymogen plasminogen (activated by urokinase or tissue-like plasminogen activator to generate plasmin) has been shown to cleave the influenza HA

proteins (40, 75, 83). The HA cleavage site of A/WSN/1933 H1N1 influenza virus governs virus spread in a plasmin-dependent manner (75). Mini-plasmin, a plasmin fragment, is distributed predominantly in the epithelial cells of the bronchioles and potentiates the replication of both plasmin-sensitive and plasmin-insensitive influenza A virus strains, suggesting a pivotal role of plasmin in the spread and pathogenicity of the influenza virus (57). Additionally, kallikreins cleave and activate HA of the influenza virus H1, H2, and H3 subtypes (27). Similar to coronavirus and influenza viruses, plasmin, trypsin, thrombin, and furin enhance RSV-induced cytopathology (17). Local fibrin and clot formation are implicated in host defense against influenza virus infections (5), thus plasminogen may affect lung injury and repair by interfering with these processes (1).

Protease cleavage may enhance or decrease the activities of various proteins. For example, prostasin increases the activity (60–80%) of ENaC, whereas TMPRSS2 markedly decreases ENaC function and protein levels (15). In neural tissues, brain-derived neurotrophic factor precursor (proBDNF) is cleaved either intracellularly by furin-like proteases or extracellularly by plasmin or matrix metalloproteinases. However, plasmin, but not related proteases, cleaves proBDNF furin sites extracellularly (4). The inhibitory effects of TMPRSS2 could be corrected by serine protease inhibitors, such as camostat mesylate that has been approved for clinical use in Japan (98). The beneficial effects of camostat mesylate, an antiprotease, may be partially due to the inhibition of plasmin (98).

C. Proteolytical Cleavage and Pathogenicity

Extracellular cleavage of virus envelope fusion glycoproteins by host cellular proteases is a prerequisite for the infectivity of respiratory viruses. The presence of a polybasic cleavage site that can be cleaved by furin-like proteases is a signature of several highly pathogenic avian influenza viruses (82). Similarly, the S protein of SARS-CoV-2 harbors a furin cleavage site at the S1/S2 boundary. The almost ubiquitous and diverse expression of furin-like proteases could lead to increasing SARS-CoV-2 cell and tissue tropism and transmissibility, and enhance its pathogenicity (84).

Four viral proteins are essential for the pathogenesis of COVID-19. The S proteins bind to ACE2 receptors after being cleaved by furin-like proteases. The RNA-dependent RNA polymerase (RdRp) is responsible for replicating SARS-CoV-2 RNA genome. 3C-like and papain-like proteases cleave two polyproteins that are important for the packing new virions. Whether plasmin and other host proteases cleave additional viral proteins is not known.

IV. ELEVATED PLASMIN(OGEN) LEVELS IN COMORBID DISEASES OF COVID-19

A. Plasmin(ogen) in Hypertensive Patients

Plasmin is generated from the cleavage of plasminogen by either urokinase (uPA) or tissue-like plasminogen activator (tPA). In general, uPA is responsible for the plasmin in body fluid (BAL, urine, tear, pleural effusion, etc.), while circulating tPA proteolytically cleaves plasminogen in the plasma. The most frequent comorbidity observed in COVID-19 patients is hypertension followed by diabetes, chronic cardiovascular conditions, cerebrovascular diseases, COPD, and chronic kidney illnesses (80, 101). The plasminogen system is a druggable target in renal hypertension (76). Plasmin, a potent protease, cleaves up to 16 sites, including the cleavage sites for trypsin, chymotrypsin, prostatic, and elastases, of the human γ ENaC subunit (99). Elevated renal plasmin results in hypertension by cleaving ENaC in the collecting tubule, which increases salt retention, causing expanded circulating volume. Urinary excretion of plasmin(ogen) and urokinase directly correlates with urine albumin in hypertensive subjects (2). On the other hand, amiloride, an inhibitor of ENaC, lowers blood pressure and urine plasminogen excretion (61). ENaC proteins are located in the apical membranes of tight epithelia, and they are the major pathways for the entry of Na^+ . As such, they play an important role in maintaining the proper depth of airway and alveolar lining fluids, the reabsorption of edema fluid in injured lungs, and the regulation of salt retention in the collecting tubules (6, 14, 18, 28, 38, 39, 50, 64). Proteolysis is an important regulatory mechanism of ENaC function (36, 63, 65, 67, 70, 99). Decreased ENaC function will result in increased fluid in body cavities (e.g., lung edema in the airspaces and increased blood volume), while increased function will cause dehydration of luminal fluid, as it occurs in cystic fibrosis (CF) and most likely dry eye syndrome.

B. Plasmin(ogen) in Cardiovascular Diseases

Significantly higher levels of urinary plasminogen and plasmin are reported in rats (71) and patients with chronic heart failure (76, 100). In addition, plasmin activity in patients with coronary artery disease is 1.7-fold greater compared with healthy subjects (16).

C. Plasmin(ogen) in Diabetes

Both types I and II diabetes are associated with higher plasmin(ogen) levels in plasma. A 25-year prospective study of type I diabetes documented an association with increased urinary plasmin(ogen), particularly in hypertensive subjects (69). Concentrations of plasmin(ogen) in urine are correlated with the development of preeclampsia late in preg-

nancy (58). Aberrant plasmin in preurine may inappropriately activate ENaC in patients with type II diabetes and microalbuminuria (7). Individuals with high plasma furin concentration have a pronounced dysmetabolic phenotype and elevated risk of diabetes mellitus and premature mortality (19).

D. Plasmin(ogen) in Other Comorbid Diseases

Higher levels of plasmin(ogen) are detected in the urine of various cancer patients as compared with healthy individuals (9). Elevated urine plasmin(ogen) levels, accompanied by increased exosomal α ENaC fragments, have been detected in pregnant women (59), a population susceptible to influenza (22). Endogenous channel-activating proteases, as well as proteases released by inflammatory cells (trypsin, elastase), activate ENaC either by cleaving critical amino acids in α and γ ENaC subunits, or by activating signaling pathways (66, 72). Aprotinin, a potent and reversible Kunitz-type inhibitor of several serine proteases, including trypsin, plasmin, and kallikreins, has been reported to inhibit sodium transport among a variety of epithelial cells (66). Other Kunitz-type serine protease inhibitors, such as hepatocyte growth factor activator inhibitor (HAI)-1 and HAI-2 (placental bikunin), have also been demonstrated to inhibit prostatic and ENaC activity (77). Finally, a1-antitrypsin, an acute-phase glycoprotein and a member of the serine protease inhibitor (SERPIN) superfamily, inhibits ENaC in vitro and in vivo by decreasing protease activity (41). Of note, SARS S protein inhibits ENaC via the protein kinase C signaling pathway (35). It is worth noting that high plasmin levels may contribute to the development of comorbid bacteremia and sepsis (26, 74). Interestingly, azithromycin, a common antibiotic for suppressing infection in the CF airways when combined with hydroxychloroquine, turns COVID-19 to SARS-CoV-2 negative in 5 days (21).

V. PLASMIN(OGEN) IN ARDS

A. Plasmin(ogen) Is Increased in ARDS

ARDS is a life-threatening disorder associated with respiratory and systemic infections, trauma, burns, inhalation of toxic gases, and aspiration of gastric contents injury (52). In addition to lung injury, patients with ARDS may develop MOF with a hallmark of excess D-dimer and other FDPs presenting in both BAL and blood biopsies. Soluble D-dimer and D-monomer are predominately produced from the proteolytic cleavage of cross-linked fibrin and fibrinogen/non-cross-linked fibrin, respectively, by plasmin (fibrinolysis). Plasmin activity in BAL is detected in healthy subjects (71). Both D-dimer and D-monomer levels are significantly increased up to 17-fold in undiluted edema fluid

in patients with ARDS (68). Significant increase in both plasminogen and cleaved plasmin protein in the BAL of ARDS patients (32) and an animal model of DAD (33) have been reported. Augmented plasmin activity contributes to elevated D-dimer in the BAL of infected lungs in a time-dependent manner during the development of ARDS (20, 87). This is further validated by the observation that plasmin-mediated fibrinolytic activity could be inhibited by 50% with α 2-antiplasmin antibody (32). Kallikrein and neutrophil elastase may contribute to the residual proteolytic activity in BAL of ARDS (53). Also, radiation-induced lung injury in mesothelioma patients is accompanied by a significant elevation in BAL plasminogen and plasmin-associated fibrinolytic activity (49).

B. Fibrinolysis in COVID-19

In comparison with patients with mild COVID-19 (such as those who did not require ICU stays, did not develop ARDS or pneumonia, and who survived), patients with severe COVID-19 have higher comorbidities, including 56% for hypertension, 21% for heart diseases, 18% for diabetes, 12% for cerebrovascular diseases, and 7% for cancer (TABLE 1) (79, 97). Some patients have more than one, even up to five preexisting conditions. Multivariate regression further links hypertension with increased incidence and fatality (85, 89, 101). Hyperfibrinolysis, reflected by elevated serum D-dimer levels, was present in 97% of COVID-19 patients at admission and increased further in all patients before death (TABLE 2) (97). FDPs were significantly increased as well (79). This is accompanied by a prolonged prothrombin time particularly in non-survivors (31, 79, 97, 101). Platelet counts were decreased significantly in severe and dead patients (79, 97, 101). 71.4% of non-survivors meet the criteria of the International Society on Thrombosis and Hemostasis (ISTH) for DIC, suggesting the coexistence of coagulation activation and hyperfibrinolysis in patients with severe COVID-19 infection (78, 85). In contrast, D-dimer levels decreased to control levels in survivors or non-ARDS patients.

The mortality rate of patients with COVID-19 who did not develop ARDS is 9 versus 49% for those who did develop ARDS (45). Of note, ARDS/respiratory failure remains the leading cause of death (70%), followed by sepsis/MOF (28%), heart failure (15%), hemorrhage (6%), and renal failure (4%) (TABLE 3). Coagulation/hemorrhage ranks among the top three leading causes of death (97). Furthermore, multivariate regression analysis identifies D-dimer and age as independent risk factors for mortality (TABLE 4) (85, 89, 101). These findings suggest that the normalization of hyperactive fibrinolysis may be a therapeutic target.

Table 1. Comorbidities of COVID-19 patients

	Epidemiology (death%) (n = 72,314)	Severe/ non-severe Pneumonia (n = 38/72)	Severe/ non-severe (n = 173/926)	ICU/ non-ICU (n = 13/28)	ARDS/ non-ARDS (n = 53/56)	Non-survivor/ survivor (n = 54/137)	Non-survivor/ survivor (n = 32/20)	Non-survivor (n = 82)
Hypertension	12.8/6	39.47/29.17 ^c	23.7/13.4	15/14	39.6/28.6	48/23 % ^c		56.1
Diabetes	5.3/7.3	21.05/9.72	16.2/5.7	8/25	20.8/1.8 ^b	31/14 ^b	22/10	18.3
Coronary heart diseases	4.2/10.5		5.8/1.8	23/11	5.7/7.1	24/1 ^d	9/10	20.7
Cerebrovascular diseases		7.89/5.56	2.3/1.2		11.3/0 ^b		22/0	12.2
COPD	2.4/6.3	10.53/2.78 ^a	3.5/0.6	8/0	3.8/3.6	7/1 ^a	6/10	14.6
Kidney diseases			1.7/0.5		15.1/3.6 ^a	4/0 ^a		4.9
Liver diseases			0.6/2.4	0/1		—		2.4
Cancer	0.5/5.6		1.7/0.8	0/1		0/1	3/5	7.3
Secondary infection			0.6/2.4					6.1
Immunodeficiency			0/0.2					17.1
Others						20/8 ^a		3.7 (surgery)
Total	26/—		38.7/21	38/29		67/40 ^c		76.8
Reference no.	80	85	24	31	45	101	95	97

ARDS, acute respiratory distress syndrome; COPD, chronic obstructive pulmonary disease; ICU, intensive care unit. ^aP < 0.05, ^bP < 0.01, ^cP < 0.001, ^dP < 0.0001.

C. Uncoordinated Coexistence of Hypercoagulation and Hyperproteolysis

The specific plasmin inhibitor α2-antiplasmin is elevated by approximately one order of magnitude in patients with ARDS, while fibrinolytic activity is reduced approximately by half, and D-dimer is elevated 50-fold in BAL (25). The nonproportional change between the expression and activity level of plasmin(ogen) and anti-plasmin indicates the stoichiometry of the plasmin-antiplasmin complexes may not be in a ratio of 1:1. Increased levels of α2-antiplasmin

and other antiproteases may not completely shield the proteolytic triad of plasmin in the complexes, suggesting that either their efficacy is inadequate or that plasmin is still able to cut fibrin to produce D-dimers and FDPs in ARDS patients. The soluble complexes of the plasmin-antiplasmin in BAL may facilitate physical interactions with the vast deposition of fibrin at the luminal surface of alveoli. Based on the pathology and laboratory results, dynamic hypercoagulation occurs as evidenced by microthrombi throughout the blood vessels of multiple organs, accompanied with extremely reduced platelets in COVID-19 patients (TABLE 2).

Table 2. Coagulation and fibrinolysis in patients with COVID-19

	Severe/ non-severe (n = 173/926)	ARDS/ non-ARDS (n = 53/56)	ICU/ non-ICU (n = 13/28)	Severe/ non-severe Pneumonia (n = 38/72)	Non-survivor/ survivor (n = 54/137)	Non-survivor/ survivor (n = 21/162)	Non-survivor/ survivor (n = 32/20)	Non-survivor (n = 82)
D-dimer, >1 mg/L (≥0.5 mg/L)	59.6/43.2	940/370 ^c	2.4/0.5 ^b	1.11/0.37 ^c	5.2/0.6 ^d	2.12/0.61 ^c , 100/		97.5–100 (>0.55 mg/L)
FDP, mg/L						7.6/4.0 ^c		
Fibrinogen, g/L		3.4/2.9 ^c				5.16/4.51, 28.6 (<1 g/L)		—
Platelets, <100 × 10 ⁹ /L (<150,000/mm ³)	57.7/31.6		8/4	144.5/179.5 ^c	20%/1% ^d	57.1/	191/164	63.2
Prothrombin time, ≥16 s			12.2/10.7 ^a		13/3 ^a	15.5/13.6 ^c		100 (>12.1)
Antithrombin activity						84/91	12.9/10.9	
APTT, s			26.2/27.7			44.8/41.2		
ISTH DIC criteria						71.4/-		
Reference no.	24	45	31	85	101	79	95	97

APTT, activated partial thromboplastin time; ARDS, acute respiratory distress syndrome; DIC, disseminated intravascular coagulation; FDP, fibrin degradation products; ICU, intensive care unit; ISTH, International Society on Thrombosis and Hemostasis. ^aP < 0.05, ^bP < 0.01, ^cP < 0.001, ^dP < 0.0001.

Table 3. Outcomes or complications (%) in patients with COVID-19

	Severe/non-severe (n = 173/926)	ICU/non-ICU (n = 13/28)	Non-survivor/survivor (n = 54/137)	Non-survivor/survivor (n = 32/20)	Injured Organs (n = 82)	Death Cause (n = 82)
Sepsis/MOF			100/42 ^d	3/0	28.0 ^e	
Respiratory failure			98/36 ^d			100
ARDS	16.5/1.1	85/4 ^d	93/7 ^d	81/45	69.5 ^e	
Septic shock	6.4/0.1	23/0 ^a	70/0 ^d			
Acute cardiac injury		31/4 ^a	59/1 ^d	28/15		89.0
Heart failure			52/12 ^d		14.6 ^e	
Coagulopathy/hemorrhage			50/7 ^d		6.1 ^e	80.5
Acute kidney injury	2.9/0.1	23/0 ^a	50/1 ^d	37.5/15		31.7
Secondary infection		31/0 ^a	50/1 ^d	9/20		
Hypoproteinemia			37/1 ^d			
Acidosis			30/1 ^d		2.4 ^e	
Renal failure	0.6/0				3.7 ^e	
Liver failure					1.2 ^e	78.0
GI failure					2.4 ^e	6.1
Reference no.	24	31	101	77	97	97

ARDS, acute respiratory distress syndrome; GI, gastrointestinal; ICU, intensive care unit; MOF, multiple organ failure. ^a*P* < 0.05, ^b*P* < 0.01, ^c*P* < 0.001, ^d*P* < 0.0001. ^ePercent of contribution to death.

On the other hand, hemorrhage and markedly elevated degraded fibrin products result from plasmin-associated hyperproteolysis. Whether patchy hemorrhage coexists with areas infected by SARS-CoV-2 is not known. Administration of anti-proteases may prove beneficial.

VI. CLINICAL RELEVANCE AND PERSPECTIVE

The cleavage of the new furin sites in the S protein of SARS-CoV-2 virus by plasmin and other proteases may enhance its infectivity by expediting entry, fusion, duplication, and release in respiratory cells. Elevated plasmin(ogen) levels are a common feature in COVID-19 patients with underlying medical conditions. The elevated plasmin(ogen) could be an independent factor for risk stratification of patients with COVID-19. Measurements of plasmin(ogen) levels and its enzymatic activity may be important biomarkers of disease severity in addition to resultant D-dimer. The ad-

ministration of antiproteases to suppress plasmin activity in the respiratory system may prevent, or at least decrease, SARS-CoV-2 entry into respiratory cells and improve the clinical outcome of patients with COVID-19. As demonstrated in vitro, a serine protease inhibitor for TMPRSS2 blocks SARS-CoV-2 S protein-driven entry into cells (30). Clinical trials conducted in China are testing various protease inhibitors (29). Currently there are no proper animal models of COVID-19 with underlying medical conditions to test new therapeutic agents. Healthy mice and monkeys infected with SARS-CoV-2 develop either mild lung injury or show no symptoms of disease (8). It remains to be seen whether mice and monkeys with preexisting comorbid conditions and higher plasmin levels develop COVID-19 when infected with SARS-CoV-2. Targeting hyperfibrinolysis with a broad spectrum or specific anti-plasmin compounds may prove to be a promising strategy for improving the clinical outcome of patients with comorbid conditions.

Table 4. Risk factors of COVID-19 associated with mortality computed with multivariate logistic regression

	Non-survivor/survivor (OR, n = 54/137)	Severe/non-severe Pneumonia (OR, n = 38/72)	ARDS (HR, n = 201)
Age	1.10 (1.03, 1.17) ^b	25.314 (1.628, 92.664) ^c >60 yr	6.17 (3.26, 11.67) ^c
Lymphocyte	0.19 (0.01, 1.62)	0.322 (0.137, 0.756) ^b	0.51 (0.22, 1.17)
D-dimer	18.42 (2.64, 128.55) ^b	17.054 (2.547, 114.171) ^b	1.02 (1.01, 1.04) ^b
Reference no.	101	85	89

ARDS, acute respiratory distress syndrome; HR, hazard ratio; OR, odd ratio. ^a*P* < 0.05, ^b*P* < 0.01, ^c*P* < 0.001.

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DISCLOSURES

No conflicts of interest, financial or otherwise, are declared by the authors.

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