



The triumvirate: why hypertension, obesity, and diabetes are risk factors for adverse effects in patients with COVID-19

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Abstract

The outbreak of coronavirus disease 2019 (COVID-19) caused by a novel coronavirus, severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), has become a pandemic. The cellular receptor for SARS-CoV-2 entry is the angiotensin-converting enzyme 2, a membrane-bound homolog of angiotensin-converting enzyme. Henceforth, this has brought the attention of the scientific community to study the interaction between COVID-19 and the renin–angiotensin system (RAS), as well as RAS inhibitors. However, these inhibitors are commonly used to treat hypertension, chronic kidney disorder, and diabetes. Obesity is a known risk factor for heart disease, diabetes, and hypertension, whereas diabetes and hypertension may be indirectly related to each other through the effects of obesity. Furthermore, people with hypertension, obesity, diabetes, and other related complications like cardiovascular and kidney diseases have a higher risk of severe COVID-19 infection than the general population and usually exhibit poor prognosis. This severity could be due to systemic inflammation and compromised immune response and RAS associated with these comorbid conditions. Therefore, there is an urgent need to develop evidence-based treatment methods that do not affect the severity of COVID-19 infection and effectively manage these chronic diseases in people with COVID-19.

Keywords SARS-CoV-2 · Viral infection · Comorbidities · ACE2

Introduction

In December 2019, a new coronavirus (2019-nCoV) causing severe acute respiratory syndrome (SARS-CoV-2) was reported in Wuhan, China, and the world has witnessed a subsequent outbreak of this disease, known as COVID-19 (Coronavirus Disease 2019). By January 30, 2020, a transition of this localized outbreak led the World Health Organization (WHO) to declare it as public health emergency of national concern (PHEIC) and later as a pandemic by March 11, 2020. As of August 5, 2020 [1], there have been over

18,000,000 of total confirmed cases of infection and more than 700,000 deaths due to COVID-19 worldwide. Countries around the world are suffering from an unprecedented burden to health care and the overall economy. COVID-19 manifestations range from asymptomatic to severe or critical type with the need for hospitalization or intensive care unit (ICU) [2, 3]. The common clinical features include fever and respiratory symptoms like cough, dyspnea [4, 5], and in severe cases, sepsis, multiple organ dysfunction including cardiac injury or kidney failure can develop [3, 6]. Clinical data have suggested a correlation between the severity of COVID-19 infection outcome with advanced age and several comorbidities including hypertension, obesity, and diabetes [5, 7–9]. The severity of complications and a need for hospitalization are more when COVID-19 infection is associated with underlying comorbid complications [5, 7, 9–11]. The prevalence of hypertension, obesity, and diabetes among hospitalized COVID-19 patients is much more than nonhospitalized COVID-19 patients [9, 12]. Hypertension, obesity, and diabetes are closely linked to each other, and these comorbidities are associated with an elevated risk of recurrent infection due to underlying impairments in the host

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defense system [13]. As an efficient and integrated immune defense system is necessary for the body's response to infection, a compromised immune response makes the host more vulnerable to infection and related severe complications.

This review is aimed at highlighting why conditions of hypertension, obesity, and diabetes increase the risk for infections, and more particularly how preexistence of these metabolic diseases in patients with COVID-19 causes severe manifestations and clinical outcomes. The available information about this novel COVID-19 pandemic is based on limited clinical data and information from similar previous infections including H1N1 influenza, severe acute respiratory syndrome-coronavirus (SARS-CoV), and Middle East respiratory syndrome (MERS). Currently, there are no vaccines or reliable treatments available for the prevention or treatment for CoV-2 infection; therefore, it is important to understand its mechanism of infection and higher risk among individuals with obesity, diabetes, and hypertension to possibly reduce the morbidity and associated mortality.

Hypertension, obesity, and diabetes: coexistence of the triumvirate with an increased risk for respiratory infections

The rising prevalence and disease burden of obesity have received significant attention globally over the past three decades. Many epidemiological studies have suggested a high body mass index (BMI) as a risk factor for many chronic diseases, including diabetes, cardiovascular disease (CVD), chronic kidney disease, hypertension, and infections. Hypertension, obesity, and diabetes are interconnected with an elevated risk of renal complications and cardiovascular diseases, and the coexistence of these diseases increases the disease-associated mortality. Considering the multifactorial and complex nature of these conditions, including the combination of environmental, genetic, behavioral, and lifestyle-related factors, we are yet to clarify all driving forces linking this triumvirate [14].

Viral infections including H1N1 influenza, SARS-CoV (severe acute respiratory syndrome-coronavirus) and MERS-CoV (middle eastern respiratory syndrome-coronavirus) cause respiratory infections, and the coexistence of various comorbidities like hypertension, obesity, and diabetes has been linked with severe illness among these patients. SARS-CoV and MERS-CoV were first reported in 2002 and 2012, respectively [15, 16]. Hyperglycemia and known history of diabetes were reported as an independent predictor of mortality and morbidity among SARS patients [17]. Similarly, systemic analysis of MERS-CoV infection cases

showed that diabetes and hypertension were equally prevalent in 50% of patients and obesity was prevalent in 16% of the patients. Additionally, these underlying comorbidities were found to be associated with severe illness and mortality in MERS-CoV patients [18]. In 2008–09, H1N1 influenza A virus transmitted from swine to human, later developing into a human to human transmission, caused a pandemic and is currently the predominant strain for seasonal human influenza [19]. During the influenza viral infection, the presence of obesity and related comorbidities showed an increased incidence of morbidity and mortality [20, 21]. A systematic review and meta-analysis study showed the association between obesity with a higher risk of ICU admission and mortality among H1N1-infected patients [22]. Taken together, these data suggest that people with obesity are more susceptible to increased risk of influenza-related hospitalization, morbidity, and mortality.

Limited availability of clinical data about the outcomes of H1N1 infection in diabetes patients makes it nonconclusive [23–25]. A study conducted with non-Hispanic Black and White US adults, to assess the impact of diabetes in influenza-related mortality, found that irrespective of race, sex, or socioeconomic status, people with diabetes are more likely to have higher medical consultation, morbidity, mortality, and hospitalization compared to the nondiabetic subjects [26]. However, the worse outcomes among diabetes patients with H1N1 infection could be a consequence of related underlying comorbidities including obesity and cardiovascular diseases but not diabetes itself alone [27].

In hypertension, the increased sympathetic nervous system (SNS) activity and decreased parasympathetic nervous system (PNS) increase the blood pressure and promote activation of innate immunity via Toll-like receptors (TLRs), which can induce oxidative stress and inflammation. These events promote disease progression and lead to organ damage [28]. Obesity is associated with low-grade systemic inflammation which leads to elevated circulatory pro-inflammatory cytokines [29]. Excessive overload of nutrients induces oxidative stress, increases infiltration of macrophages, and further leads to activation of inflammatory processes [30–32]. Moreover, in diabetes condition, immune response to infection is compromised [33]. Impaired glycemic control is also associated with alteration in various immune mechanisms against viral infections [34]. Therefore, the compromised host immune defense makes them vulnerable to infection. The increased incidence of morbidity and mortality with respiratory infections in the presence of obesity and related comorbidities suggests possible similar risks of severe outcomes for individuals with these underlying conditions during the current global pandemic with the novel SARS-CoV-2 virus.

SARS-CoV-2 infection and COVID-19

SARS-CoV-2 and SARS-CoV both belong to subgenus *Sarbecovirus*, genus *Betacoronavirus*, and family *Coronaviridae* [35]. The SARS-CoV was reported for the first time over 17 years ago in Guangdong Province, China [36], which was also responsible for causing a new disease called Severe Acute respiratory syndrome (SARS). SARS-CoV infected 8098 individuals and caused 774 deaths in 29 countries [37]. Similarly, A novel coronavirus (SARS-CoV-2) has emerged with effective human-to-human transmission and leading to pneumonia-like outbreak first reported in December 2019 in Wuhan, China [38, 39]. Both viruses can lead to life-threatening respiratory illnesses in humans.

Genomic characterization revealed around 80 percent similarity between SARS-CoV-2 and SARS-CoV [40]. Moreover, protein sequence analysis showed that both viruses share the same seven conserved nonstructural domains suggesting a relationship between these two coronaviruses [40]. The entry of coronavirus in the cell is complex. The first step in the entry process, the virus-cell fusion, requires receptor binding and proteolysis of the receptor-binding domain (RBD). Structural proteins of CoV are classified into four categories of proteins, which include nucleocapsid (*N*), envelope (*E*), membrane (*M*), and spike (*S*) proteins [41]. Ectodomain segment of spike consists of a receptor-binding subunit S1 and membrane-binding subunit S2, which together are needed for virus entry to allow viral genome to enter the cell [42–44]. Membrane fusion and binding of receptors are crucial steps for coronavirus infection. Biochemical [45, 46] and structural studies [47–49] showed that the RBD of SARS-CoV-2 interacts with ACE2 for cellular entry, similar to SARS-CoV [50, 51]. The binding affinity of S protein of virus and ACE2 appeared to be a major determinant for virus replication rate and severity of the disease [42, 46, 52]. However, RBD of SARS-CoV-2 is less exposed and has a higher affinity for its human ACE2 (hACE2) receptor compared to the RBD of SARS-CoV, suggesting efficient immune surveillance evasion and cellular entry, respectively [53]. Additionally, Priming of S protein and proteolytic cleavage of a spike by TMPRSS2, a Serine protease, and Furin, a proprotein convertase are required for the membrane fusion and subsequent viral entry via the endosomal pathway [42, 43]. After entry, the virus multiplies, and mature virions are transported to the cell surface and released via exocytosis (Fig. 1).

COVID-19 and comorbidities

Due to the lack of a well-designed cohort study for COVID-19 in patients with various comorbidities like hypertension, obesity, and diabetes, it is not clear whether these

comorbidities are risk factors for SARS-CoV-2 infections. However, currently available data indicate an association of these comorbidities with severe outcomes in patients with COVID-19 (See Table 1 for studies that show the effect of comorbidities like hypertension, obesity, and DM with SARS-CoV-2 infection).

Hypertension

Prevalence of hypertension in COVID-19 seems higher in patients with high severity, which includes the use of primary composite endpoint (i.e., intensive care unit, use of mechanical ventilation), ARDS, or death. Recently, Guan et al. [11] reported that 23.7% of subjects with hypertension as one of the coexisting illnesses had a more severe course of COVID-19 disease compared with 13.4% subjects, who had a nonsevere disease. Similarly, another study from China [10] showed that nearly 58% of COVID-19 patients who required intensive care had hypertension, whereas only 21.6% of total COVID-19 patients who did not require the use of ICU had hypertension. Two other studies [4, 54] also reported that 48% of COVID-19 patients who died had an underlying condition of hypertension. However, it is important to note that these associations did not account for age in the analysis and may be confounded by the higher incidence of hypertension in older people. As people age, they exhibit severity of disease including a high risk of acute respiratory distress syndrome (ARDS) and a high mortality rate compared to younger people [10, 54–57]. Hypertension may also present with other cardiovascular risk factors such as diabetes, hypertension-mediated heart damage, and other cardiovascular-related complications. These risk factors show an increasing prevalence with age [58]. This indicates that this association is possibly confounded by age and other comorbidities [59].

Control of blood pressure in patients with hypertension has been considered as one of the important concerns to lessen the disease burden regardless of its effect on SARS-CoV-2 infection [60]. Besides, in COVID-19 patients with coexisting hypertension, high blood pressure was independently associated with hospitalization, mortality, and heart failure [61]. The mechanistic relationship between hypertension and COVID-19 may be explained by the use of ACE2 (angiotensin converting enzyme 2) as a receptor for SARS-CoV-2 entry [42, 46]. ACE2 is an important element of the RAS (renin–angiotensin system), which regulates vasodilation and vasoconstriction and thereby plays an essential role in the pathogenesis of hypertension [62]. In severe forms of hypertension, blood angiotensin II levels are high and significantly correlated with diastolic blood pressure [63]. Angiotensin II is an essential mediator of tissue inflammation by

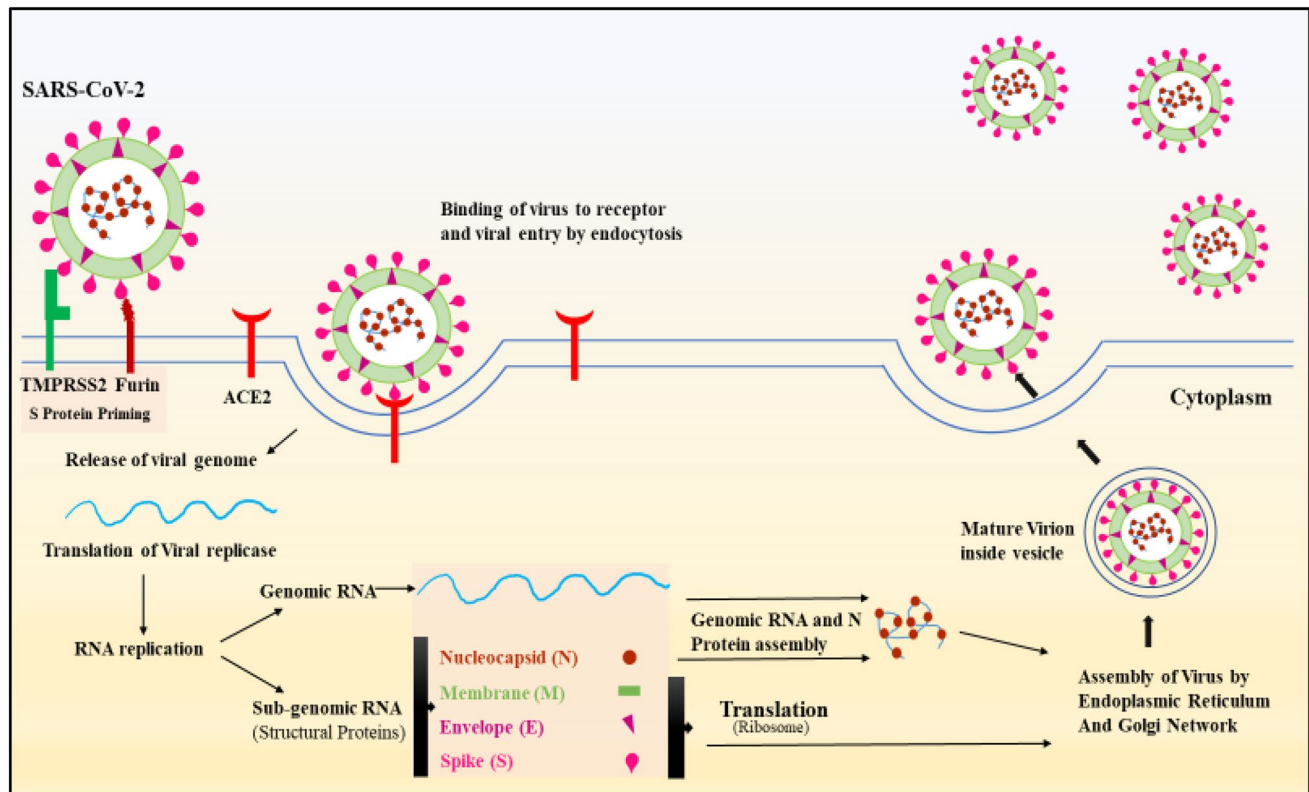


Fig. 1 Schematic diagram of SARS-CoV-2 infection cycle. S protein priming facilitates the binding of the virus with angiotensin converting enzyme—2 (ACE2) and thereby fusion of viral and cellular membrane occurs. This, in turn, releases a viral genome inside the cell. The next step is translation of the viral replicase gene followed by RNA replication to produce genomic RNA and subgenomic RNAs.

Subgenomic RNAs are then translated into proteins, and genomic RNA is packaged with N proteins and along with all other structural proteins (i.e., N, E, S, M). The assembly of the virus occurs in various steps by Endoplasmic Reticulum and Golgi network. Mature virion inside the vesicle then released from the cell

increasing vascular permeability, recruiting inflammatory cells, and oxidative stress [64, 65]. Angiotensin II has been shown to induce lung edemas, impaired lung function, and lung inflammation in pneumonia [66]. Moreover, the SARS spike protein binding to ACE2 showed elevated angiotensin II levels along with severe acid-induced pneumonia. This pathology was rescued by an angiotensin II type 1 receptor antagonist, losartan, suggesting the inflammatory role of angiotensin II [67]. ACE2 is a negative regulator of RAS that inactivates angiotensin II. Coronavirus infection causes downregulation of ACE2 [68], and probably, in the case of SARS-CoV-2, that may cause elevation of angiotensin II and tissue inflammation. Several reports suggested elevated levels of angiotensin II in COVID-19 patients with high disease severity [69, 70]. Altogether, these indicate that hypertensive individuals are more vulnerable to COVID-19 disease severity.

In early phase of COVID-19 pandemic, it was suggested that the use of RAS inhibitors (i.e., ACE inhibitors (ACEIs) and angiotensin II receptor blockers (ARBs)) may be detrimental to COVID-19 patients [71] as some experimental

studies suggested a compensatory increase in ACE2 upon the use of above-mentioned inhibitors [72–74]. A comprehensive review [75] on assessing the risk of RAS inhibitors in COVID-19 suggested inconsistency in the effect of ARB and ACEI on ACE2 expression in animals. However, human studies suggest no significant effect on ACE2 by RAS inhibitors. A study from NYU Langone system reported no significant increase in the propensity of detecting positive for SARS-CoV-2 infection or in the risk of severe infection correlated with various antihypertensive medications including ACEI and, ARB [76]. A retrospective cohort study from Denmark reported that prior use of ACEI or ARBs is neither associated with COVID-19 diagnosis nor with mortality or disease severity among patients with hypertension [77]. Another large population-based study from Italy suggested no association for the use of antihypertensive medications with the risk of developing COVID-19 and related severe outcomes [78]. Hence, although there is an underlying concern for the effect of RAS inhibitors on ACE2 and subsequent increase in susceptibility or severity of COVID-19 infection, there is no justification for the preemptive change

Table 1 List of studies that show the effect of comorbidities like hypertension, diabetes, and obesity with SARS-CoV-2 infection

Study Details [Authors, Country]	Comorbidities			* <i>p</i> values
	Hypertension (HT) n/N (%)	Diabetes (Db) n/N (%)	Obesity (Ob) n/N (%)	
	[Use of ICU or Disease Severity or Survival—n/N (%)]*			
Chen et al. [4] China	93/274 (34%) [Survivor—39/161 (24%) Non-Survivor—54/113 (48%)]	47/274 (17%) [Survivor—24/161 (21%) Non-Survivor—23/113 (14%)]	NR	NR
Huang et al. [5] China	6/41 (15%) [ICU—2/6 (33.3%) Non-ICU—4/6 (0.67%)]	8/41 (20%) [ICU—1/8 (0.13%) Non-ICU—7/8 (0.88%)]	NR	HT— <i>p</i> = 0.93 Db— <i>p</i> = 0.16
Richardson et al. [8] USA	3026 (56.6%) [#]	1808 (33.8%) [#]	1737/4170 (41.7%)	NR
Petrilli et al. [9] USA	2256/5279 (42.7%) [Not Hospitalized—557/2538 (21.9%) Hospitalized—1699/2741 (62%)]	1195/5279 (22.6%) [Not Hospitalized—245/2538 (9.7%) Hospitalized—950/2741 (34.7%)]	1865/5279 (35.3%) [Not Hospital- ized—781/2538 (30.7%) Hospital- ized—1084/2741 (39.5%)]	HT, Db, Ob— <i>p</i> < 0.001 [§]
Wang et al. [10] China	43/138 (31.2%) [ICU—21/36 (58.3%) Non-ICU—22/102 (21.6%)]	14/138 (10.1%) [ICU—8/36 (22.2%) Non-ICU—6/102 (5.9%)]	NR	HT— <i>p</i> < 0.001 Db— <i>p</i> = 0.009
Guan et al. [11] China	165/1099 (15%) [Severe—41/173 (23.7%) Non-severe—124/926 (13.4%)]	81/1099 (7.4%) [Severe—28/173 (16.2%) Non-severe—53/926 (5.7%)]	NR	NR
Grassilli et al. [12] Italy	509/1043 (49%) [Died—195/309 (63%) Discharged—84/212 (40%)]	180/1591 (17%)	NR	HT— <i>p</i> < 0.001
Zhou et al. [54] China	58/191 (30%) [Survivor—32/137 (23%) Non-survivor—26/54 (48%)]	36/191 (19%) [Survivor—19/137 (14%) Non-survivor—17/54 (31%)]	NR	HT— <i>p</i> = 0.0008 Db— <i>p</i> = 0.0051
Wu et al. [55] China	39/201 (19.4%) [ARDS—23/84 (27.4%) No ARDS—16/117 (13.7%)]	22/201 (10.9%) [ARDS—16/84 (19%) No ARDS—6/116 (0.05%)]	NR	HT— <i>p</i> = 0.02 Db— <i>p</i> = 0.002
Zhang et al. [56] China	42/140(30%) [Severe—22/58 (37.9%) Non-severe—20/82 (27.4%)]	17/140 (12.1%) [Severe—8/58 (13.8%) Non-severe—9/82 (11%)]	NR	HT— <i>p</i> = 0.85 Db— <i>p</i> = 0.615
McMicheal et al. [57] USA	74/167 (44.3%)	38/167 (22.8%)	37/167 (22.2%)	NR
Shi et al. [101] China	127/416 (31%) [With cardiac injury—49/82 (59.8%) Without cardiac injury—78/334 (23.4%)]	60/416 (14.4%) [With cardiac injury—20/82 (24.4%) Without cardiac injury—40/334 (12%)]	NR	HT— <i>p</i> < 0.001 Db— <i>p</i> = 0.008
Bean et al. [102] UK	150/205 (51.2%)	62/205 (30.2%)	NR	NR

ICU, intensive care unit, ARDS, acute respiratory distress syndrome, NR, not reported

[#]*N* value is not reported

[§]*p* value for risk of hospitalization

in using these inhibitors in people who are at risk or having COVID-19 infection, unless clinically advised [79]. Many renowned societies including the European society of hypertension, American society of hypertension, and American heart association are in agreement with this position [59,

79, 80]. Though theoretically high expression of ACE2 may suggest high chances of infection, it is also important to note that viral entry requires proteolytic degradation by TMPRSS2 [42]. Hence, increased ACE2 expression may affect the tissues that coexpress both ACE2 and TMPRSS2.

However, it is unknown whether only elevated ACE2 expression affects viral infection. Although all observational studies provide important assurance about the safety of RAS inhibitors during the COVID-19 pandemic, their confirmation in randomized controlled trials will provide a greater degree of assurance to patients and doctors.

Obesity

Several reports are suggesting an association between obesity and the severity of COVID-19 infection and mortality. Clinical data for COVID-19 infection from China found a 86% and 142% higher association for overweight and obesity, respectively, with high severity of infection compared to normal-weight individuals when adjusted for multiple potential confounders [81]. Another study from China showed increased odds of severe SARS-CoV-2 infection in obese patients [82]. A study from USA showed obesity as the second comorbidity (41.7%) after hypertension (56.6%) among hospitalized patients following COVID-19 infection [8]. Similarly, obesity was also the significant independent predictor of hospitalization, after old age, among over 5000 patients with COVID-19 infection in New York City [9]. Furthermore, a small study from France reported data from 124 patients with COVID-19, suggesting an association of the need for invasive mechanical ventilation with a BMI ≥ 35 kg/m², independent of other comorbidities [83].

Certain underlying parameters may mediate this high risk, including various pathophysiological characteristics of obesity like impaired respiratory functions, declined respiratory muscle strength, and low lung capacity [84]. Transcriptomics data from the studies of mice with diet induced obesity indicate ACE2 upregulation in the lungs of obese mice compared to lean mice [85], which may promote CoV-2 infection. Besides, adipose tissue in obese individuals may act as a reservoir for more extensive viral replication due to ACE2 expression. The higher amount of adipose tissue may lead to increased viral shedding and immune activation and subsequent severe complications [86]. Liraglutide, which is used to treat obesity [87], has also been shown to upregulate ACE2 expression in diabetic rats [88]. It remains to be determined whether using liraglutide as treatment during COVID-19 infection may be a concern. Moreover, obesity is also associated with a high risk for complications, development of ARDS, and need for hospitalization with the use of primary composite endpoint [89, 90]. During obesity, angiotensin II and angiotensinogen are elevated [91, 92]. RAS activation and angiotensin II production are involved in obesity-associated inflammation [93, 94]. Higher expression of angiotensin II leads to inflammation and fibrosis in the lung, thereby enhancing lung injury and ARDS [95, 96].

Hence, obesity-associated inflammation and ARDS may even worsen COVID-19 disease severity.

On the contrary, research on how obesity can affect outcomes of critical illness is still controversial and some reports suggest an inverse relationship between obesity and mortality [97]. This is known as the obesity survival paradox. Additionally, mortality rates are less in obese patients with pneumonia [98]. This protection could be due to obesity-mediated low-grade inflammatory processes creating protective environment to limit a detrimental effect of second hit (e.g., infection), more aggressive medication, and some other unidentified factors [98, 99]. However, obesity is found to be associated with increased disease severity in COVID-19 patients. It is possible that limited available resources during this pandemic may limit early admission of these patients to intensive care for monitoring purposes [100]. Hence, healthcare personnel should consider taking preventive measure and extra care to avoid disease severity in obese individuals.

Diabetes

Diabetes, hypertension, obesity, and age are associated with a higher risk of morbidity and mortality among COVID-19 patients [5, 8–10, 38, 54–57, 83, 101, 102]. Diabetes was also considered as a major predictor of the severity of disease during the MERS-CoV [18]. Patients with SARS also showed a strong association with DM and blood glucose levels [103]. Likewise, blood glucose is represented as a predictor of severe outcomes and death in patients with COVID-19 [104]. Recently, a retrospective study from the USA with 451 patients of COVID-19 with diabetes and/or hyperglycemia reported that uncontrolled hyperglycemia is associated with longer hospital stay and higher mortality compared to people without diabetes or uncontrolled hyperglycemia [105]. Similarly, Zhu and colleagues [106] suggested that diabetes increased the requirement of medical interventions and mortality risk for COVID-19 infected patients. A meta-analysis reported higher mortality risk and higher risk of ICU admission in COVID-19-diagnosed patients with diabetes [107]. On the contrary, well-controlled diabetes is linked with improvement in outcomes for patients with COVID-19 [106]. Previously, hyperglycemia was considered to be the link for the association between diabetes and viral infections, which influences viral growth and inflammation, thereby exacerbating mortality and morbidity in patients [108]. Severe infections may cause difficulty in managing blood glucose levels. However, the causal relationship between control of glucose level and disease severity is yet to be established.

There could be different mechanisms, which may favor COVID-19 infection in a patient with diabetes. The virus enters the cell via the ACE2 receptor, which is an important element of the system involved in blood pressure management and inflammation [42]. Various diabetic animal models have demonstrated an increased expression level of ACE2 in the pancreas, kidney, liver, and serum [109, 110]. Recently, a phenome-based mendelian randomization study on diabetes indicated that enhanced ACE2 expression in lungs may be causally linked with diseases or traits [111]. Insulin administration in nonobese diabetic (NOD) mice downregulates ACE2 expression in lungs [109], whereas pharmacological drugs for glycemic control like glucagon-like-peptide-1 (GLP-1) agonist (e.g., liraglutide), and Pioglitazone, upregulate ACE2 expression in lungs of STZ-induced diabetic rats and in insulin-sensitive tissues of high-fat diet-fed rats, respectively [88, 112]. In addition, elevated circulating furin, which is required for proteolytic cleavage of viral spike protein for cellular entry of COVID-19 [113], has been observed in patients with diabetes [114]. Delay in SARS-CoV-2 clearance was also reported in patients with diabetes [115]. A recent observational study with COVID-19 patients with diabetes suggests insulin infusion as an effective method to improve glycemic control and reduce disease severity [116]. Moreover, some metabolic complications of preexisting diabetes including ketoacidosis have been observed with COVID-19 patients [117–119]. Collectively, these studies support the hypothesis that patients with diabetes are at a higher risk of COVID-19 infection and severe outcomes.

Many leading diabetes researchers suggest a bidirectional relationship between COVID-19 and diabetes [120]. The ACE2 is expressed in key metabolic organs (i.e., muscle, liver, pancreas, and adipose tissue), which plays an important role in glucose homeostasis [121]. In vitro and humanized mice study suggests that human liver and pancreatic islets are highly permissive for SARS-CoV2 infection [122]. SARS-CoV infection damaged various metabolic organs including the endocrine part of pancreas and leads to acute insulin-dependent diabetes [123]. Similarly, few reports also indicate pancreatic injury in patients with COVID-19 [124–126]. However, careful investigation is needed to determine impairment of endocrine function of pancreas in COVID patients. Moreover, in COVID-19, elevated levels of inflammatory cytokines known as cytokine storm have been linked in multiorgan failure with severe disease condition [127]. Hence, it is possible that COVID-19 may cause impairment of insulin secretory function by damaging pancreas or by inflammation-induced acute insulin resistance to trigger pleiotropic effects that impairs glucose metabolism and exacerbate preexisting diabetes or may even cause transient diabetes.

COVID-19 and other associated complications related to triumvirate

Several reports indicated the threat of COVID-19 for patients with cardiovascular disease (CVD) and renal complications [128, 129] that are also associated with this triumvirate. Obesity is closely associated with CVD with a high propensity of mortality and morbidity [130]. Similarly, high blood pressure is one of the most critical risk factors for developing cardiovascular complications [131, 132]. Data of clinical studies indicate that outcomes of COVID-19 in CVD patients are linked with disease severity, and a high prevalence of preexisting CVD has been associated with mortality in COVID-19 patients [7, 133–138]. Also, several reports showed the development of cardiovascular disorders in COVID-19 patients [139–141]. These data suggest a bidirectional relationship between COVID-19 and cardiovascular diseases. There are two possible explanations for this relationship. First, induction of “cytokine storm” and elevated systemic inflammation in COVID-19 may accelerate the cardiovascular system’s complications. A significant linear correlation between high troponin (a marker for myocardium injury) levels and C-reactive protein (an inflammatory marker) has been observed in COVID-19 patients [141], suggesting an association between inflammation and injury of the myocardium. Second, ACE2, a receptor for SARS-CoV-2, is a part of RAS that is involved in the maintenance of blood pressure. In COVID-19, elevated levels of angiotensin II have been observed [69, 70]. High levels of angiotensin II can elevate blood pressure and damage the myocardium by inducing acute ischemia, ventricular hypertrophy, oxidative stress, and proinflammatory cascade [142, 143] that may worsen cardiovascular complications in COVID-19 patients.

Patients with diabetes are vulnerable to infection due to impaired immune function [144]. It is estimated that 30–40% of diabetes patients develop kidney infections. Also, diabetes kidney disease is associated with disease severity and mortality in diabetic patients [145]. Reports suggest that COVID-19 patients have a high incidence of kidney dysfunction (high levels of blood urea nitrogen (BUN), serum creatine, D-dimer), and acute kidney injury (AKI) [128, 146–148]. Also, AKI is associated with severe outcome and mortality in COVID-19 patients [147–150]. Kidney expresses ACE2 [151], and SARS-CoV-2 was detected in patients’ urine and kidney samples [128]. Moreover, viral structure and protein and RNA of the SARS-CoV-2 are found in kidney of COVID-19 patients [152, 153]. This suggests that SARS-CoV-2 can directly damage the kidney. However, mechanism of kidney disease in COVID-19 patient requires further investigation.

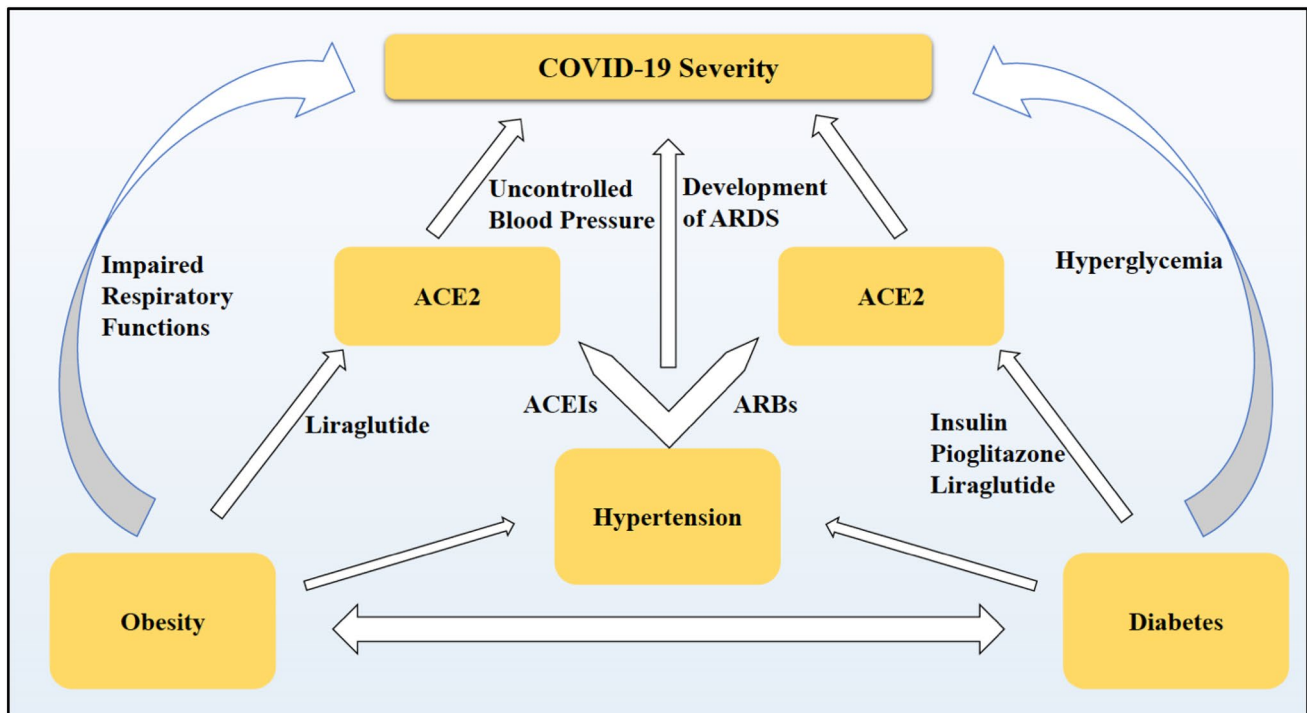


Fig. 2 Proposed mechanism of disease severity in COVID-19 patients with comorbidities like hypertension, obesity, and diabetes. Hypertension, obesity, and diabetes are associated with impaired respiratory functions, uncontrolled blood pressure and development of ARDS, and hyperglycemia, respectively, which in turn leads to increased disease burden in COVID-19 patients. Moreover, medications that are

used in these comorbidities (liraglutide for obesity, ACEIs, and ARBs for hypertension and insulin, pioglitazone, and liraglutide for diabetes) have been reported to affect ACE2 expression and hence may affect disease severity.*ARDS, acute respiratory distress syndrome, ACEIs, angiotensin-converting enzyme inhibitors, ARBs, angiotensin II receptor blockers

Conclusion and future directions

The current COVID-19 pandemic imposes an alarming condition in the world in an unprecedented way. Chronic preexisting metabolic conditions during viral infection increase disease severity and mortality. The altered metabolic environment and impaired immune system reinforced by hypertension, obesity, and diabetes may exacerbate the severity of the disease among infected patients. However, the current understanding of the connection between the triumvirate—hypertension, obesity, and diabetes—and related cardiovascular and renal complications with COVID-19 is evolving. Various studies have highlighted the role of ACE2 receptor expression and systemic inflammation in promoting SARS-CoV-2 infection and disease severity. Although several pharmacological interventions for these chronic metabolic disorders also seem to affect ACE2 expression (Fig. 2), it is not clear whether they have any role on the disease severity. Therefore, while research and rapid efforts are being made to develop treatments or vaccines for COVID-19, it is important to identify and investigate alternative clinical treatments that effectively manage the preexisting metabolic perturbations during the COVID-19 pandemic and reduce disease severity.

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Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

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