

1 **Perspective: COVID-19 Pandemic, Corona Viruses, and Diabetes Mellitus**

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32 **Abstract**

33 The pandemic of COVID-19, a disease caused by a novel coronavirus (CoV), SARS-CoV-2, is causing
34 substantial morbidity and mortality. Older age and presence of diabetes mellitus, hypertension, and
35 obesity significantly increases the risk for hospitalization and death in COVID-19 patients. In this
36 *Perspective*, informed by the studies on severe acute respiratory syndrome, SARS-CoV, and Middle East
37 respiratory syndrome, MERS-CoV, and the current literature on SARS-CoV-2, we discuss potential
38 mechanisms by which diabetes modulates the host-viral interactions and host-immune responses. We
39 hope to highlight gaps in knowledge that require further studies pertinent to COVID-19 in patients with
40 diabetes.

41 **Introduction**

42 Coronaviruses (CoV) are enveloped viruses with a single-stranded, positive-sense RNA genome
43 known to cause respiratory infections in humans (7, 38). In general, in most immunocompetent
44 individuals, human CoV infection leads to mild upper respiratory infection. However, two highly
45 pathogenic CoV have resulted in outbreaks of severe acute respiratory syndrome (SARS) in 2003 in
46 Guangdong province, China and Middle East respiratory syndrome (MERS) in Middle Eastern countries
47 a decade later. SARS-CoV and MERS-CoV were identified to cause SARS and MERS respectively (11,
48 51, 55). In December 2019, a novel coronavirus SARS-CoV-2 was identified as the pathogen causing
49 coronavirus disease of 2019 (COVID-19) in Wuhan, China (11, 51, 55). On March 11th, 2020, COVID-19
50 was declared a pandemic by the World Health Organization. As of March 27th, 2020, there have been a
51 total of 103,942 confirmed cases with 1689 deaths in the United States (1). Globally, 27,324 deaths have
52 been reported among 595,800 confirmed cases (1).

53 Individuals with diabetes mellitus (DM), hypertension, and severe obesity ($BMI \geq 40 \text{ kg/m}^2$) are
54 more likely to be infected and are at a higher risk for complications and death from COVID-19 (16, 30,
55 32, 48, 50, 52, 56). Interestingly, there was similarly an increased risk for SARS and MERS in individuals
56 with DM. In the United States, 34.2 million or 10.5% of the total population have DM (32). Among those
57 aged 65y or older, a population at higher risk for death from COVID-19, 26.8% have DM (32).
58 Hypertension and severe obesity are present in 68.4% and 15.5% of individuals diagnosed with DM,
59 respectively. Over a period of months, a substantial portion of the US population will be infected by
60 SARS-CoV-2 (12). While a significant number will remain asymptomatic and be able to transmit the
61 virus, the estimated proportion of symptomatic individuals requiring hospitalization increases with age
62 (12). In individuals older than 60y, that proportion ranges from 17-27%. Furthermore, in this older group,
63 the percentage of hospitalized patients requiring ICU care is 27-71% with an infection fatality rate (IFR)
64 ranging from 2.2-9.3% (12). While these estimates are preliminary and likely to change, considering the

65 prevalence of DM, hypertension, and severe obesity in the US and the substantial increased risk for
66 COVID-19 and its complications in patients with these conditions, it is likely the pandemic has the
67 potential to cause significant mortality and morbidity. Specialists and health care providers will be
68 providing clinical care to many patients with COVID-19 in both inpatient, outpatient, and telehealth
69 settings. Increased awareness of the clinical features, pathophysiology, and potential mechanisms that
70 increase the risk is needed to provide better care and spur new investigations, both basic and clinical, to
71 better understand COVID-19 in patients with diabetes.

72 ***Clinical Features and Natural Course of COVID-19***

73 The median age of SARS-CoV-2-infected patients is in the range of 47-56 years, males comprise
74 more than half of the cases, the average incubation period is 5.2 days, and 98% of those who develop
75 symptoms will do so within 11.5 days (5, 16, 19, 22, 42). The clinical manifestations of COVID-19 vary
76 and include the asymptomatic carrier status, acute respiratory disease (ARD), and pneumonia (16, 42).
77 The prevalence of asymptomatic cases is significant (20-86% of all infections) and are defined as
78 individuals with positive viral nucleic acid tests, but without any COVID-19 symptoms (3, 4, 23, 29, 57).
79 Transmission rates and respiratory viral load in asymptomatic carriers are similar to symptomatic patients
80 (23, 57), partially explaining the rapid spread of SARS-CoV-2. In addition to a laboratory-confirmed
81 COVID-19 diagnosis, patients with ARD manifest with fever, fatigue, respiratory (cough, dyspnea) or
82 gastrointestinal (nausea, diarrhea, vomiting) symptoms, and no significant abnormalities on chest imaging
83 (16, 42). Patients with pneumonia have respiratory symptoms and positive findings in chest imaging.
84 Severe pneumonia can present as acute respiratory distress syndrome (ARDS) leading to severe hypoxia,
85 respiratory failure, multiorgan failure, shock, and death (16, 37, 42).

86 ***The Pathophysiology of SARS-CoV-2 Infection***

87 The genetic sequence of SARS-CoV-2 showed more than 80% shared identity to SARS-CoV and
88 50% to the MERS-CoV, and both SARS-CoV and MERS-CoV originate in bats and infect humans and
89 wild animals (2, 7, 26, 38). Cellular CoV entry is a complex process that involves receptor-binding and
90 proteolysis leading to virus-cell fusion. CoV is made up of four structural proteins: spike (S), membrane
91 (M), nucleocapsid (N), and envelope (E) proteins. The S protein mediates receptor binding on the host
92 cell membrane through the receptor-binding domain (RBD) in the S1 domain and membrane fusion
93 through the S2 subunit (18, 40). Angiotensin-converting enzyme 2 (ACE2) is the cellular receptor for
94 SARS-CoV and SARS-CoV-2, in contrast to MERS-CoV, which utilizes dipeptidyl peptidase 4 (DPP4)
95 as its cellular receptor (24, 33) (Figure 1). This interaction thus determines host tropism and ultimately
96 clearance of the virus. ACE2 is expressed in the upper respiratory system, type I and II alveolar epithelial
97 cells in the lungs, the heart, endothelial cells, kidney tubular epithelium, enterocytes, and the pancreas
98 (10, 24, 25, 54). After binding to ACE2, proximal serine proteases such as TMPRSS2 are involved in S

99 protein priming and cleavage of the spike (Figure 1). Proteases such as Furin subsequently release the
100 spike fusion peptide, and the cellular virus enters through an endosomal pathway (18, 40). The low pH
101 and presence of proteases such as cathepsin-L characteristic of the endosomal microenvironment favor
102 the delivery of SARS-CoV-2 genome into the cytosol where further viral replication leads to the
103 formation of mature virions and subsequent spread.

104 Infected cells undergo apoptosis or necrosis and trigger inflammatory responses marked by the
105 activation of pro-inflammatory cytokines or chemokines, which leads to the recruitment of inflammatory
106 cells. CD4⁺ T helper (Th1) cells regulate antigen presentation and immunity against intracellular
107 pathogens such as CoV through interferon gamma (IFN- γ) production. Th17 cells induce the recruitment
108 of neutrophils and macrophages by producing interleukin-17 (IL-17), IL-21, and IL-22 (9). SARS-CoV-2
109 infects circulating immune cells and increases apoptosis of lymphocytes (CD3, CD4, and CD8 T cells)
110 leading to lymphocytopenia. Indeed, the degree of lymphocytopenia is associated with the severity of
111 SARS-CoV-2 infection (16, 45, 50, 52). Lower T cell function relieves the inhibition on innate immune
112 system leading to secretion of high amounts of inflammatory cytokines in what is known as a “cytokine
113 storm” (31). In fact, circulating levels of cytokines/chemokines (IL-6, tumor necrosis factor- α [TNF]) and
114 chemokines (CXC-chemokine ligand 10 [CXCL10] and CC-chemokine ligand 2 [CCL2]) involved in the
115 cytokine storm syndrome are elevated and may play a role in SARS-CoV-2 driven hyperinflammation
116 leading to multiorgan failure (15, 28, 41).

117 ***Potential Mechanisms that increase the risk of COVID-19 in Diabetes***

118 It is now well recognized that older age and the presence of DM, hypertension, and severe obesity
119 ($\text{BMI} \geq 40 \text{ kg/m}^2$) increases morbidity and mortality in patients with COVID-19 (16, 30, 32, 48, 50, 52,
120 56). Considering the high prevalence of cardiovascular disease (CVD), obesity, and hypertension in
121 patients with DM, it is unknown whether DM independently contributes to this increased risk. However,
122 plasma glucose levels and DM are independent predictors for mortality and morbidity in patients with
123 SARS (49). Potential mechanisms that may increase the susceptibility for COVID-19 in patients with DM
124 include: a) higher affinity cellular binding and efficient virus entry, b) decreased viral clearance, c)
125 diminished T cell function, d) increased susceptibility to hyperinflammation and cytokine storm
126 syndrome, and e) presence of CVD (Figure 2).

127 Augmented ACE2 expression in alveolar AT2 cells, myocardium, kidney, and pancreas may
128 favor increased cellular binding of SARS-CoV-2 (25, 27, 58). Increased expression of ACE2 has been
129 demonstrated in the lung, kidney, heart, and pancreas in rodent models of DM (35, 46). Insulin
130 administration attenuates ACE2 expression (35, 46), while hypoglycemic agents such as glucagon-like
131 peptide – 1 (GLP-1) agonists (liraglutide) and thiazolidinediones (TZDs; pioglitazone), antihypertensives
132 such as ACE inhibitors, and statins upregulate ACE2 (14, 36, 39, 44, 53). Until recently, whether DM

133 was causally linked to ACE2 expression levels in the lung in humans was unknown. Using a phenome-
134 wide Mendelian randomization study, Rao et al. explored diseases or traits that may be causally linked to
135 increased ACE2 expression in the lung (34). Interestingly, they found that DM was causally associated
136 with increased lung ACE2 expression (34). Circulating levels of furin, a cellular protease involved in
137 facilitating viral entry by cleaving the S1 and S2 domain of the spike protein, are elevated in patients with
138 DM (13). These studies support the hypothesis that patients with DM are susceptible to SARS-CoV-2
139 infection. Indeed, a recent study reported that clearance of SARS-CoV-2 was delayed in patients with
140 DM, a finding that needs to be confirmed in larger studies (6) (Figure 2).

141 ACE catalyzes the conversion of the prohormone, angiotensin I to the octapeptide, angiotensin II
142 (AngII), whereas ACE2 converts AngII to angiotensin₁₋₇. AngII, through the activation of Ang II type 1a
143 receptors induces vasoconstriction and proliferation, whereas angiotensin₁₋₇ stimulates vasodilatation and
144 suppresses cell growth (Figure 1). Increased ratio of pulmonary ACE/ACE2 activity as observed in
145 patients with ARDS (43), favors AngII generation. Once bound to ACE2, SARS-CoV downregulates
146 cellular expression of ACE2, and the unopposed action of AngII contributes to acute lung injury (20).
147 Binding to ACE2 alone does not lead to severe lung injury as is observed with other CoVs (NL63) (7,
148 38). Whether SARS-CoV-2 causes down-regulation of pulmonary ACE2 is unknown. Nevertheless, there
149 exists a potential for salutary, if not therapeutic effects, of Ang II receptor blockers, ACE inhibitors,
150 TZDs, GLP-1 agonists, and statins in the setting of low ACE2 expression. Lacking further evidence of
151 risk or benefit, the American College of Cardiology, the American Heart Association, and the American
152 Society of Hypertension have recommended that patients should continue treatment with their usual
153 antihypertensive therapy (8).

154 DM inhibits neutrophil chemotaxis, phagocytosis, and intracellular killing of microbes.
155 Impairments in adaptive immunity characterized by an initial delay in the activation of Th1 cell-mediated
156 immunity and a late hyper-inflammatory response is often observed in diabetics (17). In an elegant study,
157 Kulcsar et al. examined the effects of DM in a humanized mouse model of MERS-CoV infection on a
158 high-fat diet (21). Following MERS-CoV infection, the disease was more severe and prolonged in
159 diabetic male mice and was characterized by alterations in CD4⁺ T cell counts and abnormal cytokine
160 responses (such as elevated IL17a). Consistent with this finding, in patients with COVID-19, peripheral
161 counts of CD4⁺ and CD8⁺ T cells are low, but with a higher proportion of highly proinflammatory Th17
162 CD4⁺ T cells, as well as elevated cytokine levels (16, 45, 47, 50, 52). Thus, it is likely that patients with
163 DM may have blunted anti-viral IFN responses, and the delayed activation of Th1/Th17 may contribute to
164 accentuated inflammatory responses (Figure 2).

165 **Conclusion**

166 There is a paucity of data in the US regarding comorbidities and COVID-19 outcomes and
167 mechanisms that modulate viral pathogenesis. Certain racial groups such as African Americans,
168 Hispanics, Asians, and Native Americans are highly prone to develop DM and disparities in health care
169 make these groups more vulnerable. Identification of clinical and biochemical parameters using multi-
170 omics approaches that predict severity of the COVID-19 in DM using large data sets is urgently needed.
171 Studies in humanized ACE2 (hACE2) mice and non-human primates aimed at understanding how
172 hyperglycemia, hyperinsulinemia, and hypoglycemic agents affect pathogenesis of COVID-19 and how
173 DM affects the efficacy of vaccines and anti-viral investigational agents currently in trials are warranted.
174 Finally, we need to develop novel ways to deliver care to our DM patients using telehealth, remote patient
175 monitoring, and wearable technologies. As the global pandemic unfolds and rapidly spreads across the
176 US, social isolation measures will enable the transition, but there is an urgent need for basic and clinical
177 investigations to address the many important and unanswered questions.

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363 **Author Contributions**

364 RM and GS drafted and revised the manuscript

365 **Disclosure statement**

366 The authors have nothing to disclose

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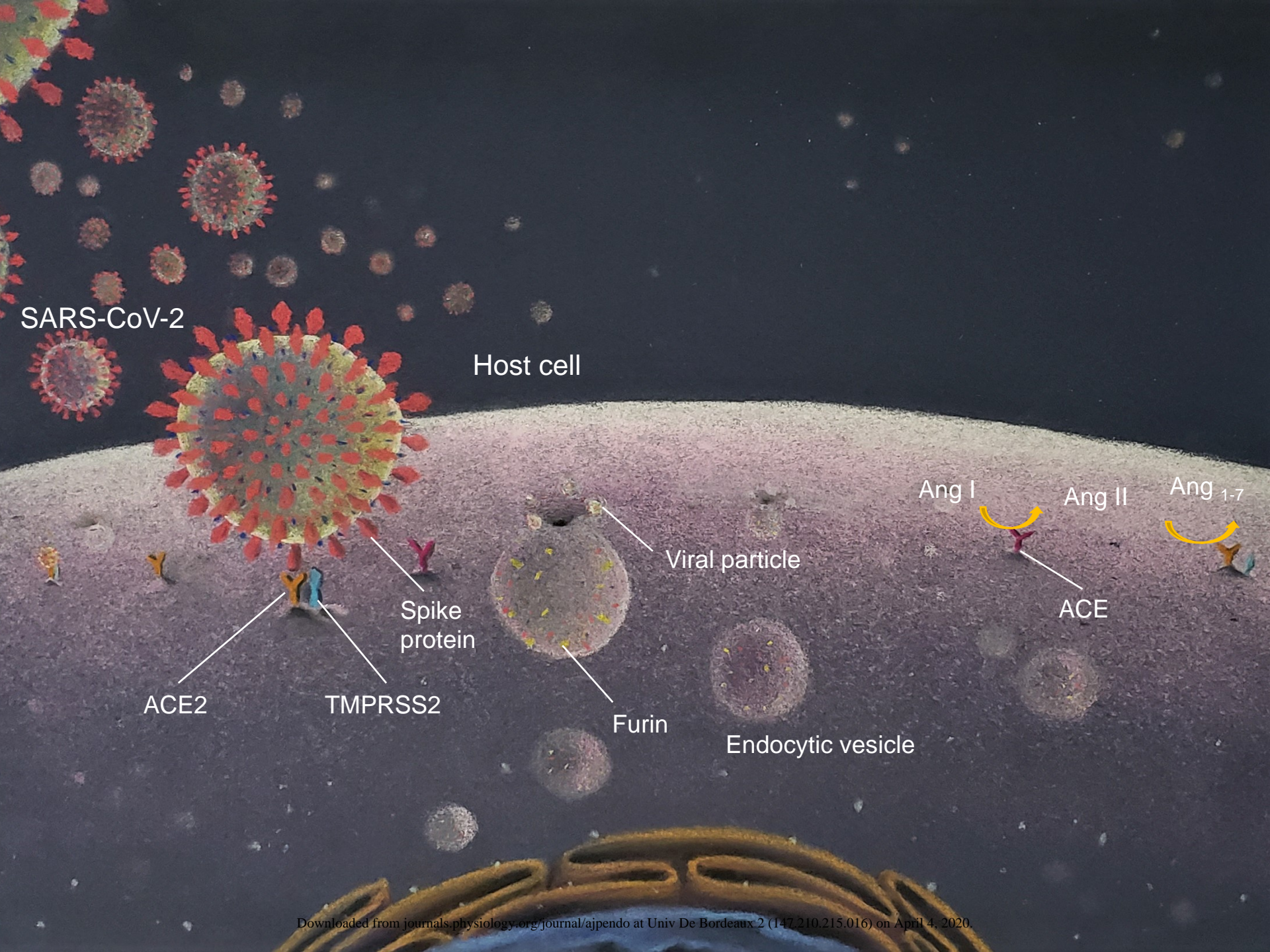
Figure Legends

372 **Figure 1. Cellular entry of SARS-CoV-2.** The initial step in cellular entry of the virus is the binding of
373 SARS-CoV-2 spike protein to cell surface angiotensin converting enzyme 2 (ACE2). Cellular proteases
374 such as TMPRSS2 and furin are involved in priming of the S protein which involves cleavage at the
375 S1/S2 domains. This allows the fusion of the virus to the cell surface. Virions are taken up into
376 endosomes, where SARS-CoV-2-S is cleaved and possibly activated by the pH-dependent cysteine
377 protease cathepsin L. Once inside the cell, SARS-CoV-2 uses the endogenous cellular machinery to
378 replicate itself. ACE catalyzes the conversion of angiotensin I to the octapeptide, angiotensin II (AngII),
379 whereas ACE2 converts AngII to angiotensin 1–7. AngII through the activation of Ang II type 1a
380 receptors induces vasoconstriction and proliferation, whereas angiotensin 1–7 stimulates vasodilatation
381 and suppresses cell growth.

Figure 2. Putative mechanisms contributing to increased susceptibility for COVID-19 in patients with diabetes mellitus (DM). Following aerosolized uptake of SARS-CoV-2, invasion of the respiratory epithelium and other target cells by SARS-CoV-2 involves binding to cell surface ACE2. Increased expression of ACE2 may favor more efficient cell binding and entry into cells. Early recruitment and function of neutrophils and macrophages are impaired in DM. Delay in the initiation of adaptive immunity and dysregulation of the cytokine response in DM may lead to the initiation of cytokine storm.

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383



SARS-CoV-2

Host cell

ACE2

TMPRSS2

Spike protein

Furin

Viral particle

Endocytic vesicle

ACE

Ang I

Ang II

Ang 1-7

