



Relationship between Diabetes and Susceptibility to COVID-19

Zainab Mohammed Jassim¹, Noor A. Neama², Thulfikar H. al-kefaei³

1AL-QASIM Green University

2College of Pharmacy, University of Alkafeel, Najaf, Iraq

3 Department of Medical Laboratory Techniques, Faculty of Medical and Health Techniques, University of Alkafeel, Iraq

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ABSTRACT:

Coronavirus disease 2019 (COVID-19), triggered by the severe acute respiratory syndrome-coronavirus 2 (SARS-CoV-2), may lead to extrapulmonary manifestations like diabetes mellitus (DM) and hyperglycemia, both predicting a poor prognosis and an increased risk of death. SARS-CoV-2 infects the pancreas through angiotensin-converting enzyme 2 (ACE2), where it is highly expressed compared to other organs, leading to pancreatic damage with subsequent impairment of insulin secretion and development of hyperglycemia even in non-DM patients. SARS-CoV-2 infection impairs glucose homeostasis and metabolism in DM and non-DM patients due to cytokine storm (CS) development, downregulation of ACE2, and direct injury of pancreatic β -cells. Therefore, the potent anti-inflammatory effect of diabetic pharmacotherapies such as metformin, pioglitazone, sodium-glucose co-transporter-2 inhibitors (SGLT2Is), and dipeptidyl peptidase-4 (DPP4) inhibitors may mitigate COVID-19 severity. In addition, some antidiabetic agents and also insulin may reduce SARS-CoV-2 infectivity and severity through the modulation of the ACE2 receptor expression. The findings presented here illustrate that insulin therapy might seem as more appropriate than other anti-DM pharmacotherapies in the management of COVID-19 patients with DM due to low risk of uncontrolled hyperglycemia and diabetic ketoacidosis (DKA). From these findings, we could not give the final conclusion about the efficacy of diabetic pharmacotherapy in COVID-19; thus, clinical trial and prospective studies are warranted to confirm this finding and concern.

Keywords: Diabetes mellitus, SARS-CoV-2, Hyperglycemia

I. INTRODUCTION:

Coronaviruses are enveloped, positive single-stranded RNA viruses widely distributed in humans and animals worldwide (Huang et al., 2020,

Isam et al., 2021). Although most human coronavirus infections are mild, major outbreaks of two betacoronaviruses, severe acute respiratory syndrome coronavirus (SARS-CoV) in 2002–2003 and Middle East respiratory syndrome coronavirus (MERS-CoV) in 2012, have caused deadly pneumonia, with mortality rates of 10% for SARS-CoV and 36% for MERS-CoV. In December 2019, clusters of pneumonia cases of unknown etiology emerged in Wuhan, Hubei Province, China. Deep sequencing analysis from lower respiratory tract samples indicated a novel coronavirus as the causative agent, which was named Severe Acute Respiratory Syndrome-Coronavirus-2 (SARS-CoV-2), and the disease it causes called COVID-19 (WHO, 2020). Although SARS-CoV-2 has shown phylogenetic and clinical similarities with SARS-CoV, the novel coronavirus appears to have a higher transmissibility and lower case fatality rates. On 30 January 2020, the World Health Organization (WHO) declared the COVID-19 outbreak a Public Health Emergency of International Concern, and on March 11, the epidemic was upgraded to pandemic (WHO, 2020). As of today (02.04.2020), 827,419 confirmed cases are officially reported in more than 200 countries or territories with 40,777 deaths. We conducted a scoping review to provide a brief summary of the general characteristics of COVID-19, as well as a more detailed description and critical assessment of the association between this new infectious disease and diabetes. We hope this review can provide meaningful information for future research and ultimately contribute to better clinical management of patients with COVID-19 and diabetes.

This new CoV infection called as COVID-19 originated in Wuhan, Hubei Province, China in December 2019 (Zhu et al., 2020). The causative agent for this respiratory illness is severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2). These epidemics have a high infection transmission rate and in general, the direct cause of death is severe atypical pneumonia (Yin and Wunderink, 2018). The disease spread rapidly from



the initial epicenter, Wuhan to rest of the world and has become a pandemic. The main risk factors include pneumonia, acute kidney failure, acute heart failure. People with underlying health conditions such as cardiovascular disease, diabetes, chronic respiratory disease and the elderly above 60 years are most susceptible to COVID-19. Hospitals announced a cluster of cases of unknown cause pneumonia in Wuhan, Hubei, China on 31 December 2019, attracting great national and worldwide attention. Coronavirus (CoV) is a wide family of single-stranded, positive-sense RNA viruses belonging to the Nidovirales order. The order includes families of the Roniviridae, Arteriviridae, and Coronaviridae. The family Coronaviridae is subdivided into subfamilies Torovirinae and Coronavirinae. Coronavirinae is further subclassified into alpha, beta, gamma and delta COVs (Fehr and Perlman, 2015). Phylogenetic clustering accounts for certain virus subtypes being named. The viral RNA genome varies in length from 26 to 32 kilobases. They can be separated from various species of animals. Those include birds, cattle and mammals such as camels, bats and masks Civets of leaves, rats, pigs, cats. The widespread COV distribution and infectivity make it a major pathogen. The moderate clinical signs are associated with human pathogenic subtypes of CoV. Yet extreme coronavirus-related acute respiratory syndrome (SARS-CoV) and the Middle East The two notable exceptions are respiratory coronavirus syndrome (MERS-CoV). In Saudi Arabia MERS-CoV was first observed in 2012. It has been responsible for 2,494 confirmed cases which have resulted in 858 deaths. In 2002, a Beta-COV subtype spread rapidly in Guangdong, China. In 37 countries this epidemic resulted in 8,000 infections and 774 deaths (Lau et al., 2020). The COVID-19 pandemic spurred a crisis that is unprecedented in modern times (Zhu et al., 2019). The disease course varies substantially among individuals, from mild or even sub-clinical infection to severe disease. Indeed, more than 1 million COVID-19-related deaths have been reported globally. There is interest in potential risk factors that affect susceptibility to infection and disease progression. Multiple medical (e.g. diabetes, hypertension) and sociodemographic (e.g. sex, age and race/ethnicity) risk factors for severe outcomes were already established (Wiersinga et al., 2019).

Association between COVID-19 and diabetes

Diabetes is one of the leading causes of morbidity and mortality throughout the world. The condition is associated with several macrovascular and microvascular complications, that ultimately

impact the overall patient's survival (Williamset al., 2020). A relationship between diabetes and infection has long been clinically recognized. Infections, particularly influenza and pneumonia, are often common and more serious in older people with type 2 diabetes mellitus (T2DM). Nevertheless, the evidence remains controversial regarding whether diabetes itself indeed increases susceptibility and impacts outcomes from infections, or the cardiovascular and renal comorbidities that are frequently associated with diabetes are the main factors involved. Diabetes and uncontrolled glycaemia were reported as significant predictors of severity and deaths in patients infected with different viruses, including the 2009 pandemic influenza A (H1N1), SARS-CoV and MERS-CoV (Banik Gouri Raniet al., 2016). In the current SARS-CoV-2 pandemic, some studies did not find a clear association between diabetes and severe disease. However, other reports from China and Italy showed that older patients with chronic diseases, including diabetes, were at higher risk for severe COVID-19 and mortality. Scarce data exist regarding glucose metabolism and development of acute complications of diabetes (e.g., ketoacidosis) in patients with COVID-19. Infection of SARS-CoV-2 in those with diabetes possibly triggers higher stress conditions, with greater release of hyperglycemic hormones, e.g., glucocorticoids and catecholamines, leading to increased blood glucose levels and abnormal glucose variability (Wang Aihong et al., 2020). On the other hand, a retrospective study from Wuhan reported that around 10% of the patients with T2DM and COVID-19 suffered at least one episode of hypoglycemia (<3.9 mmol/L). Hypoglycemia has been shown to mobilize pro-inflammatory monocytes and increase platelet reactivity, contributing to a higher cardiovascular mortality in patients with diabetes (Iqbal et al., 2019). Yet it remains largely unknown how exactly the inflammatory and immune response occurs in these patients, as well as whether hyper- or hypoglycemia may alter the SARS-CoV-2 virulence, or the virus itself interferes with insulin secretion or glycemic control. Furthermore, the impact of usual diabetes drug treatment on COVID-19 outcomes, as well as therapeutic approaches for COVID-19 on glucose regulation remains unspecified. Diabetes is a chronic inflammatory condition characterized by multiple metabolic and vascular abnormalities that can affect our response to pathogens. Hyperglycemia and insulin resistance promote increased synthesis of glycosylation end products (AGEs) and pro-inflammatory cytokines, oxidative stress, in addition



to stimulating the production of adhesion molecules that mediate tissue inflammation. This inflammatory process may compose the underlying mechanism that leads to a higher propensity to infections, with worse outcomes thereof in patients with diabetes (Knapp Sylvia, 2013). Several defects in immunity have been associated with hyperglycemia, even though the clinical relevance of some in vitro disturbances are still not fully understood. Poorly controlled diabetes has been linked to inhibited lymphocyte proliferative response to different kinds of stimuli, as well as impaired monocyte/macrophage and neutrophil functions (Knapp Sylvia, 2013). Abnormal delayed type hypersensitivity reaction and complement activation dysfunction have also been described in patients with diabetes. In vitro studies have shown that pulmonary epithelial cells exposure to high glucose concentrations significantly increases influenza virus infection and replication, indicating that hyperglycemia may enhance viral replication in vivo. In animal models, structural lung changes have been related to diabetes, such as augmented vasculature permeability and collapsed alveolar epithelium. On the other hand, patients with diabetes generally present a significant reduction in forced vital capacity (FVC) and forced expiratory volume in one second (FEV1), which is associated with raised plasma glucose levels (Popov and Simionescu, 1997).

Covid-19 and Hyperglycemia

It has been stated that COVID-19 is associated with hyperglycemia, actually considered a direct predictor of the poor prognosis of the disease and to an increased risk of death. Briefly, the binding site and entry point of SARS-CoV-2 is the ACE2 receptor, which is highly expressed in the lung, liver, brain, placenta, and pancreas. SARS-CoV-2 infects the pancreas through ACE2, being highly expressed there when compared to other organs, leading to pancreatic damage with subsequent impairment of insulin secretion and development of hyperglycemia even in non-DM patients. Similarly, SARS-CoV-2-induced pancreatic injury may worsen a preexistent DM (Wanget al., 2020). Previous data have shown that SARS-CoV, which is closely related to SARS-CoV-2, triggers transient hyperglycemia and impairment of pancreatic b-cell function during epidemic derived pneumonia. Moreover, the COVID-19-induced inflammation and cytokine storm (CS), which are characterized by profound elevations in the levels of tumor necrosis factor alpha (TNF- α) and interleukin (IL)-6, lead to peripheral insulin resistance (IR). Besides, high

TNF- α and IL-6 in CS impair pancreatic b-cell function and inhibit insulin secretion. Taken together, both IR and impairment of pancreatic b-cell function contribute to a vicious cycle in the development and progression of hyperglycemia in COVID-19 patients. Furthermore, hyperglycemia and induced oxidative stress and gluco-lipotoxicity contribute to the development of IR and impairment of pancreatic b-cell function (Mehta et al., 2020). In addition, prolonged hyperglycemia could worsen the course of COVID-19 via glycation of pancreatic ACE2, which facilitates the SARS-CoV-2 binding and entry at the pancreatic b-cell. Different reports have shown that an abnormal expression of cell ACE2 receptors in different tissues reduces the protective effect against viral entry and, consequently, exacerbates the severity and poor outcomes of SARS-CoV-2 infection. On the other hand, the systemic renin-angiotensin system (RAS) regulates pancreatic b-cell function, while local RAS of pancreatic b-cell function controls b-cell apoptosis, cell proliferation, and oxidative stress (Al-kuraishy et al., 2021). Angiotensin II (AngII) through AT1R leads to DM induction in experimental models, while inhibiting the glucose-stimulated insulin secretion. Therefore, blockade of AT1R improves pancreatic b-cell function and increases the pro-insulin and insulin biosynthesis. Besides, upregulated local pancreatic AngII induces oxidative stress that triggers b-cell damage by NADPH oxidase induction. Indeed, it has been shown that hyperglycemia upregulates AT1R leading to b-cell function impairment and insulin secretion. In COVID-19, ACE2 dysregulation by SARS-CoV-2 leads to marked elevation of vasoconstrictor AngII with a reduction in the vasodilator Ang1-7, which per se leads to pancreatic b-cell dysfunction, inhibition of insulin secretion, and hyperglycemia, which might be transient even in non-DM patients. Furthermore, elevated AngII leads to pulmonary vasoconstriction, ALI, and ARDS with the induction of inflammation cascade and oxidative stress, which together participate in the induction of pancreatic b-cell function and hyperglycemia (Dwi and Netra, 2020). Then, hyperglycemia in COVID-19 leads to ALI through the induction of pulmonary sodium-potassium chloride co-transporter 1 (NKCC1), involved in the regulation of the transport of water and ions to alveolar cells. Thus, untreated and long-standing hyperglycemia may lead to ALI through ischemic-reperfusion injury. Also noteworthy is the fact that hyperglycemia is associated with oxidative stress' induction and inflammatory mediators' overproduction, which together partake in the development of endothelial



dysfunction and thrombosis due to alterations in both function and generation of antithrombin III. Taken together, these findings reveal that both COVID-19 and hyperglycemia interact in a vicious cycle leading to more complications and worse metabolic outcomes (Al-Namiet et al., 2019).

II. CONCLUSIONS

Suggestions are made on the possible pathophysiological mechanisms of the relationship between diabetes and COVID-19, and its management. No definite conclusions can be made based on current limited evidence. Further research regarding this relationship and its clinical management is warranted. Diabetes and other comorbidities are significant predictors of morbidity and mortality in patients with COVID-19. Future research is urgently needed to provide a better understanding regarding potential differences in genetic predispositions across populations, underlying pathophysiological mechanisms of the association between COVID-19 and diabetes, and its clinical management.

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