#### **REVIEW ARTICLE**



# Association between ABO blood types and coronavirus disease 2019 (COVID-19), genetic associations, and underlying molecular mechanisms: a literature review of 23 studies

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#### Abstract

An association of various blood types and the 2019 novel coronavirus disease (COVID-19) has been found in a number of publications. The aim of this literature review is to summarize key findings related to ABO blood types and COVID-19 infection rate, symptom presentation, and outcome. Summarized findings include associations between ABO blood type and higher infection susceptibility, intubation duration, and severe outcomes, including death. The literature suggests that blood type O may serve as a protective factor, as individuals with blood type O are found COVID-19 positive at far lower rates. This could suggest that blood type O individuals are less susceptible to infection, or that they are asymptomatic at higher rates and therefore do not seek out testing. We also discuss genetic associations and potential molecular mechanisms that drive the relationship between blood type and COVID-19. Studies have found a strong association between a locus on a specific gene cluster on chromosome three (chr3p21.31) and outcome severity, such as respiratory failure. Cellular models have suggested an explanation for blood type modulation of infection, evidencing that spike protein/Angiotensin-converting enzyme 2 (ACE2)-dependent adhesion to ACE2-expressing cell lines was specifically inhibited by monoclonal or natural human anti-A antibodies, so individuals with non-A blood types, specifically O, or B blood types, which produce anti-A antibodies, may be less susceptible to severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) infection due to the inhibitory effects of anti-A antibodies.

Keywords COVID-19 · Coronavirus · 2019-nCoV · ABO blood types · SARS-CoV-2

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### Introduction

## **COVID-19 pandemic**

The World Health Organization (WHO) received several reports of unidentified pneumonia from Wuhan, China on December 31, 2019. On January 7, 2020, the cause of those reported cases was found to be the 2019 novel coronavirus (2019-nCoV). The disease was later named severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) and coronavirus disease 2019 (COVID-19). The WHO declared the outbreak a global health emergency on January 30, 2020 [1].

COVID-19 has become a global concern because it spreads easily and quickly through close contact. The main transmission mechanism is via respiratory droplets containing infectious virus particles. These droplets are transmitted during breathing, coughing, sneezing, and speaking. Airborne transmission and contact transmission are also potential vectors of infection.

Symptoms present following the incubation period, which can be 2–14 days after exposure to the virus. People infected

with COVID-19 may have mild to severe symptoms. Common mild symptoms are dry cough, fever, and fatigue. In severe cases, individuals experience acute respiratory distress syndrome, septic shock, or death [2].

Mortality varies from country to country: currently, the highest mortality is around 8.9% in Mexico, and the lowest is around 0.9% in Turkey [3]. As of December 21, 2020, there are 77,202,828 cases and 1,639,709 deaths worldwide due to COVID-19 infection [3]. Managing the spread of this pandemic is very challenging because individuals are contagious during the incubation period [2]. Moreover, new research has found that 10–30% of people are asymptomatically infected with 2019-nCoV and readily become a conduit for viral transmission [4].

#### **Defining risk factors**

In general, risk factors are variables associated with an increased risk of disease or infection [5]. There are two types of risk factors: nonmodifiable risk factors and modifiable risk factors [6]. A conclusive panel of risk factors for COVID-19 infection susceptibility and prognosis is still being explored; however, several risk factors have been proposed.

Nonmodified risk factors for COVID-19 infection are innate traits. Black and Hispanic heritage were found to be risk factors for testing positive for COVID-19, but not for 30-day mortality in a retrospective cohort study of nearly 6 million individuals seeking care from the US Department of Veteran Affairs [7]. Age over 65 has also been reported as a risk factor for severe outcome, potentially due to increased comorbidities prevalent with age, and decreased efficacy of the immune system, including delayed macrophage function and reduced T cell propagation [8].

Modifiable risk factors can be controlled, manipulated, or treated through intervention or lifestyle changes. Several meta-analyses have reported that tobacco smokers may be susceptible to increased COVID-19 severity compared with nonsmokers [9–11], including increased Intensive Care Unit (ICU) support, mechanical ventilation, and death [12]. However, some studies have also reported non-significant relationships between smoking and COVID-19 severity [13].

Identifying consistent risk factors is critical to ensure that those at higher risk of infection can take added precautions to prevent acquiring the infection. Additionally, understanding those who are at greatest risk for severe prognosis or death may help clinicians better predict patient outcome, allowing for more targeted allocation of limited critical care resources during epidemic surges.

#### Blood type as a non-modifiable risk factor

ABO blood type is an inherited, non-modifiable trait. Individuals can be A, B, AB, or O depending on the antigens present or absent on erythrocyte surfaces. Blood types may also be characterized as positive or negative depending on the presence of the Rhesus (Rh) factor protein. Numerous studies have previously found associations between ABO blood types and viral respiratory infections such as influenza A (H1N1) and acute respiratory syndrome (SARS) [14–16]. Recently, several studies have proposed relationships between blood types and susceptibility to COVID-19, its significance in the course of the disease, and outcomes [17–25].

#### Motivations and study aims

The aim of this work is to summarize the key findings of published studies on this topic to identify the relationship between blood type, COVID-19 infection rates, and outcomes, as well as the role of genetics and the underlying molecular mechanism responsible for susceptibility and severity. Although the relationship between blood type, COVID-19 infection, and severe outcome is still under debate, it is critical to understand the potential risk factor's influence to allow relevant individuals to take added precautions to prevent acquiring the infection.

#### Methodology

A comprehensive online literature search was conducted using the keywords "COVID-19," "2019-nCoV," "SARS-CoV-2," "ABO blood types," and "blood group" on PubMed, Embase (Elsevier), and Google Scholar. The identified studies were screened by title and abstracts by two independent reviewers (authors YZ and SS). After discussing the disagreements, the authors reached a consensus to include 23 studies, all published in English, for the final review. The initial data extraction, including study design, number of patients, selection criteria, limitations, and major findings, was carried out by one of the authors (YZ) and assessed for accuracy by other authors (RG, SS, MLR, and DD). We note that this study was primarily designed as a narrative literature review, hence does not entirely follow the Preferred Reporting Items for Systematic Reviews (PRISMA) [26] method which is designed for systematic reviews/meta-analysis.

# Association between ABO blood groups and viral respiratory infections

The role of blood types in bacterial and viral infections has always been an area of interest among researchers in the field [27]. The association between blood types and different viral respiratory diseases such as H1N1 and severe acute respiratory syndrome (SARS) has been previously discussed in multiple publications [14–16]. SARS, similar to COVID-19, is also caused by a member of Coronaviridae family known as SARS-associated coronavirus (SARS-CoV).

#### Blood types related to severe acute respiratory syndrome infection

In 2003, several unidentified pneumonia cases were reported in Asia, and later, the illness was confirmed to be caused by a coronavirus named SARS-associated coronavirus (SARS-CoV). This severe respiratory disease was named severe acute respiratory syndrome (SARS). Soon, SARS spread globally, including to North and South America, Europe, and Asia [28]. Researchers later found an association between blood type and SARS: individuals with blood type O were less likely to become infected with SARS compared with non-blood type O individuals [14].

#### Blood types related to influenza infection

Influenza is a common viral infection caused by the influenza virus. Type A and Type B are the two main types of influenza viruses. The flu is a contagious respiratory disease that can cause mild to severe illness, with common symptoms including fever, cough, and sore throat. Flu viruses can be easily spread by droplets that are transmitted when people talk, cough, or sneeze [29]. Studies have shown an association between flu infection and blood type. During the outbreak of flu caused by H1N1, a subtype of Type A influenza virus, patients with blood group A and B were more susceptible to infection than those with blood group O and AB [15]. Similarly, researchers found that individuals with blood group B were more susceptible to infection by H3N2, a Type A influenza subtype [16].

# Association between ABO blood type and COVID-19

In order to review evidence of associations between ABO blood types and COVID-19 infection, 23 literature papers were studied. Summarized findings include the associations between blood type and higher infection susceptibility, severe outcomes, and death, as well as genetic association and potential molecular mechanisms that drive the relationship between blood type and COVID-19 infection.

# Association of blood type with higher susceptibility of testing positive

Recent studies have shown that blood type A is associated with the highest probability of SARS-CoV-2 infection [17–21]. A summary of findings is reported in Table 1. One of the earliest studies in Wuhan, China, suggested that there was an association between blood type A and COVID-19, noting that females with blood type A were more susceptible to infection [17]. Subsequent studies also reported that blood type A patients had significantly higher odds of acquiring SARS-CoV-2 infection compared with non-A blood types [18–20].

Several studies have also reported that blood type O patients have significantly lower odds of infection, suggesting that blood type O may be a protective factor against infection [18–20, 22, 23].

In one multivariate analysis of 14,112 patients who tested positive for COVID-19 in the New York Presbyterian hospital system, investigators reported that blood type A, AB, and B had higher prevalence than blood type O after adjusting for race and ethnicity [24]. A retrospective study at the First Hospital of Changsha in Changsha, Hunan, China, similarly showed that blood type is a strong risk factor for COVID-19. Blood type O patients had decreased risk of infection compared with non-O blood group patients and blood type A patients had higher risk than all other groups [21].

While the literature generally agrees that blood type O is a protective factor [22], there is disagreement on which blood type has the highest susceptibility. Contrary to studies citing blood type A as the most at-risk group, several studies have suggested high risk for blood type B and AB [23]. One of the largest retrospective cohort studies indicated that risk ratios associated with SARS-CoV-2 infection were 0.87, 1.09, 1.06, and 1.15 for O, A, B, and AB individuals, respectively. Risk ratio compares the risk of health event among one group with another group. If the risk ratio equals to 1, it indicates the identical risk among two groups. A risk ratio is greater than 1, which means the group in the numerator is the exposed group. The risk ratio is smaller than 1; it indicates that the exposed group has lower risk and may play a protective role [32]. Padhi et al. [25] also reported that blood type B and AB were associated with higher odds of testing positive for SARS-CoV-2 infection, whereas blood type O was associated with lower odds of testing positive. However, the authors found no correlation between blood type A and positive test rates.

## ABO blood type and COVID-19 clinical outcomes of severity

Scientists found that ABO group is associated not only with COVID-19 infection rates but also with severe outcomes, such as requiring mechanical ventilation or death (Table 2). Literature regarding which blood type is most at risk of severe outcome are more variable than the literature regarding risk of infection.

For example, blood type O is regarded as a protective factor for infection, but it may not decrease risk of severe outcome. In a study conducted by Zietz et al. [24], blood type A patients had lower risk of intubation and death compared with blood type O patients, whereas blood type AB had higher risk of intubation and death. Moreover, individuals with blood type B had higher rates of intubation, but lower mortality compared

|          | Reference             | Study design                                      | Number of patients   | Selection criteria  | Limitations  | Major findings   |
|----------|-----------------------|---|--|---|--|--|
| l _      | Fan et al. [17]       | A hospital-based<br>case-control study            | Cases: <i>N</i> = 105.<br>Controls: <i>N</i> =<br>103                                  | Patients tested positive at Zhongnan<br>Hospital of Wuhan University in<br>Wuhan, China, from January 1,<br>2020, to March 5, 2020                                    | Small sample size; possible regional<br>selection bias to be considered; some<br>other respiratory diseases may<br>affect the research results   | Blood type A patients had higher risk and<br>blood type O patients had lower risk of<br>becoming infected with SARS-CoV-2  |
| 7        | Zhao et al. [18]      | A meta-analysis<br>using random effects<br>models | N = 2,173  | Patients from Jinyintan Hospital in<br>Wuhan, China; Renmin Hospital<br>of Wuhan University in Wuhan,<br>China; and Shenzhen Third<br>People's Hospital, in Shenzhen, | The number of patients is relatively<br>small; control group used lack of<br>age and sex factor for multivariate<br>analysis to adjust; the status of<br>chronic medical conditions        | Blood type A patients had higher risk<br>and blood type O patients had lower risk<br>of becoming infected with SARS-CoV-2  |
| ŝ        | Golinelli et al. [19] | A meta-analysis using<br>random effects models    | Cases: $N = 7,503$ .<br>Controls: $N = 2,962,160$                                      | Data in MEDLINE [30] and<br>LitCovid [31] databases from<br>studies published until July<br>15 <sup>th</sup> , 2020   | was mussing<br>Evidence and data relied on the<br>scientific literature to date; cases and<br>controls may be not adequately<br>representative; difficult to perform                       | Blood type A patients had higher risk and<br>blood type O patients had lower risk<br>of becoming infected with SARS-CoV-2  |
| 4        | Zietz et al. [24]     | A cohort study                                    | <i>N</i> = 14,112  | Patients above 18 years old who<br>tested positive for SARS-CoV-2<br>with known blood types in the<br>New York Presbyterian hospital                                  | a unitorn aquisatizent<br>The study mostly focuses on severely<br>ill patients and may not be<br>generalizable to all COVID-19<br>patients   | Blood type O was protective against<br>SARS-CoV-2 infection compared<br>with non-O blood type  |
| Ś        | Wu et al. [20]        | A meta-analysis                                   | N = 31,100   | system<br>Rudiabases: PubMed, MedRxiv,<br>BioRxiv, Web of Science, and<br>China National Knowledge  | The study missed some<br>demographic data, including age<br>and gender   | Blood type A patients had higher risk and<br>blood type O patients had lower risk<br>of becoming infected with SARS-CoV-2  |
| 9        | Wu et al. [21]        | A retrospective analysis                          | <i>N</i> = 187   | Patients with COVID-19 recorded<br>between January 20, 2020, and<br>March 5, 2020, at the First<br>Hosoiral of Channeda   | The sample size is relatively small;<br>most COVID-19 patients did not test<br>for their blood type; lack of normal  | Blood type A patients had higher risk<br>and blood type O patients had lower<br>risk of becoming infected with<br>SARS_CVV.7   |
| Г        | Barnkob et al. [22]   | A retrospective cohort<br>analysis                | SARS-CoV-2<br>positive: <i>N</i> = 7422.<br>SARS-CoV-2<br>negative: <i>N</i> = 466,232 | Danish people who were tested<br>for SARS-CoV-2 between<br>february 27, 2020, and<br>July 30, 2020, with<br>known hlood aroun   | population group as control<br>Some individuals lacked blood type<br>information   | Blood type O was protective against<br>SARS-CoV-2 infection compared<br>with non-O blood type  |
| $\infty$ | Latz et al. [23]      | A retrospective<br>multi-institutional<br>study   | <i>N</i> = 1289  | Patients tested positive for<br>SARS-CoV-2 with a known<br>blood type   | Sample size was relatively small<br>and there might be a lead-time bias  | Blood type B and AB were associated<br>with higher risk of testing positive for<br>COVID-19. Blood type O was<br>associated with lower risk of<br>testing positive. No correlation was<br>found between blood type A and |
| 6        | Padhi et al. [25]     | Mixed method                                      | N = 277,649  | COVID-19 patient data collected<br>from the Ministry of Health and<br>Family Welfare of India   | The study was not directly performed<br>in COVID-19 patients but based<br>on online datasets; some potential<br>covariates, such as age and gender,<br>were not considered in the analysis | positive test rates.<br>Blood type O may be protective<br>against COVID-19 infection   |

 Table 1
 Studies on associations between blood type and higher susceptibility of testing positive for COVID-19

| Tabl | e 2 Summary of find | ings for ABO blood type and CO  | <b>DVID-19</b> clinical outcon | nes of severity  |  |  |
|------|---------------------|---|--------------------------------|--|--|--|
|      | Reference           | Study design  | Number of patients             | Selection criteria   | Limitations  | Major outcomes   |
| -    | Zietz et al. [24]   | A cohort study  | <i>N</i> = 14,112              | Patients above 18 years old who tested<br>positive for SARS-CoV-2 with<br>known blood types in the<br>New York Presbyterian<br>hospital system   | The study mostly focuses on<br>severely-ill patients, so it<br>may not be generalizable to<br>all COVID-19 patients  | Blood type A patients had lower<br>risk of intubation and death<br>compared with blood type O<br>patients, whereas blood type<br>AB had higher risk of both of<br>intubation and death |
| 7    | Wu et al. [20]      | A meta-analysis   | N = 31,100                     | Studies extracted from five<br>databases: PubMed, MedRxiv,<br>BioRxiv, Web of Science, and<br>CNKI.  | The study missed some<br>demographic data, including<br>age and gender   | There is no correlation between<br>blood type and COVID-19<br>severity or mortality  |
| ε    | Latz et al. [23]    | A retrospective<br>multi-institutional study  | <i>N</i> = 1289                | Patients tested positive for<br>SARS-CoV-2 with a known<br>blood type  | Sample size was relatively<br>small and there might be a<br>lead-time bias   | There is no correlation between<br>blood type and COVID-19<br>intubation or death  |
| 4    | Padhi et al. [25]   | Mixed method  | N = 277,649                    | COVID-19 patient data collected<br>from Ministry of Health and<br>Family Welfare of India  | The study was not directly<br>performed in COVID-19<br>patients but based on online<br>datasets; some potential<br>covariates such as age<br>and gender were not<br>considered in the analysis | Blood type O was protective<br>against COVID-19 death, while<br>blood type B was more strongly<br>correlated with death.   |
| Ś    | Ray et al. [33]     | A population-based<br>retrospective<br>cohort study   | N = 225,556                    | People who had ABO blood group<br>assessed between January 2007<br>and December 2019 and who<br>also had SARS-CoV-2 testing<br>conducted between January 15,<br>2020, and June 30, 2020, in<br>Ontario, Canada | Patients who died shortly<br>after the infection   | Blood type A and AB had<br>higher risk of severe illness<br>or death   |
| 9    | Hoiland et al. [34] | A multicenter retrospective<br>analysis and nested<br>prospective observational<br>substudy | N = 125                        | ICU patients in 6 metropolitan<br>Vancouver hospitals between<br>February 21, 2020, and April<br>28, 2020  | Data were only focused on<br>critically ill patients   | Blood type A and AB patients<br>are at higher risk of requiring<br>mechanical ventilation and<br>experience more severe<br>outcomes  |

to individuals with blood type O. Conversely, a populationbased cohort study conducted in Ontario, Canada, reported that blood type O individuals have lower risk of developing severe conditions or death compared with all other blood types [23]. Similarly, in a study conducted using Spearman's rank coefficient analysis on COVID-19 patients collected from the Ministry of Health and Family Welfare in India, scientists found that blood type O was protective against COVID-19 death and that blood type B was more strongly correlated with infection and death [25].

In contrast, Latz et al. [23] proposed that blood type A had highest risk of intubation and death. Similarly, a multicenter retrospective analysis showed that patients with blood type A or AB were at higher risk of requiring mechanical ventilation when compared with blood type O or B patients after adjusting for sex, age, and other factors. Blood type A (12% of cohort) and AB patients (32% of cohort) also had higher probability of requiring continuous renal replacement therapy compared with blood type O or B patients after adjusting for sex, age, and other factors. After adjustments, there were no differences in median length of ICU stay or probability of ICU discharge or extubation. There was also no association between blood type and death [34].

### Association of Rhesus factor with susceptibility to COVID-19 infection and outcomes

As blood types can be characterized as positive or negative depending on the presence of the Rhesus (Rh) factor protein, it is also significant for researchers to investigate the association between Rh and COVID-19 infection (Table 3).

A study conducted by Latz et al. [23] reported that Rh(+) was associated with higher odds of testing positive and higher risks of intubation and death. Another study conducted by Zietz et al. [24] showed that Rh(-) patients had a 2.7% lower risk of initial infection, intubation, and death after adjusting for sex, age, and other demographics. A study conducted by Ray et al. [33] similarly indicated that individuals with Rh(-) blood type were less susceptible to infection by SARS-CoV-2, especially individuals with O-negative blood type.

However, another hematological study conducted in Sudan proposed that the O-positive blood group has the lowest risk of having severe symptoms, while the authors agreed that Apositive individuals were the most vulnerable when exposed to the virus [35].

#### Genetic associations with susceptibility and severity

ABO blood type is determined by the ABO gene, located on the 9<sup>th</sup> chromosome. It consists of 7 exons and codes for enzyme glycosyltransferases, which is responsible for the formation of antigens in blood type A and/or B [34]. In a meta-analysis, investigators sampled 1980 patients with COVID-19-related respiratory failure at seven centers in Italy and Spain and analyzed 8,582,968 single-nucleotide polymorphisms (SNPs). Researchers reported a cross-replicating association with rs11385942 at locus 3p21.31 and with rs657152 at locus 9q34. They found that the 3p21.31 gene cluster is a genetically susceptibility locus in COVID-19 patients with respiratory failure [36]. Another study conducted by Shelton et al. [37] using trans-ethnic genome-wide association studies similarly reported a strong association between blood type and COVID-19. In particular, there was strong association at a locus on chr3p21.31 with outcome severity [37].

# Underlying molecular mechanism of susceptibility and severity of ABO blood types

Several researchers have proposed potential molecular mechanisms underlying the susceptibility of particular ABO blood types to COVID-19. A review conducted by Silva-Filho et al. [38] proposed that the association between ABO blood type and COVID-19 susceptibility and infection progression may be related to the varied distribution of sialic acid-containing receptors on host cells' surfaces. This distribution is modulated by ABO antigens through carbohydrate-carbohydrate interactions (CCIs), which could maximize or minimize the binding capacity of the virus's spike protein, the structural glycoprotein that protrudes from the viral envelope, to host cell's surface receptors. Viral entry is facilitated by interaction of two subunits, S1 and S2. S1 specifically contributes to viral binding to host cell surface receptors via two domains: S1A, corresponding to the N-terminal region, which interacts with sialic-acid containing glycoproteins and glycolipids, and S1B, corresponding to the receptor-binding domain, which binds to Angiotensin-Convertor Enzyme 2 (ACE2) receptors. ACE2 is the central entry route for several coronaviruses into host cells [39].

ABH antigens are present on the erythrocyte membrane as well as many other cells, including lymphocytes, platelets, and arterial and venular capillary endothelium [40]. Antigen A especially, but also antigens AB and B, stimulate carbohydrate clustering, whereas H, the antigen characteristic of O blood type, does not induce carbohydrate promotion. Additional carbohydrate clustering facilitates CCIs, maximizing interaction, cell recognition, and aggregation [38, 41]. Increased interaction increases the probability of SARS-CoV-2 successfully binding to host cells via specific binding of spike protein domains to ACE2 and CD147, a transmembrane protein that may also facilitate the infectious process via viral anchoring to host cells [38].

|   | Reference         | Study design                                  | Number of patients | Selection criteria  | Limitations  | Major outcomes  |
|---|-------------------|---|--------------------|---|--|---|
|   | Zietz et al. [24] | A cohort study                                | <i>N</i> = 14,112  | Patients above 18 years old who tested<br>positive for SARS-CoV-2 with<br>known blood types in the New<br>York Presbyterian hospital system   | The study mostly focuses on<br>severely-ill patients, so it<br>may not be generalizable to<br>all COVID-19 patients                                  | Rh(-) patients had a 2.7% lower risk<br>of initial infection after adjustment,<br>and also lower risk for both<br>intubation and death  |
| 0 | Latz et al. [23]  | A retrospective<br>multi-institutional study  | <i>N</i> = 1289    | Patients tested positive for<br>SARS-CoV-2 with a known<br>blood type   | Sample size was relatively small<br>and there might be a<br>lead-time bias   | Individuals with Rh(-) blood type<br>were less susceptible to infection<br>by SARS-CoV-2  |
| ε | Ray et al. [33]   | A population-based retrospective cohort study | N = 225,556        | People who had ABO blood group<br>assessed between January 2007<br>and December 2019 and who also<br>had SARS-CoV-2 testing conducted<br>between January 15, 2020, and<br>June 30, 2020, in Ontario, Canada | Patients who died shortly<br>after the infection   | B-positive was associated with higher<br>odds of testing positive, whereas<br>O-negative associated with lower<br>infection rate; Rh(+) had higher<br>risk of severe illness or death |
| 4 | Taha et al. [35]  | A case-control study                          | N = 557            | Individuals surveyed online   | Potential exclusion bias due<br>to reliance on the internet for<br>survey completion (i.e., older<br>people may have less<br>access to the internet) | O-positive blood group has the<br>lowest risk of having severe<br>symptoms, and A-positive<br>individuals were the most<br>vulnerable when exposed to<br>the virus                    |

 Table 3
 Summary of findings for association of Rh with susceptibility to the COVID-19 and outcomes

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Similar work conducted on SARS-CoV proposed that natural anti-A antibodies may play a role in modulating interaction between the virus and host cells. In the cellular model, spike protein/ACE2-dependent adhesion to ACE2-expressing cell lines was specifically inhibited by monoclonal or natural human anti-A antibodies. Thus, individuals with non-A blood types, specifically O or B blood type, which produce anti-A antibodies, may be less susceptible to SARS-CoV-2 infection due to the inhibitory effects of anti-A antibodies [42].

Several studies have also suggested that host transmembrane protease serine subtype 2 (TMPRSS2) may play a significant role in ABO group modulation of infection [43, 44]. TMPRSS2 is a gene located on human chromosome 21q22.3 [45], and has been found to be essential for spike protein priming and subsequent infection of SARS-CoV [44]. However, it has not been confirmed if SARS-CoV and SARS-CoV-2 share identical genomic sequences for the serine protease. Despite this, proteolysis of viral serine by TMPRSS2 may allow for serine mobilization, a key molecule in mucin O-glycan that has been found to be critical for infection [44]. Following hybridization to yield O-N-acetyl-D-galactosamine (O-GalNAc), SARS-CoV-2 may evade innate or specific immunity by hybridizing ABO blood group, effectively mimicking self-cell presentation. A, B, and AB blood groups have upregulated immunoglobulin M (IgM) activity, while O-group has downregulated IgM activity due to glycosylation. The A, B, and AB blood groups are thus preferential targets because they possess A/B phenotypic-determining enzymes which facilitate greater viral molecular contact, whereas O blood type lacks these enzymes and only binds the virus via hybrid H-type antigen formation. Additionally, IgM downregulation in the O blood group leads to downstream anti-A and anti-B isoagglutinin activity, hallmarks of innate immune activity [43]. Arend et al. [43] and Wang et al. [46] proposed TMPRSS2 inhibition as a potential therapeutic target.

# Conclusion

This review examined published empirical papers and metaanalyses related to the relationship between blood type and COVID-19. These studies suggest that blood type may be a risk factor for COVID-19 infection and outcome. Findings related to the highest risk of infection varied among researchers. The majority of researchers report that the greatest risk for susceptibility to COVID-19 infection is among individuals with blood type A, while some others report that individuals with blood type B are the most vulnerable group to infection. Although some researchers report that there is no correlation between blood type and COVID-19 severity or mortality, most studies found that blood types A and AB had higher risk of severe illness or death, while blood type O was protective against death or severe outcomes. Genetic association and several potential mechanisms related to the relationship among blood types A/B, infection susceptibility, and severe outcome were also reported. It has been found that the 3p21.31 gene cluster is a genetically susceptibility locus in COVID-19 patients with respiratory failure. In the cellular model, ACE2 is an entry for virus into host cell. Spike protein/ACE2-dependent adhesion to ACE2-expressing cell lines was specifically inhibited by monoclonal or natural human anti-A antibodies. Thus, in individuals with non-A blood types, specifically O or B blood type, anti-A antibodies may play a protective role in SARS-CoV-2 infection due to its inhibitory effects. Moreover, A, B, and AB blood groups are preferential targets compared to O blood group because they possess A/B phenotypic-determining enzymes which facilitate greater viral molecular contact; however, blood type O lacks these enzymes and only binds the virus via hybrid Htype antigen formation.

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#### Declarations

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