



A comprehensive review of COVID-19 biology, diagnostics, therapeutics, and disease impacting the central nervous system

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Abstract

The ongoing COVID-19 pandemic, caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), is a highly transmissible disease. SARS-CoV-2 is estimated to have infected over 153 million people and to have caused over 3.2 million global deaths since its emergence in December 2019. SARS-CoV-2 is the seventh coronavirus known to infect humans, and like other coronaviruses, SARS-CoV-2 infection is characterized by a variety of symptoms including general flu-like symptoms such as a fever, sore throat, fatigue, and shortness of breath. Severe cases often display signs of pneumonia, lymphopenia, acute kidney injury, cardiac injury, cytokine storms, lung damage, acute respiratory distress syndrome (ARDS), multiple organ failure, sepsis, and death. There is evidence that around 30% of COVID-19 cases have central nervous system (CNS) or peripheral nervous system (PNS) symptoms along with or in the absence of the previously mentioned symptoms. In cases of CNS/PNS impairments, patients display dizziness, ataxia, seizure, nerve pain, and loss of taste and/or smell. This review highlights the neurological implications of SARS-CoV-2 and provides a comprehensive summary of the research done on SARS-CoV-2 pathology, diagnosis, therapeutics, and vaccines up to May 5.

Keywords SARS-CoV-2 · COVID-19 · Coronavirus · CNS

Introduction

In December 2019, a novel coronavirus was discovered in Wuhan China. Initially designated as 2019-nCoV by the World Health Organization (WHO) on January 12, 2020, this virus became the latest entrant in the family of coronaviruses able to infect humans (WHO 2020a). On February 11, 2020, the virus was renamed severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) by the International Committee on Taxonomy of Viruses (ICTV), and the WHO subsequently named the disease caused by SARS-CoV-2 as coronavirus disease 2019 (COVID-19) (Coronaviridae

Study Group of the ICTV 2020; WHO 2020a). Having been declared as the sixth public health emergency of international concern (after H1N1, polio, Ebola (West Africa), Zika, and Ebola (Democratic Republic of Congo)) by the WHO, the resulting outbreak of COVID-19 has caused a pandemic that has accelerated at an unprecedented scale (Lai et al. 2020). As of May 2021, there are an estimated 153 million reported cases and over 3.2 million global deaths associated with COVID-19, with numbers continuing to rise daily (WHO 2021).

Coronaviruses are enveloped viruses that contain a positive-sense, single-stranded RNA genome approximately 30 kb in size (Fehr and Perlman 2015). Briefly, coronaviruses belong to the Coronaviridae family of the Nidovirales order. Coronaviridae is divided into the two subfamilies Torovirinae and Coronavirinae, the latter of which is categorized into four genera including Alphacoronaviruses, Betacoronaviruses, Gammacoronaviruses, and Deltacoronaviruses (Pal et al. 2020). SARS-CoV-2, specifically, is classified in the betacoronavirus genera. Previous human betacoronaviruses have caused epidemics, namely severe acute respiratory syndrome (SARS) in 2002 and Middle East respiratory

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syndrome (MERS) in 2012, but SARS-CoV-2 has a greater transmission rate, albeit lower mortality (Liu et al. 2020; Petrosillo et al. 2020). To underscore this point, the basic reproduction number (R_0), which is a key epidemiologic metric used to indicate the transmission potential of an infectious agent, has been estimated to range between 2 and 3 for SARS-CoV-2 (Delamater et al. 2019; Lai et al. 2020; Salzberger et al. 2020). For reference, SARS had an associated R_0 of approximately 1.7 to 1.9 while the R_0 associated with MERS was reported to be < 1 (Petrosillo et al. 2020). The high transmission rate of SARS-CoV-2 may be the cause of the higher viral loads that have been observed during early infection (Hu et al. 2020; Wölfel et al. 2020; Zou et al. 2020). Moreover, it has been suggested that asymptomatic carriers of SARS-CoV-2, which in some cases have ranged from 18 to 81%, have largely contributed to the spread of the pandemic (Nikolai et al. 2020).

With the emergence of SARS-CoV-2, there are now a total of seven coronaviruses which can infect humans. As previously mentioned, the zoonotic betacoronavirus SARS-CoV-1 resulted in the SARS epidemic that first emerged in Southern China in November 2002. At the time of the last documented case in 2003, the virus had spread to 27 countries and had resulted in approximately 8000 probable cases with a mortality rate of $\sim 10\%$ (Cherry 2004; Perlman and Netland 2009). Similarly, in two separate incidents in 2012 and 2015, MERS-CoV was responsible for MERS outbreaks originating in Saudi Arabia and South Korea, respectively. To date, MERS is still considered an endemic disease in many Middle Eastern countries and there have been 2,500 confirmed cases with an associated fatality rate of $\sim 35\%$ (Chafekar and Fielding 2018; Willman et al. 2019).

Evidence suggests that the evolutionary origins of both SARS-CoV-1 and MERS-CoV involved bats and later adapted to intermediate hosts such as civets and camels, respectively, prior to humans (Corman et al. 2014). This information suggests that bats could be the natural reservoir for SARS-CoV-2; however, the intermediate host(s) have yet to be confirmed (Wu et al. 2020; Zhao et al. 2020; Zhou et al. 2020). Along these lines, the phylogenetic analysis from a recently published article showed that SARS-CoV-2 shares 88.1% identity to two bat-derived SARS-like coronaviruses (bat-SL-CoVZC45, bat-SL-CoVZXC21), whereas only 79% and 50% homology was shared with SARS-CoV-1 and MERS-CoV, respectively (Lu et al. 2020; Wu et al. 2020; Zhou et al. 2020). Another recent report has studied the interactions between the SARS-CoV-2 spike protein receptor binding domain (RBD) and the host receptor ACE2, revealing that pangolins, snakes, and turtles may also serve as potential intermediate hosts of SARS-CoV-2 (Liu et al. 2020).

Like other human coronaviruses (e.g., 229E, OC43, NL63, HRU1), SARS-CoV-2 infection is characterized by

symptoms similar to those of the common cold including fever, cough, fatigue, sore throat, and shortness of breath (Guan et al. 2020; Kumar et al. 2020; Su et al. 2016). In critical cases, patients often develop pneumonia, lymphopenia, acute kidney injury, and cardiac injury (Huang et al. 2020). The most severe cases of COVID-19 are associated with an over-activation of the immune system which can induce a cytokine storm. In the absence of therapeutic intervention, this can lead to significant pulmonary inflammation and extensive lung damage, acute respiratory distress syndrome (ARDS), secondary infection, multi-organ failure, deep venous thrombosis, and sepsis, all of which may ultimately result in death (Hojyo et al. 2020; Tang et al. 2020; Wichmann et al. 2020). Although the respiratory system is the primary target of SARS-CoV-2, it is also becoming clear that the virus may also drive dysfunction and pathology in the central nervous system (CNS). Previous reports have suggested that up to 30% of patients infected with SARS-CoV-2 suffered from neurological symptoms such as dizziness, ataxia, seizure, nerve pain, encephalitis, and impairments relating to vision, taste, and smell (Ellul et al. 2020; Li et al. 2020; Mao et al. 2020).

Since its emergence, SARS-CoV-2 has caused a significant public health crisis that has left a crippling effect not only on the medical system, but also across numerous socio-economic platforms (Ceylan et al. 2020). Thus, if left unchecked, SARS-CoV-2 could pose the greatest healthcare challenge of the century. At the time of this article, there are three FDA emergency use authorization (EUA)-approved vaccines available in the USA and a few others throughout the world. Therapeutic options currently center around preventive measures and supportive care. This review will highlight the current knowledge surrounding the biology, diagnostics, and therapeutic options relating to COVID-19. A special emphasis will also be placed on discussing the implications and potential long-term complications of SARS-CoV-2 in the context of the CNS.

COVID-19 pathology

COVID-19 is caused by the single-stranded, positive-sense RNA virus SARS-CoV-2. Its highly contagious nature, coupled with the large proportion of asymptomatic transmitters, has significantly contributed to the rapid spread of COVID-19. However, since the onset of the outbreak, substantial progress has been made in the understanding of SARS-CoV-2 transmission. This has led to the implementation of certain measures to decrease its spread. While it is widely accepted that respiratory transmission (via respiratory droplets) is the primary route of transmission, other modes of transmission such as fomites, domestic pets/farm animals, fecal, aerosol,

sexual, and bloodborne transmission cannot be conclusively ruled out at this time (Meyerowitz et al. 2020).

After exposure to the virus, symptoms typically manifest after an incubation period of 5 to 6 days; however, incubation periods of up to 14 days have also been reported (Guo et al. 2020; Shanmugam et al. 2020). It is also worth noting that in older adults, symptoms may be more severe and are often further complicated in individuals with comorbidities and other medical conditions (e.g., asthma, chronic obstructive pulmonary disease, cardiovascular disease, hypertension, diabetes) (Ahn et al. 2020; Wolff et al. 2020). Although the virus establishes infection primarily in the respiratory tract, it has also become widely appreciated that the virus can drive pathology outside of the lungs, including in the digestive tract, the circulatory system, and the central nervous system (Zhang et al. 2020).

Similar to SARS-CoV-1, SARS-CoV-2 primarily utilizes the host cellular receptor angiotensin-converting enzyme 2 (ACE2) to enter host cells and establish infection (Zhou et al. 2020). Briefly, the mechanisms of viral attachment and cell entry are mediated by the spike (S) protein located on the surface of the virus. The S protein directly associates with the ACE2 receptor via its receptor-binding motif (RBM) (Weiss and Leibowitz 2011). This is followed by membrane fusion and the release of viral RNA (Guo et al. 2020). ACE2 is mainly found on lung alveolar epithelial cells as well as small intestine enterocytes, which could explain the lower respiratory and gastrointestinal symptoms experienced by some SARS-CoV-2 patients, as well as explain the tropism of SARS-CoV-2's abundant replication in the upper respiratory epithelia (Guo et al. 2020; Wiener et al. 2007; Xu et al. 2020; V'kovski et al. 2021; Zhang et al. 2020). However, broad distribution of the ACE2 receptor has also been reported by others who have found varying levels of ACE2 mRNA or protein expression in distinct cell types (e.g., vascular endothelial and smooth muscle) and tissues (heart, kidney, thyroid, liver, colon, bladder, and brain) (Dong et al. 2020; Hamming et al. 2004; Li et al. 2020; Zou et al. 2020). Additionally, there is evidence of ACE2 expression in glial cells and neurons within the brain and upregulated on activated type 1 macrophages (M1) and on tissue-specific macrophages (Baig et al. 2020; Panariello et al. 2020; Song et al. 2020).

The ACE2 receptor has multiple potential functions during infection with SARS-CoV-1 or SARS-CoV-2, which may include the following: to mediate viral entry, initiate intracellular inflammation, and induce shedding of the ACE2 receptor. Shedding of the ACE2 ectodomain, or the downregulation of ACE2, was found to be coupled with tumor necrosis factor- α (TNF- α) activation, a trait associated with severe cases of SARS-CoV-1 infection (Haga et al. 2008). Increased levels of TNF- α could contribute to inflammation and subsequent tissue damage, increasing the severity of

the infection. Altogether, these effects may further facilitate viral spread.

Studies have shown that SARS-CoV-2 may use a variety of host proteins to enter cells as well as ACE2. The SARS-CoV-2 S protein is a homotrimer, each consisting of two subunits, S1 and S2 (Silva-Filho et al. 2020). The S1 subunit is further broken down into two parts S1A (n-terminal domain, NTD) and S1B (receptor binding domain, RBD). ACE2 is proposed to bind with the RBD while sialic acid containing glycoproteins like CD147 is proposed to bind to the NTD (Silva-Filho et al. 2020). For ACE2 binding, transmembrane proteins such as transmembrane protease serine 2 (TMPRSS2) and Cathepsin-L are important for the cleavage of the SARS-CoV-2 spike protein to facilitate host cell membrane fusion and for viral infectivity (Kumar et al. 2020).

Another proposed receptor for SARS-CoV-2 infection is type II transmembrane ectopeptidase protein dipeptidyl peptidase IV (DPP4; also known as CD26). DPP4 has been found enriched in the bronchiolar and alveolar epithelia of the lower respiratory tract, liver, and immune cells of MERS cases; and studies have found high levels of DPP4 and CD147 in SARS-CoV-2-infected cortical cells that had low ACE2 expression (Andrews et al. 2021; Boonacker and Van Noorden 2003; Chan et al. 2013; Hocke et al. 2013; Mattern et al. 1991; Meyerholz et al. 2016; Raj et al. 2013; Widagdo et al. 2019). Furthermore, Andrews et al. discovered a 30% reduction of SARS-CoV-2 N protein and a 70% reduction of a stress indicating gene, ARCN1, in astrocytes treated with a DPP4 inhibitor (Vidagliptin) compared to untreated cells, thus, indicating a potential SARS-CoV-2 tropism for cells with DPP4 (Andrews et al. 2021).

In addition to ACE2 and DPP4, there may be other receptors used by coronaviruses. For instance, GRP78 has been suggested as a possible secondary receptor for SARS-CoV-2; and research has demonstrated five peptides that bind to both the virus Spike protein and the host GRP78 binding sites to prevent viral entry (Allam et al. 2020). Furthermore, SARS-CoV-1 infection has been detected in cells that lack expression of ACE2 receptors, such as in hepatocytes, and cells that naturally express low levels of ACE2 and DPP4, such as those in the CNS, suggesting a potential role of unknown host factors in viral entry and infection (Bernstein et al. 2018; Li et al. 2020; To et al. 2004). In addition, not all ACE2 expressing human endothelial and intestinal cells become infected by SARS-CoV-1 (Chan et al. 2004; Ding et al. 2003).

To better understand the mechanics of SARS-CoV-2 infection (such as important host receptors), *in vitro* experiments are crucial. Researchers use a variety of cells such as human airway epithelial cells, brain microvascular endothelial cells, and immortal cell lines such as Vero E6 cells to mimic SARS-CoV-2 replication (Jiao et al. 2021; Takayama 2020). Furthermore, there is an increased interest in

lab grown organoids used to replicate cell systems, such as 3D brain models and human lung organoids (Ramani et al. 2020; Takayama 2020). To further understand the mechanisms of viral spread in SARS-CoV-2, it may be important to, initially, consider the symptomology of COVID-19 patients along with *in vitro* studies to determine preferentially infected systems (e.g., lungs, intestines, or CNS). These observations may eventually lead to the discovery of additional host-related factors.

Lung pathology

Earlier studies of SARS-CoV-1 reported that downregulation of the ACE2 receptor was associated with viral entry, specifically due to the interaction between the cytoplasmic domain of ACE2 and the S protein of SARS-CoV-1 (Haga et al. 2008; Kuba et al. 2005). The consequences of this deregulation are believed to be linked to intracellular inflammation and deterioration of pulmonary health in SARS-CoV-1-infected patients (Imai et al. 2005). ACE2 has been identified as a key enzyme in the regulation of inflammation related to hypertrophy, pulmonary hypertension, glomerulonephritis, lung injury, sepsis, and acute pancreatitis (Gaddam et al. 2014). ACE2, as well as DPP4, is predominantly expressed by cells of the respiratory tract. Therefore, it is important to examine the relationship between SARS-CoV-1 infection, viral entry via the ACE2/DPP4 receptors, and the associated inflammatory response in infected patients.

As mentioned above, the SARS-CoV-1 S protein induces pleiotropic effects when interacting with the recipient cell. It has been reported that binding of the S protein induces a TNF- α -converting-enzyme (TACE)-mediated loss of the ectodomain (extracellular) of the ACE2 receptor and increased production of TNF- α in Vero E6 cells (Haga et al. 2008). Deletion of the carboxyl-terminal domain (cytoplasmic tail) of the ACE2 receptor or through the use of small interfering RNA (siRNA) against TACE decreased viral infection, indicating the importance of the cytoplasmic domain and TACE to viral entry (Haga et al. 2008). A recent study has shown SARS-CoV-2 S protein shares 76% of its primary sequence with SARS-CoV-1 S protein and it is thought that SARS-CoV-2 induces a similar affect (Ou et al. 2020; Saponaro et al. 2020).

Both SARS-CoV-1 and SARS-CoV-2 are characterized by viral spread through the respiratory tract and potential transmission from person-to-person via droplets, respiratory secretions, aerosols, and direct contact (Guo et al. 2020). The lungs are among the most susceptible organs during SARS-CoV-1 and SARS-CoV-2 infection, with early signs of pneumonia in many patients (Gu et al. 2005; Huang et al. 2020; Weiss et al. 2011). The primary cell types of the alveolar walls of the lungs consist of alveolar type I and II epithelial cells (pneumocytes) and alveolar macrophages

(Gralinski and Baric 2015). However, others have reported that only type II pneumocytes, which produce pulmonary surfactant and regulate alveolar surface tension in the lungs, express ACE2 (Bombardini and Picano 2020). As such, it is not surprising that SARS-CoV-1 targets type II pneumocytes, thereby inducing the production of inflammatory cytokines and decreasing surfactant levels (Gralinski and Baric 2015). Therefore, type II pneumocytes expressing ACE2 may represent a key cell type potentially involved in early stages of SARS-CoV-2 infection via oral/nasal routes (Bombardini and Picano 2020; Hoffmann et al. 2020; Uhal et al. 2011; Wiener et al. 2007).

Interestingly, prior observations of patients infected with SARS-CoV-1 described lung pathologies consistent with scattered alveolar damage, unlike other human coronaviruses, and a tropism towards pneumocytes as confirmed by the presence of viral RNA (To et al. 2004). Furthermore, pulmonary pathological examinations of both SARS-CoV-1 and SARS-CoV-2 patients showed pneumocyte hyperplasia (i.e., atypical proliferation of pneumocytes) (Tian et al. 2020; Weiss and Leibowitz 2011). Notably, this pathological transformation marked early phases of infection as patients had not yet developed clinical symptoms (Tian et al. 2020). This implicates pneumocytes as one of the potential primary targets of SARS-CoV-2 infection. Chest computed tomography (CT) scans of COVID-19 patients have also shown abnormalities. For instance, chest CT scans in a cohort of 41 hospitalized patients revealed pneumonia in all patients, with bilateral involvement in 98% of patients; bilateral, multiple-lobular, and subsegmental areas of consolidation in intensive care unit (ICU) patients; and bilateral ground-glass opacity in non-ICU patients (Huang et al. 2020). These indicators are similar to those of SARS-CoV-1 and MERS-CoV infections and further support the potential susceptibility of pneumocytes to SARS-CoV-2 (Huang et al. 2020).

Systemic inflammation

SARS-CoV-2 infection has been linked to elevated cytokine levels in patients contributing to cytokine storms, especially in cases of severe COVID-19. The most commonly detected cytokines in the plasma of COVID-19 patients include a wide range of proinflammatory cytokines, such as IL1 β , IL6, IL12, CXCL10, IL2, IFN γ , and monocyte chemoattractant protein (MCP1) (Guo et al. 2020; Huang et al. 2020; Xiong et al. 2020). Additionally, Huang et al. noted that plasma samples obtained from ICU patients exhibited higher levels of macrophage-secreting cytokines/chemokines, including granulocyte colony-stimulating factor (GCSF), 10 kD interferon-gamma-induced protein (IP10), MCP1, macrophage inflammatory protein 1- α (MIP1 α), and TNF- α compared to non-ICU patients, thus pointing towards the potential role of cytokines in the severity of infection (Huang et al.

2020). As such, corticosteroids, routinely used to suppress inflammation, have been used in the treatment regimen of patients infected with SARS-CoV-1 or SARS-CoV-2, targeting cytokine-producing cells (e.g., endothelial cells and macrophages), reducing proinflammatory cytokine levels, and mitigating inflammatory-induced lung injury (Huang et al. 2020; Wang et al. 2020).

In addition to the aforementioned cytokines, patients suffering from severe infection with SARS-CoV-1, MERS-CoV, or SARS-CoV-2 have been reported to have high-levels of serum chemokines such as CXL9 and IL8, which may lead to the induction of a cytokine storm (Liu et al. 2020). These groups of chemokines are typically produced by monocytes/macrophages and endothelial cells and are synergistically enhanced in the presence of TNF- α , thereby leading to systemic inflammation (Tokunaga et al. 2018). For instance, transient changes in the expression of cytokines and chemokines, such as the early induction of CXCL10 and IL2, and overexpression of IL6 have been found to play a role in the immunopathological response that results in lung injury in SARS-CoV-1 infection (Chien et al. 2006; Liu et al. 2020). Furthermore, the delayed expression of type I interferon (IFN) associated with SARS-CoV-1, in conjunction with strong viral replication, stimulates the accumulation of pathogenic inflammatory monocyte-macrophages and leads to increased concentrations of cytokines/chemokines in the lungs, virus leakage, and diminished virus-specific T cell responses (Channappanavar et al. 2016). Overall, an increased production of proinflammatory cytokines has been observed in both SARS-CoV-1 and SARS-CoV-2, suggesting a significant role of proinflammatory cytokines, chemokines, and immune cells (e.g., macrophages) in the advancement and severity of the infection (Channappanavar et al. 2016; Liu et al. 2020).

Autopsy results of SARS-CoV-1 patients have shown lung injury demonstrated by diffuse alveolar damage, organizing pneumonia, squamous metaplasia, and bronchiolitis obliterans at later stages of infection, while the lungs of individuals that succumbed to the infection in the early stages featured a loss of pneumocytes, edema, hyaline membranes, small vessel thrombi, lymphocytes, macrophages, and polymorphonuclear leukocytes in the lungs (Liu et al. 2020; Weiss and Leibowitz 2011). Similar lung injury attributed to cytokine storm in the lower respiratory tract was also reported in the deceased bodies of COVID-19 patients (Guo et al. 2020). The consistent manifestation of cytokine storms in multiple severe COVID-19 cases led to the investigation of drugs used to treat malaria, such as chloroquine, due to their ability to suppress the production of proinflammatory cytokines (i.e., IL6 and TNF- α), as a treatment for COVID-19 (Guo et al. 2020; Touret and de Lamballerie 2020). Overall, the high infectivity of COVID-19 stemming in part by inflammation caused by cytokine storms urgently calls for

additional novel studies as well as treatment options to curb its global spread.

Neuroinflammation and CNS pathology

Electron microscopy (EM) images as well as reverse transcription polymerase chain reaction (RT-PCR) performed during the autopsy of SARS patients revealed a significant number of viral particles and SARS-CoV-1 genome sequences in the cytoplasm of neurons (Gu et al. 2005). This finding is suggestive of the ability of SARS-CoV-1, and by extension SARS-CoV-2, to access anatomically privileged organs such as the brain. Along these lines, certain neurological symptoms (i.e., dizziness, ataxia, seizure, nerve pain, and impairments relating to vision, taste, and smell) have been reported to affect up to 30% of patients infected with SARS-CoV-2 (Li et al. 2020; Mao et al. 2020).

SARS-CoV-1 has been found in the brains of humans and animal models where the brainstem was heavily infected (Li et al. 2020). Similarly, SARS-CoV-2 RNA has now been documented within different areas of the brain including the cerebellum and medulla oblongata (Meinhardt et al. 2020). The murine coronavirus known as mouse hepatitis virus (MHV) provides a model system for the *in vivo* study of viral tropism and pathogenesis in the CNS, including encephalitis and multiple sclerosis (Weiss and Leibowitz 2011). Following MHV infection, neutrophils, macrophages, and natural killer (NK) cells are the first to be recruited to the CNS (Bergmann et al. 2006). In the CNS, MHV infection also results in the induction of chemokines (e.g., CXCL9, CXCL10, CCL2, CCL3, and CCL5), cytokines (e.g., IL1 α , IL1 β , IL6, and IL12), and matrix metalloproteinases (MMPs) (Bergmann et al. 2006). As previously mentioned, some of these proinflammatory cytokines, including IL1 β , IL6, and IL12, have been detected in plasma samples of both SARS-CoV-1 and SARS-CoV-2 patients (Guo et al. 2020; Huang et al. 2020). Furthermore, microglia, the resident immune cells of the brain, are known to elicit the release of IL1 β , IL6, and TNF- α in response to infection in the CNS (Wang et al. 2015). Collectively, these cytokines contribute to neurodegeneration elicited by sustained inflammation. Chemokine and cytokine expression may additionally impact the permeability of the blood–brain barrier (BBB), thereby modifying the CNS-infiltrating cell populations (Bergmann et al. 2006). Production of MMP-9 (a metalloproteinase with effects on BBB permeability) by neutrophils and upregulation of adhesion molecules on CNS endothelium leads to relaxation of the BBB, facilitating the ensuing entry of additional inflammatory cells into the CNS to combat the infection (Bergmann et al. 2006). Collectively, these data imply that a secondary cytokine storm may initiate in the CNS, resulting in the neurologic symptoms that have been associated with COVID-19.

Regulatory mechanisms aimed at curbing host immune responses with the intention of maintaining CNS integrity can inadvertently lead to the lack of viral clearance. Growing evidence suggests that coronaviruses can infect the CNS possibly by first entering peripheral nerve terminals, allowing access to the CNS via a synapse-connected route (Li et al. 2020). This pathway was studied in the primary motor cortex of a rat model infected with swine hemagglutinating encephalomyelitis virus (HEV), a type of CoV (Li et al. 2013). Here, it was discovered that the virus spread from neuron to neuron via membranous-coated-mediated endocytosis using synaptic junctions, as evidenced by the presence of viral particles in axons, invaginations of presynaptic membranes, synaptic clefts, and postsynaptic cytoplasm (Li et al. 2013). This begs the critical question as to whether CNS infection by SARS-CoV-1 or SARS-CoV-2 can lead to neuronal injury. In support of this, multiple independent studies revealed the presence of SARS-CoV-1 proteins, including the viral antigen protein “N” in the brain of infected individuals (Ding et al. 2004; Gu et al. 2005; Xu et al. 2005). Another animal study using transgenic mice expressing human ACE2 receptor that were intranasally and intracranially infected with SARS-CoV-1 saw an extensive neuronal infection in the brain (measured by plaque assays and immunohistochemical staining for the viral N protein) followed by death despite minimal infection in the lungs (Netland et al. 2008). Taken together, these data suggest that neurons are susceptible to SARS-CoV-1 infection. Considering the similarities in viral structure and infection pathway of SARS-CoV-1 and SARS-CoV-2, the above data could be applicable to COVID-19. As such, neurologic symptoms observed in COVID-19 patients could generate from the invasion of the CNS by SARS-CoV-2, potentially via infection of the neurons and/or neuroinflammation, as ACE2 and other potential receptors such as neuropilin-1 expression have also been detected in the CNS (Barrantes 2020; Davies et al. 2020; Harmer et al. 2002). Likewise, proinflammatory cytokines (e.g., IL1 β , IL6, and TNF- α) primarily linked to SARS-CoV-1 immunopathology have been shown to be upregulated in the brains of transgenic mice expressing ACE2 (McCray et al. 2007).

Neurological symptoms from COVID-19 have been detected weeks past the acute phase (Groiss et al. 2020). Through the use of electrophysiological studies (e.g., electroencephalography), Groiss et al. found that both the peripheral nervous system and the central nervous system were affected by SARS-CoV-2 in 4 male patients. Delirium was seen in these patients and cognition was impaired even after delirium passed (Groiss et al. 2020). Another study showed 4 children with COVID-19 who displayed similar neurological symptoms (e.g., encephalopathy, headache, brainstems with signs of dysarthria/dysphagia, meningism, cerebellar ataxia, muscle weakness, reduced reflexes,

and lesions in the splenium of corpus callosum) had little to no respiratory symptoms (Abdel-Mannan et al. 2020). Therefore, Abdel-Mannan et al. suggested to test children for SARS-CoV-2 if neurological symptoms are displayed even with no other typical COVID-19 symptoms (Abdel-Mannan et al. 2020).

Recently, a study conducted by Mao et al. on a cohort of 214 COVID-19 hospitalized patients revealed neurological symptoms in 36.4% of patients and this percentage increased to 45.5% in more severe cases (Mao et al. 2020). Neurological manifestations included acute cerebrovascular events, dizziness, seizure, headache, impaired consciousness, nerve pain, loss of smell, loss of vision, loss of taste, muscle injury, and stroke such as cerebral venous thrombosis (blood clot in the venous sinuses of the brain) which may lead to death (Mao et al. 2020, Tu et al. 2020). Researchers also found that patients with CNS symptoms had lower lymphocyte levels, platelet counts, and higher blood urea nitrogen (Mao et al. 2020). As such, the immune system of these patients was suppressed, paving the way for a more severe infection. Therefore, the CNS manifestations associated with COVID-19 could potentially be explained by the presence of the ACE2 receptor in the CNS by providing a pathway for SARS-CoV-2 to bind to and infect cells (Hamming et al. 2004). Another theory postulates that SARS-CoV-2 enters and invades the CNS via the nasal route hence the loss of smell symptom experienced by some patients (Mao et al. 2020). This is supported by a study by Meinhardt et al. which found SARS-CoV-2 through anatomically distinct areas of the brain and nose (e.g., olfactory mucosa, olfactory bulb, olfactory tubercle, oral mucosa, trigeminal ganglion, medulla oblongata, and cerebellum), with the highest amounts being associated with the olfactory mucosa beneath the cribriform plate.

Loss of olfactory and taste functions

As mentioned earlier, an additional symptom experienced by SARS-CoV-2-infected patients has been the sudden and complete loss of olfactory function. Reports of multiple patients from around the globe have been documented (Gautier and Ravussin 2020; Gilani et al. 2020; Moein et al. 2020; Yan et al. 2020). More specifically, a woman in her 40 s diagnosed with SARS-CoV-2 (i.e., positive RT-PCR test), also presented with loss of smell, without dysgeusia and nasal obstruction, but with inflammation in the olfactory cleft confirmed by magnetic resonance imaging (MRI) (Eliezer et al. 2020). The olfactory cleft allows the flow of odors to the olfactory epithelium and subsequently to the olfactory bulb. This one patient was unable to detect odors such as phenyl-ethyl-alcohol (flower rose), cyclotene (caramel), isovaleric acid (goat cheese), and undecalactone (fruits), and skatole (manure) (Eliezer et al. 2020). The

olfactory epithelium (also known as the olfactory mucosa) is composed of epithelial cells, basal cells, and olfactory receptor neurons (Chen et al. 2017). These cells respond to cytokines, such as TNF- α and IL1 β (Chen et al. 2017). Interestingly, SARS-CoV-2 infection has been shown to cause an increase in expression of these same cytokines (Guo et al. 2020; Huang et al. 2020). Thus, the pathophysiology of SARS-CoV-2 may cause simultaneous effects on the lower respiratory tract, while also affecting neighboring cells (as present in the respiratory tract), resulting in effects on the CNS, such as in this case.

Loss of taste, like loss of smell, has been observed in COVID-19-infected individuals. Changes in saliva rate, flow, and content (e.g., hormones, enzymes, ions) are proposed to have a role in the loss of taste (Abduljabbar et al. 2020). Furthermore, the loss of smell/taste has been described as “the most discriminative symptom between seropositive and seronegative persons” in a population-based study within the Netherlands (Vos et al. 2020). This study took serological samples and used multiplex-immunoassays to look for the presence of IgG antibodies against SARS-CoV-2 spike protein in the population (3207 participants) along with a questionnaire and found that 53% of individuals with SARS-CoV-2 antibodies experienced loss of taste/smell (Vos et al. 2020). Similar findings have been reported in other studies, and there is reason to believe that self-reported loss of smell/taste is lower than reported (Gözen et al. 2020). Gözen et al. showed that out of the 59 patients tested, 83% were found to have loss of smell when Sniffin’ Sticks tests were used but only 52.5% of patients self-reported loss of smell as a symptom. Another study showed that loss of smell and loss of taste were some of the most persistent symptoms ~ 125 days of symptom onset indicating again the potential neurological impacts of COVID-19 (Petersen et al. 2020). These studies are representative of the growing interest in the effects of COVID-19 within the CNS and PNS and demonstrate the long-lasting effects even after viral clearance.

Chronic COVID syndrome

There is now a growing number of individuals who have chronic COVID syndrome (CCS; aka long-haulers) who have mostly recovered from the acute COVID-19 phase but have lingering or new symptoms months after infection (Baig 2020). As this is a developing pandemic, there is still much to learn but there already are studies that have analyzed large data sets of CCS cases which have helped to characterize the symptoms and groups most afflicted (Baig 2020; Graham et al. 2021; Huang et al. 2021; Lopez-Leon et al. 2021).

Like acute COVID-19, CCS impacts a variety of organs and displays a range of symptoms. Some of the most prevalent symptoms include sore throat, chills, tachycardia,

extreme fatigue, hair loss, inability to perform daily tasks, shortness of breath, muscle pain, chest pain, insomnia, gastrointestinal problems, and neurological symptoms (Baig 2021; Graham et al. 2021; Lopez-Leon et al. 2021). The typical neurological symptoms are as follows: brain fog, headache, attention disorder, numbness/tingling, loss of smell and/or taste, dizziness, and blurred vision (Baig 2021; Graham et al. 2021; Huang et al. 2021; Lopez-Leon et al. 2021). In one study, Graham et al. performed 100 neurological exams and determined through tests of recall and serial sevens that these CCS individuals had short-term memory deficit as well as an attention deficit (Graham et al. 2021).

There have now been a few studies that have characterized the people who are more at risk. In a study of 1407 COVID-19 cases from the University of California COVID Data Set, 27% had symptoms upwards of 60 days and of these CCS cases, 32% were asymptomatic during the acute phase (Huang et al. 2021). Huang et al. also found women to be more at risk of having CCS. This was supported in another study that found 70% the CCS in the study of 150 cases were women (Graham et al. 2021). In the aforementioned study by Graham et al., individuals with comorbidities such as such as depression, anxiety, and autoimmune disease were also at a higher risk of CCS (Graham et al. 2021). There is still much to be understood about CCS, but these first few studies have shown the importance of reducing the transmission of SARS-CoV-2, not just for acute COVID-19 but CCS as well.

Development of diagnostics

The primary goal for SARS-CoV-2 containment is to reduce transmission and, thus, reduce infection and prevent symptoms like those of the CNS. Currently, outside of vaccination, the only method available to reduce transmission is through identifying and isolating infectious persons (Cheng et al. 2020). To do this, efficient diagnosis is required, and as symptoms alone are not enough for diagnosis, diagnosis is contingent upon the availability of tests, resources, and the time to obtain diagnostic results (Cheng et al. 2020; Udugama et al. 2020).

Current diagnostic tests include nucleic acid testing, antibody testing, and protein testing. The accuracy of these tests is defined by their sensitivity and specificity, positive tests in patients with disease, and negative tests in healthy individuals, respectively (Weissleder et al. 2020). Many aspects of diagnostic tests impact their sensitivity and specificity, one of those being sample collection sites. The two most common specimens for SARS-CoV-2 are nasopharyngeal samples and blood serum (Esbin et al. 2020). Nasopharyngeal swab samples test for viral RNA and collect high viral titers starting on the first day of infection whereas blood serum samples test for host antibodies as SARS-CoV-2 levels are

low in blood (Cheng et al. 2020; Esbin et al. 2020; Udugama et al. 2020).

Nucleic acid testing

Nucleic acid testing (NAT) is the primary method for diagnosing COVID-19 as it currently has the earliest detection and is the most sensitive (Esbin et al. 2020; Udugama et al. 2020). NAT results show the presence or absence of known genetic sequences (genetic markers) within the genome of the virus under investigation (Udugama et al. 2020). The CDC suggests targeting two regions of the viral nucleocapsid (N) gene, N1 and N2, as well as human RNase P gene to confirm RNA extraction (CDC 2020a). Nucleocapsid genes, viral envelope (env), RNA-dependent RNA polymerase (RdRp), and others may also be targeted (WHO 2020b). NAT for SARS-CoV-2 is mainly performed on nasopharyngeal swabs due to their high sensitivity and ability to detect viral titers the first day of infection (Cheng et al. 2020; Esbin et al. 2020).

Many nucleic acid tests rely on RT-PCR. To date, there are over 181 commercialized RT-PCR kits approved for emergency use by the FDA (FDA 2021d; Foundation for Innovative New Diagnostics [FIND] 2021). Briefly, there are three basic steps to the CDC RT-PCR kit: sample collection/transportation, lysis and RNA purification, and amplification (FDA 2021c). Esbin et al. have an informative summary comparing some of the new commercialized kits (Esbin et al. 2020). RT-qPCR has also been used in studies such as the one mentioned above, to map the presence of SARS-CoV-2 RNA throughout the brain (Meinhardt et al. 2020).

Other NATs are under development and a few are currently in use. These include CRISPR, isothermal amplification, and next-generation sequencing (NGS) (Broughton et al. 2020; Guo et al. 2020; Nat Biotechnol 2020; Weissleder et al. 2020; Yoshikawa et al. 2020). CRISPR-Cas12- and CRISPR-Cas13-based assays simultaneously perform reverse transcription and isothermal amplification from RNA extracted from nasopharyngeal or oropharyngeal swabs; then, it goes through Cas12 or Cas13 detection of coronavirus sequences with no cross reactivity from related coronavirus strains (Broughton et al. 2020). There are currently 2 CRISPR diagnostic tests approved for commercial use by FDA EUA (FDA 2021d; FIND 2021). Isothermal amplification including a new battery powered kit is used for rapid testing in remote locations (Yoshikawa et al. 2020). There are currently 8 isothermal amplification tests approved by FDA EUA (FDA 2021d; FIND 2021). NGS is used for high-volume screening and population management, and there are now 4 approved commercially used FDA EUA (FDA 2021d; FIND 2021; Nat Biotechnol 2020).

Serological testing

Serological testing detects antibodies (e.g., immunoglobulins) or interleukins (e.g., IL6) in a patient's blood serum (Esbin et al. 2020; Weissleder et al. 2020). Serological tests are rapid, robust, and easy (as easy as a finger prick); however, this testing method lacks specificity and relies on the host's production of antibodies which can take 5–10 days after the onset of symptoms (Cheng et al. 2020; Esbin et al. 2020; Lee et al. 2020). Although the FDA does not approve the sole use of serological testing for COVID-19 diagnosis, there are over 75 serological tests FDA EUA for commercial use (including one at home test) that aid in the detection of post-infection syndrome and late presenting symptoms (CDC 2020a; FDA 2021b; FDA 2021c; FIND 2021). For example, in a case where an individual presented manic-like symptoms after recovery of vital signs, a serological test was used to find that the cerebrospinal fluid was positive for SARS-CoV-2-specific IgG (Lu et al. 2020a, b). Serological tests are also helpful for epidemiological and surveillance studies tracking antibody longevity and transmission dynamics and are also used as a tool to confirm negative results (CDC 2020a; Esbin et al. 2020; Weissleder et al. 2020). For these reasons, there are many companies working to create point-of-care serological tests (Lisboa Bastos et al. 2020).

Immunoassays are the main method used for antibody testing. Most infections incite the production of IgM first and then IgG, but in SARS-CoV-2 infections they are both produced around the same time (Lee et al. 2020). Lee et al. suggest that both IgG and IgM levels may correlate to disease severity and have reported that IgG3 may be the dominant subtype in SARS-CoV-2 infection (Lee et al. 2020). While there are different types of immunoassays, the most common are CLIA (chemiluminescence immunoassay) and ELISA (enzyme-linked immunosorbent assay) due to their simple readouts (Lee et al. 2020). In addition to the wait time for antibody production, another main issue is cross-reactivity with past infections from other coronaviruses (e.g., SARS-CoV-1, MERS-CoV), or even current infections with endemic coronaviruses like MERS-CoV (Lee et al. 2020; Weissleder et al. 2020).

Protein testing

Viral antigen tests (VATs) currently probe for nucleocapsid (N) or spike (S) proteins on SARS-CoV-2 through nasopharyngeal or other respiratory tract samples (Weissleder et al. 2020). VATs are most accurate in the early phase of infection as intermediate phase and late phase have varied to consistently negative results, and current tests have too low sensitivity to rule out infection (Cheng et al. 2020; Hirotsu et al. 2020a). However, VATs are useful in tracking

viral clearance in infected persons as there is a gradual loss of viral antigens over time as opposed to the positive/negative results seen with RT-qPCR, although the persistence of antigens does not mean infectious host (Hirotzu et al. 2020a).

There are a variety of tests for detecting viral antigens. The most common are through the use of lateral flow assays or single antigen ELISA tests and take around an hour to complete (Steiner et al. 2020; Weissleder et al. 2020). LUMIPULSE is a chemiluminescence enzyme immunoassay that is high throughput with the ability to run 60–120 samples in 30 min on automated machines and has the potential for hospitals to track viral clearance in patients (Hirotzu et al. 2020b). There are also monoclonal antibody tests in development that use monoclonal antibodies to test against the N proteins of SARS-CoV-2 and may create a rapid antigen test (Cheng et al. 2020). Other antigen tests include western blots, virus neutralization test (VNT) that uses antibodies against specific virus proteins to identify the virus, and immunofluorescence microscopy (IFM) which uses antibody interaction with virus proteins to create fluorescent glow that can be visualized with microscopy to identify the virus (Cheng et al. 2020; Weissleder et al. 2020). Protein testing can also be used for understanding the SARS-CoV-2 infection. For example, Song et al. used immunolabeling against the N protein in their transgenic mice (expressing human ACE2) to show the viral infection of SARS-CoV-2. This experiment showed the presence of SARS-CoV-2 throughout the brain of the infected mice, specifically in the neural cells of the fore brain, as well as the S protein in human brains (Song et al. 2021).

Point-of-care testing

The need for rapid point-of-care testing (on-site diagnosis) is crucial for infected individuals to isolate themselves. Many tests have been developed for point-of-care (POC) diagnosis of COVID-19, but POC tests tend to have lower throughput, are more expensive, or are not automated (Weissleder et al. 2020). Although there are issues with POC testing, quick diagnosis in hospitals allows for improved infection control measures including the following: patient flow, enrollment of patients into clinical trials, admission to secondary care, reduction of nosocomial infection in non-COVID-19 positive patients who might otherwise be exposed, and reduction in bed movement which reduces exposure to cleaning staff and healthcare staff (Brendish et al. 2020). This can be especially useful with individuals who display CNS symptoms without the tell-tale respiratory symptoms that may otherwise fly under the radar (Abdel-Mannan et al. 2020; Mao et al. 2020).

There are now a variety of POC tests. LAMPs are popular POC tests as there is no need for thermocyclers and results can be ready in 20 min (Orooji et al. 2021). There have been

advances in PCR kits which can produce results in less than 30 min via cartridges or lateral flow technology, and 15-min immunoassays (Cheng et al. 2020; Orooji et al. 2021; Weissleder et al. 2020). Other POC tests that are in development now include microarrays which are cost effective, non-fluorescent, low-density oligonucleotide assay tools that have been used for whole genomic detection of past coronaviruses, and mobile analysis platforms (MAPS) which have been used to diagnose past acute respiratory illnesses (de Souza Luna et al. 2007; Hardick et al. 2018; Prabhakar & Lakhanpal 2020).

Ancillary tests and programs

On top of these diagnostic tests, there are a variety of tools that epidemiologists, healthcare workers, businesses, and individuals may use that help to determine possible exposure, detect early symptoms, and confirm COVID-19 infections. Health care professionals may use other tests to help determine if a patient is COVID-19 positive along with the abovementioned tests including CT scans and ultrasounds (Cheng et al. 2020; Taylor et al. 2020). Healthcare professionals may also use a variety of non-contact tools to determine abnormal breathing and/or heart rate with less exposure to themselves. For example, RGB-thermal cameras can be used to diagnose COVID-19 on a mask wearing individual with 83.7% accuracy (Taylor et al. 2020).

Temperature monitoring has become a useful tool used to detect infected individuals with low-grade fevers and little to no symptoms. This period typically occurs in the early stage of COVID-19 where viral shed is highest and so detecting a low-grade fever can encourage testing and isolation (Zhou et al. 2020a, b, c). Infrared thermography especially has become useful as it can detect body temperature without physical touch and can screen for potential infected people in high-risk areas like airports. All these diagnostic methods help to identify infected individuals which leads to isolation and a reduction of viral transmission, helping to prevent infection and thus CNS symptoms.

Therapeutics for the treatment of COVID-19

Due to the novelty of SARS-CoV-2 and the spread of the virus, scientists and clinicians have worked together to discover therapeutics for the treatment of COVID-19 and alleviants for the disease symptoms. As described above, there are a variety of systems impacted including the respiratory tract, gastrointestinal tract, central nervous system, and systemic inflammation. Due to the heterogeneity in symptoms and disease severity, as well as the heterogeneity across populations, a variety of treatments are needed. The rapid spread of the virus has resulted in many drugs and therapeutics

that are already approved for other similar infections like SARS and MERS or for general antiviral activities to be used for potential COVID-19 treatment (Reddy et al. 2020). For example, the S proteins of SARS-CoV-2 and SARS-CoV-1 share 77.5% identical amino acid sequence and use the ACE2 receptor to enter host cells (Li et al. 2003; Walls et al. 2020; Wrapp et al. 2020; Zhou et al. 2020a, b, c). As such, the interaction of the S protein of SARS-CoV-2 with the host cellular receptor(s) (e.g., ACE2) is the main target for inhibiting viral entry into cells (Li et al. 2005). Additionally, other therapies like stem cell therapy are also being considered (Harrell et al. 2020; Kim et al. 2020).

Host protein–targeted approaches

One method of viral suppression is through targeting host proteins that play a role in viral replication. Human receptor ACE2 and TMPRSS2 are the main host proteins targeted by therapeutics with 115 and 32 clinical trials currently underway, respectively (Clinical Trials 2021). Vitamin D and calcitriol (which have known anti-inflammatory effects) are studied for their preemptive effects to help regulate angiotensin-renin mechanisms, which increase lipopolysaccharide permeability into lungs and reduce severe lung damage (Sarwar et al. 2020; Xu et al. 2017). Typical ACE2 inhibitors and receptor antagonists Captopril and enalapril, and losartan and valsartan, respectively, are under study along with new inhibitory methods including single-chain variable fragments, VHH domains, nanobodies, fusion of human recombinant soluble ACE2 (hrsACE2) to an immune-adhesion molecule, and even the use of inhibitory peptides isolated from marine organisms (Festa et al. 2020; Zhou et al. 2020a, b, c).

The primary focus of TMPRSS2 inhibition is on camostat mesylate. Camostat mesylate is a serine protease inhibitor that has been shown to reduce inflammatory markers IL6, TNF- α , and TGF- β as well as inhibit TMPRSS2 and thus block viral entry and replication (Breining et al. 2021). Other drugs under study for TMPRSS2 inhibition are aprotinin and nafamostat mesylate (Bojkova et al. 2020; Zhou et al. 2020a, b, c). Nafamostat mesylate has shown neuroprotective effects with neurovascular ischemia and has been shown to reduce the production of inflammatory cytokines, which could be another benefit when used by COVID-19 individuals afflicted with CNS cytokine storm brain injury (Duan et al. 2018; Ghali and Ghali 2020; Li et al. 2016).

While the main host proteins involved with SARS-CoV-2 entry (ACE2 and TMPRSS2) are important targets for therapeutics, there are also other proteins that may play a role in SARS-CoV-2 replication including GRP78 and CD147. As discussed above, GRP78 may be a secondary receptor, and a study showed an increase in the inhibitory effect of viral entry through the use of four polyphenols which bind

to GRP78 (Allam et al. 2020). CD147, another potential secondary receptor, has been shown to be blocked by meplazumab, a humanized anti-CD147 antibody that is used to treat malaria (Wang et al. 2020a, b, c).

Monoclonal antibodies

In recent years, monoclonal antibodies have emerged as powerful therapeutic tools for a variety of pathologies, including those related to cancer, immunology, and cardiovascular and respiratory diseases (Singh et al. 2018). In general, this form of immunotherapy relies on the manufacturing of monoclonal antibodies that are able to bind specifically to certain antigens. This binding consequently results in the induction of an immune response that will eliminate the targeted cell(s) (Zhou et al. 2020a, b, c).

Antibody-based treatments are a useful option for immediate effects or disease prevention for people at high risk of infection, as well as in the treatment of infected patients to prevent disease progression (Nie et al. 2004). To date, there is FDA EUA of bamlanivimab, and the combinations of bamlanivimab/etesevimab, and casirivimab/imdevimab for the treatment of COVID-19 (FDA 2020b, FDA 2021a). Similar to these EUA treatments, most monoclonal antibodies under consideration work by targeting the SARS-CoV-2 S protein (e.g., 80R, CR3013, CR3014, CR3022, and m396) (Chung et al. 2020; National Center for Biotechnology Information [NCBI] 2020; Sui et al. 2004; ter Meulen et al. 2006; van den Brink et al. 2005). Recently, Wang et al. reported the first human monoclonal antibody (47D11) against the spike receptor of SARS-CoV-2, as well as SARS-CoV-1, which could potentially be used to control COVID-19 spread (Wang et al. 2020a, b, c). Several other monoclonal antibodies have been identified as potential candidates to neutralize SARS-CoV-1 and have been reviewed elsewhere (Jin et al. 2017; Liu et al. 2020a, b, c, d; Nie et al. 2004; Prabhakaran et al. 2009; Zhou and Zhao 2020).

Antiviral monoclonal antibodies have also been developed to neutralize proteins associated with SARS pathogenesis such as antibodies against IL6/IL6R, TLR3, CD16, immunoreceptor tyrosine-based activation motif (ITAM), dendritic cell-specific intercellular adhesion molecule-grabbing nonintegrin (DC-SIGN), intercellular adhesion molecule 3 (ICAM-3), or interferon γ -inducible protein 10 (IP10/CXCL10) (Liu et al. 2020a, b, c, d). Antibody-dependent opsonization or complement activation is the primary mechanism of virus clearance after neutralization (Coughlin and Prabhakar 2012).

Currently, monoclonal antibody treatment of COVID-19 has focused on respiratory tract, but due to their anti-inflammatory effects, monoclonal antibodies should be studied to help mediate CNS distress. Previous studies have found monoclonal antibodies to be effective in the treatment of a

variety of CNS and PNS diseases in animal models such as stroke, brain injury, and epilepsy (Liu et al. 2007; Nishibori et al. 2019; Wang et al. 2017; Zhang et al. 2011). Due to the similarity of CNS damage from SARS-CoV-2 infection, it would be beneficial to further investigate these effects.

Suppression of immune response

The release of cytokines results in the induction of an immune response that aids in viral clearance. As described above, with respect to SARS-CoV-2, the most severe infections are often associated with the development of cytokine storms that may lead to lung damage, incite CNS inflammation, and may ultimately lead to death (Chen et al. 2020a, b). Elevated levels of pro-inflammatory cytokines (e.g., TNF α , IL1 β , IL2, IL6, IFN α , IFN β , IFN γ , and MCP-1) are observed in many patients with severe infections. Therefore, the use of existing approved treatments coupled with the development of anti-inflammatory therapies may help reduce the mortality rate of SARS-CoV-2. Intravenous immunoglobulin (IVIG), tocilizumab, and methylprednisolone have been used to manage cytokine storms and to halt respiratory failure in severe SARS-CoV-1 patients, and could potentially be utilized as treatments for the current COVID-19 outbreak (Chen et al. 2020a, b; Lee 2014). Currently, there are 7 completed clinical trials that investigated IVIG therapy against COVID-19 that measured the ability of IVIG to reduce the immune response through providing an anti-inflammatory, immunomodulatory effect in severe cases but none is approved by FDA EUA at this time (Clinical Trials 2021).

Tocilizumab is marketed as a rheumatoid arthritis drug and can bind to the membrane bound and soluble forms of IL6 receptor (Kaly and Rosner 2012; Zhang et al. 2020a, b, c, d). In a recent study, patients with SARS-CoV-2-induced ARDS displayed clinical improvements after taking tocilizumab for 5 days (Xu et al. 2020a, b). Additionally, siltuximab, emapalumab, and anakinra are being tested in clinical trials to demonstrate the efficacy of these drugs in alleviating severe symptoms in SARS-CoV-2 patients (Clinical Trials 2021).

Despite the efficient use of corticosteroids (e.g., methylprednisolone) in the reduction of early acute phase viral infection, their use is not routinely recommended to treat SARS-CoV-2 patients (Russell et al. 2020a; Russell et al. 2020b; Russell et al. 2020a, b, c). Corticosteroids may not be effective and may even produce adverse effects as seen in prior cases (e.g., avascular osteonecrosis in SARS, a decreased ability to eradicate virus in later stages of MERS; and increased mortality rate and secondary infections in influenza patients) (Arabi et al. 2018; Russell et al. 2020a; Russell et al. 2020b; Russell

et al. 2020b; Chang 2005; Griffith et al. 2005; Lee et al. 2015; Rodrigo et al. 2016).

Cytokine signaling pathways are largely dependent on the JAK-STAT pathway. Therefore, Janus kinase (JAK1/JAK2) inhibitors (e.g., baricitinib or tofacitinib) could be used to block cytokine release in order to dampen inflammation. Baricitinib is used as an anti-rheumatic drug and may reduce SARS-CoV-2 entry to the host cells via interference with the viral endocytosis moderators: adaptor protein-2 complex (AP2) and adaptor-associated kinase-1 (AAK1) (Richardson et al. 2020a). Baricitinib has also been used in the CNS of an animal model, and resolved hallucinations in a COVID-19 patient, potentially indicating its ability to reduce neurological deficits (Hoang et al. 2021; Richardson et al. 2020b). Since then, baricitinib along with antiviral remdesivir has received FDA EUA for patients over 2 years of age who require supplemental oxygen, invasive mechanical ventilation, or extracorporeal membrane oxygenation (FDA 2020a).

Convalescent plasma therapy

Convalescent plasma (CP) from recovered SARS-CoV-2-infected patients could be potentially beneficial to treat SARS-CoV-2-infected patients (Hoffmann et al. 2020). In this sense, CP containing antibodies developed against the previous infection (i.e., against viral proteins) could be used to neutralize viral infection and lead to viral clearance (Marano et al. 2016). CP has been used as a therapeutic option in a variety of viral infections, such as Ebola virus, SARS-CoV-1, avian influenza A(H5N1) virus, and influenza A(H1N1) virus (Chen et al. 2020; Hung et al. 2013; Luke et al. 2006). SARS patients who received CP from SARS-recovered donors observed a shorter length of hospital stay and a lower mortality rate relative to the untreated group (Soo et al. 2004; Cheng et al. 2005).

Many patients recovering from SARS-CoV-2 infection have high titers of neutralizing antibodies (< 1:640) against several SARS-CoV-2 proteins within 2–3 weeks following the onset of the symptoms (Shen et al. 2020). These neutralizing antibodies can be detected by ELISA and other quantitative tests (Shen et al. 2020). Therefore, plasma transfusion could be given to high-risk cases in order to reduce the severity of the disease in combination with antiviral therapies.

A recent study reported improvement of clinical symptoms in 6 patients after CP treatment including a reduction in body temperature, viral loads, and C-reactive protein (Zhang et al. 2020). Additional studies are required for large testing groups with a proper control group, to evaluate whether the observed improvement is caused by the CP or antiviral drugs. Moreover, it is crucial to determine whether CP should only be used for the critical patient, as well as the

efficient time that is needed to administer plasma transfusion (Shen et al. 2020).

It is also worth noting that patients that receive CP transfusions may develop severe allergies, transfusion-associated circulatory overload (TACO), and transfusion-related acute lung injury (TRALI) (Chun et al. 2016). For instance, a MERS patient who received CP from a cured MERS-CoV donor displayed TRALI after transfusion (Chun et al. 2016). Adverse reactions can occur due to failure of the antibodies being neutralized in the host. Ultimately, this may result in entry of SARS-CoV-2 into the host cells and increase inflammation which can lead to severe pulmonary and/or CNS disease (de Alwis et al. 2020). Therefore, the use of CP should be cautiously approached. Currently, the FDA has not approved CP as a treatment method but does provide guidelines on CP use as an investigational product (FDA 2021e).

Stem cell-based therapy

Stem cell therapy has received attention as an alternative supportive treatment for patients infected with COVID-19. Early reports emerging from China have provided evidence that transplantation of allogeneic mesenchymal stem cells (MSCs) is associated with improved clinical outcomes of patients with varying stages of COVID-19-associated pneumonia (Liang et al. 2020; Leng et al. 2020). The use of MSCs as a therapeutic strategy for critical cases of COVID-19 has been the focus of a recent review which emphasizes their reparative functions (Atluri et al. 2020).

MSCs represent one of the most widely studied types of stem cells, and in recent decades, it has been well established that MSCs possess broad regenerative and restorative properties. These properties are believed to be modulated through the secretion of paracrine factors (i.e., growth factors, cytokines, extracellular vesicles) which regulate immunomodulatory and inflammatory processes in damaged or diseased tissues (Baraniak and McDevitt 2010; Phinney and Pittenger 2017; Samsonraj et al. 2017; Wang et al. 2016). Specifically, MSCs have demonstrated the ability to regulate proliferation, activation, and function of key immune cells (e.g., macrophages, B cells, effector/regulatory T-cells). Additionally, MSCs have been shown to exert anti-inflammatory effects through the secretion of various immunosuppressive molecules such as TGF- β , PGE2, IDO, and NO, as well as through the polarization of macrophages from an M1 to M2 phenotype which reduces the expression of pro-inflammatory cytokines (Gao et al. 2016; Harrell et al. 2019; Volarevic et al. 2017; Weiss and Dahlke 2019). Accordingly, numerous clinical trials have evaluated the therapeutic potential of MSCs in various diseases such as cardiac, bone, neurodegenerative, autoimmune, kidney, and lung (Marquez-Curtis et al. 2015; Trounson and McDonald 2015). Due to these holistic

effects, MSCs hold great promise for the future of tissue engineering and regenerative medicine.

With respect to COVID-19 infection, an overactivation of the immune system can induce a cytokine storm in severe and critical severe cases (Zhang et al. 2020). As discussed above, in the absence of therapeutic intervention, this cytokine storm can lead to significant pulmonary inflammation and extensive lung damage, ARDS, secondary infection, enhanced neurological symptoms, and sepsis that may ultimately result in death (Atluri et al. 2020). Therefore, preventing and/or reversing the cytokine storm may be paramount to treating critically ill COVID-19 patients (Atluri et al. 2020; Leng et al. 2020). Thus, due to their potent immunomodulatory and anti-inflammatory properties, MSCs may be well-suited for therapeutic intervention in these cases.

Throughout the last decade, MSCs have been utilized in a number of pre-clinical and clinical experiments relating to CNS and lung pathologies ranging from ARDS and pneumonia to Parkinson's disease and multiple sclerosis. Accordingly, there is accumulating evidence to support their potential therapeutic use in these conditions (Harrell et al. 2019; Wilson et al. 2018). For example, several *in vivo* studies have shown that MSCs can reduce lung injury and promote repair in animal models of ARDS and pneumonia. Here, some of the key efficacy outcomes were related to the modulation of inflammatory cytokine expression and neutrophil influx, regeneration of alveolar type II cells, regeneration of the alveolar-epithelial barrier, and reduction of cellular apoptosis (Asmussen et al. 2014; Curley et al. 2017; Gupta et al. 2007; Harrell et al. 2019; Lee et al. 2009; Kotani et al. 2017; Zhu et al. 2017). In the CNS, MSCs have shown tissue repair, modulation of inflamed tissue, and improved neurological functions in clinical and disease models (Branscome et al. 2020; Harris et al. 2012; Simorgh et al. 2019; Wilson et al. 2018). Specifically, in a mouse model of Parkinson's disease, MSCs were shown to improve behavior, increase dopamine transporters, among other benefits (Simorgh et al. 2019). Since ARDS, pneumonia, and neuronal damage can arise from COVID-19 infection, it stands to reason that COVID-19 patients may benefit from MSC therapy.

As mentioned above, two recent reports from China have already evaluated the therapeutic potential of clinical grade allogeneic MSCs against COVID-19-associated pneumonia. These studies showed improvements in vital signs, reduction of pneumonia, decrease in TNF, and an increase in IL10, VEGF, and IP10, as well as cellular indications of the reversal of the cytokine storm (Leng et al. 2020; Liang et al. 2020). Importantly, Leng et al. showed that transplanted MSCs were ACE2 and TMPRSS2 negative via RNA-seq analysis, confirming their natural immunity against SARS-CoV-2 (Leng et al. 2020). Collectively, these results suggest that MSCs have the potential to

reverse immune system over-activation to promote endogenous tissue repair. While these studies are promising, further research is warranted to validate these findings.

It is also crucial for future studies to better define the mechanisms of action through which MSCs act in damaged tissue. In recent years there has been a surge of interest in extracellular vesicles (EVs), especially exosomes, and it is now widely recognized that these vesicles are responsible for mediating many of the reparative and regenerative effects of MSCs and this topic has been extensively reviewed elsewhere (Joo et al. 2020; Marote et al. 2016; Mendt et al. 2019). Briefly, EVs are nano-sized vesicles released by virtually all cell types and, furthermore, are enriched with a rich assortment of bio-active cargo including RNAs (both small and long non-coding), cytokines, and proteins. These molecules can be horizontally transferred to recipient cells and, thus, have the ability to significantly affect cellular activity (Deng et al. 2018; Katsuda and Ochiya 2015). It is also worth noting that, from a therapeutic perspective, EVs offer certain advantages over stem cells due to their increased potency and stability, longer shelf life, and lower immunogenicity (Phinney and Pittenger 2017; Seo et al. 2019). In this sense, EVs represent a relatively novel paradigm for the field of regenerative medicine.

Similar to the previously discussed literature surrounding MSC-based therapy for lung and neurodegenerative pathologies, complementary studies involving the use of MSC EVs have revealed that these vesicles also possess the immunomodulatory, anti-inflammatory, antifibrotic, and antibacterial properties associated with their donor stem cells. A recent review nicely summarizes these experimental results in the context of asthma, ARDS, pneumonia, COPD, and idiopathic pulmonary fibrosis (IPF) (Cruz and Rocco 2017). In the context of the CNS, EVs provide additional benefits through their ability to cross the blood–brain barrier to deliver the therapeutic and regenerative qualities of MSCs (Branscome et al. 2020; Gorabi et al. 2019). It therefore stands to reason that an EV or exosome-based approach should also be considered for COVID-19 patients.

Due to the severity of the COVID-19 pandemic, it is imperative to investigate and exhaust all therapeutic options. Preliminary data strongly suggests that severely ill COVID-19 patients may benefit from MSC transplantation and regulators have been called upon to minimize the regulatory burdens governing access to its use (Atluri et al. 2020). Meanwhile, next-generation strategies employing the use of MSC EVs should also be rigorously pursued. Overall, the therapeutic methods offered and under research are vast but there is little information on if treatments help mitigate the neurological symptoms seen with COVID-19, especially for CCS individuals. This highlights an important area to study as we move forward.

Vaccines

The goal of a vaccine is to stimulate the immune system to produce antibodies and gain immunity against SARS-CoV-2 infection without contracting COVID-19. Currently, the S protein is the main candidate for vaccine development as the S protein mediates entry into host cells. As such, antibodies against the S protein can block virus-host cell interactions, ultimately neutralizing the virus (Li et al. 2005). The CDC explains that vaccine development follows the following prescribed steps: Exploratory stage, Pre-clinical stage, Clinical Development, Regulatory review and approval, Manufacturing, and Quality control (CDC 2020b). Due to the many stages that a new vaccine has to follow, WHO estimates that vaccine development can take as long as 10 years. However, the current COVID-19 crisis has substantially accelerated this process and it has taken less than a year to develop the current COVID-19 vaccines. To date, there are 3 COVID-19 vaccines with FDA EUA and several in clinical trials.

Two of the vaccines currently available in the USA are RNA vaccines. RNA vaccines are a relatively new type of vaccine that works through mRNA-encoded viral antigens, thus removing the genome integration step and providing an advantage over conventional vaccine approaches (Zhang et al. 2019). Furthermore, RNA vaccines can be developed faster and are more easily manufactured in large quantities than traditional vaccines (Amanat and Krammer 2020; Zhang et al. 2019). The two RNA vaccines with FDA EUA are mRNA-1273 (developed by the National Institute of Allergy and Infectious Diseases in collaboration with Moderna) and mRNA BNT 162b2 (developed by Pfizer) (Clinical Trials 2021). Both are double-dose vaccines that use an encoded mRNA SARS-CoV-2 S protein (Moderna 2020, Walsh et al. 2020). Once injected, the host immune cells produce and display the viral S protein which results in the activation of the host immune system and creation of antigens that target the S protein of the coronavirus and protect the host from future infection.

The third vaccine with FDA EUA is an adenoviral vector vaccine, which is one of the most common approaches in the development of vaccines, along with lentiviral vectors. These approaches have foreign genes which transduce the dividing cells, as well as the non-dividing cells. The lentiviral vectors target dendritic cells, as well as T-cells, to prompt an immune reaction (Pincha et al. 2010). The Janssen COVID-19 vaccine is a single-dose vaccine that encodes the SARS-CoV-2 S protein in an adenovirus vector to stimulate an immune response similar to that of the mRNA vaccines discussed above (FDA 2021a; Johnson and Johnson 2021).

There are a few other types of vaccines that are also undergoing clinical trials including recombinant subunit vaccines, DNA vaccines, live attenuated vaccines, and inactivated virus vaccines summarized by Amanat and Krammer

(2020). Lastly, there are some studies that have not yet been peer reviewed that show that vaccination has improved symptoms of brain fog and GI distress in cases of CCS, but as these are preliminary results, further research must be done (CDC 2021).

Conclusion

As discussed in this review, there have been great strides in SARS-CoV-2 research this last year. There are over 122,000 scientific papers in PubMed that discuss COVID-19, and publications increase daily along with our understanding of the disease. Here we try to summarize the major findings in SARS-CoV-2 biology, transmission, disease symptoms, diagnostic techniques, vaccines, and therapeutics.

SARS-CoV-2 is one of the seven known viruses within the Coronaviridae family that can infect humans. SARS-CoV-2 is a single-stranded positive sense RNA virus similar to other coronaviruses like SARS-CoV-1 and MERS-CoV. The main route of transmission of SARS-CoV-2 is through the respiratory tract via droplets, but other routes of infection may include pet, fomite, aerosol, sexual, and blood-borne transmission. Entry of SARS-CoV-2 into host cells is thought to be through the interaction between the viral spike protein and host receptors like ACE2, DPP4, CD147, and GRP78. Once in the host cell, SARS-CoV-2 replicates in the cytosol and infects the host. ACE2, proposed as (one of) the main receptor(s) for initial infection, is found in high concentrations in the lung alveolar epithelial cells and small intestine enterocytes which logically follows the respiratory and gastrointestinal symptoms seen in COVID-19 cases. Although mainly found in the above locations, ACE2 is also found throughout the body including nasal and CNS cells.

Once SARS-CoV-2 infects the body, the host can display a wide variety of symptoms that are associated with COVID-19. COVID-19 symptoms appear to worsen in older adults and individuals with comorbidities, but anybody can be infected. Symptoms usually occur within 5 to 6 days of infection (but may take up to 14 days) and impact many areas of the body. Typical symptoms include fever, cough, fatigue, sore throat, and shortness of breath. In more severe cases, pneumonia, lymphopenia, acute kidney injury, cardiac injury, lung damage, overactive immune system, ARDS, multi-organ failure, sepsis, and even death have been observed. CNS and PNS symptoms have also been observed including dizziness, ataxia, seizure, nerve pain, vision, taste, and smell impairments. The presence of SARS-CoV-2 has been shown to cause cytokine storms in the lower respiratory tract as well as a secondary cytokine storm within the brain which may result in additional lung damage as well as induce/increase neurological symptoms. In addition to acute COVID-19, there is now mounting evidence of chronic

COVID syndrome (CCS) which may impact individuals for months post exposure with symptoms such as fatigue, memory loss, attention deficit, brain fog, tachycardia, and shortness of breath.

The wide variety of symptoms associated with COVID-19 make diagnosis through symptoms alone ill-advised, and thus, there are now many tests to determine if an individual has COVID-19. The most common are NATs via oropharyngeal samples as they are currently the most sensitive diagnostic tests and provide the earliest detection. The CDC has advised the use of the viral N1, N2, genes as well as human RNase P gene for accurate results. These RT-qPCR tests can be done in as little as 3 h, but due to reagent and staffing shortages may take up to multiple days. Point-of-care test such as LAMPs, PCR kits, and immunoassays are also used as rapid tests, and can produce results in as little as 15–30 min. Other typical diagnostic methods have been unable to meet CDC standards for COVID-19 diagnosis but are used for other purposes. For example, serological tests and virus antigen tests are used to monitor antibody longevity and for tracking viral clearance in infected persons respectively, as well as provide information for epidemiological studies. On top of these tests, healthcare providers may use other ancillary tests, like CT scans, to provide insight for treatment plans for patients.

Most treatment approaches alleviate disease symptoms, and as the virus is still new with vast symptoms, there are a variety of therapeutics and ongoing clinical trials. Currently the FDA has approved remdesivir as well as given EUA to the use of the combination of remdesivir/baricitinib, and monoclonal antibodies bamlanivimab, bamlanivimab/etesevimab, and casirivimab/imdevimab. Other therapeutic methods in trial include the suppression of host receptors like ACE2 and GRP78. ACE2 levels are regulated through vitamin D and calcitriol and can be suppressed with typical ACE2 inhibitors like enalapril. TMPRSS2 is a co-protein utilized by SARS-CoV-2 alongside ACE2 in infection, and so drugs that target TMPRSS2 like camostat mesylate are also under study. Monoclonal antibodies similar to bamlanivimab, bamlanivimab/etesevimab, and casirivimab/imdevimab are undergoing clinical trials and are of interest as they can work both as treatment as well as preventive therapeutics through their ability to recognize SARS-CoV-2 proteins, but also host proteins that are associated with SARS-CoV-2 pathogenesis. Following this train of thought, some studies are looking at suppressing the immune response to prevent cytokine storms and the damage they cause. Anti-inflammatory baricitinib has been approved for emergency use in combination with anti-viral remdesivir. Other anti-inflammatory therapeutics are under study including some that have been approved for other diseases (i.e., tocilizumab) as well as new anti-inflammatory therapies (i.e., IVIG). Convalescent plasma from recovered

SARS-CoV-2-infected patients is also under investigation as it provides antibodies that can help neutralize the viral infection and lead to viral clearance. Finally, stem cell therapy could also aid in recovery as transplanted MSCs have been shown to improve patient outcomes.

To prevent disease transmission, there are currently 3 vaccines approved by the FDA for emergency use in the USA, produced by Johnson and Johnson, Pfizer, and Moderna. Both the Pfizer and Moderna vaccines require a double-dose injection of mRNA, while the Johnson and Johnson is a single-dose adenovirus vaccine, but all three target SARS-CoV-2 spike protein. Although these three vaccines are across headlines as they are approved, there are many other vaccines that are under trial including DNA vaccines, live attenuated vaccines, and inactivated vaccines.

Although there is still much to be studied about this virus, scientists have come a long way in a year. However, with new strains emerging and transmission rates rising, it is clear that the fight is not over, and that studies investigating all aspects of SARS-CoV-2 must continue.

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Declarations

Competing interests The authors declare no competing interests.

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