REVIEW

www.neurosci.cn <www.springer.com/12264>

Brain Pathology in COVID‑19: Clinical Manifestations and Potential Mechanisms

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Received: 12 January 2023 / Accepted: 25 May 2023 / Published online: 16 September 2023 © Center for Excellence in Brain Science and Intelligence Technology, Chinese Academy of Sciences 2023

Abstract Neurological manifestations of coronavirus disease 2019 (COVID-19) are less noticeable than the respiratory symptoms, but they may be associated with disability and mortality in COVID-19. Even though Omicron caused less severe disease than Delta, the incidence of neurological manifestations is similar. More than 30% of patients experienced "brain fog", delirium, stroke, and cognitive impairment, and over half of these patients presented abnormal neuroimaging outcomes. In this review, we summarize current advances in the clinical fndings of neurological manifestations in COVID-19 patients and compare them with those in patients with infuenza infection. We also illustrate

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the structure and cellular invasion mechanisms of SARS-CoV-2 and describe the pathway for central SARS-CoV-2 invasion. In addition, we discuss direct damage and other pathological conditions caused by SARS-CoV-2, such as an aberrant interferon response, cytokine storm, lymphopenia, and hypercoagulation, to provide treatment ideas. This review may offer new insights into preventing or treating brain damage in COVID-19.

Keywords COVID-19 · SARS-CoV-2 · Brain · Neurological pathology

Introduction

As of Jan 5, 2023, the global pandemic coronavirus disease 2019 (COVID-19) caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) has resulted in over 600 million confrmed cases and 6 million deaths ([https://covid](https://covid19.who.int/) [19.who.int/\)](https://covid19.who.int/). The majority of COVID-19 patients initially suffer respiratory symptoms, which can progress to severe substantial pulmonary diseases, such as pneumonia or acute respiratory distress syndrome (ARDS) [[1\]](#page-11-0). With increasing clinical evidence, researchers have discovered that SARS-CoV-2 causes neurological manifestations in \sim 30% of patients who recovered from COVID-19. The most common neurological manifestations include cognitive impairment, depression, and psychosis [\[2](#page-12-0), [3](#page-12-1)]. Further investigations have revealed brain pathology in COVID-19 patients with neurological manifestations [[4\]](#page-12-2). Although Omicron causes less severe disease than Delta, it still leads to brain pathology in many patients. However, the mechanisms of brain disease in COVID-19 continue to be a mystery because of the complexity of brain structure and function, the uncertain course of central SARS-CoV-2 invasion, and the diverse responses to the virus. In this review, we summarize the clinical manifestations and brain pathological features of COVID-19 and discuss the mechanisms of brain damage caused by SARS-CoV-2.

Clinical Neurological Manifestations in COVID‑19 Patients

Neurological manifestations have been reported in COVID-19 patients of all ages. For instance, in a study conducted in Wuhan, China, more than a third of COVID-19 patients developed new neurological symptoms after being infected, as shown in Table [1](#page-2-0) [\[5](#page-12-3)]. Another retrospective cohort study that included 236,379 patients also found a similar rate (33.62%) of patients with neurological manifestations [\[6](#page-12-4)]. During the acute phase of infection, patients most frequently experience headache, anosmia, and stroke [[7\]](#page-12-5), while "brain fog", headache, anxiety, anosmia, ageusia, and cognitive impairment are common post-acute neurological sequelae of SARS-CoV-2 infection [\[8](#page-12-6)]. Many patients had anosmia, and further investigations showed that their nasal components [[9\]](#page-12-7) and olfactory gyrus [\[10](#page-12-8)] had been damaged. A recent study suggested that sensory and neurological disorders are more evident in younger patients, whereas mental health, musculoskeletal, and episodic disorders are more noticeable in older patients [[11\]](#page-12-9). While most of these symptoms gradually disappear within a few months after COVID-19 recovery, more research is required to confrm the duration of these symptoms and subsequent changes.

Compared to the patients infected by infuenza, COVID-19 patients with neurological manifestations are older and exhibit a higher incidence of altered mental status, headache, anosmia, dysgeusia, and ischemic stroke [[12,](#page-12-10) [13](#page-12-11)]. In addition, COVID-19 patients develop symptoms more quickly. As shown in Tables [3](#page-4-0) and [5,](#page-6-0) fu patients may sufer from influenza-associated encephalitis/encephalopathy (IAE) with symptoms such as altered consciousness and seizures. Flu patients may also experience post-infuenza encephalitis, Guillain-Barre syndrome (GBS), Reye's syndrome, and Parkinsonian symptoms (PD) [\[14\]](#page-12-12).

Neuroimaging Findings of COVID‑19 Patients

Neuroimaging applies quantitative techniques to visualize the brain's structure and function. As listed in Table [2](#page-3-0), medical imaging examinations, particularly magnetic resonance imaging (MRI) and positron emission computed tomography (PET) reveal the brain structure and function of patients with neurological manifestations [\[15\]](#page-12-13). The most frequent imaging features (afecting >60% of patients) include ischemic infarcts, intracerebral hemorrhages, perfusion abnormalities,

and leptomeningeal enhancement. Hypometabolism in the pons, cerebellum, bilateral insula, bilateral medial lobes, and prefrontal cortex indicates brain function dysregulation in COVID patients. The high incidence of cerebrovascular events such as ischemic stroke and intracerebral hemorrhage, indicates potential endothelial injury and coagulation dysfunction. Other investigations have also reported hypoxic alterations in the cerebellum and cerebrum, metabolic alterations of astrocytes, microglial activation [[16\]](#page-12-14), neuro-axonal damage, and neuronal loss [[17](#page-12-15)], which provide additional evidence for brain damage in COVID patients.

As for fu patients, the most prevalent imaging features include brain lesions, edema, rapid and fulminant demyelination, and infammation. These abnormalities are mainly in the cortex, white matter, or brainstem (Tables [4](#page-5-0) and [5\)](#page-6-0). In addition, glial activation, metabolic disorders, and genetic factors may be involved [[18,](#page-12-16) [19\]](#page-12-17).

Variant and Neurological Complications in Patients with COVID‑19

After the appearance of the Alpha variant, there was no discernible diference in the prevalence of neurological manifestations in patients with COVID. However, the risk of anxiety disorders, insomnia, cognitive defcit, and ischemic stroke were signifcantly higher after the appearance of the Delta variant. Patients diagnosed with COVID-19 after the emergence of the Omicron variant have a lower death rate, but their neurological outcomes still carry a similar risk to those with the Alpha variant [[20](#page-12-18)]. Therefore, despite the reduced rate of severe cases of Omicron, we cannot underestimate the high incidence of neurological sequelae. It is important to note that this aspect has not been adequately studied because of the difficulty of differentiating the variants of viruses that infect patients in those published clinical studies.

Patients' Characteristics and Brain Pathology in COVID‑19

Cohort studies on the neurological manifestations of COVID-19 have shown that older age and pre-existing neurological disorders are risk factors for neuropathy and mortality $[21]$ $[21]$. It is known that aging results in immunosenescence, which is a progressive weakening of the ability to mount efficient immune responses against infections [[22\]](#page-12-20). In addition, older people are more likely to have preexisting neurodegenerative diseases, including Alzheimer's disease (AD) and PD [\[23](#page-12-21)]. Previous studies have suggested that patients with AD are vulnerable to experiencing severe conditions and passing away during COVID-19 [[24](#page-12-22)]. It is

explicit inclusion criteria and an experimental process to ensure the validity of the results.

Search strategy: search "(brain) AND (infuenza)" in PUBMED and flter articles with specifc neurological manifestations. Notably, the study should have explicit inclusion criteria and an experimental process to ensure the validity of the results.

worth noticing that the apolipoprotein E4 (APOE4) allele, which is strongly associated with AD, is also a risk factor for severe COVID-19 [[25](#page-12-30)]. Neuropathy in COVID-19 may be exacerbated by APOE4 because it can increase fbrinogenesis, blood-brain barrier (BBB) permeability, and cerebral amyloid angiopathy in the brain [[26\]](#page-12-31). In contrast to female patients, aged male patients have fewer activated and diferentiated T cells, as well as poorer $CD8⁺ T$ cell activation and IFNγ production [\[27](#page-12-32)]. However, the incidence of COVID-19 sequelae does not difer signifcantly by gender or exhibits a modest overrepresentation of female patients [[28\]](#page-12-33). Patients with pre-existing diseases, such as hypertension, diabetes, and cardiac diseases, are at a higher risk of infection and brain pathology [\[29](#page-12-34)]. Pre-existing pathological conditions,

such as arteriosclerosis, contribute to acute ischemia, stroke, dementia, and other neurological manifestations [\[30](#page-12-35)]. Moreover, drugs used to treat pre-existing diseases must also be considered for their potential impact on the incidence and severity of infection. One study has suggested that patients treated with drugs that increase angiotensin-converting enzyme 2 (ACE2) expression may be more vulnerable to severe COVID-19 infection [\[31](#page-12-36)]. Therefore, patient characteristics, including age, pre-existing conditions, and medication use, should be carefully evaluated when making clinical decisions.

Table 4 Neuroimaging fndings in infuenza patients

Search strategy: search "(brain) AND (COVID-19) AND ((MRI) OR (EEG))" in PUBMED and flter articles with detailed and concrete outcomes. Notably, the study should have explicit inclusion criteria and an experimental process to ensure the validity of the results

Evidence for Central Invasion of SARS‑CoV‑2

In several post-mortem studies, small amounts of SARS-CoV-2 have been detected in the brains of patients, but in almost all cases [[60](#page-13-13)], the patients had at least one site in the brain with low but positive amounts of SARS-CoV-2 RNA, with the cerebellum being the most frequently affected [[61](#page-13-14)]. The astrocyte has been identified to be the preferred target for SARS-CoV-2 infection and replication. However, SARS-CoV-2 is difficult to detect in microglia, although the activation of microglia is prominent in the brain of severe COVID patients and is similar to the state of human neurodegenerative disease [[62](#page-13-15)]. In addition, SARS-CoV-2 has also been detected in brain capillary endothelial cells [[63\]](#page-13-16). Notably, SARS-CoV-2 has been found to infect and kill neurons in human brain organoids [[64\]](#page-13-17). A recent study identifed virus-specifc protein expression in the hypothalamus and spinal ganglia neurons by immunofuorescence. Furthermore, this study demonstrated that SARS-CoV-2 RNA persists in the brain for months, even after the virus has been eliminated from the plasma [\[65\]](#page-13-18). However, additional research is required to confrm the virus's capacity for replication in the brain. It should be emphasized that the existing studies are insufficient to demonstrate direct viral invasion of neurons or detect low virus levels in the brain.

Table 5 COVID-19 versus infuenza virus

Currently, no conclusive information about the brain invasion of SARS-CoV-2 is available in human post-mortem studies. Besides, unlike animal studies with a good perfusion process before sampling, the SARS-CoV-2 detected in the human brain might originate in the blood. Thus, it is more likely that the virus indirectly causes damage to the brain by triggering a series of responses. The importance (even the existence) of direct viral brain invasion of SARS-CoV-2 still needs further study.

Virus Structure and Cellular Invasion Mechanisms

SARS-CoV-2, known as the main culprit of this global pandemic, possesses single-stranded positive-sense RNA (+ssRNA), four structural proteins (N, E, M, and S), and sixteen non-structural proteins (nsp1−16) [\[66](#page-13-24), [67\]](#page-13-25). The spike (S) protein, which consists of S1 and S2 subunits, plays a prominent role in viral invasion and is regarded as a crucial target for antiviral treatment [\[68](#page-13-26), [69](#page-13-27)] (Fig. [1\)](#page-8-0). In addition, the nucleocapsid (N), envelope (E), and membrane (M) proteins are all involved in viral production and combine with host cell organelles to cause various dysregulation of physiological processes [[70,](#page-13-28) [71](#page-13-29)]. ACE2 is a signifcant receptor for SARS-CoV-2 entering host cells [\[72](#page-13-30)]. After binding to ACE2 *via* the S1 unit, SARS-CoV-2 can initiate membrane fusion depending on a furin-like cleavage site on the S1/S2 and S2 units [\[73](#page-13-31)]. The S2 unit is cleaved by the endosomal proteases cathepsin B and L (CatB/L) [\[74](#page-14-0)], or the transmembrane protease serine 2 (TMPRSS2) [[75\]](#page-14-1), to form a fusion pore, which allows the viral genome to enter cells [\[76](#page-14-2)]. The replication and transcription complex discontinuously processes the viral RNA, mainly formed by nsp12 cooperating with nsp7 and nsp8 [[77\]](#page-14-3). Viral transcription, replication, and translation depend on host cell organelles, including the endoplasmic reticulum and Golgi [\[78](#page-14-4)]. It is worth noting that the S protein of Omicron has a higher ACE2 binding affinity than the S protein of other variants, which might explain the higher infection of Omicron [\[79](#page-14-5)]. In addition, since different receptors are highly expressed on diferent types of cells, it is important to consider the impact of additional receptors, such as neuropilin-1, in bringing about pathological changes.

As SARS-CoV-2 has been shown to cause damage and dysregulation in other systems [[80](#page-14-6)], its detection in the brain by polymerase chain reaction [[81](#page-14-7)] and immunohistochemistry [\[82](#page-14-8)] suggests that it may also harm brain cells and tissues. Typically, the virus causes damage through two mechanisms: (1) replication in cells to produce daughter viruses and breaking the cell to release more viruses to invade other cells [\[83](#page-14-9)]; (2) the body's response against the virus inevitably results in cell destruction. Thus, SARS-CoV-2 might also enter the brain and directly or indirectly damage brain cells. In direct damage, while the daughter virus is released by secretory vesicles and causes light damage to cell structure, interference by viral protein in the cell cycle can result in severe cell damage and death [\[84\]](#page-14-10). The nonstructural proteins (nsps) of SARS-CoV-2 disturb host DNA replication, protein synthesis, and transport [[85](#page-14-11)], leading to some pathological brain changes. Besides, the virus can directly kill cells by inducing apoptosis, necroptosis, and autophagy in infected cells [[86\]](#page-14-12).

Possible Brain Entry Mechanisms of SARS‑CoV‑2

Via **the BBB**

The BBB typically serves as a barrier and a transporter. It transports ions, macromolecules, nutrients, and toxins into or out of the brain [\[87](#page-14-13)]. The BBB is mostly composed of vascular endothelial cells, pericytes, and astrocytes. The tight junctions (Tjs) and adhesion junctions between endothelial cells act as signifcant parts of the barrier [[88\]](#page-14-14). Pericytes, embedded in the basement membrane of microvessels, regulate BBB formation and permeability and serve in CNS immune surveillance, such as the leukocyte migration across endothelial cells [\[89](#page-14-15)]. Astrocytes exhibit a strong link with BBB formation and dominantly enhance the frequency, length, and complexity of Tjs [[90,](#page-14-16) [91\]](#page-14-17).

As remarked above, ACE2 is the main receptor of SARS-CoV-2 and has been detected in several tissues, especially in arterial and venous endothelial cells [[92](#page-14-18)]. In the human brain, the substantial nigra, paraventricular nuclei of the thalamus, raphe nuclei, tuberomammillary nucleus, central glial substance (an area of gray matter surrounding the central canal), and choroid plexus (ChP) of the lateral ventricle may have high ACE2 expression [\[93](#page-14-19)]. SARS-CoV-2 might infect brain microvascular endothelial cells through ACE2 and replicate in them [[94\]](#page-14-20). However, entry through endothelial cells may be limited, because numerous studies have not yet demonstrated that infection and replication can occur there effectively. Besides, the basement membrane disruption of the BBB induced by SARS-CoV-2 through matrix metalloproteinase-9 is considered a signifcant factor for virus entry [\[95](#page-14-21)]. Indeed, BBB disruption and leakage have also been reported in COVID-19 patients with neurological manifestations [[96\]](#page-14-22).

The SRAS-CoV-2 may also use host cells to convey traffic across the BBB $[97]$ $[97]$ $[97]$. The virus is believed to preferentially invade lipid-secreting cells and use lipid metabolism for replication and spread [[98](#page-14-24)]. A study has also shown replication of SRAS-CoV-2 in infected monocytes and macrophages [\[99\]](#page-14-25). Since the immune cells can enter the brain through the BBB and the ChP, SARS-CoV-2 may get a ride and get out after brain entry [[100](#page-14-26), [101\]](#page-14-27). However, it is not yet known whether viral replication in monocytes produces infectious viruses [[102\]](#page-14-28). Further research is needed to investigate this mechanism

Via **the CSF**

As some cases have reported the presence of SARS-CoV-2 in cerebrospinal fuid (CSF), it is worth discussing the possibility of the CSF route [[103\]](#page-14-29). The blood-cerebrospinal fuid barrier (BCSFB) is one of the selectively permeable barriers which regulates the transportation of substances between the blood and CSF. The BCSFB is formed by epithelial cells of the ChP, which separate the fenestrated capillaries from the CSF [[104](#page-14-30)] (Fig. [2](#page-9-0)A and [B](#page-9-0)). Similar to the BBB, The Tjs between ChP endothelial cells function as the barrier [[105](#page-14-31)]. The ChP also secretes pro-infammatory cytokines and provides space for transporting immune cells into the stroma [[106\]](#page-14-32). ACE2 and its co-receptors TMPRSS2 and TMPRSS4 are highly expressed in the ChP, through which SARS-CoV-2 destroys the structure and function of the BCSFB in organoids [\[107](#page-14-33), [108](#page-14-34)]. The circumventricular organs (CVOs) are located in the third and fourth ventricles and have continually fenestrated and highly permeable vessels. ACE2 has been detected at a high level in CVOs, which provide a facility for virus entry [[109](#page-14-35), [110\]](#page-14-36). However, due to the limited detection of SARS-CoV-2 in CSF, the efficiency of virus entry through this route must be further studied.

Via **Retrograde Nerve Transmission**

Olfactory dysfunction has been frequently reported in COVID-19 patients [\[111\]](#page-14-37) and may be correlated with the invasion of olfactory cells by viruses (leading to the downregulation of odor detection pathways) [[112](#page-15-0)]. Olfactory mucosa is in contact with droplets by ciliated cells and perceives them through the olfactory sensory nerves (OSNs) in the basal cells, which are supported and nourished by sustentacular cells. Signals are then transferred to the olfactory bulb (OB) where they reach the mitral cells. The virus might invade the peripheral nerves, move retrogradely along them, and fnally enter the CNS [[113\]](#page-15-1). Given that many viruses enter the brain in this way, it might also be a possible route for SARS-CoV-2 to invade the brain [\[114](#page-15-2)] (Fig. [2C](#page-9-0)). Meinhardt *et al.* analyzed regional mapping of the consecutive olfactory nervous tracts and defned CNS regions in 33 individuals, suggesting that neuroinvasion can occur by transmucosal entry *via* regional nervous structures [[115](#page-15-3)]. Besides, the invasion of OSNs by SARS-CoV and SARS-CoV-2 has also been revealed in hamsters [[116\]](#page-15-4). However, many other studies conversely showed a low possibility of the OB route in humans. Based on 88 post-mortem cases of COVID-19 patients, Khan *et al.* found that SARS-CoV-2 infects ciliated cells, sustentacular cells, and even leptomeningeal layers surrounding the OB, while the OSNs and OB are uninfected [\[117](#page-15-5)]. Another post-mortem research also obtained the same result and suggested that infammation might cause axon and vascular injury [\[118\]](#page-15-6). Butowt *et al.* pointed out that a lack of quantitative analysis and false positives might explain the contradictory reports on OSN infection [[119\]](#page-15-7). In addition to the OSNs, Bowman's gland, a branching tubular gland in the olfactory mucosa, has been found to show both CatB/L and furin expression [[120\]](#page-15-8). Thus, the nerve terminals of neurons

Fig. 1 The structure of SARS-CoV-2. SARS-CoV-2 consists of 4 structural proteins (S, M, E, and N) and 16 non-structural proteins (nsp1−16). After the S1 component binds to ACE2, the S1/S2 cleavage primes the fusion and creates extra cell surface receptors, like NRP1. Then, SARS-CoV-2 starts entry by CatB/L and TMPRSS2. The viral RNA is released after the entry and translated into corresponding viral structures. The open reading frame 1a (ORF1a) and

might receive SARS-CoV-2 from the infected Bowman's gland. Overall, retrograde nerve transmission at least is a theoretically possible means of SARS-CoV-2 invasion but needs further determination. Besides, retrograde nerve transmission is also being considered in the infection of the eye, intestine, and lung through the vagus nerve [[121](#page-15-9)] or optic nerve [\[122](#page-15-10)].

Other Mechanisms for Brain Pathology in COVID‑19

Apart from the direct damage caused by the cellular invasion, several pathological conditions, such as an aberrant IFN response, cytokine storm, lymphopenia, and hypercoagulation in the brain, may result in more severe and widespread damage to the brain. In this section, we elucidate the periphery-related potential mechanisms of central pathological conditions (Fig. [3\)](#page-10-0).

Aberrant IFN Response

The innate immune system plays a signifcant role in defending against infection and determining the disease severity [[123](#page-15-11)]. Dysregulation of type I and II IFN is the main

ORF1b of RNA are translated into polyprotein 1a (pp1a) and pp1ab, further being translated into nsp1−11 and nsp12−16. Host cell organelles, such as the Golgi, are involved in virus replication. SARS-CoV-2: severe acute respiratory syndrome coronavirus 2; ACE2: angiotensin-converting enzyme 2; NRP-1: neuropilin-1; TMPRSS2: transmembrane protease serine 2.

cause of the insufficient innate immune response. In the early stages of infection, a large amount of type I IFN may be present in the patient's body to fght the virus, and the ChP is the most vulnerable area to infection in the brain. A study by Suzzi *et al.* has demonstrated that the increased IFN response may disrupt the ChP and be involved in the process of brain damage [\[124](#page-15-12)]. After a peak of type I IFN in the early stage of infection, a delayed aberrant IFN response has been reported in most severe cases [[125\]](#page-15-13). For example, in the later period of COVID-19, a decrease of IFN occurs in the CSF of severe patients with neurological manifestations, and this may be related to the defcient innate immune response [[126](#page-15-14)]. SARS-CoV-2 interferes the IFN production and activation in two ways: (1) the nsp1 of SARS-CoV-2 inhibits the gene expression and translation of IFN, and (2) pattern-recognition receptor (PRR)-mediated signaling pathways are suppressed to downregulate the IFN signaling. PRRs, including membrane-bound C-type lectin receptors, NOD-like receptors (NLRs), retinoic acid-inducible gene I (RIG-I)-like receptors, and Toll-like receptors (TLRs), are commonly expressed on sensory neurons, astrocytes, and microglia [\[127,](#page-15-15) [128\]](#page-15-16). SARS-CoV-2 RNA and M proteins can bind to RIG-I and melanoma diferentiation-associated gene 5, blocking the activation of interferon regulatory factor 3 and the type I IFN signaling [[129](#page-15-17)]. In addition, it has

been demonstrated that several SARS-CoV-2 components, including nsp6, nsp13, and open reading frame 3a, inhibit the type I IFN response *in vitro* [[130](#page-15-18)]. As the balance of the innate immune response is closely related to the disease severity of COVID-19 patients with neurological manifestations, the potential strategy to rectify the IFN response is worth discussing.

Cytokine Storm

The cytokine storm is a dangerous sharp increase of cytokines in body fuid that is also a concern in COVID-19 patients, especially in severe cases. As infammation has been considered to be involved in several CNS diseases such as AD and PD, the elevated cytokines induced by SARS-CoV-2, including IL-6, TNF- α , IL-1, and IL-10 [\[131](#page-15-19)], might also be related to the pathological brain changes in COVID-19. It is worth mentioning that the increase of cytokines not only occurs in the acute phase but also may persist in patients for 3 months after recovery from COVID-19. Lowgrade chronic infammation may be associated with brain injury and vascular injury-related stroke in patients [[132](#page-15-20)].

The generation of inflammation in the brain may be closely relevant to the brain barriers. SRAS-CoV-2 can infect components of barriers such as endothelial cells and astrocytes to process pro-inflammatory cytokines [[133\]](#page-15-21) and infuence BBB permeability [[134](#page-15-22)]. The neuroflament light chain protein and glial fbrillary acidic protein are at high levels in both the plasma serum and CSF of COVID-19 patients, indicating that astrocytic and neuronal injury indeed occurs in these patients [[135](#page-15-23)]. Notably, C-C motif chemokine ligand 11, which can specifcally cause hippocampal microglial reactivity and impaired neurogenesis, has been found to be elevated in CSF only in COVID-19 patients with cognitive syndrome [\[136\]](#page-15-24). In addition to elevated CSF cytokines [\[137\]](#page-15-25), the choroid-to-cortex network across infammatory pathways also increases in COVID-19 patients. This indicates that the infammatory signals are sent from the choroid plexus into the brain more efectively to activate the glia. Microglia, the innate immune cells in the CNS, respond to the virus and form clusters with infltrating T cells, which are associated with the brain and perivascular infammation [\[138](#page-15-26)]. Elevated complements like C1q and C3, combined with microglia, are linked to the synaptic loss and toxicity of soluble β-amyloid $(Aβ)$ oligomers [[139](#page-15-27)]. Moreover, the mast cells, as innate immune cells participating in the adaptive immune response, can be activated by SARS-CoV-2 and release pro-infammatory cytokines, including IL-1β, C-C motif chemokine ligand 2, IL-6, granulocytemacrophage colony-stimulating factor, and TNF- α , which makes them signifcant components in neuroinfammation [[140,](#page-15-28) [141\]](#page-15-29).

SARS-CoV-2 may bind to receptors on host cells, including ACE2 and PRRs, to promote infammation. ACE2 plays a key part in the renin-angiotensin system (RAS) [[142\]](#page-15-30). As SARS-CoV-2 interacts with ACE2 and infuences its catalytic activity, it can further regulate the RAS and contribute to the cytokine storm [[143\]](#page-15-31). Besides, the reduction of ACE2 leads to an increase of serum Ang II, which activates the production of reactive oxygen species (ROS) and nuclear

Fig. 2 The CSF and retrograde nerve transmission pathways for virus entry. **A**, **B** The CSF pathway. SARS-CoV-2 traverses the permeable vessels of the choroid plexus (ChP) to reach the stroma. In the stroma, it binds to ACE2 and TMPRSS2/4 to enter endothelial cells and then be released into the CSF. **C** The retrograde nerve transmission pathway. The virus invades the olfactory sensory nerves (OSNs), moves retrogradely along nerves, and fnally enters the neurons of the CNS. Besides, the virus can invade nervous terminal neurons that contact Bowman's gland through cathepsin B/L (CatB/L). Tj, tight junction; ACE2, angiotensin-converting enzyme 2; TMPRSS2, transmembrane protease serine 2; OB, olfactory bulb.

factor-kappa B pathways. These activations accordingly promote the expression of pro-infammatory cytokines and inactive NOD-like receptor family pyrin domain containing 3 (NLRP3) [[144,](#page-15-32) [145\]](#page-15-33). Excessive ROS can trigger apoptosis through the mitochondria, which causes morphological and functional changes in the host cell [[146](#page-15-34)]. ROS also leads to the overactivation of the immune response and other patho-logical reactions [[147](#page-16-0)]. NLRP3 inflammasomes, activated by ROS production and mitochondrial dysfunction, show a high correlation with apoptosis, dysregulation of tau protein phosphorylation, and neurofbrillary tangles [\[148\]](#page-16-1). Uncontrolled activation of NLRP3 has also been found in endothelial progenitor cells of COVID-19 patients [\[149,](#page-16-2) [150\]](#page-16-3).

As discussed above, PRRs, including TLRs, NLRs, and RIG-I, participate in the signaling pathways of IFN. PRRs in the CNS, especially TLRs, are also involved in the production and release of several cytokines including IL-1, IL-6, and TNF- α [[151\]](#page-16-4). The S protein of SARS-CoV-2 can bind to TLR2 and TLR4 and activate the myeloid diferentiation factor 88 signaling pathway to promote the cytokine storm [\[152](#page-16-5)]. Besides, TLR4-mediated inflammatory cytokines are reported to be upregulated in peripheral blood mononuclear cells and CSF [[153\]](#page-16-6).

Lymphopenia

Lymphopenia has been found in severe COVID-19 patients, and several factors may account for this phenomenon [\[154,](#page-16-7) [155](#page-16-8)]. Firstly, ACE2 has remarkable expression in lymphocyte cells, through which the SARS-CoV-2 can directly destroy them. SRAS-CoV-2 RNA has also been detected in

Fig. 3 Pathological conditions in the brain. SARS-CoV-2 may cause several pathological conditions, including an aberrant IFN response, cytokine storm, lymphopenia, and hypercoagulation, resulting in pathological brain changes. NETs, neutrophil extracellular traps.

macrophages, neutrophils, B cells, T cells, and natural killer cells [\[156](#page-16-9)]. Secondly, the elevated cytokines, especially TNF and IFNγ, can contribute to apoptosis and pyroptosis, which leads to the death of lymphocytes and atrophy of the spleen [\[157,](#page-16-10) [158](#page-16-11)]. The systemic reduction of lymphocytes also results in lymphopenia in the brain [[60\]](#page-13-13), which directly decreases the body's defense against viruses and leads to pathological changes. Several studies have suggested that changes in immune cells might induce neural and neuroendocrine dysregulation, such as anxiety [[159](#page-16-12)], depression $[160]$ $[160]$ $[160]$, and neurodegenerative diseases $[161]$ $[161]$. Meanwhile, these psychiatric sequelae have been reported in COVID-19 patients with an aberrant immune system and include significant post-traumatic stress disorder [[162\]](#page-16-15). In addition, lymphopenia is also considered a hallmark of poor prognosis in COVID-19 patients, further highlighting the importance of lymphopenia [\[163](#page-16-16), [164\]](#page-16-17).

Hypercoagulation

Hypercoagulation-related neurological manifestations, such as ischemic stroke, have been reported in COVID-19 patients [[165](#page-16-18)]. In addition, risk factors of coagulation are also considerably elevated in COVID-19 patients, including D-dimer, von Willebrand Factor (vWF), vWF antigen, and FVIII [[1](#page-11-0), [166\]](#page-16-19). Both white and red thrombi have been observed in microvascular, macrovascular, and venous sys-tems [[167](#page-16-20)]. Affected individuals may experience neuropsychiatric diseases, neurocognitive (dementia-like) syndrome, and altered mental status as a result of the thrombi in the cerebrovascular, system; they obstruct the bloodstream and kill endothelial cells [\[21](#page-12-19)]. The term "coagulation cascade" refers to the sequence of events that leads to the formation of clots during the coagulation process.

The coagulation process includes the extrinsic, intrinsic, and complement pathways. (1) The extrinsic pathway starts with tissue factor, which is expressed on neutrophils and monocytes and can be activated by cytokines and neutrophil extracellular traps (NETs) [[168](#page-16-21)]. NETs consist of cytosolic and granule proteins and mitochondrial DNA. Triggered by ROS against infection, NETs regulate the activation of the extrinsic pathway [[169\]](#page-16-22). An elevated number of neutrophils and myeloperoxidase /DNA complexes, a welldefned marker of NETosis, indicates a high incidence of NET formation in COVID-19 patients [[170\]](#page-16-23). (2) The intrinsic pathway begins with factor 12 and ultimately participates in fbrin formation. The intrinsic pathway is triggered by an endothelial injury, which can be caused by infection with SRAS-CoV-2. Factor 8 and vWF, a carrier of factor 8, and fbrinogen are highly expressed in COVID-19 patients [[171](#page-16-24)]. In addition, patients with COVID-19 have remarkably elevated levels of the plasminogen activator inhibitor-1 (PAI-1), which downregulates the plasminogen activation to plasmin by inhibiting tissue-type plasminogen activator (tPA). Contradictory opinions have been generated by the fact that some other COVID-19 patients have elevated tPA. That is, the PAI-1 could counteract the local tPA efect to create a net prothrombotic hypofbrinolytic state, while the fbrinolysis led by tPA is predominant [[172,](#page-16-25) [173](#page-16-26)]. Furthermore, despite the decrease or maintenance of platelet numbers in COVID-19 patients, the plasma thrombopoietin levels, platelet surface P-selectin expression (a marker of platelet activation), and circulating platelet-leukocyte aggregations are prominently high. These results may partly apply to the hypercoagulation state in COVID-19 patients and also suggest that the JAK3-MAPK pathway is involved in the process [[174](#page-16-27), [175](#page-16-28)]. The platelet-to-lymphocyte ratio level is signifcantly higher in severe patients than in non-severe patients with COVID-19 and shows a correlation with the cytokine storm [[176\]](#page-16-29). The D-dimer, a protein produced when clots dissolve, is elevated in COVID-19 patients, which also suggests a serious hypercoagulation syndrome [\[177\]](#page-16-30). (3) The complement pathway is a cascade of events that leads to hemostasis. It has been found that the N protein of SARS-CoV-2 can cause abnormal complement activation [\[178\]](#page-16-31). Indeed, complements such as C5, C6, C5a, and C8 also increase in severe COVID-19 patients [[179](#page-16-32)]. The complement dysregulation might impel thrombosis and endothelial injury, ultimately leading to brain damage. In summary, the "coagulation cascade" might be started by SARS-CoV-2 and result in a pathological hypercoagulation state, which induces endothelial injury and subsequently causes pathological brain changes.

Other Potential Mechanisms

Several studies have detected anti-neuronal and anti-glial autoantibodies in the serum or CSF of COVID-19 patients with neurological symptoms, suggesting that autoantibodies may be involved in the brain injury of COVID-19 [\[180,](#page-16-33) [181](#page-16-34)]. Autoantibodies may directly damage neurons and can also lead to immune abnormalities. In addition, Franke *et al.* found that autoantibodies have undetermined antigenic epitopes, which may be related to the molecular mimicry of the SARS-CoV-2 virus $[182]$ $[182]$.

Hypoxia due to pulmonary fbrosis can cause multi-organ damage, which is more prominent in patients with severe disease [\[183\]](#page-17-1). Hypoxia in COVID-19 may be correlated with the expression of ACE2 and TMPRSS2 as well as $Aβ$ deposits in the neocortex [[184,](#page-17-2) [185](#page-17-3)]. Zilberman-Itskovich *et al.* found a lower incidence of neurological manifestations in COVID-19 patients subjected to hyperbaric oxygen therapy, suggesting that improving hypoxia may reduce neurological pathology by improving brain perfusion and neuroplasticity [[186\]](#page-17-4).

Conclusion

COVID-19 is still a problem that afects the entire world and interferes with people's daily lives. Approximately 30% of COVID-19 patients experience new neurological manifestations or exhibit pathological brain changes. However, the current treatments have hitherto been insufficient. In this review, we emphasize the need for clinical practice to consider patient features, such as pre-existing diseases, when treating COVID-19 patients with neurological symptoms. As for the mechanisms, in this review, we indicate that SARS-CoV-2 can infect host cells through ACE2 (accompanied by TMPRSS2 or CatB/L) and may further enter the brain through the BBB, CSF, and retrograde nerve transmission. SARS-CoV-2 may cause cell and tissue damage once it has entered the brain by preventing replication and promoting apoptosis. Apart from these direct efects, several systemic or peripheral pathological conditions, including an aberrant IFN response, cytokine storm, and hypercoagulation, may also contribute to brain pathology. The aberrant IFN response can result in a delayed innate immune response, rendering the viral defense inefective. The cytokine storm produces excessive tissue damage and cell death, even in the brain, despite its original function as a defense mechanism against the virus. Besides, severe COVID-19 patients exhibit remarkable lymphopenia, indicating a high risk of neurological pathology. Hypercoagulation may be related to endothelial injuries, infarcts, and hemorrhages in the brain. Thus, in addition to host receptors (such as ACE2 and TMPRSS2) and virus structures (such as the S protein and nsps) as potential targets, we suggest that protecting the brain barrier, regulating immune responses, and ameliorating the hypercoagulation state *via* specifc targets also potentially prevent or suggest treatment strategies for pathological brain changes in patients with COVID-19.

Acknowledgements We disclosed receipt of the following fnancial support for the research, authorship, and/or publication of this article: This review was supported by the National Natural Science Foundation of China (82174005) and The Natural Science Foundation of Zhejiang Province (LY22H280007 and LEZ20H190001).

Confict of interest Zhong Chen is an Editorial Board member of Neuroscience Bulletin and a co-author of this article. To minimize bias, he was excluded from all editorial decision-making related to the acceptance of this article for publication.

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