REVIEW



Tangled quest of post-COVID-19 infection-caused neuropathology and what 3P nano-bio-medicine can solve?

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Abstract

COVID-19-caused neurological problems are the important post-CoV-2 infection complications, which are recorded in ~ 40% of critically ill COVID-19 patients. Neurodegeneration (ND) is one of the most serious complications. It is necessary to understand its molecular mechanism(s), define research gaps to direct research to, hopefully, design new treatment modalities, for predictive diagnosis, patient stratification, targeted prevention, prognostic assessment, and personalized medical services for this type of complication. Individualized nanobio-medicine combines nano-medicine (NM) with clinical and molecular biomarkers based on omics data to improve during- and post-illness management or post-infection prognosis, in addition to personalized dosage profiling and drug selection for maximum treatment efficacy, safety with least side-effects. This review will enumerate proteins, receptors, and enzymes involved in CoV-2 entrance into the central nervous system (CNS) via the blood-brain barrier (BBB), and list the repercussions after that entry, ranging from neuroinflammation to neurological symptoms disruption mechanism. Moreover, molecular mechanisms that mediate the host effect or viral detrimental effect on the host are discussed here, including autophagy, non-coding RNAs, inflammasome, and other molecular mechanisms of CoV-2 infection neuro-affection that are defined here as hallmarks of neuropathology related to COVID-19 infection. Thus, a couple of questions are raised; for example, "What are the hallmarks of neurodegeneration during COVID-19 infection?" and "Are epigenetics promising solution against post-COVID-19 neurodegeneration?" In addition, nano-formulas might be a better novel treatment for COVID-19 neurological complications, which raises one more question, "What are the challenges of nano-bio-based nanocarriers pre- or post-COVID-19 infection?" especially in the light of omics-based changes/challenges, research, and clinical practice in the framework of predictive preventive personalized medicine (PPPM / 3P medicine).

Keywords SARS-CoV2 · Post-CoV-2 infection · Post-COVID-19 complications · Nano-medicine (NM) · Hallmarks of neuropathology · Neurodegeneration · Central nervous system (CNS) · Blood-brain barrier (BBB) · Autophagy · Non-coding RNAs (ncRNA) · Inflammasome · Nanotechnology · Biomarker · Multi-omics · Predictive preventive personalized medicine (PPPM / 3P medicine) · Predictive diagnosis · Patient stratification · Targeted prevention · Prognostic assessment · Personalized medical service · 3P nano-bio-medicine

Introduction

COVID-19-caused neurological problems

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Severe acute respiratory syndrome coronavirus-2 (SARS-CoV-2; CoV-2) pandemic, coronavirus disease 2019 (COVID-19) named by the World Health Organization (WHO) in February 2020, continues to have a far-reaching impact on nearly all aspects of health, inducing respiratory distress, renal and cardiac failure, and eventually death [1]. In addition to these, post-COVID-19 neurological

manifestations have been reported in nearly 40% of critically ill patients [2]. COVID-19 patient's neurological symptoms are ranging from minor as headache and anosmia to more severe complications such as stroke, seizure, cognitive dysfunction, depression, psychosis, and delirium [1]. The global prevalence of COVID-19 and neurodegenerative disorders adds urgency to the study of their potential relationships, especially at the molecular level.

Importance of PPPM approach in improvement of the overall management of COVID-19-caused neurological problems

COVID-19-caused neurological problems are a gradually developing process since CoV-2 infection. Studies on the routes for CoV-2 entry to the central nervous system (CNS), the cellular machinery and consequences after CoV-2 entry to CNS, molecular hallmarks and signaling pathways of CoV-2-mediated neuropathology, and the molecular mechanisms of COVID-19 as a trigger for the neurological Parkinson-like symptoms will clarify in-depth the molecular mechanisms and signaling pathways of COVID-19-caused neurological problems; establish effective molecular biomarkers for predictive diagnosis, patient stratification, and prognostic assessment; and discover effective therapeutic targets for targeted prevention and personalized medical services of COVID-19-caused neurological problems in the framework of predictive, preventive, and personalized medicine (PPPM; 3P medicine). Additionally, molecular biotechnology, multi-omics, and modern drug delivery tools have great roles in highlighting post-infection studies/ trials to address nano-bio-medicine/nanotechnology (NT) potential/advances in targeted prevention and personalized therapy of COVID-19-caused neuropathology. It emphasizes the important scientific merit of the PPPM approach in the management of this type of neurological complication.

Working hypothesis in the PPPM framework of COVID-19-caused neurological complications

We hypothesize a series of molecular events and signaling pathway alterations occur in the pathophysiological processes of COVID-19-caused neurological complications. This review discusses the underlying molecular mechanisms of COVID-19 infection accompanied by CNS invasive pathway(s) and/or associated CNS complications, and what NT can add, in the following three aspects: (i) we will list proteins/receptors/enzymes that are responsible for CoV-2 entry to CNS and also list consequences after that entry from neuroinflammation to neurological manifestations via the blood-brain barrier (BBB) disruption mechanisms; (ii) we will address the hallmarks of molecular mechanisms mediating the host effect or the viral deleterious effect towards the host and highlight if SARS-CoV-2 would replicate in cells of human BBB in the presence of an accompanied neurodegenerative condition and the underlying molecular or omics-based mechanisms; and (iii) we will enumerate nanobio-medicine advancement in the light of epigenetics and/ or omics, for post-COVID-19 infection with neurological complications. These discussions will much benefit the 3P medicine research and clinical practice of post-COVID-19 infection-caused neuropathology.

Sars-CoV-2 AND the central nervous system

Infectivity is host-determined per the availability of viral receptors together with the host cell entry cofactors or coreceptors [3] (Fig. 1). Here, we will discuss the mechanism(s) for SARS-CoV-2 entry to CNS (Fig. 1–01) and neuronal target(s) for SARS-CoV-2 post entry (Fig. 1–02).

Route(s) for CoV-2 entry to the CNS

The SARS-CoV-2 virus could enter the CNS via the hematogenous or neuronal routes.

The hematogenous route

It is either pleomorphic or spherical, SARS-CoV-2 has a bear's club–shaped glycoprotein projection known as spike (S) protein that is composed of 2 subunits (S1 and S2) and has a high affinity with angiotensin-converting enzyme-2 (ACE2) receptors of the host cells [4]. Host(human)-ACE2 protein expressed in neurons is a possible target for SARS-CoV-2 infection, facilitating the virus to bind and invade the cerebral neurons and glia [5].

Another type II transmembrane domain, a receptor class A domain, a scavenger receptor cysteine-rich domain and a protease domain, which is present in the host cells, is the transmembrane protease serine 2 (TMPRSS2). Host-TMPRSS2 facilitates viral entry to the host cell by proteolytically cleaving and activating viral envelope glycoproteins [3]. The co-receptors neuropilins 1 and 2 (NRP1 and NRP2) at the host neuronal cell surface also co-facilitate viral entry [3]. The virus enters the brain via infecting the epithelial cells of the blood-cerebrospinal fluid (CSF) barrier and/or the endothelial cells of the BBB, or via the inflammatory cells of the cytokine storm as "trojan horses" to attain access into the CNS, where the virus is transferred intracellularly and is concealed by the host immune cells [6]. One study supported the hematogenous-endothelia-based neuro-invasion by SARS-CoV-2, which clearly shows the virus in the neural and capillary endothelial cells of the frontal lobe tissue from a patient infected with SARS-CoV2 [7]. Blood pressure imbalance,



Fig. 1 Mechanism(s) for SARS-CoV-2 entry to CNS (01) and neuronal target(s) for SARS-CoV-2 post entry (02). SARS-CoV-2 entry to the CNS occurs via BBB endothelial cells, viral spike proteins bind the host ACE2 receptors, with facilitated viral entry activating the viral envelope glycoproteins, and finally more interaction occurs via the host cofactor proteins neuropilins, MMPs, tyrosine kinase receptors, basigin, CD147, RAGE, and AGTR2. Post SARS-CoV-2

hypotension, or hypertension is one of the mechanism(s) by which post-COVID-19 viral infection neurological affection is signified. The membrane-bound co-receptor tyrosine kinase receptor (TKR) for vascular endothelial growth factor (VEGF), NRP1, basigin (BSG), and extracellular matrix metalloproteinase (MMPs) inducers and a cluster of differentiation 147 (CD147), the receptor for advanced glycation end products (RAGE), angiotensin II receptor type 2 (AGTR2), are identified as host cofactors/ coreceptor factors that enhance COVID-19 entry to hostcell with the possibility of ischemic/thrombotic manifestations occurring post-COVID-19 infection [8]. Hopefully, these co-receptors could be new potential targets for personalized nano-carrier anti-COVID-19 treatment. As previously stated, the virus can spread through the bloodstream and cross the BBB via attaching the viral S protein with host-ACE2 expression in the capillary endothelial lining to promote viral entry into the brain, followed by a viral infection, replication, and neuro-inflammation to

entry to the host cell, viral replication promotion occurs, induced host cell apoptosis plus cytokines and chemokines overproduction, with reduced host interferon formation, leads to host-cell necrosis, host mitochondrial and lysosomal dysfunction, and inflammasome and autophagy activation and caspase-independent host cell death, all are triggered via various viral-*ORF-genes*. v = virus

raise the possibility that SARS-CoV-2 is associated with potential neuronal complications such as encephalitis and/or stroke [9, 10]. Also, the release of inflammatory cytokines in response to the virus could disrupt the BBB's normal function and increase its permeability leading to more CNS infection [11].

The neuronal route

For this route, COVID-19 could initially infect peripheral nerve endings and then travel in a retrograde way to reach the CNS [6]. Viral components or molecules, which are released following SARS-CoV-2 invasion, interact with enteric neurons, glia, vagus, facial, glossopharyngeal, and olfactory nerve fibers to trigger an altered-neural signaling or alter neurotransmission, leading to cerebral dysfunctions [7, 12]. A prior in vivo investigation revealed that when the olfactory bulbs were removed, COVID-19's entry into the CNS was reduced, which demonstrates that the virus invades the CNS through this pathway [12].

SARS-CoV-2 interaction with the cellular machinery after entering the CNS

Viral S protein accumulates in the host cell endoplasmic reticulum (ER) and directly modulates the host unfolded protein response (UPR) to promote viral replication [13]. After entering the host cell, the COVID-19 virus genome is then released inside the host cell and the virus replicates its polyproteins 1a and 1ab that are coded by open reading frame genes (ORF-1a and ORF-1b) [14]. This helps to hijack the host's cellular machinery and take control of the host ribosomes, for their own replication and translation processes [15]. Besides, the viral proteins ORF-3a, ORF-3b, ORF-6, and ORF-7a have the ability to induce host cell apoptosis [13]. Of them, viral-ORF-7a protein activates the expression of nuclear factor-kappa B-cell (NF-kB), ORF-3b increases the host expression level of cytokines and chemokines, ORF-6 inhibits host-interferon (IFN) production, and ORF-3a induces host-cell necrosis [16]. Additionally, the viral-ORF-9b protein targets host mitochondria, which leads to mitochondrial elongation and alteration of redox homeostasis, mitochondrial dysfunction, and host cell autophagy induction [10]. Moreover, viral-ORF-8b activates the inflamma complex and interleukin-1 beta (IL-1 β) release, ER stress-response, lysosomal dysfunction, mitochondrial impairments, and caspase-independent cell death [6, 7].

Consequences of CoV-2 entry to the CNS

Anosmia affects more than 40% of the post-infected patients, which was evidenced by cranial nerve involvement in COVID-19 neurovirulence and neurotropism [17]. Interestingly, resonance image findings revealed cranial nerve, cranial root, and meningeal involvement, and reflected neuroinvasion or an edematous neuro-inflammatory mechanism(s) for neurological post-COVID-19 manifestations [18]. Neurological manifestations and BBB disruption mechanism(s) are the major complications in long-COVID-19 [19]. The proposed interaction of COVID-19, after entry, within the nervous system tissues, is sketched (Fig. 2).

Neuro-inflammation

Neuro-inflammation is one of the major hallmarks associated with a wide range of CNS disorders [20]. During infection, *IFNs* and *tumor necrosis factor-\alpha (TNF-\alpha)* genes trigger initial responses of the innate immune system. Upon activation, IFNs induce the expression of IFN-stimulated gene (ISG) proteins contributing to the cytokine storm syndrome (CSS) during severe COVID-19 [21]. CSS development by COVID-19 causes an increase in IL-6 production by glial cells in the brain [22]. Chemokines, which are produced from the choroid plexus epithelium to brain astrocytes C-C motif chemokine ligand (CCL) and C-X-C motif chemokine ligand (CXCL) family, also have roles in neuro-inflammation [23]. Primary glial cells cultured in vitro, after being infected with COVID-19, proved to secrete a considerable amount of inflammatory mediators such as IL-6, IL-12, IL-15, and TNF- α [12]. Several inflammatory genes such as *NAD*(*P*) H quinone dehydrogenase1 (NQO1) and zinc finger protein 36 (ZFP36) are upregulated during post-COVID-19 neuroaffection [23]. On the contrary, choroid and glia limitans barrier cells in patients with COVID-19 express the first-line antiviral defense gene interferon-induced transmembrane protein 3 (IFITM3) [23]. Host coreceptor cofactor CD147 activation increases the proinflammatory transcripts lectin galactoside-binding soluble 3 binding protein (LGALS3BP) and the S100 calcium-binding protein A9 (S100A9) known as migration inhibitory factor-related protein 14 (MRP14) or calgranulin B [8].

Neurological manifestation(s)

Neurological manifestations and disorders may be the first indication of SARS-CoV-2-infection-related neuro-invasion, such as Guillain–Barré syndrome, Miller Fisher syndrome, polyneuritis crania's, epilepsy, and cerebral stroke [7, 22]. COVID-19-related neurological manifestations such as fatigue, headache, altered consciousness, ataxia, generalized tonic–clonic seizures, hyposmia, and hypogeusia in 36% of infected patients are responsible for peripheral organ dysfunctions in some patients [24]. SARS-CoV-2 brain infiltration results in cerebral vascular/endothelial dysfunction with cerebral circulatory deficits such as frontotemporal hypoperfusion [25].

Neurotransmission-mediating gene(s) downregulation

Deregulation of excitatory neuron synaptic VAMP2, *SNAP25*, and *ATP6V0C* genes with a concomitant overexpression in the proximal inhibitory neurons suggests disrupted neuro-function, with cognitive deficits to occur post-COVID-19 infection [23].

BBB disruption mechanism(s)

The primary function of the BBB endothelial cells is to form a protective barrier against particles/pathogens not to pass from the arterial blood into the cerebral extracellular fluid and reach the nerve tissues, so the BBB should be ensured without damage/harm [26]. Viral molecules in the



Fig. 2 Neurological post-COVID-19 manifestations, addressing mechanism(s) of long-COVID-19 complications within the nervous system tissues. Neuro-inflammation from CSS, ILs, TNF, and more

respiratory, gastrointestinal, or other areas of the body could be transported retrogradely to the brain via the systemic circulation, interact with the cerebral-vascular-endothelium to mediate neuroinflammation, followed by BBB endothelial cell disruption [10] and encephalitis [27].

It is worth mentioning that the BBB damage by direct viral invasion differs from that caused by cytokines. In this latter situation, the BBB involvement has more diffuse damage to affect several brain regions. In other words, the integrity of the BBB is compromised by neuroinflammation induction rather than the viral attack [28]. Activation of the G-protein-coupled receptor C5a receptor 1 (C5aR1) initiates inflammatory processes in the brain via increasing BBB fluid permeability [29]. Therefore, inhibiting C5aR1 with an anti-C5aR1 antibody effectively suppressed the expressions of caspase-1 and IL-1 β , which implied that the C5a/C5aR1 signaling pathway was a potential target for SARS-CoV-2 treatment [30].

Moreover, interestingly, the aging brain BBB is weaker than the adult one. Therefore, upon CoV-2 infection, infection-related neuro-invasion, cytokine storm syndrome increases IL-6 production and the BBB permeability leading to BBB endothelial cells disruption mechanisms

macrophage activation, cerebral inflammation, and fluid accumulation responses are more severe in the former condition [19] (Fig. 3).

Hallmarks of sars-CoV-2-mediated neuropathology

Candidate mechanisms/hallmarks that mediate COVID-19 infection-related neuropathology are shown in detail (Table 1 and Fig. 4). The global pandemic mandates the need to understand *the neurotropic potential of COVID-19 and the mechanisms by which that virus triggers post-infection neuropathology*.

The molecular mimicry—human heat shock proteins

Human heat shock protein (HSP) family is now recognized as one of the mechanisms that can evoke autoimmunity [31].



Fig. 3 Young vs aging BBB in response to SARS-CoV-2 infection

SARS-CoV-2 shares sequence similarity with HSPs, which is implicated in several immune-mediated clinical conditions [32]. Research demonstrated that immunological targeting by HSPs could be a potential pathogenic mechanism of neuropathy following SARS-CoV-2 infection [33]. Accordingly, there is a possibility of cross-reactivity between epitopes within the COVID-19 spike–bearing gangliosides and the host sugar residues of surface peripheral nerve glycolipid [18], which is responsible for viral infection entry into the host body. The Guillain-Barré syndrome (GBS) postinfection nephropathy is a result of molecular mimicry from direct IgG-induced damage to nociceptive fibers [34].

Cytokine storm—the deadly systemic inflammation

The hyperinflammatory state with the elevated cytokine levels [19], CSS, results in an overwhelming systemic inflammation to exacerbate viral pathogenesis, sepsis, acute respiratory distress syndrome (ARDS), and multiorgan failure/damage [35]. This cytokine storm drives and predicts disease mortality and/or severity [36]. A patient with COVID-19-induced encephalitis was reported to have altered consciousness; akinetic syndrome with mutism with an increased IL-6, IL-8, and TNF- α ; and

moderate nuchal rigidity [37, 38]. Another case reported that encephalopathy and seizures are the initial presentation of COVID-19, along with high levels of IL-17A, IL-6, IL-8, and IFN- γ -inducible protein (IP-10), as well as a unique monocyte chemoattractant protein-1 (MCP-1) signature in CSF [39].

Cytokines have been implicated in hyposmia symptoms and olfactory nerve impairment [40], ad IL-6 was higher in the plasma, saliva, and nasal mucus of hyposmia patients [41]. Lipocalin-2, which is a glycoprotein secreted by activated neutrophils, cerebral endothelial cells, and perivascular astrocytes in response to pathogen-derived inflammation [10], is considered a biomarker of neural inflammation and is highly expressed in the CNS and contributes to the innate immunity [42].

The inflammasome NLRP3 molecular signaling

The NOD-like receptor protein 3 (NLRP3) inflammasome and IL-6 are the primary cause of clinical and pathological manifestations in COVID-19 patients [43, 44]. IL-6 stimulates downstream signaling pathways including Janus kinase/ signal transducers and transcription activators (JAK/ STAT) [45] to promote B lymphocytes proliferation/differentiation

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allmark	Marker	Mechanism	Consequence(s)
folecular mimicry; hHSPs	SARS-CoV-2 shares sequence similarity with hHSPs, direct 1gG-induced damage	Immunological targeting byhHSPs	Neuropathy post-covid ini nation, Guillain-Barré s
ytokine storm systemic inflammation	Surge of IL-6, IL-8, TNF-α, IL-17A, IL-8, IP-10, MCP-1, lipocalin-2, etc	Multi-organ failure, hyposmia symptoms, olfactory nerve impairment	Sepsis, seizures, and ence
nflammasome NLRP3 signaling	NLRP3 activation, increased IL-16/IL-18/	v-PAMPs alert the host innate immune	Neuropathy

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Table 1 Hallmarks of COVID-19 infection-	related neuropathology			
Hallmark	Marker	Mechanism	Consequence(s)	Ref
Molecular mimicry; hHSPs	SARS-CoV-2 shares sequence similarity with hHSPs, direct IgG-induced damage	Immunological targeting byhHSPs	Neuropathy post-covid infection, demyeli- nation, Guillain-Barré syndrome	[31–34]
Cytokine storm systemic inflammation	Surge of IL-6, IL-8, TNF-α, IL-17A, IL-8, IP-10, MCP-1, lipocalin-2, etc	Multi-organ failure, hyposmia symptoms, olfactory nerve impairment	Sepsis, seizures, and encephalitis	[35-42]
Inflammasome NLRP3 signaling	NLRP3 activation, increased IL-1β/IL-18/ caspase-1, B lymphocytes proliferation/ differentiation, antibody production, activated TGF-β, JAK/STAT	v-PAMPs alert the host innate immune cells, releasing IFN- $\alpha\beta$, NK CSS	Neuropathy	[43-49]
Neurodegeneration	Altered <i>PTBP1</i> , <i>YWHAZ</i> , and <i>YWHAE</i> genes expressions	CSS	Risk of AD, PD	[50-52]
Oxidative stress	Decreased glutathione and increased LDH, promotes PARP1, ROS levels	Inflammatory stress, ER stress, microbial, viral invasion	Trigger autophagy, damage to the brain, and nervous tissues	[53–59]
Cell self-eating; Autophagy	ATG5, 7; dysregulation of the autophagy machinery, decreased NLRP3 activation factors, TLR activation, IRE1 activation, ORF-9b-mediated	Autophagy negatively regulates NLRP3 activation, CSS	Neuropathy	[6067]
Amyloid neurofibrillary tangling plaques deposition	Activation of NLRP3, increased IL-1β, BBB increased permeability, amyloid beta plaques deposition, endothelial dysfunction, pericyte loss	Distorts microglia phagocytic function, failure of brain Aβ clearance, immune cell infiltration	Progression of AD, common CNS comor- bidities of COVID-19 infection	[68–71]
Prion-disease	Abnormally folded protease-resistant isoforms of host cellular sialo-glycopro- teins; prion proteins	Progressive dementia, cerebellar dysfunc- tion	Development of CJD	[73–75]
Hypercoagulation	Increased D-dimer, Factor VIII IFN, IL-2, IL-6 elevation	CSS	Pulmonary and cerebral venous thrombo- embolism	[76, 77]
Post-translational epigenetic modulation	DNA hypomethylation, histone modifica- tion, miRs, lncRNAs	ACE2 modulation, dysregulated immunomodulation	Increased susceptibility to infection	[21, 78]
Parkinson-like symptoms	APOE4 alleles, α-synuclein misfolding and aggregation, mitochondrial dysfunc- tion, impairment of protein clearance, neuroinflammation, oxidative stress, the loss of dopaminergic neurons in the substantia nigra pars in the brain	Neuronal lysis, microglia activation, pro- inflammatory factor production, vascular injury, hypoxic brain	Development and progression of PD	[27, 106–110]
CNS demyelination	IL-6 induction, activation of TLR signal- ing, production of antibodies against myelin	Neurotropic feature	Development of MS symptoms	[111-115]
Vitamin D, VDR Neurotransmission-mediating gene(s)	Vitamin D deficiency, activation of VDR VAMP2, SNAP25, ATP6V0C deregula- tion	Increased severity of infection Overexpression in the proximal inhibitory neurons	SNHG6 and 16 IncRNAs influence VDR Disrupted neuro-function, with cognitive deficits	[116, 117] [23]



Fig. 4 List of hallmarks of SARS-CoV-2-infection-related neuropathology

and induce antibody production via recruiting the transforming growth factor-beta (TGF- β) [43].

The multiprotein complex—inflammasome is a key inflammatory component and participates in both the innate immunity and inflammation response when it is activated by various stimuli [46]. NLRP3 inflammasome is activated via sensing virus surface ligands such as pathogen-associated molecular patterns (PAMPs), deathassociated molecular patterns (DAMPs), and cytokines [47]. v-PAMPs alert the hosts' innate immune cells to the invading viral molecule through releasing IFN- α/β , natural killer (NK) cell activation and various CSS molecule productions [8]. Moreover, v-PAMPs are recognized by host cell pattern recognition receptors (PRRs) as retinoic acid-inducible gene I-like receptors (RIGs), toll-like receptors (TLRs), and nucleotide-binding and oligomerization domain-like receptors (NOD) [48]. NLRP3 inflammasome activation induces caspase-1 activation and maturation of pro-IL-1β and pro-IL-18 [49]. SARS-CoV envelope M2 protein ion channel activity is linked to IL-1ß secretion with the formation of the NLRP3 inflammasome compartment. Whereas, SARS-CoV-3a protein acts as a viroporin during the formation of the NLRP3 inflammasome compartment [46]. Accordingly, owing to the potential role of NLRP3 inflammasome in SARS-CoV-2--induced immune response, therefore, an attempt to block the inflammasome component for CoV-2 CSS post-infection treatment would be recommended.

Neurodegeneration—cerebral structural damage

A recent translatomics study revealed that SARS-CoV-2 infection altered the expressions of three genes, essential for host cell neuronal survival, namely, *polypyrimidine tract binding protein1 (PTBP1)* and *tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein Zeta and Epsilon (YWHAZ and YWHAE)*, despite the fact that either gene's aberrant expression can lead to neurodegeneration [50]. For idiopathic Parkinson's disease (PD), *PTBP1* knockdown or depletion stimulates neurogenesis and reverts disease phenotypes [51]. Also, the expressions of YWHAZ and YWHAE are related directly to neurodegenerative diseases such as Alzheimer's disease (AD) [52].

Oxidative stress—the reactive spark destructive process

SARS-CoV invasion increased expressions of oxidative stress genes, prooxidants [53], and lactate dehydrogenase (LDH) [54], and downregulated glutathione antioxidant enzymes capacity [55]. The increased levels of age-related-lower-NAD⁺ and reactive oxygen species (ROS) are major predictors of SARS-CoV-2-associated in-hospital mortality [56]. Oxidative stress promotes poly (ADP)-ribose polymerase-1 (PARP1) that has been proposed as a potential therapeutic target for COVID-19 due to its impacts on the viral life cycle [57] in the brain and nerve tissue injury [58]. Inflammatory, oxidative and ER stresses, and microbial/viral invasion [59] trigger autophagy.

Autophagy—the self-degradation process

Autophagy, self-eating of cellular components, is characterized by the sequestration of cell organelles or cytosolic components into autophagosomes in combination with lysosomes for degradation [60]. Autophagy is modulated by a wide range of proteins encoded by autophagy-related genes (ATGs) in several consecutive stages [61]. Although autophagy is a survival mechanism that helps the cell clear out damaged components [62], defective autophagy signaling is involved in the pathogenesis of several diseases, including infectious and inflammatory CNS diseases [10].

Crosstalk between the inflammasome machinery and autophagy is characterized by the fact that autophagy negatively regulates NLRP3 inflammasome activation through the removal of endogenous inflammasome activators [60]. Aside from sequestering inflammasome components, autophagosomes target IL-1 β in macrophages after TLR activation, which highlights autophagy's modulatory involvement during inflammation [60].

Therefore, autophagy is a fine-tuning molecular mechanism that is directly related to the inflammatory response/inflammasome. Autophagy inhibition is required to avoid COVID-19 inflammatory response [63]. ATG5 or ATG7 controls autophagosome formation and is not necessary for viral replication in SARS-CoV-infected cells [64]. According to a recent study, Middle East Respiratory Syndrome (MERS)-CoV inhibits the fusion of autophagosomes and lysosomes. However, the reverse stimulation of autophagy lowers MERS-CoV replication [64]. Autophagosome formation is induced in the infectious bronchitis virus (IBV) through modulation of ATG5 and ER stress sensor inositol requiring enzyme 1 (IRE1) [65]. Moreover, SARS-CoV genome encodes ORF-9b that interacts with mitochondrial antiviral signaling (MAVS); further, ORF-9b-mediated MAVS deletion reduces host innate responses by lowering NLRP3 inflammasome activity [66]. Finally, ORF-9b is known to play a significant role in autophagy via a series of signaling reactions, depending upon ATG5 [67]. Now we can implicate that there is a certain type of interplay between the autophagy molecular machinery and CoVs. However, the exact nature of such interplay/interaction remains to be further elucidated.

Amyloid plaques—the neurofibrillary tangling process

One of the most prevalent neurodegenerative disorders and the leading cause of dementia is AD, which is the most common CNS comorbidity of COVID-19 infection [68]. In AD patients, deposition of amyloid beta (A β) plaques or neurofibrillary tangles can damage memory and learning brain regions [69]. Again, COVID-19 activates the NLRP3 inflammasome to distort the normal phagocytic function of microglia, with brain A β clearance failure [70]. Increased IL-1ß results in an increased Aß production, COVID-19-induced inflammation and cognitive dysfunction, and finally neurodegeneration [71]. Furthermore, COVID-19 contributes to an increased BBB permeability with endothelial dysfunction, immune cell infiltration, and pericyte loss, to hinder clearance of brain metabolites like $A\beta$ peptides [72]. The accumulation of Aß proteins in senile plaques, particularly, in the hippocampus, is the primary pathophysiological mechanism behind AD [72].

Prion-disease—the protein misfolding disorder

A transmissible protein-misfolding disorder is a prion disease, which is caused by misfolding of a host-encoded prion protein. Prion disease comes in two forms, a normal cellular prion protein or pathogenic misfolded conformer [73]. The cascade of systemic inflammatory mediators, which is characteristic of COVID-19, accelerates the prion disease pathogenesis and neurodegeneration [74]. Fatal neurodegenerative conditions may result from the accumulation of abnormally folded protease-resistant isoforms of host cellular sialo-glycoproteins (i.e., prion proteins) [74]. Creutzfeldt-Jacob disease (CJD) is a kind of human prion disease with progressive dementia and cerebellar dysfunction appearing as muscular incoordination, visual, speech, and gait clinical symptoms [73]. The CoV-2 disease might induce an exacerbated, faster pathogenesis and manifestations of CJD, as well as decreased overall survival, due to a sudden surge in the systemic inflammatory response to COVID-19 viral load [75].

Hypercoagulation—the microvascular thrombosis

IFN, IL-2, and IL-6 are elevated in COVID-19 patients to promote a hypercoagulable state in favor of thrombus formation and CSS [76]. A recent study provided a correlation between the hypercoagulable condition and severity of COVID-19, to reveal poor prognosis [77]. COVID-19-infected patients are at risk of experiencing venous, arterial, and microvascular thrombosis, from an excessive [17] immune-thrombogenic response, with an elevated serum D-dimer, factor VIII, hyperfibrinogenemia, and potential vasculitis [77].

Epigenetic modulation—the post-translational modifications

Epigenetic modulations such as DNA methylation, posttranslational modifications of histone tail, and non-coding RNAs, help regulate all biological processes and are implicated in a wide range of human disorders [78]. DNA methylation and histone modifications are associated with inflammatory complexes such as TNF- α , INF, and CSS [21]. Epigenetic modifications are implicated in the IL-6 and inflammasome pathways activation and can improve the efficacy of potential medications used for SARS-CoV-2 [43]. Host ACE2 receptors can be epigenetically modulated to increase host susceptibility to COVID-19 infection through DNA hypomethylation [21] and histone modifications [79]. Micro-RNAs (miRs) and long non-coding RNAs (lncRNAs) are genetic regulatory elements to enable the regulation of gene expressions implicated in multiple immune and cellular activities/processes (Table 2). It is worth mentioning that the host epigenetics constitute the bases of molecular biomarkers' uniqueness for predictive, preventive, and personalized (3P) intervention.

Micro-RNAs (miRs)

Epigenetic modulations may occur via miRs that regulate the immune system gene expression post-transcriptionally at the untranslated region (UTR) of genes [80]. The functionality of homo sapiens (hsa)-miRs plays an important role in the control of pro-inflammatory responses that are involved in pathophysiological condition of COVID-19 [81]. Accordingly, the tumor suppressor hsa-miR lethal-7 (Let-7) reduces the expression of IL-6 [82]; hsa-miR-124 stimulates anti-inflammatory actions via downregulating TLR-6 [83]; hsa-miR-132, hsa-miR-145, and hsa-miR-146 target NF- κ B, IL-6, IL-8, TNF- α , and MCP-1, through repression of SIRT1 (sirtuin1; silent mating type information regulation 2 homolog1) which is a member of seven protein family that function in the cellular response to inflammatory, metabolic, and oxidative stressors [84]. However, hsa-miR-146, hsa-miR-187, hsa-miR-221, and hsa-miR-155 regulate the TLR/IL-16 pathway [80]. Moreover, hsa-miR-223 inhibits IL-1ß secretion from the inflammasome with NLRP3 protein hypo-accumulation [85]. It is worth mentioning the role of hsa-miR-155 in orchestrating signaling molecule expressions to control the viral-induced immune response, namely suppressor of cytokine signaling 1 (SOCS1) [86], together with viral transcriptional modulators such as Jumonji and AT-Rich interaction domain-containing 2 (Jarid2) [87], proto-oncoprotein1 of the Ets family of transcription factors (Ets1) [88], and FOS Like 2, AP-1 transcription factor subunit (Fosl2) [89], repress T-regulatory cells (Tregs) as well as promote vascular endothelial growth factor (VEGF)independent angiogenesis [90], leading to more systemic/ neuro-inflammation and hence neuro-affection. Furthermore, has-miRs might target the genes encoding SARS-CoV-2 structural proteins. For example, hsa-miR-98, hsa-miR-744-3p, hsa-miR-410, hsa-miR-23b, and others are capable of binding to S protein-coding gene [81]. hsa-miR-126 and hsa-miR- 378 could target the viral nucleocapsid (N) protein [91], a multifunctional protein participates in the viral core formation, viral components, and genomic v-RNA synthesis [92]. hsa-miR-367, hsa-miR-6751, hsa-miR-203b-3p, hsa-miR-3132 target ORF-3a, however, ORF-6 might be bound by hsa-miR-190a. ORF-7a is targeted by hsa-miR-4436a, hsa-miR-1910-3p, hsa-miR-6866, and hsa-miR-6731. hsa-miR-4732, hsa-miR-23b, hsa-miR-3190-3p, and hsamiR-5011-3p bind to ORF-8, whereas hsa-miR-3682, and hsa-miR-411 target ORF-10 [81, 93, 94]. Interestingly, hsamiR-574-5p attenuates COVID-19-related CSS, suppresses TLR4/NF-kB signaling, and decreases proinflammatory cytokine production in ARDS patients [95].

On the other hand, viral (v)-miRs might influence the expressions of vital molecules during antiviral and proinflammatory signaling pathways. For example, SARS-CoV-2-encoded v-miR-19 and v-miR-62 can target some subunits of the RNA polymerase II enzyme, and v-miR-61 is able to target host STAT1 to decrease host-related IFNmediated signaling pathways [93]. v-miR-MR147-3p and v-miR-MR66-3p target TMPRSS2 and TNF- α , respectively, to effect an adverse clinical symptom via elevating the expression levels of TMPRSS2 and TNF- α [81]. The hsa-miRs are sketched to target the virus particle/protein itself (Fig. 5).

Long non-coding RNAs

A novel class of non-coding transcripts (longer than 200 nucleotides in length) plays an important role in cell proliferation, differentiation, apoptosis, autophagy, tissue repair, and tissue remodeling, development, and progression of disorders such as cancer and infectious diseases [20, 92]. LncRNAs have roles in viral transformation, viral latency

ncRNA		Target(s)	Effect(s)	Ref
miRNA	hsa-miR Let-7	IL-6	Reduces IL-6 expression	[82]
	hsa-miR-124	TLR-6	Induces the anti-inflammatory actions by downregulating TLR-6	[83]
	hsa-miR-132, hsa-miR-145, hsa- miR-146	Sirtuin1	Modulate NF-κB expression, IL-6, IL-8, TNF-α, MCP-1	[84]
	hsa-miR-146, hsa-miR-187, hsa- miR-221, hsa-miR-155	TLR/ IL-1β	Regulate TLR/IL-1β pathway	[80]
	hsa-miR-223	IL-1β/ NLRP3	Inhibits IL-1β secretion from the inflammasome, decrease NLRP3 protein	[85]
	hsa-miR-155	SOCS1/ Jarid2/ Ets1/ Fosl2/Tregs, VEGF	Systemic/neuro-inflammation, neuro- affection	[87, 90, 91]
	hsa-miRNA 126, hsa-miRNA 378	Nucleocapsid protein	Influence viral-host interaction	[91]
	hsa-miR-98, hsa-miR-744-3p, hsa- miR-410, hsa-miR-23b	S protein-coding gene	Influence expression of genes involved in viral pathogenesis, attachment with ACE2	[81]
	hsa-miR-367, hsa-miR-6751, hsa-miR- 203b-3p, hsa-miR-3132	ORF-3a	Influence viral replication, pathogen- esis	[81] [93]
	hsa-miR-190a	ORF-6	Influences nuclear import of STAT1	[94]
	hsa-miR-4436a, hsa-miR-1910-3p, hsa-miR-6866, hsa-miR-6731	ORF-7a	Influence viral accessory protein, composed of a type I transmembrane protein, modulate caspase-dependent apoptosis pathway	
	hsa-miR-4732, hsa-miR-23b,hsa-miR-3190-3p,hsa- miR-5011-3p	ORF-8	Influence viral replication, interspecies transmission	
	hsa-miR-3682, hsa-miR-411	ORF-10	Influence viral trans-species transmis- sion	
	hsa-miR-574–5p	TLR4/NF-kB	Suppress TLR4/NF-kB signal- ing, decreases proinflammatory cytokines, attenuates the COVID-19- caused CSS	[95]
	v- miR-61	STAT1	Decreases IFNs-mediated signaling pathways	[93]
	v-miR-19, v-miR-62	RNA polymerase II	Disruption to hsa-miRs biogenesis	[93]
	v-miR-MR147-3p v-miR-MR66-3p	TMPRSS2, TNF-α	Elevate their expressions, elevate inflammation to worsen the overall clinical outcome	[81]
LncRNA	MALAT1	NF-κB/ MAPK	Key role during SARS-CoV-2 inflam- matory immune response	[99]
	NEAT1	NLRP3 inflammasome	Activating caspase-1, inflammatory cytokine production	[100]
	NORAD	IL-6, TNFα	Implicated in modulating viral- induced inflammatory responses	[98, 102]
	ANRIL	hsa-miR-122 /BRCC3 axis	Inducing NLRP3 expression, develop- ment of cytokine storm	[104]
	XIST	NF-κB/NLRP3 inflammasome	Mediates the inflammatory response	[103]
	MEG3	hsa-miR-122, hsa-miR-22, hsa-miR-223	Inhibiting hsa-miR-223 function, increasing NLRP3 expression with virus replication progression, cytokine storm	[81]

maintenance, viral replication promotion, and viral-gene expression augmentation [96]; implicate in modulating host cell immunity and/or inflammation, host cell energy/

metabolism, host cell cycle, and apoptosis; and thus negatively impact the normal physiological activities of host cells [97]. Multiple lncRNAs were differentially expressed during



Fig. 5 The hsa-miRs that target the virus particle/protein itself. v-miR-MR147-3p and v-miR-MR66-3p target host TMPRSS2, and the reverse human miRs target the viral envelope glycoprotein. Some

host miRs affect the hACE2 receptor which binds viral-spike protein. Host miR-126 and -378 target the viral nucleocapsid protein

SARS-CoV-2 infection, which are considered potential disease biomarkers [98]. Metastasis-associated lung adenocarcinoma transcript 1 (MALAT1) and lncRNA X-inactive specific transcript (XIST) have been reported to play a key role during the SARS-CoV-2 inflammatory immune response [99]. Non-coding nuclear-enriched abundant transcript 2 (NEAT2) has dual roles in the inflammatory response and in the secretions of IL-6, TNF- α , and IL-1 β cytokines, during different signaling pathways [43]. NEAT1, the oncogenic lncRNA, has been shown to translocate from the nucleus to the cytoplasm, which induces inflammasome NLRP3 complex formation, and activates caspase-1 and inflammatory cytokines production [100, 101]. ncRNA activated by DNA damage (NORAD) has been identified in modulating viral pathogenesis via targeting CSS cytokines [98, 102]. Whereas XIST mediates inflammation via generating a negative feedback loop to regulate NF-kB/NLRP3 inflammasome pathway [103]. Maternally expressed gene 3 (MEG3) via sponging hsa-miR-122, hsa-miR-22, and hsa-miR-223 increases NLRP3 expression, affects pyroptosis, increases inflammation intensity, and worsens the overall clinical outcome [81]. Antisense ncRNA in the INK4 locus (ANRIL) was found to be involved in the NF-kB-mediated inflammatory response via sponging miR-122-5p, thus activates the NLRP3 inflammasome [104].

Therefore, these ncRNAs might constitute potential target(s) to interevent SARS-CoV-2 invasion [102]. Nearly 40% of lncRNAs identified so far are expressed, particularly in the brain, and are distributed in the olfactory bulb and cerebellum of the hippocampus cortex [105]. Viruses stimulate host-lncRNAs near the olfactory bulb to participate in viral infection, replication, latent infection, and immunological responses, with an increased IL-2, IL-7, INF, and TNF- α production [102].

Future research is warranted to link more, epigenetics; lncRNAs to post-CoV-2 cellular-stress-response function, which will aid insight into the research gap at the molecular mechanistic roles of lncRNAs, pointing to them as potential therapeutic new modalities/targets to treat post-SARS-CoV viral infection with neuro-affection.

COVID-19 as a trigger for the neurological Parkinson-like symptoms

Parkinson's disease (PD) is a chronic neurological disorder that primarily affects the elderly and impairs mobility. Given their vulnerability, it is probable that PD patients are more vulnerable to COVID-19 and COVID-19-post infection complications [106]. PD is attributed to a combination of molecular pathogenic mechanisms of α -synuclein misfolding and aggregation, with mitochondrial dysfunction, and impairment of protein clearance, neuroinflammation, oxidative stress, and the loss of dopaminergic neurons in the substantia nigra pars in the brain [107].

The link between SARS-CoV-2 and Parkinson's disease PD

COVID-19 can exacerbate PD symptoms, which indicates the neurotropic nature of COVID-19 infection, either directly by virus tropism and host neuronal cell lysis in the basal ganglia as well as indirectly via microglia activation, increased proinflammatory mediators production, with vascular injury in the stressful hypoxic brain [108].

Human ACE2 receptor has been found in the substantia nigra, ventricles, middle temporal gyrus, posterior cingulate cortex, and olfactory bulb. Accordingly, the possibility of the substantia nigra to be infected by the COVID-19 virus, through binding ACE2, might contribute to the acute parkinsonism condition in COVID-19-infected patients [109]. A recent case of Parkinsonism associated with COVID-19 in the later stages of the disease has been reported with hyposmia, the generalized myoclonus, asymmetric hypokinetic-rigid syndrome symptoms, the altered consciousness, and a bilateral decrease in presynaptic dopamine uptake [110]. Another COVID-19 patient, who experienced parkinsonism with neurological symptoms such as rigidity, tremors, and bradykinesia, is resolved after convalescent plasma treatment [108].

Following SARS-CoV-2 infection, α -synuclein, a natural antiviral neuronal protein that regulates synaptic vesicle trafficking, is implicated in PD. α -Synuclein aggregation pushes sensitive neurons over the edge of neurodegeneration [106, 107]. Apolipoprotein E4 (APOE4) alleles are considered significant risk factors, not only for tauopathies such as AD but also for the development of dementia in synucleinopathies; PD strongly impacts the neurotropism of the SARS-CoV-2 virus (glial cells and neurons expressing ACE2 receptors) and the severity of COVID-19-post-infection complications [27].

CNS demyelination

CNS demyelination has been reported with MERS-CoV, SARS-CoV-1, and SARS-CoV-2 infections, as well as neurotropic features [111]. SARS-CoV-2 has neurotropic and neuro-invasive features, inducing direct neurological damage, which are mediated via IL-6 and undesirable immune response, antibody production surge against myelin [112, 113] as well as via TLRs production following SARS-CoV-2 acute infection, resulting in CNS demyelination diseases as multiple sclerosis [114, 115].

The hormone-like vitamin D

The cholesterol-derived steroid hormone, vitamin D, regulates the expression of 5% of human genes involved in the immunological response to pathogens [116]. Post-COVID-19 complications were reported to be triggered by vitamin D deficiency and being mitigated via vitamin D receptor (VDR) activation. The expression of VDR has been found to be influenced by small nucleolar RNA host genes 6 and 16 (SNHG6 and 16) lncRNAs [117].

Thereby, it is noteworthy mentioning that Parkinson-like symptoms, CNS demyelination and vitamin D deficiency, VDR activation, all occurring in cases of SARS-CoV-2 infection, could be considered among COVID-19 infectionmediated neuropathology hallmarks, which could be beneficial markers for better prediction, prevention of COVID-19 infection–mediated neuropathology complications.

Nano-bio-medicine advancement(s) in the era of COVID-19 infection

Advancement of magnetism-based nanoparticles for COVID-19 molecular diagnosis

NT contributes to SARS-CoVs-2 RNA extraction and/ or qRT-PCR detection. This is a nanoparticle (NP)-based one-step SARS-CoVs-2 RNA extraction technique using amino-modified magnetic NPs (MNPs) decorated with polycarboxyl groups (PC-coated NH2-MNPs) to bind solely to the virus RNA. Upon applying a magnetic field, the nucleic acids are eventually collected for viral RNA release from the MNPs, an elution buffer is used [54].

Nano-medicine assembly-based structure advancement for COVID-19 vaccination

Assembly-based structure is used for fabrication of CoV-2 nanovaccines (NVs), via encapsulation of CoV antigens or exposing them on a NP surface, with an advantage of less aggregation and an improved *vaccine efficacy*, via (i) protection of COVID-19 antigens from degradation, during the quest of vaccination, (ii) controlling the amount of antigens to be delivered within the NV from the NP-matrix, and (iii) regulation of antigen uptake and processing [118]. Gold NPs are antigen carriers, used in NVs, acting as adjuvants during immunization [119].

Nanostructure advancement for post-COVID-19 infection immune-related complication(s)

Besides nano-dots and nano-fibers, for point-of-care testing (POCT) diagnosis, as well as NP usage for vaccine development, re-purposed NT is now used to engineer repurposed antiviral drugs within targeted-NPs, as "reformulated repurposed" drugs, the "2Re" approach, delivered within a nanocarrier, protected and potentiated in action (immunemodulatory effect) [120].

To use and globalize the concept of "nano-by-design" (NbD) [121] and design or engineer a/- treatment cargo(s), as *exosomes*, for immune modulation, either enhancing/ potentiating or repressing the immune response, for counteracting the replicative capabilities of SARS-CoV-2 or compacting post-CoV-2 complications as CSS and hence guard against neuro-affection.

Nano-bio-based therapy or regeneration

Effective therapeutic anti-SARS-CoV-2 small interfering RNA (siRNA) activators would be an effective strategy to inhibit the highly replicating RNA virus, COVID-19, in a specific and rapid manner [8]. One study reported blocking the replication of the ORF-1b sequence of the virus genome with plasmid-mediated siRNAs [122]. However, the use of plasmid-mediated siRNAs has limitations of rapid enzymatic degradation or unwarranted hyper-stimulated immune reaction and the most serious drawback was the random siRNA insertion into chromosomes [123]. Therefore, the use of the FDA-approved, non-toxic, biocompatible nanocarriers would be a better approach to protect siRNA and ensure their in vivo delivery. These nanocarriers are prepared as either nano-capsules, nano-spheres, nano-emulsions or soluble lipid nano-carriers (SLN), lipid nano-carriers (LNC), or from polymeric micelles and polymer-based as poly(lactic acid) (PLA), polycaprolactone, poly(glycolic acid) (PGA), poly(lactic-co-glycolic acid) (PLGA) or hybrid (polymer/ lipid) NPs [124].

Anti-viral-drug-loaded exosomes as nano-bio-materials can cross the BBB to modulate neuro-affection, modulate immunity, enhance neural regeneration, and deliver treatment post-CoV-2 neuropathies [125]. Moreover, cytokine receptor–loaded exosomes deliver cytokine receptors as an immune-modulatory approach to neutralize or remove proinflammatory cytokines or chemokines [126], so compact CSS. Enveloped nano-complexes with an amphiphilic polymer poly(glutamic acid)-PEG (PGA-PEG) to transport miRs to the brain and modulate its mRNA targets [127]. BBB endothelial cells are tightly connected to block any polar compounds from passing, unless being lipophilic, or coming from systemic absorption or via the nasal route (noseto-brain), not intranasal, across the olfactory mucosa [128], with the recommended use of cell penetration peptides onto the NPs as dextran nano-micelles or bio-muco-adhesive nano-carrier.

NPs to be formulated using polymers, metals, nanogel and colloidal systems, or vesicular systems as liposomes organized from double-chain phospholipids, ethosomes, transfersomes, and bilosomes, or niosomes for efficient drug delivery to the brain [129]. Niosomes are composed of cholesterol or phosphatidylcholine and uncharged non-ionic single-chain surfactant as spans, tweens 20, 40, 60, and 80, and Brij, incorporated into an aqueous phase [130]. Proniosomes consist of water-soluble carriers and surfactants, being dehydrated, hydrogels for biomolecules delivery as protein, monoclonal Ab, self-assembling peptides, or vaccine delivery as inhaler or I.V, parenteral [131].

Nano-carriers are synthesized using biomaterials like protein polymers (collagen) and lipids (micelles), synthesized using ceramics (silica), metals (gold and silver), oxides and salts, and polymers (proteins and lipids). Nanoenzymes that are said to be functionalized, could be used for controlling and repairing/regenerative, at a molecular level, prion, amyloid protein misfolding with limited clearance, cytokines release inhibitors or enhancers, enzymes activators or depressors, antibodies, signaling molecules, RNA, enhancer or repressor factors, stem cells, as a challenge. Cerium oxide NPs (CeONPs) can mimic the effects of antioxidant enzymes to treat oxidative stress [132]. NPs loaded with either enzymatic or non-enzymatic antioxidant vitamins, minerals, metabolites, and several phytochemical compounds (nutraceuticals) to protect the neurons from oxidative stress are an acceptable approach nowadays [133], like platinum, gold NPs, titanium dioxide NPs, silver, iron, copper oxide, and zinc oxide have an anti-inflammatory effect [134]. Gold NPs can suppress Aβ aggregation [135]. NPs loaded with complement diminished the microglial neurotoxicity after injury via decreasing the levels and activity of pro-inflammatory factors, inflammatory cells, and neuronal apoptosis [136]. Selenium NPs provided a neuroprotective effect via diminishing the oxidative-inflammatory-apoptotic cascade [137]. SLNs may serve to target amyloid plaques, prion, tau, as anti-amyloid, anti-protein aggregation, antitangling, accelerating and facilitating autophagy and to prevent neurofibrillary tangles formation [133]. Gold NPs and small liposomes could pass through BBB passively [138]. Some NPs use carrier-mediated transport pathways such as the glucose transporter 1 (GLUT1) protein and amino acid transporter (ASCT2) for delivery across BBB [133].

Challenges of nanotechnology pre-COVID-19 infection

First, NP formulations would be a promising platform for hostimmune cells enrichment with IFN- α /- γ /I prior to infection as a prophylactic approach. This is to (i) enhance the expressions of host IFN-related genes and signaling pathways like protein kinase receptor (PKR) and 2'-5'-oligoadenylate synthase (OAS)/RNase L; and (ii) stimulate apoptosis of the COVID-19-infected cells [8], with an overall anti-viral replicative effect. Moreover, nanomaterials used for formulating the nano-carriers by themselves can have an immune-modulatory effect(s) activating or inhibiting CDs such as PLGA. Second, NP-based nose-to-brain smart delivery of the formulated and characterized intranasal NPs carrying immuno-modulatory drugs could be used, which is an approach that our group is working on currently.

Limitation/challenges of nanotechnology post-COVID-19 infection

First, the minimum literature information reporting bio-nano experiments is summarized. Second, the use of NP-based immunotherapy against CoV-2 infection clinically is still challenging and is limited by (i) NP therapy side effects or possible toxicity, (ii) NM degradation throughout their delivery journey to the target immune cells or receptors. Therefore, the ability to create and evaluate massive biomedical datasets [139] could aid in answering how can nanoscale chemical structure affect medicine distribution in vivo? And what biological processes are in-charge of NPs delivery in vivo? and (iii) the genetic diversity derived from biomolecular modifications.

Genetic diversity is derived from biomolecular modifications where cellular function/dysfunction depends on biological complexity, and because of the intricacy, determining how individual biomolecules contribute to a phenotype [139] as post-CoV-2 infection neuro-complications is difficult. This task is made more complex by the magnitude of the biological system involvement or activation/deactivation.

The chemical composition, shape, and size of NPs affect their interactions with different molecules. Many intriguing concerns remain unresolved, given that individual genes, systemic physiology, and NM chemical structure all interact to determine NM behavior. Whether there are master regulatory genes impact many different types of NPs, or the reverse, if the NPs as LNC loaded with drugs based on proteins would affect target genes in a different manner between infected and normal cells, or an effect different from the free mRNA effect on this same gene, have yet to be discovered [140]. Fortunately, complexity is a biological norm, and new sequencing tools (omics) are to be positioned to aid in the research of nano-bio-medicine interaction(s). Omics is the study of biological molecules on a large scale in order to comprehend their whole genetic or molecular profiles [141]. In other words, multi-omics are required to aid more in elucidating how cells respond to NPs.

Drugs based on siRNA, miR, mRNA, lncRNA, zinc fingers, and CRISPR-Cas proteins have joined traditional small molecule therapies, where each drug class will bring multiple options [139] as the biological response to NPs loaded with nucleic acids differs from those loaded with RNA.

Predictive, preventive, and personalized nano-bio-medicine in COVID-19-caused neuropathology

To personalize nano-bio-medicine therapeutic interventions based on the growing understanding of the human multiple omics [142], genomics, epigenomics, transcriptomics, proteomics, and metabolomics have led to the discovery of various biomarkers. Similarly, the same approach can be used to detect early-stage or post-infection neuro-affection and predict post-infection neurodegenerative progression, drug response, and the overall clinical outcome.

Challenges

The heterogeneity of RNA viruses poses a significant challenge for future pandemic infection personalized therapy. However, the use of multi-parameter omics data for specific molecular biomarker recognition in combination with the versatile drug delivery nano-carriers (Table 3), which can specifically target multiple immune cell subpopulations or host-affected receptors, might herald a promising future for predictive, preventive, and personalized post-pandemic management. First, figure out what kind of omics data you will need, including genomics, epigenomics, miRNomics, transcriptomics, proteomics, ubiquitinomics, acetylomics, phosphoproteomics, nitroproteomics, and metabolomics; second, determine whether it is critical to comprehend the transcriptomic, epigenetic, proteomic, and metabolomic responses, or a mix of responses. However, any single omics method has limited insights into interrelated molecular pathways and complicated biological activities in cells and organisms [141].

Expert recommendation in the framework of 3P medicine

It is recommended to strengthen the awareness and research of post-COVID-19 infection-caused neuropathology and take the corresponding early medical services to resolve this complication that might cause long-term side effects for COVID-19-infected patients. The reality is that neuropathology has been confirmed with the presence of SARS-CoV-2 virus particle positive qRT-PCR. From the multi-parameter systematic angle [143, 144], it is necessary to assess the brainimmune microenvironment via a brain-specific panel

Bio-nano materials		Role	Applications	Ref
Magnetic NPs; amino-modified n	nagnetic NPs decorated with polycarboxyls	Contribute SARS-CoVs-2 RNA extraction and detection	Molecular diagnosis	[54]
Nano-dots/nano-fibers		Point-of-care testing	Diagnosis	[120]
Nano vaccine	Antigen's encapsulation	Of less aggregation and an improved vaccine efficacy	NVs	[118]
	Gold NPs	Antigen carrier	Vaccine, and adjuvants in immunization	[119]
NPs and re-purposed NT		Antiviral repurposed drugs	Treatment	[120]
Nano-carrier	exosomes	Protection of immune-modulatory drugs	Counteracting the replicative capabilities of SARS- CoV-2 /counteracting post-CoV-2 complications	[121]
		Anti-viral drugs, cross the BBB	Modulate neuro-affection, modulate immunity, enhance neural regeneration, to treat post CoV-2 neuropathies	[125]
		Cytokine receptor-loaded	Neutralize or remove pro-inflammatory cytokines or chemokines, compact CSS	[126]
	Capsules, spheres, emulsions, LNC, PLA, PLGA, liposomes	Plasmid-mediated siRNAs protection	Block the replication of the ORF-1b sequence of the virus genome	[122–124, 129, 138]
	Enveloped nanocomplexes with PGA-PEG	Transport miRs to the brain	Modulates miRs mRNA targets	[127]
	Nose-to-brain route with dextran nanomicelles, bio-muco-adhesive NC	Cross the olfactory mucosa using cell penetration peptides onto NPs	Pass BBB	[128]
	Proniosomes, Niosomes are cholesterol or phosphatidylcholine, uncharged non-ionic single-chain surfactant	For protein, monoclonal Ab, self-assembling pep- tides brain delivery	for BBB delivery of biomolecules or vaccine delivery as inhaler or I.V, parenteral	[130, 131]
	Cerium oxide NPs	Mimic antioxidant enzymes	Treat oxidative stress contributing to neuropathology	[132]
	enzymatic, non-enzymatic antioxidant vitamins, metabolites loaded NPs	Phytochemical compounds (nutraceuticals) metabolites loaded NPs	Protect the neurons from oxidative stress so modulate neuro-affection, enhance neural regeneration, to treat post CoV-2 neuropathies	[133]
	Platinum, titanium dioxide NPs, silver, iron, copper oxide, zinc oxide	Phytochemical compounds (nutraceuticals) to protect the neurons from oxidative stress	Anti-inflammatory effect, modulate neuro-affection, modulate immunity, to treat post CoV-2 neuropathies	[134]
	Gold NPs	Suppress A β aggregation, cross BBB passively	Modulate neuro-affection, modulate immunity, enhance neural regeneration, to treat post CoV-2 neuropathies	[135, 138]
	Complement loaded NPs	Diminished microglial neurotoxicity after injury	Decreasing levels, activity of pro-inflammatory factors, inflammatory cells, neuronal apoptosis	[136]
	Selenium NPs	Neuroprotective effect	Diminishing oxidative-inflammatory-apoptotic cascade	[137]
	SLNs	Target amyloid plaques, prion, tau, as anti-amyloid, anti-protein aggregation, anti-tangling, accelerat- ing	Facilitating autophagy, prevent neurofibrillary tangles formation	[133]
	Carrier-mediated transport GLUT1 protein ASCT2	NPs for delivery across BBB		[133]



Fig. 6 Predictive, preventive, and personalized 3P nano-bio-medicine implementation during and post-pandemic era

of molecular biomarkers, including T-cell markers and neuronal biomarkers, as well as the viral entry receptors into host cells; CD147, hACE2, and hTMPRSS2 (Fig. 6). This panel of biomarkers can be taken to understand the molecular mechanism of this type of neuropathology, evaluate and score neuro-affection status of the patient, stratify the patients, predict the occurrence of neurodegeneration, assess its prognosis, and even discover



Fig. 7 COVID-19-infection caused cytokine storm and current or future CoV-2 infection treatment attempts



Fig. 8 Post COVID-19-related neuro-affection manifestations summary

therapeutic targets for personalized prevention and earlystage therapy of this type of neurodegeneration in the framework of 3P medicine.

Conclusions

The review unraveled SARS-CoV-2 disease molecular pathophysiology, cellular and molecular mechanisms of viral infection, cellular/BBB penetration, and reproduction. Figure 7 illustrates COVID-19-infection-caused cytokine storm and the current as well as the future CoV-2 infection treatment attempts. Hallmarks of the disease post-affection are summarized in Fig. 8 as post-COVID-19-related neuro-affection manifestations. Systemic and neuro-inflammation, neuro-apoptosis/ autophagy balance, viral miRs and has miRs as well as lncRNAs epigenetics, neuronal oxidative stress, and protein misfolding with limited clearance; tau, amyloid, and prion, a-synucleino-pathy are deregulated signaling mechanism(s) that synchronize to affect neuronal damage. COVID-19-related neurological manifestations are either mild or severe ischemic affection, stroke, or inflammation. Underlying neuro-affection causative processes are either an indirect arm from thrombosis, generalized rapid aggressive inflammatory response, hypoxia, or others. Direct neurotropic dysfunction related to COVID-19-infection may be the other causative arm. Though neurological manifestations are not synonymous

with early COVID-19 detection and treatment, their prevalence is lower than that of respiratory symptoms. Being not uncommon, the neurological signs should not be overlooked because they occur at any time during the infection course or post-COVID-19 infection and, hence, can lead to serious problems if it is not detected and treated properly early. Nonetheless, given the speed with which the pandemic is spreading and the likelihood that 50-70% of the world's population would be COVID-19 infected, before gaining herd immunity, the overall number of people with neurological symptoms and/or experiencing CNS harm is growing exponentially. ND being caused by cytokine storm or direct COVID-19-infection and neuro-related mechanism(s), fully addressed in this review are the current or future CoV-2 infection treatment attempts. Scientists that embrace omics data in the nano-bio-medicine, NbD, have a strong meaningful chance to address global heterogenic highly mutating current or future infection challenge(s) and will be at the forefront of creating new treatments as well as boost precision medicine and 2030 good health initiative via predictive, preventive, and personalized medicine.

In summary, novel repurposed NT approaches, engineering and fabricating one or more combined drugs with nano-carriers' materials ensuring multi-drug delivery system with least or no toxicity might solve the problem of "lack of treatment" availability of the highly mutating heterogenous SARS-CoV-2 virus and further provide a promise to apply the PPPM concept of post-COVID neuro-complication management, to this pandemic.

Abbreviations A_β: Amyloid beta; ACE2: Angiotensin-converting enzyme-2; AD: Alzheimer's disease; ANRIL: Antisense non-coding RNA in the INK4 locus; ARDS: Acute respiratory distress syndrome; ATG: Autophagy-related-genes; BBB: Blood-brain barrier; C5aR: C5a receptor; CD: Cluster of differentiation; CJD: Cruetzfeldt-Jacob disease; CNS: Central nervous system; COVID-19: Coronavirus disease-2019; CSF: Cerebrospinal fluid; CSS: Cytokine storm syndrome;; DNA: Deoxyribonucleic acid; ER: Endoplasmic reticulum; GBS: Guillain-Barré syndrome; HCoV: Human corona viruses; Has: Homo sapiens; HSPs: Heat shock proteins; IFN: Interferon; IL: Interleukin; IncRNA: Long non-coding RNA; LNC: Lipid nanocarriers; MALAT1: Metastasis associated lung adenocarcinoma transcript 1; MAVS: Mitochondrial antiviral signaling; MCP-1: Monocyte chemoattractant protein-1; MERS: Middle east respiratory syndrome; miRs: Micro-RNA; MMPs: Matrix metalloproteinases; MNPs: Aminomodified magnetic NPs; MS: Multiple sclerosis; NAD: Nicotinamide adenine dinucleotide; NbD: Nano by design; ND: Neurodegeneration; ncRNA: Non-coding RNA; NEAT: Non-coding nuclear-enriched abundant transcript; NF-KB: Nuclear factor-Kappa BM; NLRP3: NOD-like receptor protein; NM: Nano-medicine; NORAD: Non-coding RNA activated by DNA damage; NP: Nanoparticle; NRP: Neuropilin; NT: Nanotechnology; NV: Nanovaccine; ORF: Open reading frame; PAMPs: Pathogen-associated molecular patterns; PD: Parkinson's disease; PLGA: Poly lactic-co- glycolic acid; PTBP1: Polypyrimidine tract binding protein1; 3P medicine: Prediction, prevention, personalized medicine; RNA: Ribo-nucleic acid; S: Spike; SARS-CoV-2: Severe acute respiratory syndrome - coronavirus - 2; siRNA: Small interfering RNA; SLNs: Soluble lipid nanocarriers; STAT: Signal transducers and transcription activators; TLR: Toll-like receptors; TMPRSS: Transmembrane protease serine; TNF-α: Tumor necrosis factor-a; VDR: Vitamin D receptor; V-miR: Viral-miR; XIST: X-inactive specific transcript; YWHAE: Tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein Epsilon; YWHAZ: Tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein Zeta

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