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# Food and Chemical Toxicology



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# Recent advances in potential drug therapies combating COVID-19 and related coronaviruses-A perspective



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### ABSTRACT

Coronaviruses (CoVs) are a large family of viruses responsible for the severe pathophysiological effects on human health. The most severe outbreak includes Severe Acute Respiratory Syndrome (SARS-CoV), Middle East Respiratory Syndrome (MERS-CoV) and Coronavirus disease 2019 (COVID-19) caused by Severe Acute Respiratory Syndrome coronavirus 2 (SARS-CoV-2). The COVID-19 poses major challenges to clinical management because no specific FDA-approved therapy yet to be available. Thus, the existing therapies are being used for the treatment of COVID-19, which are under clinical trials and compassionate use, based on *in vitro* and *in silico*  studies. In this review, we summarize the potential therapies utilizing small molecules, bioactive compounds, nucleoside and nucleotide analogs, peptides, antibodies, natural products, and synthetic compounds targeting the complex molecular signaling network involved in COVID-19. In this review>230 natural and chemically synthesized drug therapies are described with their recent advances in research and development being done in terms of their chemical, structural and functional properties. This review focuses on possible targets for viral cells, viral proteins, viral replication, and different molecular pathways for the discovery of novel viral- and hostbased therapeutic targets against SARS-CoV-2.

### **1. Introduction**

Coronaviruses (CoVs) are enveloped viruses having non-segmented, positive sense single-stranded RNA genome rather than DNA, belonging to the family *Coronaviridae* and contain the largest genomic RNA among any viruses broadly distributed in humans and other mammals [\(Pillaiyar](#page-16-0)  [et al., 2020;](#page-16-0) [Zumla et al., 2016](#page-17-0)). CoVs are named from crown-like spikes protruding from their outer surface and grouped in four main sub-groups, mainly alpha, beta, gamma, and delta ([Sheahan et al.,](#page-16-0)  [2020a,](#page-16-0) [2020b](#page-16-0), [2020b](#page-16-0)). CoVs were first identified in the mid-1960s, seven of which infect human beings. These are MERS-CoV (the beta

coronavirus that causes Middle East Respiratory Syndrome, or MERS), SARS-CoV (the beta coronavirus that causes severe acute respiratory syndrome, or SARS), NL63 (alpha coronavirus), 229E (alpha coronavirus), OC43 (beta coronavirus), HKU1 (beta coronavirus), and severe acute respiratory syndrome-related coronavirus (SARS-CoV-2, novel coronavirus responsible for COVID-19). People around the world commonly get infected by human CoVs like HCoV-229E, HCoV-NL63, HCoV-OC43, and HCoV-HKU1 (B. [Chen et al., 2020;](#page-14-0) [Shen et al., 2019](#page-16-0)). Sometimes CoVs that infect animals can evolve, make people sick and become a new human coronavirus. Recent examples are SARS-CoV, MERS-CoV and SARS-CoV-2 [\(Pillaiyar et al., 2020](#page-16-0); [Zumla et al.,](#page-17-0) 

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*Abbreviations:* COVID-19, Coronavirus disease 2019; SARS-CoV, severe acute respiratory syndrome coronavirus; SARS-CoV-2, severe acute respiratory syndrome coronavirus 2; MERS, Middle East respiratory syndrome; HCoV, human coronavirus; CHIKV, Chikungunya virus; DHODH, dihydroorotate dehydrogenase; HBV, hepatitis B virus; IAV, influenza A virus; HCV, hepatitis C virus; JEV, Japanese encephalitis virus; PEDV, porcine epidemic diarrhea virus; PLpro, papain-like protease; 3CLpro, 3 chymotrypsin-like proteases; RdRp, RNA-dependent RNA polymerase; SAH, S-adenosyl-l-homocysteine; RBD, receptor-binding domain; RSV, respiratory syncytial virus; ZIKV, Zika virus; IMPDH, inosine-monophosphate dehydrogenase; PPIase, peptidyl-prolyl isomerase; IMPTH, inosine-5′ -monophosphate dehydrogenase; NS3, non-structural protein 3; VEGF, Vascular Endothelial Growth Factor; ALI, acute lung injury; ARDS, acute respiratory distress syndrome; HTCC, N- (2hydroxypropyl)-3-trimethylammoniumchitosan chloride; SCV, SARS-associated coronavirus; HCMV, Human cytomegalovirus; COX, cyclooxygenase; JAK, Janusassociated kinase; NAK, Numb-associated kinase; HCMV, Human Cytomegalovirus; NS, Not studied; S protein, Spike (S) protein; E, Enveloped protein; ACE2, angiotensin-converting enzyme 2 (ACE2) blockers; TCM, traditional Chinese medicine.

<span id="page-2-0"></span>[2016\)](#page-17-0). The detail taxonomical classification of coronaviruses (according to the International Committee on Taxonomy of Viruses) illustrated in Fig. 1.

In late December 2019, several cases of unexplained pneumonia have been reported in Wuhan, China. Most of the infected or confirmed patients live near the local Huanan seafood wholesale where live animals are widely sold, where live animals are widely sold. In the early stages of pneumonia, severe acute respiratory infections occur, and some patients develop rapidly into acute respiratory distress syndrome (ARDS), acute respiratory failure and other serious complications ([Huang et al., 2020\)](#page-15-0). The Chinese Centers for Disease Control and Prevention identified a new type of coronavirus from a patient's throat swab sample on January 7, 2020. Subsequently, on February 7, 2020, a notice issued by the National Health Committee of China temporarily named the coronavirus-infected pneumonia a New/Novel Coronavirus Pneumonia, referred to as "New Crown Pneumonia" (NCP). On January 13, 2020, the World Health Organization temporarily referred to the coronavirus that caused the disease as 2019 new coronavirus ("2019-nCoV). On January 30, 2020, the disease caused by the virus was temporarily named "2019-nCoV acute respiratory disease" (2019 new type of coronavirus acute respiratory disease). On February 11, 2020, the World Health Organization officially named it "Coronavirus Disease 2019", abbreviated as "COVID-19". On the same day, the International Viral Classification Commission officially named the disease-causing coronavirus "severe acute respiratory syndrome coronavirus 2", abbreviated as SARS-CoV-2. According to WHO, the disease caused by Novel Coronavirus (2019-nCoV), or SARS-Cov-2 is now officially called COVID-19 [\(Huang et al., 2020a](#page-15-0); [Shen et al., 2019;](#page-16-0) [Zhang and Liu, 2020](#page-17-0)). By February 25, 2021, more than **>**120000000 cases of COVID-19 have been confirmed, with an estimated mortality risk of  $\sim$ 3.4%, which was comparatively less than that of major viral outbreaks that occurred in past years (Table 1). So far, the infection keeps spreading and more and more exported cases were confirmed in many countries worldwide, posing great pressure on public health security.

Regarding COVID-19 treatment and its spread, it is currently unclear; current knowledge is mainly based on known similar coronaviruses. CoVs are a large series of viruses that are common in many different animal species, including camels, cows, cats, and bats. Animal coronaviruses rarely infect people and then spread from person to person, such



**Table 1** 

Comparative detail on major outbreaks with fatality rate of epi- and pandemics.

Viral outbreaks	Year identified	Number infected cases	Number of deaths	Number of countries affected	Fatality rate $(\% )$
Marburg	1967	466	373	11	80
Ebola***	1976	33577	13562	9	40.4
Hendra	1994	7	$\overline{4}$	1	57
<b>H5N1</b>	1997	861	455	18	52.8
(Bird					
f <sub>1u</sub>					
Nipah	1998	513	398	$\overline{2}$	77.6
<b>SARS-CoV</b>	2002	8096	774	29	9.6
H1N1	2009	$\,>\,$	284500	214	17.4
(Swine		762630000			
flu) **					
MERS-	2012	2494	858	28	34.4
$CoV***$					
<b>H7N9</b>	2013	1568	616	3	39.3
(Bird					
f <sub>1u</sub>					
SARS-	2019	>	>	>219	$-3.4$
$CoV-2*$		120000000	2600000		

\*As of 25 February 2021, \*\* Between 2009 and 2010, \*\*\*As of November 2019 [\(CDC, 2020;](#page-14-0) [WHO, 2020](#page-17-0)).

as the respiratory system related diseases MERS, SARS, and now with SARS-CoV-2 [\(Pillaiyar et al., 2020;](#page-16-0) [Zumla et al., 2016](#page-17-0)). The most common case is transmission between close contacts (about 6 feet). Human-to-human transmission is believed to occur mainly through respiratory droplets produced when an infected person coughs or sneezes, similar to the way influenza and other respiratory pathogens spread. These water droplets can land on the mouth or nose of nearby people, or they can be inhaled into the lungs. It is unclear whether a person can contract COVID-19 by touching a surface or object, and then touching their mouth, nose or eyes. Generally, for most respiratory viruses, when patients have the serious symptoms (most sick), they are considered most infectious. It should be noted that how easy it is for the virus to spread from person to person varies depending on the type of virus. Some viruses are highly contagious (such as measles), while others are less common ([CDC, 2020](#page-14-0); [WHO, 2020](#page-17-0)). There is more to be



**Fig. 1.** Schematic taxonomical classification of coronaviruses (according to the International Committee on Taxonomy of Viruses).

<span id="page-3-0"></span>understood about the transmissibility, severity and other characteristics related to SARS-CoV-2, and the investigation is ongoing.

## **2. Signs and symptoms**

Common symptoms of COVID-19 infection include fever, cough, shortness of breath, and respiratory symptoms (Fig. 2). In more severe cases, the infection can cause pneumonia, severe acute respiratory syndrome, kidney failure, and even death. An infected person may be asymptomatic or has symptoms such as fever, cough and shortness of breath, also having diarrhea or upper respiratory symptoms, including sneezing, runny nose and sore throat [\(CDC, 2020;](#page-14-0) [WHO, 2020](#page-17-0)). According to WHO, the estimated incubation period for development of symptom after infection ranges from 1 to 14 days, with the median incubation period being 5–6 days. A study found some rare cases with an incubation period of up to 27 days ([CDC, 2020;](#page-14-0) [WHO, 2020](#page-17-0)).

# **3. SARS-COV-2 structural details**

SARS-CoV-2 (2019-nCoV) is an enveloped, single-stranded RNA, positive-sense, β-coronavirus, similar to SARS and MERS. The SARS-CoV-2 genome encodes non-structural proteins, like papain-like protease, helicase, 3-chymotrypsin-like protease, and RNA-dependent RNA polymerase, structural proteins, mainly spike glycoprotein and other accessory proteins (Fig. 3) ([McKee et al., 2020](#page-16-0)). From this point of view, the Spike (S), Envelope (E) and Membrane (M) proteins, which are located on the outer surface of the particles are also identified under electron microscope ( $Dömling$  [and Gao, 2020](#page-15-0)). A novel type of coronavirus called "Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) was identified as the cause of the respiratory disease outbreak that was first detected in Wuhan, China in 2019. The disease caused by this virus was named as Coronavirus Disease 2019 (COVID-19) ([Nile and Kai, 2021](#page-16-0)).



**Fig. 3.** The illustration is created with schematic structural details of the SARS-CoV-2 virion and its major structural proteins. Note that when observed under an electron microscope, the spikes adorned with the outer surface of the virus give rise to the corona like appearance around the virus body.

### **4. SARS-COV-2 genomic details**

Research evidence shows that SARS-CoV, MERS-CoV and SARS-CoV-2 all originated from bats. The sequence of SARS-CoV-2 is similar to that of the β-coronavirus found in bats, and the virus is genetically different from other coronaviruses, such as severe acute respiratory syndromeassociated coronavirus (SARS), member of Beta-CoV lineage B (that is, the subspecies Sabeco virus) and the Middle East respiratory system Syndrome-associated coronavirus (MERS). The genome of CoVs is a single-stranded sense RNA (+ssRNA) (~30 kb) with a 5'-cap structure and a 3′ -poly-A tail. The genome size of CoV (~30 kb) is the largest of all RNA viruses and almost twice the size of the second largest RNA virus. The maintenance of the giant genome size of CoV may be related to the



**Fig. 2.** Symptoms of COVID-19 caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2).

special characteristics of CoV RTC, which contains several RNA processing enzymes, such as the 3′ -5′ exoribonuclease enzyme of nsp14, which is unique to CoV among all RNA viruses, and has been proven to be used as a proofreading part of RTC ([Chen et al., 2020](#page-14-0)). Fig. 4 shows the schematic structure of SARS-CoV-2 in the perfusion conformation. Sequence analysis shows that SARS-CoV-2 has a typical genome structure, similar to the β-coronavirus group, including bat-SL ZXC21, bat-SARS (SL)-ZC45, SARS-CoV and MERS-CoV. Based on the phylogenetic tree of CoV, SARS-CoV-2 is more closely related to bat-SL-CoV ZC45 and bat-SL-CoV ZXC21, and is further related to SARS-CoV ([Pil](#page-16-0)[laiyar et al., 2020](#page-16-0); [Zumla et al., 2016\)](#page-17-0).

# **5. SARS-COV-2 infection and life cycle**

The spike proteins present on viral outer surface act as a key that allows the virus to enter the cells of a specific host human body. The binding of viral particle to the surface of host human cells through receptors constitutes the first step in the life cycle of coronaviruses. The steps and events involved in the life cycle of SARS-Cov-2 in human cells are shown in [Fig. 5](#page-5-0). SARS-CoV-2 virion can enter human cells through endosome or plasma membrane fusion, and the spike protein of SARS-CoV-2 mediates attachment to the host cell membrane and engages angiotensin-converting enzyme 2 (ACE2) as the cellular entry receptor ([Shereen et al., 2020](#page-16-0)). Once the virion enters the complete endosome, cathepsin L activates the spike protein, which is also activated by the cellular serine protease TMPRSS2 in close proximity to the ACE2 receptor, thereby initiating the fusion of the viral membrane and plasma membranes ([Hoffmann et al., 2020](#page-15-0)). Plasma membrane fusion entry is unlikely to trigger host cell anti-viral immunity, so it is more effective for virus replication. Once the virus enters the cell, the gene is translated from the viral genome RNA, and the virus replicates by using viral enzymes such as RNA polymerase. These enzymes are induced by the release of virus from endosomal viral RNA. In addition, the virus hijacks the host machinery, brakes transcription, replicates, and

reverse-transcribes its RNA genome for integration into host chromosome, and then reassembles, encapsulates and replicates in infected human cells ([Fehr and Perlman, 2015\)](#page-15-0). 5′ end two-thirds of the viral genome encode the polyproteins PP1a and PP1ab, which are cleaved by 3C-like protease (3CLPro) and papain (PLPro) into non-structural protein replicas. An important part of these nonstructural proteins is the RNA-dependent RNA polymerase (RdRp) that forms the replication complex [\(Fehr and Perlman, 2015;](#page-15-0) [Hoffmann et al., 2020](#page-15-0)). This replication complex performs transcription of the full-length negative strand. Then, the 3'end of the virus genome encodes four structural proteins, called spike protein (S) envelope (E) protein, nucleocapsid (N) protein and matrix/membrane (M) protein, and a set of accessory proteins ([Perlman and Netland, 2009\)](#page-16-0). When the transcription and replication of the viral RNA genome and accessory proteins are completed, the newly synthesized viral protein is trafficked from the endoplasmic reticulum to the Golgi apparatus, and then the mature virion is assembled in the budding vesicles and finally, mature virions are released through the process of exocytosis and release viral replicas outside the host cell and infect nearby cells [\(Shereen et al., 2020](#page-16-0)).

### **6. Prevention & potential therapies**

Currently, there are no any specific drugs or vaccines to prevent or treat 2019 coronavirus disease (COVID-19), as for the majority of other diseases; prevention of the infection by avoiding exposure or close contact to infected persons is the best way in the management of COVID-19. The Centers for Disease Control and Prevention recommended preventive actions to prevent the spread of COVID-19, including; avoiding close contact with infected people, touching eye, nose, mouth, and covering mouth during coughing and sneezing, staying at home in case of illness, cleaning or disinfecting objects and surfaces that are regularly touched. and CDC also recommends people with COVID-19 symptoms should use a mask to prevent the disease from spreading to others ([CDC,](#page-14-0)  [2020;](#page-14-0) [WHO, 2020\)](#page-17-0). The use of masks is also important for healthcare



**Fig. 4.** Schematic structures of SARS-CoV-2 S in the prefusion conformation. (A) SARS-CoV-2 genomic structure, with the un-translated region (UTR), open reading frame regions ORF1a and ORF1b, spike (S), envelope (E), membrane (M), and nucleocapsid (N) genes. (B) Select 2D class averages of the particles that were used to calculate the SARS-CoV-2 S reconstruction (left). Side and top views of the prefusion structure of the SARS-CoV-2 S protein with a single RBD in the "up" conformation (right). The two RBD "down" protomers are shown as cryo-EM density in either white or gray and the RBD "up" protomer is shown in ribbons (C) Schematic of SARS-CoV-2 S primary structures, colored by domain. Domains that were excluded from the ectodomain expression construct or could not be visualized in the final map are colored white.  $SS = signal$  sequence, NTD= N-terminal domain, RBD = receptor-binding domain, SD1 = subdomain 1, SD2 = subdomain 2, S1/  $S2 = S1/S2$  protease cleavage site,  $S2' = S2'$  protease cleavage site, FP = fusion peptide, HR1 = heptad repeat 1, CH = central helix, CD = connector domain, HR2 = heptad repeat 2, TM = transmembrane domain, CT = cytoplasmic tail. Arrows denote protease cleavage sites (isolate Wuhan-Hu-1, GenBank Acc MN908947).

<span id="page-5-0"></span>

**Fig. 5.** Life cycle of SARS-Cov-2 in human cell (1. Binding of spike protein to ACE2, 2: TMPRSS2 helps the virion entry, 3: The virion releases its genomic RNA 4: RNA is translated into proteins by the cell's machinery 5: Proteins forms a replication complex to make more RNA 6: Translation and RNA replication 7: Proteins and RNA are assembled into a new virion in the Golgi and released 8: Packaging of RNA synthesized 9: Virion release).

professionals and those who take care of infected individuals in a closed environment (home or medical institution). Washing hands with soap and water for at least 20 s after coughing or squeezing, use at least 60% alcohol-containing hand sanitizer. In severe cases, treatment should include care that supports vital organ functions [\(CDC, 2020](#page-14-0); [WHO,](#page-17-0)  [2020\)](#page-17-0).

Researchers, clinicians and virologists have been exploring and gaining some experience since the outbreaks of SARS-2003 and MERS-2012. The study of coronaviruses such as SARS and MERS have provided us with several potentially effective drugs. Researchers are using MERS-CoV and SARS-CoV as prototypes to evaluate COVID-19 countermeasures. Broad-spectrum antiviral drugs, such as remdesivir, lopinavir/ritonavir and interferon beta, have shown promising therapeutic effects against MERS-CoV in animal models are currently being used for treatment and prevention of SARS -CoV-2 developed COVID-19 ([Pil](#page-16-0)[laiyar et al., 2020](#page-16-0); [Zumla et al., 2016](#page-17-0)). Based on previous studies, angiotensin-converting enzyme 2 (ACE2), trans membrane protease serine 2 (TMPRSS2), spike (S) protein, RNA-dependent RNA polymerase (RdRp), angiotensin AT2 receptor, chymotripsin-like protease (3CLpro) and papain-like protease (PLpro) are considered as major targets for development of antiviral drugs against SARS-CoV-2 and another infectious coronavirus ([Zumla et al., 2016\)](#page-17-0). Doctors and scientists form different countries, are trying to use different pharmacological strategies to fight COVID-19, which include currently established antiviral drugs, different modes of oxygen therapy or mechanical aeration. Development of vaccines is crucial factor to prevent and control this COVID-19 pandemic as it plays an important role in controlling replication and spread SARS-CoV-2 through production of antibodies against virus and reducing mortality. Currently about 35 vaccine candidates have been entered into a clinical trial, few of them already approved and used against covid treatment and 145 vaccines are in the preclinical phase ([Kaur and Gupta, 2020](#page-15-0); [Rawat et al., 2021\)](#page-16-0). The COVID-19 pandemic requires rapid development of effective treatment strategies in pursuit of three concepts being applied: (1) the first method relies on testing currently known antiviral drugs and verifying their clinical effectiveness. (2) Another model is based on molecular libraries and databases, allowing high computing power and simultaneous verification of millions of potential drugs at the same time. (3) Finally, the third strategy involves targeted treatments aimed at disrupting the genome and function of the virus (Drozdzal [et al., 2020;](#page-15-0) [Lu, 2020\)](#page-16-0).

Scientists and physicians around the world have been carrying out an important campaign to understand this emerging disease and its epidemiology to in the context of identifying possible treatment options, finding effective therapeutic agents and developing vaccines. The development of a vaccine may take at least 12–18 months, and the typical schedule for approval of new antiviral therapies may exceed 10 years. Therefore, the reuse of known drugs currently being used for MERS and SARS can significantly accelerate the deployment of new COVID-19 therapies as described in this article. Here are some examples of synthetic ([Table 2\)](#page-6-0) and natural [\(Table 3](#page-11-0)) compounds used to treat SARS-CoV and related coronaviruses infection. Their chemical structures details provided in supplementary file (S1).

### *6.1. Promising antiviral, antimalarial and anti-HIV agents*

Various antiviral, antimalarial and anti-HIV agents are currently being evaluated for use to treat or prevent COVID -19 infections. Currently, several previously available drugs such as Nafamostat, Chloroquine, Hydroxychloroquine, Lopinavir; Ritonavir, Remdesivir, Favipiravir, Lopinavir/Ritonavir, Darunavir/Umifenovir, Nitazoxanide, Ribavirin, Penciclovir, Tocilizumab, Baricitinib, Arbidol, and other antiviral, antimalarial and anti-HIV agents as discussed in [Table 1](#page-2-0), with structural details provided in supplementary file (Supplementary file S1), Some of these compounds have exhibited promising results in patients and *in-vitro* clinical studies [\(Costanzo et al., 2020;](#page-15-0) [Shereen et al.,](#page-16-0)  [2020\)](#page-16-0). One of the most common treatments available for SARS-CoV-2 consists of 'cocktail therapies' based on various antivirals which are mainly protease inhibitors, the binding of which to the SARS-CoV-1 protease was predicted *in silico* and *in vitro* ([Costanzo et al., 2020](#page-15-0)). Various combinational therapies have been used by doctors and researchers for treatment of COVID-19. Thus, the previously approved drugs against MERS, SARS, Malaria and HIV were used as target agent against to block viral protease, clathrin-mediated endocytosis, inhibit the inflammatory cytokine surge, regulate immunity, reduce lung viral loads and improve pulmonary function ([Nile et al., 2020\)](#page-16-0). The anti-HIV protease inhibitory drug Kaletra, composed of ritonavir and lopinavir, showed a promising antiviral effect on SARS-CoV and SARS-CoV-2. The other anti-HIV drugs like lopinavir, ritonavir, niclosamide, promazine, and two other HIV inhibitors, PNU and UC2 were also studied as 3CLpro inhibitors of SARS-CoV, demonstrating their potential as templates for

# <span id="page-6-0"></span>**Table 2**

Commercially available remedies and drugs as possible targets for SARS-CoV-2 and related human coronavirus.





# **Table 2** (*continued* )







# <span id="page-11-0"></span>**Table 3**

Different types of natural compounds as possible targets for SARS-CoV-2 and related human coronavirus.



# **Table 3** (*continued* )



### **Table 3** (*continued* )



designing promising drug against SARS-CoV replication [\(Ghosh et al.,](#page-15-0)  [2020\)](#page-15-0).

Although there have been some preliminary positive reports on use of preexisting antiviral, antimalarial and anti-HIV drugs against treatment of COVID-19 infection, well-designed randomized, controlled clinical trials for evaluating their safety and efficacy will be necessary for the proper treatment of patients diagnosed with COVID-19 in comparison with controls who did not receive the same treatment ([Costanzo](#page-15-0)  [et al., 2020](#page-15-0)). The details on potential therapeutic remedies against COVID-19 and related human coronavirus were discussed and presented in [Tables 1 and 2](#page-2-0), with structural details provided as supplementary file (S1).

## *6.2. Nucleoside and nucleotide analogs (NAs)*

Nucleoside and nucleotide analogs (NAs) are chemically synthesized of purines and pyrimidines analogs having a heterocyclic ring or a sugar moiety. NAs are essential building blocks for nucleic acid biosynthesis and represents as the largest class of anti-inflammatory and antiviral drugs for the treatment of cancer and different viral infections ([Pruijssers and Denison, 2019\)](#page-16-0). Some NAs, including amivudine, sofosbuvir, adefovir, telbivudine, entecavir, and tenofovir (Supplementary file S1), have strong antiviral activity and have been used for the treatment of immunodeficiency virus type 1 (HIV-1), hepatitis C (HCV) and hepatitis B (HBV) infection provided proof that these class of compounds used as strong antiviral agents [\(Fung et al., 2011; Jordheim](#page-15-0)  [et al., 2013\)](#page-15-0). Over twenty NAs were approved by US FDA as antiviral drugs for use against various viral infections like; immunodeficiency virus type 1 (HIV-1), hepatitis C (HCV) and hepatitis B (HBV), human cytomegalovirus (HCMV), herpes simplex virus (HSV), varicella zoster virus (VZV)[\(Mahmoud et al., 2018\)](#page-16-0). These NAs are used to treat both acute and chronic viral infections are delivered as nucleoside and nucleotide precursors or pro-drugs, which are metabolized by host or viral kinases to their active triphosphate once inside the cell and inhibits the viral replication by non-mutually exclusive mechanisms [\(Pruijssers](#page-16-0)  [and Denison, 2019](#page-16-0)). In this review we summarized the antiviral effects of NAs, mainly remdesivir, lamivudine, amivudine, sofosbuvir, adefovir, entecavir, telbivudine, ribavirin, velpatasvir, and tenofovir against SARS-CoV-2 and related coronaviruses [\(Table 1](#page-2-0), the drug structural details provided as supplementary file (S1).

## *6.3. Protein (enzyme) inhibitors*

The SARS-CoV-2 containing positive-strand RNA causes severe respiratory syndrome in humans and responsible for COVID-19. This virus contains four structural proteins: Spike (S), Envelope (E), Membrane (M), and Nucleocapsid (N) protein ( $Fig. 3$ ). S protein plays a role in viral attachment to host cell, E and M proteins are involved in viral assembly, and N protein is needed for RNA synthesis (Dömling [and Gao, 2020](#page-15-0)). Angiotensin-converting enzyme 2 (ACE2), transmembrane protease serine 2 (TMPRSS2), spike (S) protein, RNA-dependent RNA polymerase (RdRp), angiotensin AT2 receptor, chymotripsin-like protease (3CLpro) and papain-like protease (PLpro) are considered as major targets for antiviral drugs against SARS-CoV-2 and another infectious coronavirus ([McKee et al., 2020](#page-16-0); [Zumla et al., 2016](#page-17-0)). We summarized all the synthetic and natural protein (enzyme) inhibitors used to treat SARS-CoV and related coronaviruses infection in [Tables 1 and 2](#page-2-0), respectively. The structural details of these synthetic and natural protein (enzyme) inhibitors were provided as supplementary file (S1).

### *6.4. Corticosteroids*

Corticosteroids are a class of drugs used to treat illnesses that result from inflammation and reduces immune system activity by mounting an exaggerated response to something or attacks its own cells [\(Singh et al.,](#page-16-0)  [2020\)](#page-16-0). The study reported by RECOVERY Collaborative Group showed the benefit of dexamethasone for patients with COVID-19 who were receiving mechanical ventilation at the time of randomization. Corticosteroids might be effective in preventing acute respiratory distress syndrome and death for patients having shortness of breath or requires oxygen therapy [\(The RECOVERY Collaborative Group, 2020\)](#page-16-0). Also, World Health Organization has confirmed that the corticosteroids as a potentially effective for the treatment of COVID-19, and patients' survival rates were improved significantly through the application of dexamethasone and other corticosteroids [\(Table 1\)](#page-2-0). Interestingly, most of earlier studies conducted on SARS-CoV and MERS-CoV showed adverse outcomes for use of corticosteroid in treatment ([Singh et al.,](#page-16-0)  [2020\)](#page-16-0). Indeed, the Lancet study also reported that corticosteroids should be avoided for the treatment of COVID19. However, such warnings are mainly based on the experiences in a similar viral illness but not on COVID-19 specifically ([Russell et al., 2020](#page-16-0)). Debates are continuing on potential use of corticosteroids as therapy for the treatment of acute respiratory distress syndrome (ARDS) and COVID -19. Indeed, corticosteroids have been speculated to be used as a potential therapy for ARDS as they have ability to reduce inflammation and fibrosis ([Reddy et al.,](#page-16-0)  [2020\)](#page-16-0). The various corticosteroids which are used as potential drug candidates were discussed in [Table 1](#page-2-0) and structural formulas provided as supplementary file (S1).

### *6.5. Natural products and traditional medicines*

Plant based natural products and various traditional medicines have been used as an excellent source for discovery of natural/herbal drugs, as they display great diversity among their chemical structures and wide range of biological activities [\(Wang et al., 2020](#page-17-0)). Many natural compounds are widely used as antiviral drugs shown to possess promising antiviral effects against influenza viruses, coronaviruses, herpes simplex virus, human immunodeficiency virus, severe acute respiratory syndrome (SARS), Middle East respiratory syndrome (MERS), and hepatitis B and C viruses [\(Mani et al., 2020;](#page-16-0) [Xian et al., 2020\)](#page-17-0). Numerous natural compounds have been screened *in silico* targeting various viral proteins; main protease (3CLpro, also named 3-chymotrypsin-like protease), papain like protease (PLpro), helicase, RNA-dependent RNA polymerase (RdRp), and spike protein([Mani et al., 2020;](#page-16-0) [Wu et al., 2020\)](#page-17-0). Various traditional medications based on indigenous theories and experiences are currently being used in the prevention and treatment various microbial diseases, these medicines mainly includes traditional Chinese medicine (TCM), Indian ayurvedic medicine, ancient Iranian medicine, traditional African medicine and Islamic medicine ([Wang et al., 2020](#page-17-0)). Naturally occurring agents that have potential for prevention of COVID-19 include various alkaloids, anthraquinones, terpenoids, fatty acids, flavonoids, glucosinolates, lignans, peptides, phenolics, proteins,

<span id="page-14-0"></span>saponins, and vitamins ([Wang et al., 2020;](#page-17-0) [Zhang and Liu, 2020](#page-17-0)). ([Table 3](#page-11-0) and Supplementary file S1), details about these natural products provided in [Table 2](#page-6-0) and structural details provided as supplementary file (S1). This comprehensive review provides details insights on some active natural products which are being proposed for COVID-19 drug development and prevention.

### *6.6. Convalescent plasma*

Convalescent (immune) plasma therapy refers to use of antibodies obtained from individual who has been recently recovered from particular resolution of infection and disease (Bloch et al., 2020). Convalescent plasma therapy is a passive immunization used to prevent and manage of infectious diseases and considered to be an emergency intervention in controlling several pandemics like SARS-CoV, West Nile virus, Spanish flu, Ebola virus, and recently emerged COVID-19 ([Chen](#page-15-0)  [et al., 2020](#page-15-0)). Food and Drug Administration has recently suggested that administration and study of investigational convalescent plasma therapy may provide effective clinical treatment against COVID-19 ([Rajendran](#page-16-0)  [et al., 2020\)](#page-16-0). Hence, convalescent plasma transfusion therapy has been the subject of increasing attention, especially in the wake of large-scale epidemics like COVID-19.

## **7. Conclusions**

To date, there is no any approved therapy for prevention or treatment of COVID-19, thus many scientists working on possible drug repurposing by using available different Therefore, therapies preexisting drugs including antivirals, antimalarial, immunosuppressive, antipsychotic, antidiarrheal, antidiabetic, anticancer, antifungal, antibacterial, anticoagulant, and antihelminthic agents have been suggested as potential targets preventives or therapeutics against COVID-19. However, the factors like small sample size, poor quality of drug and long completion period are not allowing obtaining reliable and there is paucity of clinical evidence for the therapeutic efficacy as well as safety of aforementioned agents for COVID-19 treatments. Development of effective therapeutic agents is subordinated to the understanding of molecular mechanisms underlying SARS-CoV-2 replication, pathogenesis and virus-host interaction.

The current available knowledge on the safety and efficacy on various therapies needs proper research, like *in vitro* studies, animal studies and clinical trials for use as potential drug against COVID-19. Several drugs currently being used and which are under clinical trials are remdesivir, umifenovir, oseltamivir, favipiravir, lopinavir/ritonavir, danoprevir/ritonavir, darunavir/cobicistat, triazavirin, hydroxychloroquine, ASC09F, baloxavir marboxil, azvudine, sofosbuvir/ledipasvir, sofosbuvir/daclatasvir, and emtricitabine/tenofovir [\(Table 1](#page-2-0) 2). Also, clinical trials are undergoing for various natural compounds like heparin and vitamin C as therapeutic agents or immune boosters in against COVID-19 infection [\(Table 2](#page-6-0) 3). Thus, application of the existing potential candidate therapies may represent an effective strategy for the identification of new pathways and targets for intervention of SARS-CoV-2 infection and pathogenesis. In order to effectively deal with the current strategies, needs further exploration to determine the effective agent/therapies for modifying research conduct for this COVID -19. The safety and efficacy of various suggested COVID-19 therapies needs proper systematic research, coordinated by both preclinical studies and clinical trials.

### **CRediT authorship contribution statement**

**Shivraj Hariram Nile:** colleted data, Writing – original draft, Writing – review & editing. **Arti Nile:** collected information on this topic and provided all chemical structural details. **Shivkumar Jalde:**  collected information on this topic and provided all chemical structural details. **Guoyin Kai:** Writing – review & editing, All authors read and approved the final version of this manuscript.

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## **Declaration of competing interest**

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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### **Appendix A. Supplementary data**

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