

Phytotherapeutic Evidence Against Coronaviruses and Prospects for COVID-19

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ABSTRACT

The emergence of the novel β -coronavirus (SARS-CoV-2) and subsequent outbreak of COVID-19, is a global health challenge with no known treatment to date and has culminated in significant morbidity and mortality. This article highlights current understanding on SARS-CoV-2 based on the available scientific evidence on human coronavirus (HCoV) infections, which could offer novel insights and therapeutic targets for SARS-CoV-2, the causative agent of COVID-19. Specifically, the paper presents available phytotherapeutic evidence against pathogenic HCoVs with a view to identifying potent plant-derived antiviral agents that could be developed to aid the fight against coronaviruses and the current COVID-19. Evidently, elucidation of CoV integral proteins such as the spike protein, angiotensin-converting enzyme 2, 3C-like cysteine protease and papain-like protease, as good targets for drug developments has lent credence to the use of medicinal plants or their metabolites as prophylaxis or treatment interventions in CoV infections and holds promising ground for SARS-CoV-2. While some promising phytocompounds are currently under clinical trials for COVID-19, increased research into plants and in-depth characterization of their metabolites could reveal more interesting results that would benefit humanity in its fight against emerging and re-emerging viral infections including the current COVID-19. Overall, given the current body of evidence on the potential development of phytotherapeutics for COVID-19, fears need to be allayed while clinical trials continue. Conclusively, the lockdown and other preventive measures which have been implemented in most parts of the world should be humanely exercised and supported to ensure compliance and safety of lives.

Key words: Coronavirus; COVID-19; Antivirals; Drug target; Natural products; Plants; Plant metabolites; SARS-CoV-2.

INTRODUCTION

Globally, viruses are integral groups of etiological agents associated with significant morbidity and mortality.^{1, 2} Interventions such as the use of antiviral agents including drugs and vaccines are normally employed to reduce the rate and extent of comorbidities due to viral infections in animals and humans.³ Such interventions, however, remain under trial for use as palliatives to fight the current severe acute respiratory syndrome (SARS) infection called COVID-19. COVID-19 is caused by a novel coronavirus (CoV) named SARS-CoV-2.⁴ However, CoVs are not new in the emerging infections terrain and have been generally recognized as a large group of related viruses of medical and veterinary significance. The first Coronavirus-Infectious Bronchitis Virus (IBV) - was isolated in 1931 by Schalk and Hawn, who thought it to be a respiratory disease of chicks.⁵ The studies that led to their characterization and acknowledgment as an agent of medical importance started in 1965, with the work of Tyrrell and Bynoe. Whilst working with IBV, mouse hepatitis virus and transmissible gastroenteritis virus of swine, the viruses were found to be morphologically similar under electron microscope. They were, thus, dubbed a new group of viruses in 1968, with the name 'Coronaviruses',⁶

but only accepted as a new genus of viruses in 1975.⁷ While the initial classification of the genus was mainly based on microscopic examination,⁸ the advent of advanced molecular methods, such as nucleic acid amplification technologies, automated DNA sequencing and bioinformatics, have now elucidated CoVs as pleomorphic, enveloped viruses with a positive-sense single-stranded-RNA genome and a nucleocapsid of helical symmetry.⁹ Compared to other RNA viruses, CoV has the largest viral genome of about 27 to 32 kb in length, with about 65-125 nm in diameter,^{10,11} and can diffuse among mammals and humans.¹² This large genome confers them with high recombination rates by constantly developing transcription error and RNA-Dependent-RNA-polymerase (RdRP) jumps.¹³ Structurally, CoVs are made up of nucleocapsid protein (N) which forms a complex with the RNA, the surface glycoprotein (S) which forms the petal-shaped surface projection (spike) that is responsible for virus entry into host cells, the membrane glycoprotein (M) which gives the virion its shape, and the envelop glycoprotein (E) that is involved in virion assembly¹⁴ (Figure 1).

Just like the recent emergence of SARS-CoV-2 in December 2019 in Wuhan, a similar occurrence in the later part of the year 2002 also witnessed SARS infection that emerged in the Southern China

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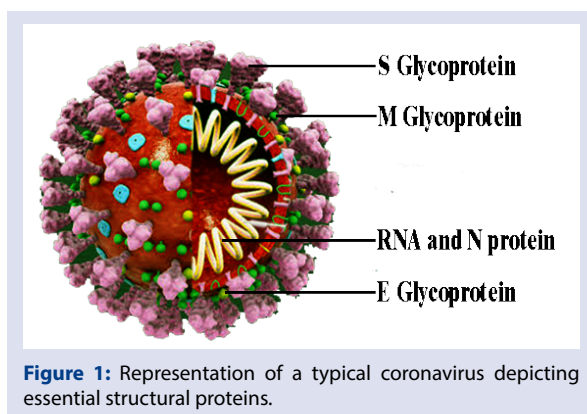


Figure 1: Representation of a typical coronavirus depicting essential structural proteins.

territory. This was reportedly caused by the CoV called SARS-CoV that caused significant epidemic claiming almost a thousand deaths and lasted till 2003.^{15, 16} Generally, CoVs constitute a genus within the family Coronaviridae, order Nidovirales and subfamily Orthocoronavirinae.¹⁷ To date, four CoV genera (alpha, beta, delta and gamma) have been elucidated, with human CoVs (HCoVs) identified in the alpha-CoV (HCoV-229E and HCoV-NL63) and beta-CoV (MERS-CoV, SARS-CoV, HCoV-OC43 and HCoV-HKU1) genera.¹⁸ The HCoV-229E, HCoVNL63, HCoV-OC43 and HCoV-HKU1 strains cause respiratory tract infections (RTIs) that are mostly non-virulent, mild and self-limiting like the common cold, while SARS-CoV or SARS-CoV-2 and MERS-CoV are associated with severe RTIs which may either cause significant morbidity or lead to death.¹⁹ The severe respiratory diseases caused by these viruses could also culminate into enteric infections or ‘cytokine storm’, an event which is indicative of an overreaction in immune responses.^{20,21}

For antiviral drug development, viral penetration and its subsequent replication are usually the targets and these processes have been well established on integral proteins of CoVs. The most studied of the CoV integral proteins which make good targets for drug developments are the spike (S) protein, 3C-like protease (3CLpro) and papain-like protease (PLpro). Notably, studies have shown the capability of natural compounds to inhibit these proteins associated with SARS or MERS CoV infections. Following the release of the SARS-CoV-2 gene sequence which showed high similarities with SARS or MERS proteins, it was theorized that existing effective anti-MERS or anti-SARS interventions could be invaluable in the race to find a drug that is anti-SARS-CoV-2 to fight the COVID-19 infection.¹⁷ SARS-CoV-2 and SARS-CoV also bind to the same host cell receptor, but SARS-CoV-2 binds more easily and tenaciously, and this could give an insight into why COVID-19 seems to be more effectively spread through human-to-human interaction. However, more research data is still required to support this insight.²²

Generally, although, CoV infections are controllable, there remains the possibility of sporadic new cases if humans continue to encounter their animal hosts.^{23, 24} Therefore, to ensure some level of preparedness, active antiviral moieties which abound in medicinal plants could form a formidable force to fight such outbreaks, especially when control measures fail. Hence, science may have to resort to nature, plants, their metabolites and/or extracts for some level of succour.^{25,26} While, ethnomedicinal plants have been shown to possess antiviral activity over the years,²⁷⁻³⁰ the demand for new antivirals continue to exist and increase given the increasing prevalence and emergence of novel viral infections like human immunodeficiency virus (HIV) and SARS-CoV. This article, therefore, presents available phytotherapeutic evidence against pathogenic HCoVs with a view to identifying potent plant-derived antiviral agents that could be developed to aid the fight against CoVs and the current global challenge of COVID-19.

MATERIALS AND METHODS

The online resources and database range for this article included PubMed, Google Scholar, MeSH, ScienceDirect, National Institute of Health (NIH) and Nation Centre for Biotechnology Information (NCBI) web resources. Word combinations and phrases pertinent to the subject under review were used. Some of these include coronaviruses, emerging topics on COVID-19, plant-based antivirals against coronaviruses, plant antiviral agents, plant metabolites for CoV prophylaxis, potent plant metabolites for CoV treatment, potential plants and their antiviral activity against CoVs, among others. Records and reports included relevant information from inception till May 2020 to help fine-tune results for thorough discussion on the appraised topic. Following the gathering of data, a mechanistic model (Figure 2) was generated to show inclusion and exclusion basis used for screening research records.

RESULTS AND DISCUSSION

Mode of transmission and life cycle of SARS-CoVs and MERS-CoV

Different strains of CoVs infect different hosts, and they normally deploy species-specific approaches for attachment and entry into the host cell. Studies have identified how CoVs enter the host cells and the definite manner through which each species of CoVs interact with the unique cellular receptors of the host. Such efforts have found that CoVs demonstrate a complex pattern for receptor recognition.¹⁴ Even when different species of CoVs infect similar hosts, they usually vary in the degree of diseases they cause, including acute, persistent, severe and highly lethal infections.³¹ While bats have been recognized as the reservoir for SARS-CoVs and MERS-CoV, their intermediate hosts differ considerably ranging from palm civets, dromedary camels to Malayan pangolin for SARS-CoV, MERS-CoV and SARS-CoV-2, respectively.³²⁻³⁴

For CoVs, the S glycoprotein has two subunits (S1 and S2) and has been reportedly responsible for binding to host-receptor on the cell surface, and consequently entry into the cell. While the S1 subunit contains N- and C-terminal domains (S1-NTD and S1-CTD, respectively),

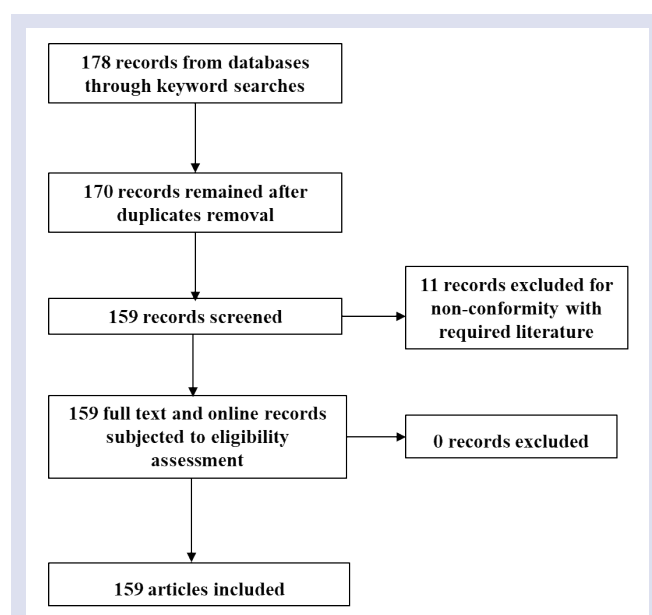


Figure 2: PRISMA flow chart showing total records, screened, included and excluded materials for this review.

both of which are receptor-binding domains (RBD), the S2 subunit, an elongated structure which forms the stalk, is mainly involved in ensuring attachment of viral envelop to the target cell membrane.³⁵ To facilitate entry, the S1-CTD of MERS-CoV and SARS-CoVs interact with dipeptidyl peptidase 4 (DPP4)³⁶ and angiotensin-converting enzyme 2 (ACE2), respectively.³⁷ The ACE2 is an ectoenzyme anchored to the plasma membrane of the cells of several tissues, especially the lower respiratory tract, heart, kidney and gastrointestinal tract.³⁶ Following S1-CTD's interaction with the receptor, the viral nucleocapsid gets deposited into the cytoplasm, where consecutive replication, assembly and release of a new viral particle occurs.³⁸ RNA replication in MERS-CoV and SARS-CoVs takes advantage of their open reading frames (ORFs) with the involvement of replicase genes (rep1a and rep1ab), a slippery sequence (5'-UUUAAAC-3') and polyproteins (pp1a and pp1ab). The polyproteins encode important non-structural proteins (NSP1-11 and NSP1-16) of the beta-CoVs. Specifically, the RNA replication which occurs on double-membrane vesicles (DMVs),³⁷ involves the positive stranded RNA genome as a template which facilitates the production of the negative strand RNA. Using the replicase gene encoded enzymes, the negative stranded RNA genome was used to produce overlapping mRNA molecules which, subsequently, gets translated into the four structural proteins (N, M, E and S). While the NSPs serve to assemble the RNA into a helical twisted structure,³² the

membrane-bound structural proteins get packed into the endoplasmic reticulum (ER) before translocation into endoplasmic reticulum-Golgi intermediate compartment (ERGIC). The nucleocapsids formed from the encapsidation of progeny genomes by N protein are then merged with the membrane-bound components, forming virions by budding into the ERGIC. Subsequently, the new virions get exported by Golgi bodies and are exocytosed into the extracellular space of the host cell,³¹ that allows attack by virions and possible transmission to other individuals (Figure 3).

Person-to-person transmission of SARS-CoV is through respiratory droplets, close contact with infected persons, faecal-oral route and aerosols. MERS-CoV is transmitted through respiratory droplets, close contact with diseased patients/camels and ingestion of camel milk. For SARS-CoV-2, current data shows major transmission routes are droplets transmission, contact transmission, with possibility of faecal-oral and aerosol transmission.³⁴ Table 1 presents an overview of the characteristic clinical features of MERS-CoV and SARS-CoVs.

Some potent drugs, metabolites and interventions against SARS-CoV-2

The recent outbreaks of viral diseases such as Ebola, Zika and SARS-CoV-2, among others called for the development of new host-targeted

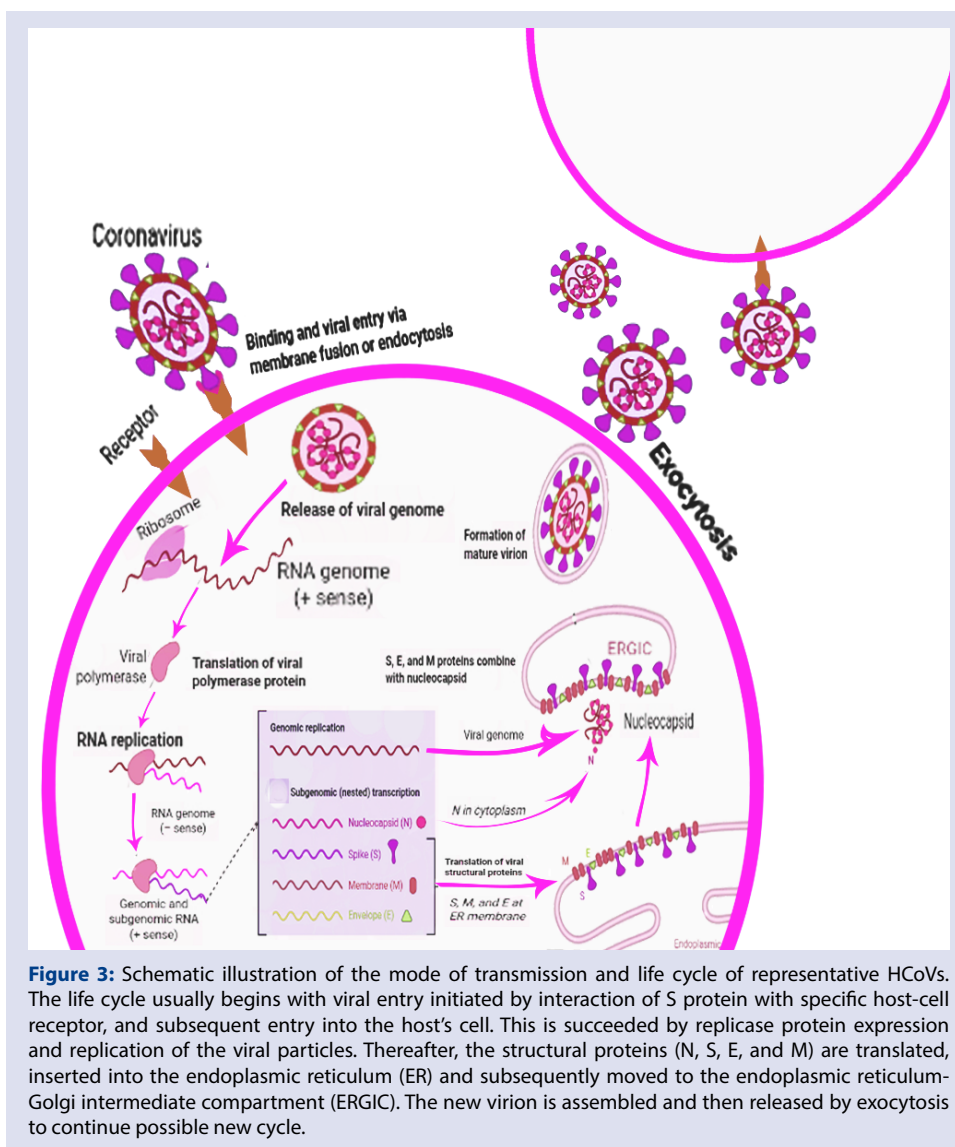


Figure 3: Schematic illustration of the mode of transmission and life cycle of representative HCoVs. The life cycle usually begins with viral entry initiated by interaction of S protein with specific host-cell receptor, and subsequent entry into the host's cell. This is succeeded by replicase protein expression and replication of the viral particles. Thereafter, the structural proteins (N, S, E, and M) are translated, inserted into the endoplasmic reticulum (ER) and subsequently moved to the endoplasmic reticulum-Golgi intermediate compartment (ERGIC). The new virion is assembled and then released by exocytosis to continue possible new cycle.

Table 1: Comparative representation of biological features of SARS-CoVs and MERS-CoV.

Virus	SARS-CoV	MERS-CoV	SARS-CoV-2
Disease caused	SARS	MERS	COVID-19
Emergence	2002	2012	2019
Reservoir host	Bat	Bat	Bat
Intermediate host	Palm Civet	Dromedary Camel	Malayan Pangolin?
Incubation period	2-7 days	5-6 days	2-14 days
Host receptor	ACE2	DDP4	ACE2
Case fatality rate	9.50%	34.40%	7.10%
Symptoms	Fever, dry cough, headache, difficulty in breathing, muscle aches, loss of appetite, diarrhoea	Fever, chills, diarrhoea, nausea, vomiting, congestion, sneezing, sore throat	Fever, cough, shortness of breath, fatigue
Complications	Heart, liver and respiratory failure in adverse condition.	Acute pneumonia and kidney failure in adverse condition.	Acute pneumonia, septic shock and respiratory failure in adverse condition

therapeutics, as well as a revisit of the drug reprofiling/repositioning methodology (DRM). DRM implies producing an added value from pre-existing medications and is usually achieved by channelling known drugs to another disease/infection besides that for which it was originally targeted.^{39,40} The major advantages of DRM over new drug formulations are that the pre-existing drug production process is already established, safe, reliable, and there is also reduced cost and timeline to clinical availability, plus the significantly improved success to market probability. In addition, early drug developmental phases and drug-interaction properties from animal models used in pre-clinical trials are readily available.⁴¹⁻⁴²

As part of DRM, compounds which target viruses belonging to two or more viral families known as broad-spectrum antiviral agents (BSAAs) which are safe for use in humans have been proposed to provide herd protection against re-emerging and emerging viral infections while also broadening possible antiviral options.¹⁹ Some notable examples of BSAAs include emetine (an antiprotozoal capable of inhibiting HCoV-OC43, Zika and Ebola viruses and HIV-1 infections), enoxacin (anti-Zika and anti-HIV-1),⁴³ amodiaquine (antimalarial and anti-Zika),⁴⁴ and niclosamide (anthelminthic and anti-Zika).⁴⁵⁻⁴⁸ Several other BSAAs including chloroquine, azithromycin, cyclosporine, rapamycin, mycophenolic acid, ezetimibe and nitazoxanide are currently undergoing phase IV surveillance studies under this concept.¹⁹ A wide assortment of safe BSAAs which could be subjected to further studies as potential SARS-CoV-2 drug candidates are available from the database at <https://drugvirus.info/>. Study data must however be fused and harmonized for greater efficiency.¹⁹

Another therapeutic is the cyclooxygenase (COX) mediator/inhibitor, indomethacin. Indomethacin/indocin has been shown to be an effective inhibitor of SARS-CoV-2 *in vitro*.⁴⁹ It is a non-steroidal anti-inflammatory drug (NSAID) with good antiviral potential against canine CoV *in vivo*.⁴⁹ It also showed similar potential against human SARS-CoV.¹⁶ In a study involving canine models, the faecal shedding of the control dog group increased steadily and peaked on the seventh day post-infection, while the viral titre in the indomethacin-treated group reduced drastically after resumption of treatment post-infection. Viral shedding in the treated dog group reached minimal levels at day seven post-infection.¹⁶

Again, human monoclonal antibodies (MAb) targeting the viral spike protein may be useful prophylactically against SARS. Fusion inhibitors like enfuvirtide which is used in HIV may be redesigned for SARS-CoV. Nonpeptide inhibitors and peptide mimics which target viral 3C-like cysteine protease (3CLpro) in CoVs with high selectivity index may show promise for SARS-CoV inhibition. Short interfering RNAs (ribavirin), α -interferon and β -interferon could also show promise for SARS control. Also, compounds with known

in vitro inhibition of SARS such as calpain inhibitors, valinomycin,⁵⁰ glycopeptide antibiotics, nelfinavir,⁵¹ plant lectins, aurintricarboxylic acid and hesperetin⁵² could be subjected to further studies targeting SARS-CoV-2. Other identified anti-SARS-CoV agents include HIV protease inhibitors,⁵³ anthraquinone-based compound,⁵⁴ carbohydrate-binding agents,⁵⁵ an antipsychotic⁵⁴ and a nucleoside analogue.⁵⁶ Hsieh et al.⁵⁷ further expanded the list from their report on the successful test of a combination therapy that involved nelfinavir and agglutinin from *Galanthus nivalis* which showed synergistic antiviral efficacy, although used in the treatment of feline SARS-CoV. Nelfinavir is a known safe anti-HIV-1 protease inhibitor with remarkable activity *in vivo*⁵⁸ and its efficacy in SARS-CoV infection has been demonstrated since 2004.⁵³

Agglutinin from *G. nivalis* belongs to a class of antivirals called carbohydrate-binding agent and has the ability to bind to both the membrane and spike proteins of CoVs.⁵⁹ As the foremost study that showed anti-HIV-1 protease inhibitors as effective blockers of feline CoV replication, it serves as a classical and successful example of DRM.⁵⁴ A CoV-membrane active compound denoted as K22 was reported to actively interfere with the DMVs formation, a process which is integral to the establishment of CoV replication. This effect was followed by a significant inhibition of viral RNA synthesis in MERS and HCoV-229E. This research finding points future studies toward a new drug development target for the treatment of pathogenic HCoVs including SARS-CoVs.⁶⁰

With clinical trials underway, a reprofiled Ebola medication, remdesivir, seems to show great promise for use in COVID-19 therapy compared to other SARS-CoV-2 test drugs. Remdesivir is a nucleotide analogue thought to act by halting the ability of the virus to replicate, thus reducing the infection of healthy body cells. Nonetheless, a study in China showed remdesivir as being more effective against SARS-CoV-2 when combined with chloroquine. Additional data on the drug efficacy and safety are however expected soonest.^{61,62} Another oral pill intervention being tested is favipiravir known as Avigan and is favoured over remdesivir which is intravenously injected.⁶³ Other promising antiviral drugs undergoing test and clinical trials include SPL7013 from Starpharm, an antiviral dendrimer,⁶⁴ Kaletra (a combination of lopinavir and ritonavir), EIDD-2801 (similar action as remdesivir and may be available as a pill), ivermectin, Actemra, Kevzara and Calquence. The use of stem cells, blood plasma from recovered COVID-19 patients and pluristems are also being investigated for potential use in COVID-19 prophylaxis or treatment.⁶⁵

Antibody-based treatments capable of preventing viral entry and infection like SAB-301 have also shown great promise. Again, MABs such as REGN3048 and REGN3051 were shown in initial trials to be well tolerated and are at advanced stages of clinical trial.⁶⁶ With regards to vaccines for SARS-CoV-2, the National Institute of Allergy

and Infectious Diseases (NIAID) in conjunction with other research institutes would be conducting clinical trials for a vaccine named mRNA-1273. The vaccine would however not be publicly available till 2021.⁶⁷ Another vaccine in form of a microneedle patch called PittCoVacc, developed by the University of Pittsburgh School of Medicine has been shown to evoke the production of anti-SARS-CoV2 antibodies within two weeks of the patch prick.⁶⁸ Several other vaccines are also being designed with possible clinical trials underway.

Plants and plant secondary metabolites as antiviral agents and prospects for COVID-19

Antiviral agents and prospects for COVID-19

Natural products from plants including plant extracts and plant-derived compounds have wide applications as nutraceuticals and with potential use in the prevention and treatment of several communicable and non-communicable diseases.⁶⁹ Unsurprisingly, synthesis of conventional drugs is greatly dependent on medicinal plants and a staggering one-quarter of the commonly used conventional drugs were originally synthesised from plant-derived compounds.⁷⁰ A typical example is chloroquine phosphate, a structural analogue of quinine derived from the bark of the *Cinchona* tree. This drug is traditionally used for the treatment of malaria however, it has been reported to exert considerable antiviral and immunomodulating properties.^{69,71} Another antimalarial drug, artemisinin is obtained from *Artemisia annua*.⁷² Again, while emetine, an amoebicidal drug is obtained from *Cephaelis ipecacuanha*, others such as quinidine, topotecan, taxol, morphine, aspirin, digitalis and colchicine are derived and developed from herbal plants.^{69,73-75} Although, modern pharmaceuticals are effective for the management and treatment of various viral diseases, the emergence of antiviral-resistant mutants and occurrence of novel viral strains (e.g. SARS-CoV-2) with no known effective cure and vaccine, have undermined their applications.^{38,76} Moreover, some antiviral drugs are expensive, toxic and not broad-spectrum based. Hence, there is a continuous demand for novel, highly potent, less toxic and cost-effective antivirals which could be proficiently fulfilled by medicinal plants.

Medicinal plants have rich bioactive components which could be explored for therapeutic leads. In fact, in recent times medicinal plants and their products have gained greater attention and a variety of herbs have been investigated for their antiviral potential. Extracts of plants such as *Lycoris radiata*, *Artemisia annua*, *Pyrrhosia lingua*, and plant derived compounds such as apigenin, berbamine, lycorine and glycyrrhizin are not only potent antiviral agents, but they also possess remarkable anti-human coronaviral (HCoV) properties. These make them excellent lead compounds for novel antiviral drug development, especially amidst the current COVID-19 pandemic.⁷⁷⁻⁸⁰ Natural compounds may confer antiviral activity via the modulation of immune system against the virus or through direct inhibition or blockage of viral entry, replication, infectivity, reverse transcription, protein expression, assembly, release or host-specific interactions.^{32,79}

Interestingly, traditional or herbal medicine has been previously employed to treat HCoVs, and studies have confirmed its efficacy either as a holistic intervention or as combined therapy with conventional medicine.^{78, 81, 82} For instance, during the SARS-CoV outbreak in 2002, Cinatl and co-workers⁷⁷ evaluated the antiviral potential of liquorice roots-derived compounds such as ribavirin, 6-azauridine, pyrazofurin, mycophenolic acid and glycyrrhizin against two clinical isolates of SARS-CoV (FFM-1 and FFM-2) isolated from German patients in Vero cell cultures. Of the five compounds, glycyrrhizin had the most potent activity with 50% effective concentration (EC_{50}) of 300 mg/L and selectivity index of 67. The mode of action of glycyrrhizin against SARS-CoV was obscure; however, it could be due to its induction of nitrous oxide synthase since nitrous oxide is a known inhibitor of replication in several viruses (e.g. Japanese encephalitis virus).⁸³ The *Camellia*

sinensis plant has been reported to contain theaflavin,⁸⁴ water soluble tannic acid and theaflavin-3-gallate⁸⁵ which contribute to its antiviral function against rotavirus and SARS-CoV. *Eleutherococcus senticosus* containing theaflavin and catechin has also been shown to be effective against rotavirus and HCoVs.^{84,86} Glycyrrhizin is also a good inhibitor of hepatitis B virus; it is capable of stimulating endogenous production of interferons and with remarkable antioxidant activity.⁸⁷ Similarly, a further study by Hoefer and co-workers⁸⁸ reported the increased anti-SARS-CoV activity of chemically modified glycyrrhizin derivatives, however the modified derivatives had increased cytotoxicity and reduced selective index compared to the original compound.⁸⁸

Ginsenoside-Rb1, a steroid from *Panax Ginseng*, a traditional Chinese medicine, shows considerable activity against SARS-CoV at a concentration of 100 μ M. The same study also demonstrated the anti-SARS-CoV potential of aescin and reserpine, with EC_{50} values of 6.0 μ M and 3.4 μ M, respectively.⁵⁰ Similarly, baicalin, a flavonoid from another traditional Chinese medicine, *Scutellaria baicalensis* has also been reported to possess anti-SARS-CoV activity.⁸⁹ Furthermore, the anti-HCoV-229E activity of glucosidic compounds, saikosaponins (A, B₂, C and D) has been shown. The strongest potency was displayed by saikosaponin B₂ and its mechanism of action was attributed to the interference of early stages of viral replication such as viral attachment, adsorption and penetration.⁹⁰ In another study by Yi and co-workers⁹¹ that involved the screening of small molecules from 121 Chinese herbs extracts resulted in the identification of two molecules, tetra-O-galloyl- β -D-glucose (TGG) and luteolin from *Galla chinensis* and *Rhodiola kirilowii*, respectively with substantial effects on SARS-CoV. The proposed mechanism of action of these two compounds was via the blockage of viral entry. The high selective index (SI) value of TGG (SI: 240) compared to luteolin (SI: 24) might make it a better lead compound, since this means that it can be used at high concentration with no significant cytotoxic effects.⁹¹ A high throughput screening of 200 Chinese medicinal herb extracts has demonstrated the significant anti-SARS-CoV activity of *Lycoris radiata* (ethanolic extract), *Artemisia annua* (ethanolic extract), *Pyrrhosia lingua* (chloroform extract), and *Lindera aggregata* (ethanolic extract) with EC_{50} ranging from 2.4 ± 0.2 to 88.2 ± 7.7 μ g/ml (Table 2). Further fractionation and purification of the alkaloid components of the most potent extract, *Lycoris radiata* (EC_{50} : 2.4 ± 0.2 μ g/ml) resulted in the isolation of the active ingredient of the herb, lycorine with EC_{50} of 15.7 ± 1.2 mM. Its remarkable SI value greater than 900 also makes it a good lead compound for future drug design.⁷⁸

With the use of a Vero E6 cell-based cytopathogenic effect (CPE) assay, the anti-SARS-CoV potential of 221 compounds was determined by Wen and co-workers.⁹² The study demonstrated that ten diterpenoids which include ferruginol, dehydroabieta-7-one, sugiol, cryptojaponol, 8 β -hydroxyabieta-9(11),13-dien-12-one, 7 β -hydroxydeoxycryptojaponol, 6,7-dehydroroyleanone, 3 β -,12-diacetoxyabieta-6,8,11,13-tetraene, pinusolidic acid, forskolin; two sesquiterpenoids viz., cedrane-3 β ,12-diol, α -cadinol; two triterpenoids namely betulonic acid, betulonic acid; five lignoids hinokinin, savinin, 4,4'-O-benzoylolariciresinol, honokiol, magnolol; and a phenolic compound, curcumin had remarkable anti-SARS-CoV activity at concentrations between 3.3 and 10 μ M. All of these phytochemicals, except sugiol and 4,4'-O-benzoylolariciresinol markedly inhibited SARS-CoV replication. Moreover, the SI values of ferruginol (SI: 58), dehydroabieta-7-one (SI: 76.3), 8 β -hydroxyabieta-9(11),13-dien-12-one (SI: >510), 7 β -hydroxydeoxycryptojaponol (SI: 111), 3 β -,12-diacetoxyabieta-6,8,11,13-tetraene (SI: 193), pinusolidic acid (SI: >159), forskolin (SI: 89.8), betulonic acid (SI: 180), and savinin (SI: >667) were higher than that of the positive control (valinomycin, SI: 41.4) used in the study, thus highlighting their potential as good lead compounds.

In some study, an aqueous extract of the leaf of *Toona sinensis*

Table 2: Medicinal plants with anti-coronaviral potential.

SN	Plant	Family	Plant part	Extract	Metabolite or extract (Dose)	EC ₅₀	Proposed mechanism	HCoV	Reference
1.	<i>Artemisia annua</i>	Asteraceae	Whole plant	Ethanol	Artemisinin (10 ⁻¹ - 10 ⁻⁴ mg/ml)	34.5 ± 2.6 µg/ml	Unclear	SARS-CoV	78
2	Cassiae Semen extract (<i>Cassia tora</i>)	Fabaceae	Seed	n-hexane	Na (0 - 10 µg/ml)	8.43 µg/ml	Inhibition of SARS-CoV 3CL protease activity	SARS-CoV	96
3.	Dioscoreae Rhizoma extract (<i>Dioscorea batatas</i>)	Dioscoreaceae	Tuber	Methanol	Na (0 - 10 µg/ml)	8.03 µg/ml	Inhibition of SARS-CoV 3CL protease activity	SARS-CoV	96
4.	Gentianae Radix extract (<i>Gentiana scabra</i>)	Gentianaceae	Rhizome	n-hexane	Na (0 - 10 µg/ml)	8.70 µg/ml	Inhibition of viral replication	SARS-CoV	96
5.	<i>Lindera aggregata</i>	Lauraceae	Root	Ethanol	Na (10 ⁻¹ - 10 ⁻⁴ mg/ml)	88.2 ± 7.7 µg/ml	Unclear	SARS-CoV	96
6.	Loranthi Ramus extract (<i>Taxillus chinensis</i>)	Loranthaceae	Stem, leaf	n-hexane	Na (0 - 10 µg/ml)	5.39 µg/ml	Inhibition of viral replication	SARS-CoV	96
7.	<i>Lycoris radiata</i>	Amaryllidaceae	Stem cortex	Ethanol	Lycorine (10 ⁻¹ - 10 ⁻⁴ mg/ml)	2.4 ± 0.2 µg/ml	Unclear	SARS-CoV	78
					Lycorine (0 - 20 µM)	0.15 to 1.63 µM		HCoV-OC43, HCoV-NL63, MERS-CoV, MHV-A59	99
8.	<i>Pyrrosia lingua</i>	Polypodiaceae	Leaf	Chloroform	Na (10 ⁻¹ - 10 ⁻⁴ mg/ml)	43.2 ± 14.1 µg/ml	Unclear	SARS-CoV	78
9.	Rhizoma Cibotii extracts (<i>Cibotium barometz</i>)	Cibotiaceae	Rhizome	Ethanol and methanol	Na (0 - 10 µg/ml)	8.42 and >10 µg/ml	Inhibition of SARS-CoV 3CL protease activity	SARS-CoV	92
10.	<i>Strobilanthes cusia</i>	Acanthaceae	Leaf	Methanol	Na (0 - 10 µg/ml)	0.64 ± 0.43 µg/ml	Virucidal activity	HCoV-NL63	100
11.	<i>Glycyrrhiza glabra</i> L., <i>Glycyrrhiza uralensis</i>	Fabaceae	Root	Na	Glycyrrhizin (300 mg/L)	Na	Virucidal activity	SARS-CoV	77,101,102
12.	<i>Houttuynia cordata</i> Thunb.	Saururaceae	Whole plant	Aqueous	Na	Na	Increase the activity CD4+ and CD8+ of T cells; inhibition of protease (3CLpro); RNA dependent RNA polymerase (RdRp) activity.	SARS-CoV	103
13.	<i>Toona sinensis</i> Roem	Meliaceae	Leaf	Aqueous	Quercetin (Na)	Na	Unclear	SARS-CoV	91,93
14.	<i>Pelargonium sidoides</i> DC.	Geraniaceae	Root	Na	Catechin and gallic acid (Na)	Na	Inhibit replication of H1N1, H3N2 virus strains, respiratory syncytial virus, and human coronavirus	SARS-CoV	95,104
15.	<i>Rheum officinale</i> Baill	Taxaceae	Root tubers	Aqueous	Emodin (Na)	Na	Inhibited the interaction of SARS-CoV S protein and ACE2	SARS-CoV	54
16.	<i>Polygonum multiflorum</i> Thunb.	Polygonaceae	Root tuber and vine	Aqueous	Emodin (Na)	200 µM	Inhibited the interaction of SARS-CoV S protein and ACE2	SARS-CoV	54
					Juglanin (10 - 40 µM)	2.3 µM			105
17.	<i>Anthemis hyalina</i> DC.	Asteraceae	Flower and bud	Ethanol	Flavanoids. (Na)	Na	Inhibit the replication and expression of coronavirus and TRP gene	SARS-CoV	98
18.	<i>Nigella sativa</i> L.	Ranunculaceae,	Seed	Ethanol	Thymoquinone, α-Hederin and Nigellidine (Na)	Na	Inhibit the replication and expression of coronavirus and TRP gene	SARS-CoV	98,106

19.	<i>Citrus sinensis</i> (L.) Osbeck	Rutaceae	Peel	Ethanol	Carvacrol and α -pinene (Na)	Na	Inhibit the replication and expression of coronavirus and TRP gene	SARS-CoV	98
20.	<i>Torreya nucifera</i> (L.) Siebold & Zucc.	Taxaceae	Leaf	Ethanol	Luteolin, Quercetin and Apigenin (Na)	20, 23.8 and 280.8 μ M	Inhibition of 3CLpro	SARS-CoV	107
21.	<i>Euphorbia neriifolia</i> L.	Euphorbiaceae	Leaf	Ethanol	Triterpenoids (0 - 25 μ M), flavonoid glycoside (Na). Hinokinin (0 --10 μ M)	Na	Virucidal activity	HCoV	97, 108
	<i>Phyllanthus urinaria</i>	Phyllanthaceae							92
22.	<i>Isatidis indigotica</i>	Brassicaceae	Root	Aqueous	Sinigrin, Indigo, β -sitosterol and Aloe-emodin (Na), Hesperetin (Na)	Na	anti-SARS-CoV 3CLpro	SARS-CoV	78, 79
23.	<i>Scrophularia scorodonia</i>	Scrophulariaceae	Na	Na	Saikosaponin A, B2, C and D (0 - 25 μ M)	Na	Virucidal activity	HCoV-22E9	90
24.	<i>Panax Ginseng</i> <i>Aesculus hippocastanum</i> <i>Rauwolfia species</i>	Araliaceae	Root	Na	Ginsenoside-Rb1. (Na) Aescin (0 - 20 μ M) Reserpine (0 - 20 μ M)	100 μ M 6.0 μ M 3.4 μ M	Inhibition of viral replication	SARS-CoV	50
25.	<i>Cinnamomum</i> sp.	Lauraceae	Cortex	Na	Procyanidin B1 (0 - 500 μ M) Procyanidin A2 (0 - 500 μ M) Cinnamtannin B1 (0 - 500 μ M)	41.3 \pm 3.4 μ M 29.9 \pm 3.3 μ M 32.9 \pm 3.9 μ M	Inhibition of pseudovirus infection	SARS-CoV	109
26.	<i>Curcuma longa</i> (Turmeric)	Zingiberaceae	Rhizome	Na	Curcumin (0 - 10 μ M)	>10 μ M	Viral replication inhibition	SARS-CoV	92

Na = Not available

inhibited SARS-COV *in vitro*, though the elicited mechanism is yet to be elucidated, its activity was attributed to the presence of quercetin, a known plant flavonol with reported antiviral activity against HIV-luc/SARS.^{91,93} Furthermore, a species from the family Geraniaceae in southern Africa that has been formulated into an herbal drug for the treatment of respiratory infection, *Pelargonium sidoides* was also found to hinder the replication of H1N1, H3N2 virus strains, respiratory syncytial virus, and HCoVs.^{94,95} Another study that involves the anti-SARS-CoV activities of more than 200 Chinese medicinal herbs' extracts was conducted by Wen and co-workers.⁹⁶ The study reported significant anti-SARS-CoV activity of six extracts from *Cassia tora*, *Cibotium barometz*, *Gentiana scabra*, *Dioscorea batatas*, and *Taxillus chinensis* with 50% effective concentration (EC_{50}) values ranging from 5 to 10 μ g/ml. While all the extracts showed good inhibitory activity against Vero E6 cells, the methanolic extracts of *Cibotium barometz* and *Dioscorea batatas* elicited the most significant effect and thus confirming their probable therapeutic potential in the treatment of SARS-COVs.

In 2012, Chang *et al.* reported the presence of triterpenoids in *Euphorbia neriifolia* L. and had anti-HCoV potential.⁹⁷ Found in Pingtung, Taiwan in the South-eastern region of Asia, the ethanolic leaf extract of *Euphorbia neriifolia* revealed the presence of 22 triterpenoids and a flavonoid glycoside, and thirteen of these compounds were identified for the first time in *E. neriifolia*. Of the triterpenoids, 3 β -Friedelanol and friedelane derivatives demonstrated the highest antiviral potential against HCoV. The study concluded that, due to the structure-activity relationship, the friedelane skeleton could serve a potential scaffold in the evolution and synthesis of novel anti-HCoV-229E medications. Ulasli *et al.*⁹⁸ also reported the efficacy of *Anthemis hyalina*, *Citrus sinensis*, and *Nigella sativa* in the management of HCoV infection. The extracts from the three investigated plants inhibited the replication and expression of CoV and TRP gene, respectively.

Recently, the broad-spectrum inhibitory effects of phytochemicals such as lycorine, emetine, monensin sodium, and mycophenolic acid against a panel of HCoVs (HCoV-OC43, HCoV-NL63, MERS-CoV, and MHV-A59) have also been demonstrated.¹¹⁰ Moreover, while emetine and lycorine are potent inhibitors of dengue virus replication, lycorine has also been reported as a good inhibitor of replication in several viruses such as poliomyelitis virus, herpes simplex virus 1, Bunyamwera virus, and West Nile virus.¹¹⁰⁻¹¹² Very recently, the anti-HCoV-NL63 activity of a traditional Chinese medicine, *Strobilanthes cusia* (IC_{50} : 0.64 μ g/ml) has been reported. Its derivatives such as tryptanthrin and indigodole B are also potent inhibitors of HCoV-NL63 replication with IC_{50} values of 1.52 and 2.60 μ M, respectively.¹⁰⁰ The plant and its two derivatives also exhibited significant virucidal effects on HCoV-NL63. The mechanistic effects of tryptanthrin was via the alteration of spike proteins and inference with viral enzymes (e.g. RNA-dependent RNA polymerase, papain-like protease 2) that are involved in viral (HCoV-NL63) RNA genome synthesis, late stages of replication, and production of virus progeny.¹⁰⁰ Furthermore, this phytochemical has also been reported to possess remarkable anti-neuroinflammatory, hepatoprotective, antitumor (leukemia, breast and colon cancer cells), and antimicrobial properties.¹¹³⁻¹¹⁵

A recent meta-analysis study of medicinal plants has also lent credence and suggested that further assays aimed at tackling SAR-CoV-2 and other CoVs could be done on thistle, barley, sundew, and *Ficus* sp.¹¹⁶ Another docking study showed the potential of *Citrus* sp., galangal (*G. nivalis*), sappan wood (*Biancaea sappan*), and *Curcuma* sp. for use in prophylaxis against SAR-CoV-2.¹¹⁷ Also, *in silico* studies have further showed that 13 natural compounds from Chinese traditional medicine databases had anti-SARS-CoV-2 potential. The compounds were sugiol, kaempferol, cryptotanshinone, moupinamide, coumaroyltyramine, quercetin, dihomogamma-linolenic acid, dihydrotanshinone, desmethoxyreserpine, lignan, tanshinone IIa, N-cis feruloyltyramine and betulinic acid.¹⁷

Saposhnikovia divaricata coumarins, phloroglucinols of *Dryopteris crassirhizoma* and oleanane triterpenes derived from *Camellia japonica* flowers also demonstrated antiviral potential against porcine epidemic diarrhoea virus (PEDV), a member of the Coronaviridae family.^{118,119} The replication pathway in PEDV are similar to human CoVs. In other words, vaccines for MERS and PEDV could be based on similar theory^{120, 121} such that the compounds reported in medicinal plant species studied by Yang et al.¹¹⁹ could become promising candidates for further research to better tackle fatal HCoV. Also, anti-HCoV-22E9 potential of saikosaponin A, B2, C and D expressed from medicinal *Scrophularia scorodonia*, *Heteromorpha* and *Bupleurum* species have been demonstrated to prevent viral attachment and penetration *in vitro*. These compounds could be repurposed for SARS-CoV-2.⁹⁰ Other potential plants believed to fight viral respiratory infections include *Pelargonium sidoides* (African geranium), *Andrographis paniculata* (kalmegh) and fruit of *Sambucus nigra* (black elder).²¹ A list of prospective plants and their metabolites for COVID-19 are presented in Table 2.

Plants and their metabolites as specific inhibitors of HCoV target proteins

As earlier mentioned, the HCoVs encode proteins including the 3-chymotrypsin-like protease (3CL^{pro}) – a cysteine protease, which is important for viral replication, transcription and subsequent maturation; papain-like protease (PL^{pro}) – essential for translation and deubiquitination; and spike protein (S) – which is indispensable for host cell entry.¹²²⁻¹²⁵ These proteins provide possible targets for drug development and screening of traditional medicines with anti-coronaviral potential. In fact, some structure-based analyses and high-throughput studies have highlighted plant metabolites as potent inhibitors of these proteins. Phytocompounds such as hesperetin, sinigrin, indigo, β -sitosterol, hirsutenone, emodin, myricetin and tannic acid have been reported to exhibit anti-coronaviral properties through their modulatory effect on HCoV target proteins.^{85,126}

A study by Lin and co-workers¹²⁶ reported the anti-SARS-CoV 3CL^{pro} potential of aqueous extract of *Isatis indigotica*. The same study was further expanded to screen five major compounds of *I. indigotica* root extract (indigo, indirubin, indican, sinigrin, and β -sitosterol) and seven other plant-derived phenolic compounds including aloe-emodin, hesperetin, quercetin, naringenin, daidzein, emodin, and chrysohanol) against the same protein. Although, hesperetin (a phenolic compound) was the most efficient in blocking the cleavage processing of the protein with 50% inhibitory concentration (IC₅₀) of 8.3 μ M, other compounds such as sinigrin, indigo, β -sitosterol, and aloe-emodin also exerted remarkable inhibitory effects with IC₅₀ values ranging from 217 to 1210 μ M. Chen and co-workers⁸⁵ have also highlighted the anti-SARS CoV 3CL^{pro} properties of three phenolic compounds from black tea which include tannic acid, 3-isothaflavin-3-gallate, and theaflavin-3,3'-digallate with IC₅₀ values of 3, 7 and 9.5 μ M, respectively. The study noted the better inhibitory effect of 3-isothaflavin-3-gallate and theaflavin-3,3'-digallate against the protease than theaflavin; this observation was attributed to the absence of gallate group in theaflavin. Findings of this study also demonstrated the significant SARS CoV 3CL^{pro} inhibitory activity of black tea (IC₅₀: 70 μ g/ml) and Puer tea (IC₅₀: 25 μ g/ml) compared to oolong and green tea (IC₅₀: 125 μ g/ml).⁸⁵

In another study, the anti-SARS CoV 3CL^{pro} activity of a phenolic compound, curcumin (IC₅₀: 40 μ M); two triterpenoids, betulinic acid (IC₅₀: 10 μ M), betulonic acid (IC₅₀: >100 μ M); and two lignoids, hinokinin (IC₅₀: >100 μ M), savinin (IC₅₀: 25 μ M) has been demonstrated.⁹² Another phenolic compound, quercetin-3- β -galactoside and some of its derivatives have been identified as potent inhibitors of SARS-CoV 3CL^{pro} through a series of molecular docking and enzyme inhibition assays. The findings further demonstrated that

although residue Gln189 of SARS-CoV 3CL^{pro} plays indispensable role in its interaction with quercetin-3- β -galactoside, its mutation does not affect the enzymatic activity of the protease.¹²⁷ Quercetin also possesses significant anti-murine coronavirus activity.¹²⁸ Findings of Wen and co-workers⁹⁶ have shown the anti-SARS CoV 3CL^{pro} activity of extracts of *Cibotium barometz* (Rhizoma Cibotii) and *Dioscorea batatas* (Dioscoreae Rhizoma) with IC₅₀ values of 39 μ g/ml and 44 μ g/ml, respectively. The aqueous extract of *Houttuynia cordata* have also been reported to possess significant anti-SARS-CoV 3CL^{pro} property as well as considerable immune-stimulatory effect via the increment of CD4⁺ and CD8⁺.¹⁰³

Out of the 312 Chinese medicinal herbs screened by Ho and co-workers,⁵⁴ only three herbs, *Rheum officinale* Baill. (root tuber), *Polygonum multiflorum* Thunb. (root tuber) and *P. multiflorum* Thunb. (vine) displayed considerable inhibition of SARS-CoV S protein and angiotensin converting enzyme type 2 (ACE2) interaction with IC₅₀ values ranging from 1 to 10 μ g/ml. Furthermore, to confirm the phytochemical component responsible for the inhibitory effect, three previously characterised biomolecules of the plants were screened. Emodin, but not Rhein and chrysin significantly blocked SARS-CoV S-protein and ACE2 interaction in a dose-dependent fashion with IC₅₀ value of 200 μ M. It is noteworthy that ACE2 is the entry receptor for SARS-CoV and blockage of its interaction with SARS-CoV S protein would prevent virus entry.⁵⁴ The antiviral activity of emodin against enveloped viruses including influenza virus and varicella-zoster virus, via the disruption of lipid layer has been reported by an earlier study.¹²⁹

The anti-SARS-CoV 3CL^{pro} property of ethanol extract of *Torreya nucifera* leaves has been reported.¹⁰⁷ A bioassay guided fractionation and purification led to the identification of eight diterpenoids [18-hydroxyferruginol (1), hinokiol (2), ferruginol (3), 18-oxoferruginol (4), O-acetyl-18-hydroxyferruginol (5), methyl dehydroabietate (6), isopimaric acid (7), and kayadiol (8)] and four biflavonoids [amentoflavone (9), bilobetin (10), ginkgetin (11), and sciadopitysin (12)] with anti-SARS-CoV 3CL^{pro} activity from the plant's n-hexane and ethyl acetate fractions, respectively. Although the bioflavonoids were the most potent inhibitors of the protease with IC₅₀ values ranging from 8.3 to 72.3 μ M, the diterpenoids also exerted considerable inhibitory effects with IC₅₀ values ranging from 49.6 to 283.5 μ M. Three other flavonoids viz., luteolin, quercetin and apigenin anti-SARS CoV 3CL^{pro} activity was also demonstrated with IC₅₀ values of 20, 23.8 and 280.8 μ M, respectively.¹⁰⁷ In a similar study, four triterpenes from *Tripterygium regelii* namely celastrol, pristimerin, tingenone, and iguesterin inhibited 3CL^{pro} activity with IC₅₀ values of 10.3, 5.5, 9.9, and 2.6 μ M, respectively.¹³⁰ Similarly, a study by Yu and co-workers¹³¹ has reported the antiviral activity (via the inhibition of SARS-CoV helicase protein) of two flavonoids, myricetin and scutellarein with IC₅₀ values of 2.71 \pm 0.19 μ M and 0.86 \pm 0.48 μ M, respectively.

A bioactivity guided fractionation and spectroscopic analysis of the ethanolic extract of *Alnus japonica* led to the isolation of nine small diarylheptanoid's molecules (polyphenols). Six out of the polyphenolic compounds namely hirsutenone, hirsutanonol, oregonin, rubranol, rubranoside B and rubranoside A exerted significant dose-dependent inhibitory effect against SARS-CoV PL^{pro}. It is worthwhile that the biological activity of these molecules was greatly impacted by their chemical structures with most remarkable effect (IC₅₀: 4.1 μ M) exerted by hirsutenone which contains an α , β -unsaturated carbonyl group with a catechol moiety.¹³² The remarkable interactions of natural myricetin and scutellarein flavonoids targeting viral helicase [SARS CoV (SCV) helicase-nsP13]¹³³ could be seen as another promising lead for future anti-SARS-CoV medications.

A flavonoid scaffold of rhoifolin, herbacetin and pectolarin have been shown to significantly inhibit the 3CL^{pro} of SARS-CoV. Molecular

docking studies indicated that the flavonoids bind with the spike S1, S2 and S3' protein sites.¹³⁴ Antiviral flavonoids found in vegetables and fruits such as kaempferol^{105, 135} and daidzein, quercetin, puerarin, epigallocatechin, gallic acid, epigallocatechin gallate¹³⁶ have shown anti-SARS-CoV 3CLpro activity. Phenolics in *Isatis indigotica*,^{78, 79} and the aqueous extract of *Houttuynia cordata*¹⁰³ also showed SARS-CoV 3CLpro inhibition. A recent study by Runfeng and co-workers¹³⁷ noted the inhibitory activity of Lianhuaqingwen, a traditional Chinese medicine formula composing of 13 herbs against the most recent SARS-CoV-2 with IC₅₀ value of 411.2 µg/ml. The exerted anti-SARS-CoV-2 activity was through the deformation of viral morphology, inhibition of viral replication and reduction of cytokines release from host cells.¹³⁷ It is worthwhile that the formula has also been reported to exert broad-spectrum effects on influenza viruses via inhibition of viral propagation and immunomodulation.¹³⁸

Similarly, a study on medicinal plant metabolites such as diallyl disulfide from *Allium sativum* (garlic), capsaicin from *Capsicum* (pepper), limonene from *Elettaria* (cardamom), thymol from *Mentha pulegium* (pennyroyal), coumarin from liquorice, verbascoside from *Stachys schtschegleevi* (hedge nettle), curcumin from *Curcuma longa* (turmeric) and glucuronic acid from *Astragalus gossypinus* (Tragacanth) was recently conducted to detect potential compounds with activity against SARS-CoV-2 using protease enzyme interaction.¹⁰² Of these compounds which showed good interaction, curcumin showed the highest protease inhibiting activity against SARS-CoV-2 and may be useful in antiprotease-based medication for treatment of COVID-19 infection.¹⁰²

Worthy of mention is a CoV infection intervention (no. EP19990203128) put forward by Mas Pharmaceutical. The intervention includes the glucopyranoside analogues found in Ginseng *Panax* species, methylpanaxadiol and 1-protopanaxatriol, in combination with a protodioscin derivative associated with *Dioscorea* plant, dimethylprotobioside. The latter is an anti-mitotic agent and the former showed anti-mutagenic and pro-apoptotic activities. In combination, the intervention targets the total functional and structural destruction of the virus through their RNA-dependent RNA polymerase and DNA gyrase inhibitive characteristics.¹³⁹ In another development with the recent advancement in bioinformatics, Salim and Noureddine¹⁰⁶ studied the molecular docking of the metabolites isolated from *Nigella sativa* as a potential COVID-19 inhibitor. They found out that α-hederin and nigellidine, when compared with chloroquine, have better energy scores thus touting the metabolites as good candidates for the treatment of COVID-19. Table 3 presents some of the specific plant-derived inhibitors of HCoV target proteins.

Overall, an insight into recognition of the class of compounds studied so far against HCoVs revealed that, phenolics (31.78%) and terpenoids (28.04%) are the most investigated phytochemicals with flavonoids (13.08%) and alkaloids (12.15%) also finding some promising applications (Table 3, Figure 4). The seldom application of lignoids (6.54%), saponins (5.61%) and steroids (0.93%) (Table 3, Fig. 4) and several other classes of phytonutrients of therapeutic significance not even being studied against HCoVs is a further promising ground calling for in-depth research targeting the prime proteins of these viruses and especially the current SARS-CoV-2 posing significant global challenge.

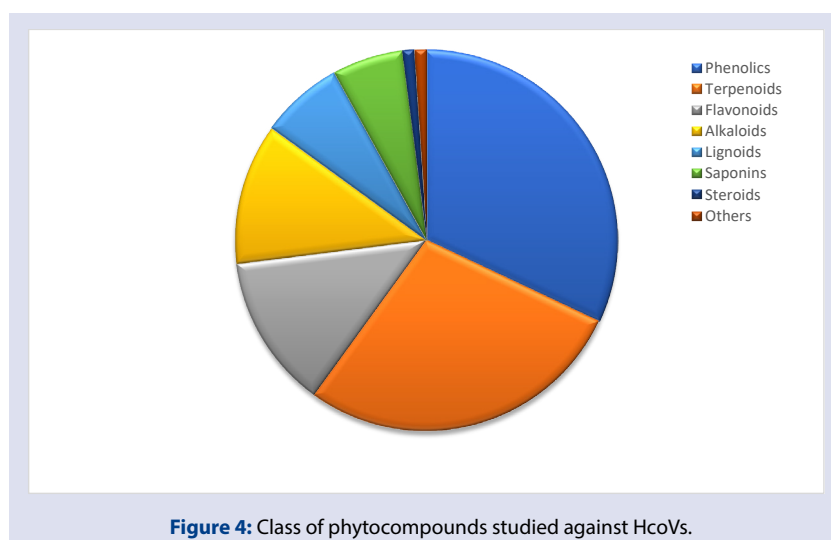
Table 3: Plant-derived specific inhibitors of HCoVs.

SN	Compound	Type	IC ⁵⁰ or EC ₅₀	Proposed mechanism of action	HCoV	Reference
1.	Lycorine (<i>Lycoris radiata</i>)	Alkaloid	15.7 ± 1.2 nM	Unclear	SARS-CoV	78
2.	Glycyrrhizin (<i>Glycyrrhiza glabra</i>) (licorice root)	Saponin	300 mg/L	May inhibit 3CL ^{pro}	SARS-CoV	77
3.	Reserpine (<i>Aesculus hippocastanum</i>)	Alkaloid	3.4 µM	May inhibit 3CL ^{pro}	SARS-CoV	50
4.	Aescin (<i>Rauwolfia</i> species)	Saponin	6.0 µM	May inhibit 3CL ^{pro}	SARS-CoV	50
5.	Lianhuaqingwen	Chinese medicine formula	411.2 µg/ml	May inhibit 3CL ^{pro}	SARS-CoV-2	137
6.	Tetra-O-galloyl-β-d-glucose (<i>Galla chinensis</i>)	Phenolic	4.5 µM	May inhibit S protein	SARS-CoV	91
7.	Luteolin (<i>Rhodiola kirilowii</i>)	Flavonoid	10.6 µM	May inhibit S protein	SARS-CoV	91
8.	Ferruginol (<i>Chamaecyparis obtusa</i> var. <i>formosana</i>)	Terpenoid	1.39 µM	May inhibit 3CL ^{pro}	SARS-CoV	92
9.	Dehydroabieta-7-one (<i>Chamaecyparis obtusa</i> var. <i>formosana</i>)	Terpenoid	4.00 µM	May inhibit 3CL ^{pro}	SARS-CoV	92
10.	Sugiol (<i>Chamaecyparis obtusa</i> var. <i>formosana</i>)	Terpenoid	-	-	SARS-CoV	92
11.	Cryptojaponol (<i>Cryptomeria japonica</i>)	Terpenoid	>10 µM	May inhibit 3CL ^{pro}	SARS-CoV	92
12.	8β-hydroxyabieta-9(11),13-dien-12-one (<i>Chamaecyparis obtusa</i> var. <i>formosana</i>)	Terpenoid	1.47 µM	May inhibit 3CL ^{pro}	SARS-CoV	92
13.	7β-hydroxydeoxycryptojaponol (<i>Cryptomeria japonica</i>)	Terpenoid	1.15 µM	May inhibit 3CL ^{pro}	SARS-CoV	92
14.	6,7-dehydroroyleanone (<i>Chamaecyparis obtusa</i> var. <i>formosana</i>)	Terpenoid	5.55 µM	May inhibit 3CL ^{pro}	SARS-CoV	92
15.	3β-,12-diacetoxyabieta-6,8,11,13-tetraene (<i>Juniperus formosana</i>)	Terpenoid	1.57 µM	May inhibit 3CL ^{pro}	SARS-CoV	92
16.	Pinusolidic acid (<i>Chamaecyparis obtusa</i> var. <i>formosana</i>)	Terpenoid	4.71 µM	May inhibit 3CL ^{pro}	SARS-CoV	92
17.	Forskolin (<i>Coleus forskohlii</i>)	Terpenoid	7.5 µM	May inhibit 3CL ^{pro}	SARS-CoV	92
18.	Cedrane-3β,12-diol (<i>Juniperus formosana</i>)	Terpenoid	>10 µM	May inhibit 3CL ^{pro}	SARS-CoV	92

19.	α -cadinol (<i>Chamaecyparis obtusa</i> var. <i>formosana</i>)	Terpenoid	4.44 μ M	May inhibit 3CL ^{pro}	SARS-CoV	92
20.	Betulonic acid (<i>Betula pubescens</i>)	Terpenoid	>10 μ M	May inhibit 3CL ^{pro} and S protease	SARS-CoV	92
21.	Betulonic acid (<i>Juniperus formosana</i>)	Terpenoid	0.63 μ M	May inhibit 3CL ^{pro}	SARS-CoV	92
22.	Hinokinin (<i>Chamaecyparis obtusa</i> var. <i>formosana</i>)	Lignoid	>10 μ M	May inhibit 3CL ^{pro}	SARS-CoV	92
23.	Savinin (<i>Chamaecyparis obtusa</i> var. <i>formosana</i>)	Lignoid	1.13 μ M	May inhibit 3CL ^{pro} and S protease	SARS-CoV	92
24.	4,4'-O-benzoylisolariciresinol (Synthetic lignoid)	Lignoid	-	-	SARS-CoV	92
25.	Honokiol (<i>Magnolia</i> spp.)	Lignoid	6.50 μ M	May inhibit 3CL ^{pro}	SARS-CoV	92
26.	Magnolol (<i>Magnolia</i> spp.)	Lignoid	3.80 μ M	May inhibit 3CL ^{pro}	SARS-CoV	92
27.	Curcumin (<i>Curcuma longa</i>)	Phenolic	>10 μ M	May inhibit 3CL ^{pro}	SARS-CoV	92
28.	Baicalin (<i>Scutellaria baicalensis</i>)	Flavonoid	12.5 to 25 μ g/ml	-	SARS-CoV	89
29.	Saikosaponin A (<i>Bupleurum</i> spp.)	Saponin	8.6 \pm 0.3 μ M	May inhibit 3CL ^{pro}	HCoV-229E	90
30.	Saikosaponin B ₂ (<i>Bupleurum</i> spp.)	Saponin	1.7 \pm 0.1 μ M	May inhibit 3CL ^{pro} and S protease	HCoV-229E	90
31.	Saikosaponin C (<i>Bupleurum</i> spp.)	Saponin	19.9 \pm 0.1 μ M	May inhibit 3CL ^{pro}	HCoV-229E	90
32.	Saikosaponin D (<i>Bupleurum</i> spp.)	Saponin	13.2 \pm 0.3 μ M	May inhibit 3CL ^{pro}	HCoV-229E	90
33.	Lycorine (<i>Lycoris radiata</i>)	Alkaloid	0.15 to 1.63 μ M	-	HCoV-OC43, HCoV-NL63, MERS-CoV, MHV-A59	110
34.	Emetine (<i>Cephaelis ipecacuanha</i>)	Alkaloid	0.12 to 1.43 μ M	-	HCoV-OC43, HCoV-NL63, MERS-CoV, MHV-A59	110
35.	Berberamine (<i>Berberis</i> spp.)	Alkaloid	1.48 to 13.14 μ M	-	HCoV-OC43, HCoV-NL63, MERS-CoV, MHV-A59	110
36.	Tetrandrine (<i>Stephania tetrandra</i>)	Alkaloid	0.29 to 12.68 μ M	-	HCoV-OC43, HCoV-NL63, MERS-CoV, MHV-A59	110
37.	Pristimerin (<i>Tripterygium wilfordii</i>)	Terpenoid	1.63 to 13.87 μ M	-	HCoV-OC43, HCoV-NL63, MERS-CoV, MHV-A59	110
38.	Harmine (<i>Peganum harmala</i>)	Alkaloid	1.90 to 13.77 μ M	-	HCoV-OC43, HCoV-NL63, MERS-CoV, MHV-A59	110
39.	Conessine (<i>Holarhena floribunda</i>)	Alkaloid	2.34 to 11.46 μ M	-	HCoV-OC43, HCoV-NL63, MERS-CoV, MHV-A59	110
40.	Tryptanthrin (<i>Strobilanthes cusia</i>)	Alkaloid	1.52 \pm 0.13 μ M	Alteration of spike proteins; Inhibition of RNA-dependent RNA polymerase; papain-like protease 2 inhibition; inhibition of viral replication	HCoV-NL63	100
41.	Chrysin	Phenolic	200 μ M	Inhibition of (S) protein and ACE2 interaction	SARS-CoV	54
42.	Indigodole B (<i>Strobilanthes cusia</i>)	Alkaloid	2.60 \pm 0.11 μ M	May inhibit 3CL ^{pro}	HCoV-NL63	100
43.	Tetrandrine (<i>Stephania tetrandra</i>)	Alkaloid	0.33 \pm 0.03 μ M	Inhibition of viral S and N protein	HCoV-OC43	140
44.	Fangchinoline (<i>Stephania tetrandra</i>)	Alkaloid	1.01 \pm 0.07 μ M	Inhibition of viral S and N protein	HCoV-OC43	140
45.	Cepharanthine (<i>Stephania tetrandra</i>)	Alkaloid	0.83 \pm 0.07 μ M	Suppression of viral replication; inhibition of viral S and N protein	HCoV-OC43	140
46.	Procyanidin B1 (Cinnamomi Cortex)	Flavonoid	41.3 \pm 3.4 μ M	Inhibition of pseudovirus infection	SARS-CoV	109

47.	Procyanidin A2 (Cinnamomi Cortex)	Flavonoid	29.9 ± 3.3 μM	Inhibition of pseudovirus infection	SARS-CoV	109
48.	Cinnamtannin B1 (<i>Cinnamomum verum</i>)	Flavonoid	32.9 ± 3.9 μM	Inhibition of pseudovirus infection	SARS-CoV	109
49.	Silvestrol (<i>Aglaia</i> spp.)	Phenolic	3 nM	Inhibition of cap-dependent viral mRNA translation	HCoV-229E	141
50.	Juglanin (<i>Polygonum aviculare</i>)	Phenolic	2.3 μM	Blockage of 3a channel	SARS-CoV	105
51.	Silvestrol (<i>Aglaia</i> spp.)	Phenolic	1.3 nM	Inhibition of cap-dependent viral mRNA translation	MERS-CoV	141
52.	Sinigrin (<i>Isatis indigotica</i>)	Glucoside	217 μM	SARS-CoV 3CL ^{pro} inhibition	SARS-CoV	126
53.	Beta-sitosterol (<i>Isatis indigotica</i>)	steroid	1210 μM	SARS-CoV 3CL ^{pro} inhibition	SARS-CoV	126
54.	Indigo (<i>Isatis indigotica</i>)		752 μM	SARS-CoV 3CL ^{pro} inhibition	SARS-CoV	126
55.	Aloe-emodin	Phenolic	366 μM	SARS-CoV 3CL ^{pro} inhibition	SARS-CoV	126
56.	Hesperetin	Phenolic	8.3 μM	SARS-CoV 3CL ^{pro} inhibition	SARS-CoV	126
57.	Kazinol A	Phenolic	66.2–88.5 μM	Inhibition of 3CL ^{pro} and PL ^{pro}	SARS-CoV and MERS-CoV	142
58.	Emodin (<i>Rheum</i> and <i>Polygonum</i> genera)	Phenolic	200 μM	Inhibition of SARS-CoV S protein and ACE2 interaction	SARS-CoV	54
59.	Myricetin	Flavonoid	2.71 ± 0.19 μM	Inhibition of SARS-CoV helicase	SARS-CoV	131
60.	Scutellarein	Flavonoid	0.86 ± 0.48 μM	Inhibition of SARS-CoV helicase	SARS-CoV	131
61.	Hirsutenone (<i>Alnus japonica</i>)	Phenolic	4.1 ± 0.3 μM	Inhibition of SARS-CoV PL ^{pro}	SARS-CoV	132
62.	Hirsutanonol (<i>Alnus japonica</i>)	Phenolic	7.8 ± 1.7 μM	Inhibition of SARS-CoV PL ^{pro}	SARS-CoV	132
63.	Oregonin (<i>Alnus japonica</i>)	Phenolic	20.1 ± 2.2 μM	Inhibition of SARS-CoV PL ^{pro}	SARS-CoV	132
64.	Rubranol (<i>Alnus japonica</i>)	Phenolic	12.3 ± 0.9 μM	Inhibition of SARS-CoV PL ^{pro}	SARS-CoV	132
65.	Rubranoside B (<i>Alnus japonica</i>)	Phenolic	8.0 ± 0.2 μM	Inhibition of SARS-CoV PL ^{pro}	SARS-CoV	132
66.	Rubranoside A (<i>Alnus japonica</i>)	Phenolic	9.1 ± 1.0 μM	Inhibition of SARS-CoV PL ^{pro}	SARS-CoV	132
67.	Quercetin-3-β-galactoside	Flavonoid	42.79 ± 4.97 μM	Competitive inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	127
68.	Quercetin	Flavonoid	23.8 ± 1.9 μM	Inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	107
69.	Betulinic acid	Terpenoid	10 μM	Competitive inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	92
70.	Betulonic acid	Terpenoid	>100 μM	Inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	92
71.	Hinokinin	Lignoid	>100 μM	SARS-CoV 3CL ^{pro} inhibition	SARS-CoV	92
72.	Savinin	Lignoid	25 μM	Competitive inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	92
73.	Curcumin	Phenolic	40 μM	SARS-CoV 3CL ^{pro} inhibition	SARS-CoV	92
74.	Broussoualchone B	Phenolic	11.6–112.9 μM	Inhibition of 3CL ^{pro} and PL ^{pro}	SARS-CoV and MERS-CoV	142
75.	Broussoualchone A	Phenolic	9.2–88.1 μM	Inhibition of 3CL ^{pro} and PL ^{pro}	SARS-CoV and MERS-CoV	142
76.	4-hydroxyisolonchocarpin	Phenolic	35.4–202.7 μM	Inhibition of 3CL ^{pro} and PL ^{pro}	SARS-CoV and MERS-CoV	142
77.	Papyriflavonol A	Phenolic	3.7–112.5 μM	Inhibition of 3CL ^{pro} and PL ^{pro}	SARS-CoV and MERS-CoV	142
78.	Tannic acid	Phenolic	3 μM	SARS-CoV 3CL ^{pro} inhibition	SARS-CoV	85
79.	3-isothaeafavin-3-gallate	Phenolic	7 μM	SARS-CoV 3CL ^{pro} inhibition	SARS-CoV	85
80.	Theaflavin-3,3'-digallate	Phenolic	9.5 μM	SARS-CoV 3CL ^{pro} inhibition	SARS-CoV	85
81.	Theaflavin	Phenolic	56 μM	SARS-CoV 3CL ^{pro} inhibition	SARS-CoV	85
82.	Theaflavin-3-gallate and Theaflavin-3'-gallate	Phenolics	43 μM	SARS-CoV 3CL ^{pro} inhibition	SARS-CoV	85
83.	Apigenin	Phenolic	280.8 ± 21.4 μM	Inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	107
84.	Kazinol J	Phenolic	15.2–109.2 μM	Inhibition of 3CL ^{pro} and PL ^{pro}	SARS-CoV and MERS-CoV	142

85.	18-hydroxyferruginol	Terpenoid	220.8 ± 10.4 μM	Inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	107
86.	Hinokiol	Terpenoid	233.4 ± 22.2 μM	Inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	107
87.	Ferruginol	Terpenoid	49.6 ± 1.5 μM	Inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	107
88.	18-oxoferruginol	Terpenoid	163.2 ± 13.8 μM	Inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	107
89.	O-acetyl-18-hydroxyferruginol	Terpenoid	128.9 ± 25.2 μM	Inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	107
90.	Methyl dehydroabietate	Terpenoid	207.0 ± 14.3 μM	Inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	107
91.	Isopimaric acid	Terpenoid	283.5 ± 18.4 μM	Inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	107
92.	Kayadiol	Terpenoid	137.7 ± 12.5 μM	Inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	107
93.	Amentoflavone	Flavonoid	8.3 ± 1.2 μM	Non-competitive inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	107
94.	Bilobetin	Flavonoid	72.3 ± 4.5 μM	Non-competitive inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	107
95.	Ginkgetin	Flavonoid	32.0 ± 1.7 μM	Non-competitive inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	107
96.	Sciadopitysin	Flavonoid	38.4 ± 0.2 μM	Non-competitive inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	107
97.	Abietic acid	Terpenoid	189.1 ± 15.5 μM	Inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	107
98.	Kazinol F	Phenolic	39.5–135.0 μM	Inhibition of 3CL ^{pro} and PL ^{pro}	SARS-CoV and MERS-CoV	142
99.	Luteolin	Flavonoid	20.0 ± 2.2 μM	Inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	107
100.	Kazinol B	Phenolic	31.4–233.3 μM	Inhibition of 3CL ^{pro} and PL ^{pro}	SARS-CoV and MERS-CoV	142
101.	Brousoflavan A	Phenolic	49.1–125.7 μM	Inhibition of 3CL ^{pro} and PL ^{pro}	SARS-CoV and MERS-CoV	142
102.	Celastrol	Terpenoid	10.3 ± 0.2 μM	Competitive inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	130
103.	Pristimerin	Terpenoid	5.5 ± 0.7 μM	Competitive inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	130
104.	Tingenone	Terpenoid	9.9 ± 0.1 μM	Competitive inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	130
105.	Iguesterin	Terpenoid	2.6 ± 0.3 μM	Competitive inhibition of SARS-CoV 3CL ^{pro}	SARS-CoV	130
106.	3'-(3-methylbut-2-enyl)-3',4,7-trihydroxyflavane	Phenolic	30.2–48.8 μM	Inhibition of 3CL ^{pro} and PL ^{pro}	SARS-CoV and MERS-CoV	142
107.	Corylifol	Phenolic	32.3 ± 3.2 μM	Inhibition of PL ^{pro}	SARS-CoV	143



CONCLUSION AND PERSPECTIVES

Quite an array of plants, either whole, extract or plant metabolites, have shown great potential as antiviral agents and moieties. Potent antiviral phytochemical groups identified include saponins, tannins, lignans, alkaloids, flavonoids, lectins, coumarins, terpenoids, peptides and proteins. It therefore seems reasonable to suggest that the world should look to plants for novel natural compounds and antivirals. In addition to the possibility of harnessing natural phytochemicals for new pharmaceuticals development, plant metabolites could also become more efficacious when utilized in form of combination therapies with existing drug interventions. Increased research into plants and in-depth characterization of their metabolites could uncover more interesting results that would benefit humanity in its fight against emerging and re-emerging viral coronavirus infections such as the current COVID-19. Even though some promising compounds remain under clinical trials, the use of medicinal plants as prophylaxis or treatment interventions in viral respiratory infections should not be undermined. Also, with the recent speculation that, some of the target proteins of SARS-CoV-2 bind to and displaces oxygen from β -chain of the hemoglobin which subsequently results in hemotoxicity and inflammation of the alveolar macrophages,¹⁴⁴ it could be logically suggested that blood purifiers including those of plant origin may offer complementary benefits against COVID-19. This speculation could support the touted efficacy of chloroquine for COVID-19 as it could compete for binding on the porphyrin of the hemoglobin in a manner that may prevent SARS-CoV-2 protein from binding. While this calls for further submission to substantiate the claim, metabolites with the ability to purify the blood and inhibit inflammation and oxidative stress could also be possible interventions for the prevention of COVID-19. Of recent, Van Vuuren and Frank¹⁴⁵ reported Southern African medicinal plants that can be used for blood purifications. Interestingly, some of the plants reported have also been validated for antiviral, inflammatory, and antioxidant activities which make them probable candidates for the management of COVID-19. Some of the plants are *Bridelia micrantha*, *Bulbine latifolia* var. *latifolia*, *Burchellia bubalina*, *Crinum moorei*, *Cymbopogon validus*, *Euclea natalensis*, *Polygala virgate*, *Polygonum hystriculium*, *Salix mucronate*, *Scadoxus puniceus*, *Schotia brachypetala*, *Tropaeolum majus*, *Vitellariopsis marginata*, and *Zanthoxylum capense*.¹⁴⁶⁻¹⁵⁷ Overall, given the current body of evidence on the potential development of phytodrugs or phytomedicines for COVID-19, fears need to be allayed while clinical trials continue. The lockdown and other preventive measures which have been implemented in most parts of the world should be humanely exercised and supported with palliatives to ensure effectiveness and compliance. Increased awareness and update on developments should also be engaged in through all possible and genuine media. However, after aligning with all possible measures, one can only hope for the best possible outcome in a not-so-distant future. But will the world ever remain the same again? Time will tell.

CONFLICTS OF INTEREST

No conflict of interest exists among authors.

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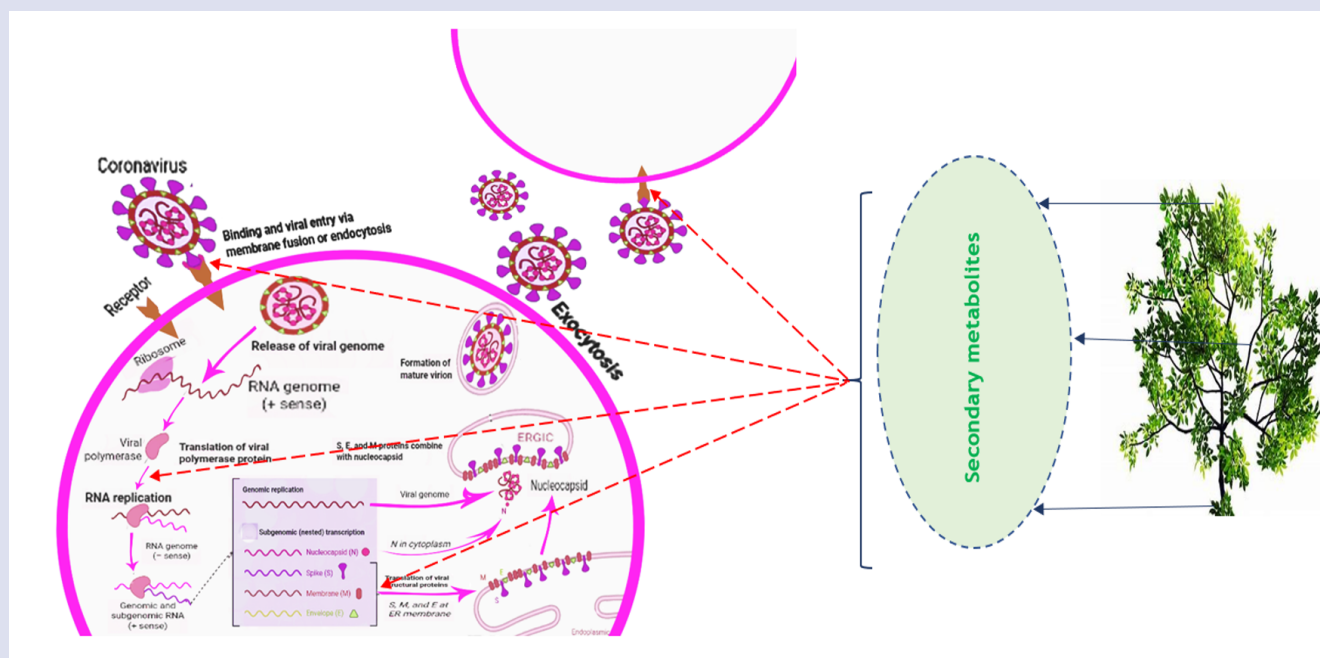
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GRAPHICAL ABSTRACT



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