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## Review

## Natural product-derived phytochemicals as potential agents against coronaviruses: A review

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## ABSTRACT

Coronaviruses are responsible for a growing economic, social and mortality burden, as the causative agent of diseases such as severe acute respiratory syndrome (SARS), Middle East respiratory syndrome (MERS), avian infectious bronchitis virus (IBV) and COVID-19. However, there is a lack of effective antiviral agents for many coronavirus strains. Naturally existing compounds provide a wealth of chemical diversity, including antiviral activity, and thus may have utility as therapeutic agents against coronaviral infections. The PubMed database was searched for papers including the keywords coronavirus, SARS or MERS, as well as traditional medicine, herbal, remedy or plants, with 55 primary research articles identified. The overwhelming majority of publications focussed on polar compounds. Compounds that show promise for the inhibition of coronavirus in humans include scutellarein, silvestrol, tryptanthrin, saikosaponin B<sub>2</sub>, quercetin, myricetin, caffeic acid, psoralidin, isobavachalcone, and lectins such as griffithsin. Other compounds such as lycorine may be suitable if a therapeutic level of antiviral activity can be achieved without exceeding toxic plasma concentrations. It was noted that the most promising small molecules identified as coronavirus inhibitors contained a conjugated fused ring structure with the majority being classified as being polyphenols.

## 1. Introduction

The dramatic change of events with the recent unprecedented coronavirus pandemic declared by the World Health Organisation (WHO) has prompted an exponential increase of scientific interest in coronaviruses globally. As of April 22nd 2020, the pandemic has resulted in 2,553,112 infections, with 177,286 deaths worldwide, which continues to drastically increase as we write (<https://www.who.int/emergencies/diseases/novel-coronavirus-2019>).

Coronaviruses (CoVs) belong to the family Coronaviridae, subfamily *Coronavirinae* and are large (genome size 26–32 kb; Wu et al., 2020a), enveloped, positive-sense single-stranded ribonucleic acid (RNA) viruses that can infect both animals and humans (Fig. 1). Based on their genotypic and serological characteristics, the viruses are subdivided into four genera: *Alpha*-, *Beta*-, *Gamma*-, and *Deltacoronavirus* (Chu et al., 2020; Lu et al., 2015). At present, all identified CoVs that are capable of infecting humans belong to the first two genera. These include the alphacoronaviruses ( $\alpha$ CoVs) HCoV-NL63 (Human CoV-NL63)

and HCoV-229E and the betacoronaviruses ( $\beta$ CoVs) HCoV-OC43 (Human CoV-OC43), HKU1 (Human CoV), SARS-CoV (Severe Acute Respiratory Syndrome CoV), and MERS-CoV (Middle Eastern Respiratory Syndrome CoV) (Lu et al., 2015). In the past two decades there have been three epidemics caused by the betaCoVs, namely SARS in 2002–03, MERS in 2012 and COVID-19, first identified in 2019 (Yang et al., 2020b).

SARS-CoV emerged in 2002–03 in Southern China, causing a global threat and infecting more than 8000 people, with approximately 800 fatalities recorded, largely in China and the surrounding regions (Lu et al., 2015; Paraskevis et al., 2020). MERS-CoV emerged in the Middle East, spreading to several countries to infect close to 2300 individuals, resulting in 845 deaths as of July 2019 (World Health Organization, 2019). The present CoV pandemic resulting from SARS-CoV-2, which causes COVID-19 (coronavirus disease), was identified in Wuhan City, in the Hubei province of southern mainland China on the 31st December 2019 (Sohrabi et al., 2020). The genome of SARS-CoV-2 is approximately 70 % identical to that of SARS-CoV (Hui et al., 2020),

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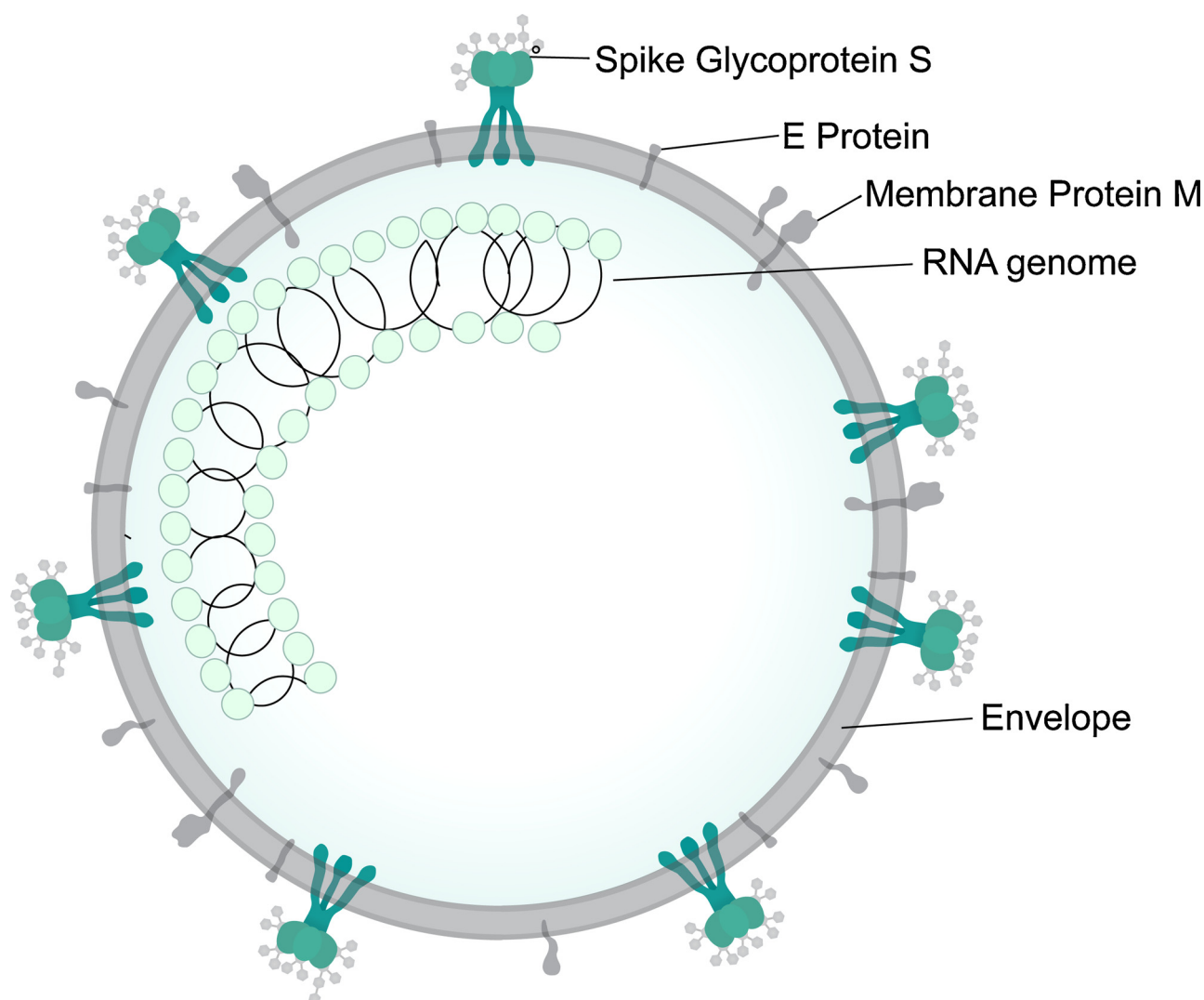


Fig. 1. The general structure of a coronavirus (reproduced from Wikipedia under CC licence 4.0). E protein = envelope protein.

hence leading to its current name.

The major druggable targets of SARS-CoV-2 include 3-chymotrypsin-like protease (3CLpro), papain like protease (PLpro), RNA-dependent RNA polymerase, and spike (S) proteins (Wu et al., 2020b). The S proteins interact directly with human angiotensin-converting enzyme (ACE) 2, allowing the virus to enter the cells. At present, no preventive vaccines or established antiviral therapies are available for coronaviruses (Sohrabi et al., 2020). However, several synthetic compounds have shown promise, including hydroxychloroquine and chloroquine phosphate (Cortegiani et al., 2020; Gao et al., 2020), which act through several mechanisms, including alkalinisation of the host cell phagolysosomes. Newer antiviral medications such as lopinavir (Yao et al., 2020), remdesivir (Holshue et al., 2020; Wang et al., 2020), and arbidol (Khamitov et al., 2008) also show promise. Other suggested treatment options include lopinavir/ritonavir, nucleoside analogues, neuraminidase inhibitors, and peptide EK1 (Lu, 2020). A detailed list of current and planned clinical trials investigating various drugs for the treatment of SARS-CoV-2 was provided by Pang et al. (2020), with updated results available from ClinicalTrials.gov (2020).

In addition, traditional herbal medicines and purified natural products may guide the development of novel antiviral drugs. In other words, more efficient drugs can often be designed based on the structure of natural compounds that exhibit the desired activity. Classic examples of this drug discovery pathway include emetine, an

isoquinoline alkaloid isolated from *Cephaelis ipecacuanha* and used as an amoebicidal drug; quinine, derived from the bark of *Cinchona* trees; and numerous other drugs modified from natural compounds, such as aspirin, morphine and paclitaxel, an antineoplastic drug used for the treatment of cancer (Ganjhu et al., 2015). Indeed, half of all drugs approved between 1981 and 2014 were derived from or mimicked a natural compound (Newman and Cragg, 2016). Furthermore, in the current outbreak of COVID-19, many patients appear to be turning to complementary or traditional medicinal therapies, albeit using them almost exclusively in conjunction with western medicine. For example, one study suggested that almost 92 % of 135 hospitalised patients in northeast Chongqing (China) received traditional Chinese medicine in addition to western medicine (Wan et al., 2020). However, based on the many studies conducted on this topic, it is hard to separate the potential effects of, and interaction between, traditional Chinese herbal medicine and western medicine. Recent reviews have suggested that traditional Chinese medicine could be used for the prevention (Luo et al., 2020) or treatment (Yang et al., 2020a) of COVID-19; while still acknowledging that many studies involving clinical trials are poorly designed or controlled, and the choice of treatments is largely empirically based. As previous work has highlighted the potential of traditional Chinese medicines as a source of potential novel drugs (Ling, 2020), we have not included details on such studies investigating the antiviral activity of remedies comprising portions of numerous plant species in this

review. Rather, our aim is to collate data on the broad spectrum of natural phytochemicals from individual plant species that may have therapeutic potential.

Naturally occurring antiviral agents acting against general coronaviruses were briefly reviewed by Lin et al. (2014) six years ago, while more recent reviews by Pang et al. (2020) and Lu (2020) on therapies for COVID-19 made only brief mention of natural therapeutics and did not explore the active compounds or their mechanism of action. In light of the current COVID-19 pandemic, this review aims to gather and consolidate information on extracts and compound(s) derived from natural products which show potential antiviral bioactivity for the inhibition of coronaviruses. It is hoped that the information presented may guide the naturally-derived drug discovery process in finding a treatment for SARS-CoV-2.

## 2. Methods

The PubMed database ([www.ncbi.nlm.nih.gov/pubmed/](http://www.ncbi.nlm.nih.gov/pubmed/)) was used to locate articles including the following combination of terms: (coronavirus, SARS OR MERS) AND (traditional medicine, herbal, remedy OR plants). Papers primarily focussed on the antiviral activity of prepared Chinese traditional medicines, which typically comprise multiple plant species, were considered out of scope of this review. All articles up to and including 25 March 2020 were considered, yielding a total of 659 results. Two of the authors independently screened the results and identified relevant articles, yielding a total of 35 primary articles on human coronaviruses and 22 on animal coronaviruses were found to be pertinent and thus included in this review (total = 58 papers). Of these, two papers (6 %) were on SARS-CoV-2 (COVID-19). The majority of studies on human coronaviruses (69 %;  $n = 24$ ) included SARS-CoV, with only 3 (9 %) including MERS-CoV and 8 (23 %) other human coronaviruses. It should be noted that one study included both SARS-CoV and MERS-CoV (O'Keefe et al., 2010) while another included both MERS-CoV and HCoV-229E (Müller et al., 2018), hence these percentages do not add to 100 %.

## 3.

Table 1 summarises the studies reporting the inhibition of various human coronavirus strains using compounds derived from plant sources. The table is arranged by viral strain in order to better compare the bioactivity of compounds from different studies upon the same viral genotypes. Where identified, the key compounds responsible for the antiviral activity and their identified mechanisms of action are presented. It should be noted the term  $EC_{50}$  (effective concentration) applies to cell-based assays, while  $IC_{50}$  (inhibitory concentration) applies to enzyme- or biochemical-based assays.

### 3.1. Inhibitors of SARS-CoV-2 (COVID-19)

Few studies report on SARS-CoV-2, as expected given the short time since its emergence. However, a number of studies report on use of computer modelling for screening purposes (Liu and Zhou, 2005; Lung et al., 2020; Toney et al., 2004; Wang et al., 2007; Zhang et al., 2020). Typically, these models determine the free binding of energy between a ligand and a receptor (Forli et al., 2016), with a lower free binding energy indicating a weaker bond between the ligand and receptor. Although obtaining consistent results via different modelling approaches can be challenging (Aldeghi et al., 2016), computer modelling nevertheless allows for comparison of the relative binding affinity of bank of molecules toward the receptor in question.

In addition to reducing the high costs and length of time associated with physically screening large banks of compounds or plant extracts for bioactivity (Chen et al., 2017), the speed and versatility of this method may be particularly valuable for rapidly finding a potent inhibitor of SARS-CoV-2. Compounds that are highlighted through this

method can then be forwarded on cell-based assays to assess their *in vitro* effectiveness and toxicity, before continuing to animal and clinical trials.

Lung et al. (2020) virtually screened 83 compounds found in Chinese traditional medicines for activity against the RNA-dependent RNA polymerase of SARS-CoV-2, identifying theaflavin, an antioxidant polyphenol, as a potential inhibitor. Similarly, Zhang et al. (2020) virtually screened 115 compounds found in Chinese traditional medicines, highlighting 13 for further studies. Several of these were naturally occurring polyphenolic compounds such as quercetin and kaempferol, which have already received considerable interest for the treatment of other disease types (Cassidy et al., 2019; Khan et al., 2019; Tome-Carneiro and Visioli, 2016).

### 3.2. Inhibitors of SARS-CoV

Given the relatively large amount of research that has been performed searching for inhibitors of SARS-CoV, antiviral agents that successfully inhibit this viral strain may provide a good starting point for identifying compounds that are active against SARS-CoV-2.

#### 3.2.1. Virtual screening

Several authors have utilised virtual computer docking models to screen for potential compounds that could bind to and inhibit key proteins present in SARS-CoV (Liu and Zhou, 2005; Toney et al., 2004; Wang et al., 2007), highlighting the potential antiviral activity of compounds such as sabadinine and aurantiamide acetate. Compounds may be screened against a number of binding sites in order to test for potential inhibition of coronaviruses; the main sites that are typically used are the chymotrypsin-like protease (3CLpro), papain like protease (PLpro), spike proteins and RNA-dependent RNA polymerase.

#### 3.2.2. Compound library screening

Several large *in vitro* screening studies searching for inhibitory activity of naturally occurring compounds against SARS-CoV have been performed, mainly on Chinese medicinal herbs (Li et al., 2005; Wang et al., 2003). While the results highlight the potential of selected plant extracts against SARS-CoV, they also demonstrate that such work can be akin to searching for a 'needle in a haystack'. For example, Li et al. (2005) screened over 200 ethanol/chloroform extracts of Chinese medicinal herbs and found only four (*Lycoris radiata*, *Artemisia annua*, *Pyrrhosia lingua* and *Lindera aggregata*) with moderate to high antiviral activity using a CPE assay ( $EC_{50}$  values ranging from  $2.4 \pm 0.2$ – $88.2 \pm 7.7$   $\mu\text{g/mL}$ ). Of these, a single compound (lycorine) from one plant species (*L. radiata*) was earmarked as a potential drug candidate against SARS-CoV. The antiviral efficacy of lycorine was quite high ( $EC_{50}$  of  $15.7 \pm 1.2$  nM), with a selectivity index greater than 900. Although the authors did not make mention of this fact, lycorine can cause toxic effects at low dosage levels (around 1 mg/kg in dogs) (Kretzing et al., 2011), hence great caution would be required for further development of this compound as a drug candidate.

Yu et al. (2012) screened the activity of a library of 64 naturally occurring compounds against SARS-CoV helicase, which plays a key role in the viral genome replication, transcription, and translation. The polyphenolics myricetin and scutellarein (Fig. 2) were identified as the most promising candidates ( $IC_{50}$  values of  $2.71 \pm 0.19$  and  $0.86 \pm 0.48$   $\mu\text{M}$ , respectively). Although the antiviral activity of the compounds was not assessed in cell-based assays, the authors did report that neither compound was toxic to normal (non-tumorigenic) breast epithelial cells. Myricetin is found in reasonably high concentrations in fruits such as cranberry (Häkkinen et al., 1999) as well as in several vegetables such as *Calamus scipionum* and garlic (Miean and Mohamed, 2001). Scutellarein was isolated from *Scutellaria baicalensis* (Chinese Skullcap), which has been traditionally used in the treatment of inflammation and respiratory infections, amongst other uses (Zhao et al., 2016). Both compounds were found to inhibit SARS-CoV helicase (nsP13) through

**Table 1**  
Studies reporting antiviral activity of natural products or isolates against human coronavirus strains.

Viral strain	Assay method	Plant species	Plant part/ isolate	EC <sub>50</sub> or IC <sub>50</sub> (μM unless otherwise stated)	SI	Key compounds present (if identified)	Biological action	Reference
SARS-CoV-2	Computer modelling	Present in: <i>Camellia sinensis</i>	–	ND	ND	Theaflavin	Binding to RNA-dependent RNA polymerase	(Lung et al., 2020)
SARS-CoV-2	Computer modelling	Compounds previously identified in a range of Chinese traditional medicines	n/a	ND	ND	Betulinic acid Coumaroyltyramine Cryptotanshinone Desmethoxyreserpine Dihomo-c-linolenic Dihydrotanshinone Kaempferol Lignan Moupinamide N-cis-feruloyltyramine Quercetin Sugiol Tanshinone IIa Leptodactylone	Replication & 3CLpro PIpro & 3CLpro Replication, 3CLpro & entry 3CLpro Entry & spike protein PIpro & 3CLpro Replication & 3CLpro PIpro PIpro & 3CLpro Replication & 3CLpro PIpro & 3CLpro Not determined	(Zhang et al., 2020)
SARS-CoV	CPE assay	<i>Boenninghausenia sessilicarpa</i>	Isolated compound from ethanol extract	~ 450	ND			(Yang et al., 2007)
SARS-CoV	CPE assay	<i>Hippeastrum hybrid</i> <i>Galanthus nivalis</i> <i>Narcissus pseudonarcissus</i> <i>Lycoris radiata</i> <i>Allium porrum</i> <i>Allium ursinum</i> <i>Cymbidium hybrid</i> <i>Listera ovata</i> <i>Epipactis helleborine</i> <i>Tulipa hybrid</i> <i>Morus nigra</i> <i>Nicotiana tabacum</i> <i>Urtica dioica</i> <i>Morus nigra</i> <i>Cladastis lutea</i> <i>Polygonatum multiflorum</i> <i>Iris hybrid</i>	Lectins isolated from diaminopropane extracts	3.2 ± 2.8 6.2 ± 0.6 5.7 ± 4.4 48 0.45 ± 0.0 8 18 ± 4 4.9 ± 0.8 2.2 ± 1.3 1.8 ± 0.3 22 ± 6 1.6 ± 0.5 1.7 ± 0.3 1.3 ± 0.1 50 ± 13 7.4 ± 0.2 18 ± 13 28 ± 11	> 31.3 > 16.1 > 17.5 > 2.1 > 222.2 > 5.5 > 20 > 45.5 > 55.5 > 2.3 > 62.5 > 58.8 > 76.9 > 2 > 13.5 > 5.5 > 12.6 22.7	Agglutinins: mannose-specific	Inhibit viral attachment and another target at end of replication cycle	(Keyaerts et al., 2007)
SARS-CoV (BJ001 and BJ006)	CPE/MTS assay	<i>Tulipa hybrid</i> <i>Artemisia annua</i> <i>Pyrrosia lingua</i> <i>Lindera aggregata</i> <i>Lycoris radiata</i> Isolated lycorine Commercial lycorine	95 % EtOH extract Chloroform 95 % EtOH 95 % EtOH	38 ± 0 (all μg/mL) 34.5–39.2 40.5–43.2 80.6–88.2 2.1–2.4 (all μg/mL) 48.8 ± 3.6 nM 15.7 ± 1.2 nM	27–31 55–59 16–17 370–422 954 885 14.622 40.0	In <i>L. radiata</i> : lycorine	Not determined	(Li et al., 2005)
SARS-CoV BJ01	MTT cytotoxicity assay	<i>Galla chinensis</i>	Isolated compounds from 85 % ethanol extract	10.6 4.5		Luteolin Tetra-O-galloyl-β-D-glucose	Binds with S2 subunit and preventing entry	(Yi et al., 2004)
SARS-CoV FFM1	CPE assay	<i>Toona sinensis</i>	Bolled water extract of leaves	30–43 μg/mL	12–17	Not determined	Not determined	(Chen et al., 2008)

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Table 1 (continued)

Viral strain	Assay method	Plant species	Plant part/ isolate	EC <sub>50</sub> or IC <sub>50</sub> (μM unless otherwise stated)	SI	Key compounds present (if identified)	Biological action	reference
SARS-CoV FFM1	CPE assay	Glycyrrhizin and glycyrrhetic acid found in: <i>Glycyrrhiza radix</i>	Chemical standards	365 ± 12 > 20	> 65 –	Glycyrrhizin 18β-glycyrrhetic acid  Selected synthetic derivatives	Not determined	(Hoeve et al., 2005)
SARS-CoV FFM1	CPE assay	<i>Laurus nobilis</i>	Essential oil	40 ± 13 35 ± 7 139 ± 20 8 ± 2 50 ± 10 5 ± 3 16 ± 1	> 75 41 2 6 5 3 4	<i>L. nobilis</i> : β-cymene, 1,8-cineole, α-pinene, β-pinene <i>T. orientalis</i> : α-pinene, δ-3-carene, α-cedrol	Inhibition of viral replication	(Loizzo et al., 2008)
SARS-CoV (Hong Kong strain)	CPE assay	<i>Thuja orientalis</i>	75 % ethanol extract	120 ± 1.2 μg/mL 130 ± 0.4 μg/mL	4.2 3.8	Not determined Secoiridoid & glycosides? Polysaccharides? Emodin? Quercetin? Procyanidin A2	Not determined	(Wen et al., 2011)
SARS-CoV PUMC01 F5	Plaque reduction assay	<i>Cibotium barometz</i> <i>Gentiana scabra</i> <i>Dioscorea batatas</i> <i>Cassia tora</i> <i>Taxillus chinensis</i> <i>Cinnamomi</i> sp.	Water extraction followed by phase extraction	8.42 > 10 8.70 8.06 8.43 5.39 (all μg/mL) 10.7 ± 0.4 μg/mL (EtOH fraction)	> 59.4 > 57.5 > 62.0 > 59.3 > 92.8 16.9		Early stage inhibition of viral entry (clathrin-dependent endocytosis pathway)	(Zhuang et al., 2009)
SARS-CoV <i>urbani</i> strain (200,300,592)	Neutral red uptake assay SARS-CoV-infected BALB/c mouse model	Found in: <i>Urtica dioica</i>	Isolated compound Chemical standard used	7.8 ± 0.3 μg/mL (butanol fraction) 29.9 ± 3.3 μM 2.6 ± 3.7 μg/mL	23.1 37.35 10.2 ± 5.6	<i>Urtica dioica</i> agglutinin	Dose-dependent inhibition of viral replication, likely in adsorption or penetration stages. Binds to SARS-CoV spike glycoprotein and N-acetylglucosamine-like residues on the glycosylated envelope. Direct binding to surface envelope glycoprotein spike	(Kumaki et al., 2011)
SARS-CoV <i>Urban</i> strain <i>Tor-II</i> strain <i>CuHK</i> strain <i>Frank</i> strain SARS-CoV helicase nsP13 SARS-CoV S protein	CPE assay	Found in <i>Griffithsia</i> sp.	Chemical standard used	(μg/mL) 0.61 0.61 0.78 1.19	> 164 > 164 > 128 > 83	Griffithsin	Inhibit ATPase activity of SARS-CoV helicase nsP13 Inhibited binding of S protein to ACE2	(Yu et al., 2012) (Ho et al., 2007)
SARS-CoV 3CLpro	Computer modelling	n/a <i>Scutellaria baicalensis</i> <i>Rheum officinale</i> <i>Polygonum multiflorum</i>	Chemical standard Isolated compounds Water extracts (at 40 °C) of roots Synthetic emodin standard n/a	2.71 ± 0.19 0.86 ± 0.48 ~5 μg/mL 1–5 μg/mL 200 μM n/a	ND ND ND ND n/a	Myricetin Scutellarein Emodin  18 compounds identified: M3927, M4367, M4890, M5410, M5789, M6601, M6602, T1434, T1441, T2826, T2831, T4744, T537, T5656, T6791, T8593, T3091, T5242 Sabadanine	Inhibition of 3CLpro	(Liu and Zhou, 2005)
SARS-CoV 3CLpro	Computer modelling	Compounds from marine natural products database and traditional Chinese medicines database Identified via computer modelling. Found in: <i>Veratrum sabadilla</i>	n/a	ND	ND	Aurantiamide acetate	Inhibition of CoV protease	(Toney et al., 2004)
SARS-CoV 3CLpro	Computer modelling for compounds docking in cathepsin-L protease	Found in: <i>Artemisia annua</i>	n/a	ND	ND		Inhibition of active pocket of CoV protease	(Wang et al., 2007)

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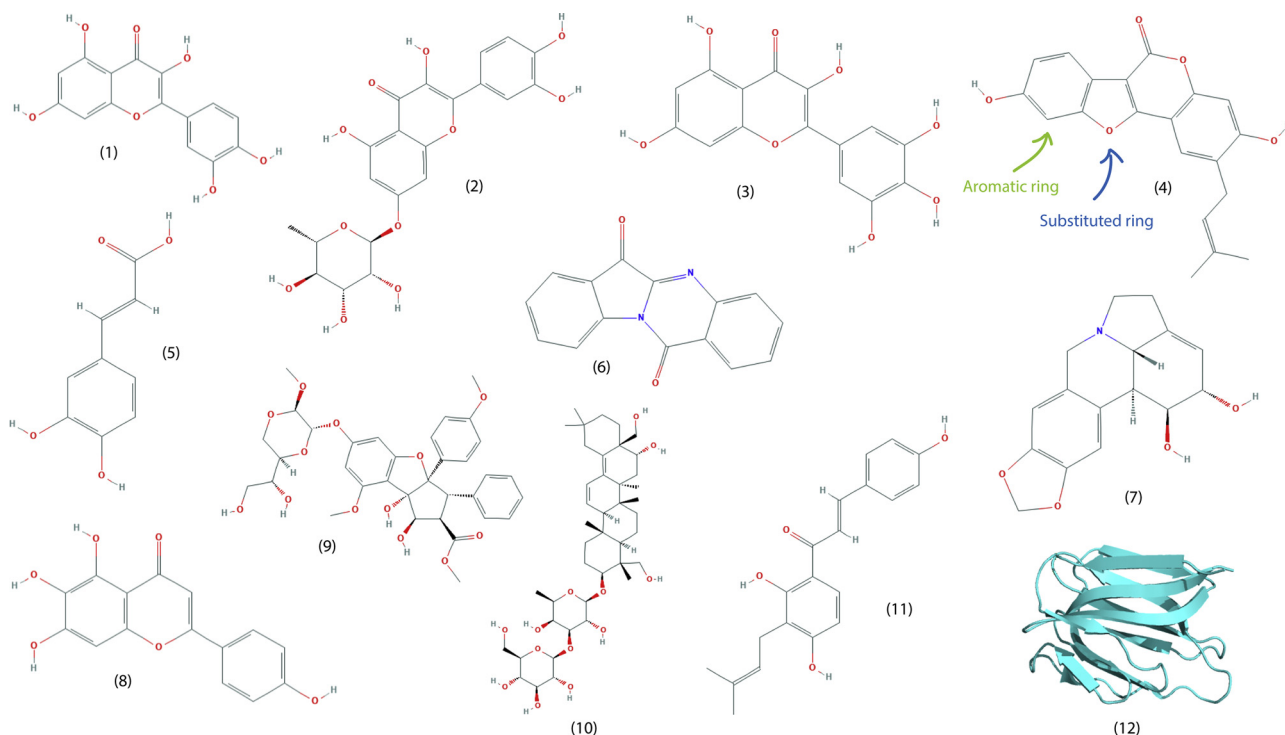


Table 1 (continued)

Viral strain	Assay method	Plant species	Plant part/ isolate	EC <sub>50</sub> or IC <sub>50</sub> (μM unless otherwise stated)	SI	Key compounds present (if identified)	Biological action	Reference
HCoV-229E	XTT assay	<i>Calophyllum blancoi</i>	Isolated compounds from acetone extract of roots	3 15	ND	Blancoxanthone Pyranojacareubin	Not determined	(Shen et al., 2005)
HCoV-229E	XTT assay	Found in: <i>Bupleurum</i> spp., <i>Heteromorpha</i> spp. and <i>Scrophularia scorodonia</i>	Chemical standards used	8.6 ± 0.3 1.7 ± 0.1 19.9 ± 0.1 13.2 ± 0.3	26.6 221.9 19.2 13.3	Saikosaponin A Saikosaponin B <sub>2</sub> Saikosaponin C Saikosaponin D	Possible interference in early stage of viral replication, e.g. absorption and penetration	(Cheng et al., 2006)
HCoV-229E	CPE assay	<i>Pelargonium sidoides</i>	EPs® 7630 (proprietary extract using 11 % ethanol)	44.50 ± 15.84 μg/mL	> 2.3	Not determined	Possibly interference of virus surface resulting in viral inactivation	(Michaelis et al., 2011)
HCoV-229E	Cellular dual luciferase reporter assay	Found in <i>Aglaia</i> sp.	Chemical standard used	3	> 3330	Silvestrol	Specific inhibitor of RNA helicase eIF4A	(Müller et al., 2018)
HCoV-NL63	Plaque viricidal assay	<i>Strobilanthes cusia</i> leaf	Methanol extract	0.64 μg/mL	> 156	Tryptanthrin	Blocking viral RNA genome synthesis and papain-like protease 2 activity	(Tsay et al., 2020)
HCoV-NL63	Virus yield reduction assay	<i>Sambucus formosana</i>	Isolated compounds	2.09	> 191	Indigodole B		
HCoV-NL63			Ethanol extract of stem	1.17 ± 0.75 (μg/mL)	~154			(Weng et al., 2019)
HCoV-OC43			Isolated compounds	3.54 ± 0.77 43.5 ± 6.0	> 141 > 11	Caffeic acid Chlorogenic acid	Inhibits cell docking	
HCoV-299E				71.5 ± 18.4	> 7	Galic acid	Not determined	
HCoV-NL63	CPE assay and neutral red assay	Found in <i>Griffithsia</i> sp.	Chemical standard used	0.048–0.16 0.18–0.33	320–> 2100 > 30–56	Griffithsin	Direct binding to surface envelope glycoprotein spike	(O'Keefe et al., 2010)
HCoV-OC43	MTS assay and qRT-PCR	Found in: <i>Stephania tetrandra</i> and related species	Chemical standards used	< 0.0032 (all μg/mL) 0.33 ± 0.03 1.01 ± 0.07 0.83 ± 0.07	> 3100 > 40.2 11.5 13.6	Terrandrine Fangchinoline Cepharanthine	Inhibit viral replication and expression of viral S and N protein	(Kim et al., 2019)

CLpro = chymotrypsin-like protease; CPE assay = cytopathogenic effect assay; n/a = not applicable to this study; ND = no data; PLpro = papain-like protease.





**Fig. 2.** The structure of selected naturally occurring compounds that demonstrate promising anti-coronavirus activity. (1) Quercetin (2) Quercetin 7-rhamnoside (3) Myricetin (4) Psoralidin (5) Caffeic acid (6) Tryptanthrin (7) Lycorine (8) Scutellarein (9) Silvestrol (10) Saikosaponin B2 (11) Isobavachalcone (12) Griffithsin. As annotated on (4), note the aromatic rings and substituted fused rings present in most compounds.

the inhibition of ATPase activity, but did not directly inhibit helicase activity. This appeared to be the only publication identifying naturally occurring compounds as inhibitors of SARS-CoV helicase.

The need to isolate and synthesise more active agents as part of the development pipeline was highlighted in a study by Runfeng et al. (2020). The authors reported that the Chinese herbal medicine Lianhuaqingwen (comprised of a mixture of plant species) showed antiviral activity against SARS-CoV-2; however, the  $EC_{50}$  was quite high ( $\sim 411 \mu\text{g/mL}$ ). For comparison, the commercial drug remdesivir showed an  $EC_{50}$  of  $0.651 \mu\text{M}$  (approx.  $0.39 \mu\text{g/mL}$ ) using the same assay (Runfeng et al., 2020). This also underscores the potential lack of potency in some traditional medicines promoted for the treatment of coronavirus symptoms.

### 3.2.3. Polyphenols

One family of compounds that demonstrate antiviral activity across a number of studies is the polyphenols. For example, quercetin (Fig. 3) showed an  $IC_{50}$  of  $8.6 \pm 3.2 \mu\text{M}$  against SARS-CoV PLpro (Park et al., 2017). No cell-based assay of antiviral activity was performed. Quercetin (Fig. 2) is a flavonoid found in many foods, but in particularly high levels in certain berries and herbs (Justesen and Knuthsen, 2001; Kaack and Austed, 1998). As previously mentioned (section 3.2.2), the structurally similar polyphenolics myricetin and scutellarein (Fig. 2) display reasonable levels of inhibitory activity against SARS-CoV helicase (Yu et al., 2012).

Bioassay-guided fractionation of the ethanolic extract obtained from *Psoralea corylifolia* seeds has also identified polyphenolics as the bioactive compounds responsible for the activity of this plant species against SARS-CoV PLpro (Kim et al., 2014). Furthermore, six phenolic phytochemicals were isolated from the ethanolic extracts - identified as bavachinin, neobavaisoflavone, isobavachalcone, 4'-O-methylbavachalcone, psoralidin and corylifol A - with their antiviral activity varying widely ( $IC_{50}$  values between  $4.2\text{--}38.4 \mu\text{M}$ ). Again, no cell-based antiviral assays were performed. Isobavachalcone and psoralidin (Fig. 2) showed the greatest antiviral activity, with both found to be

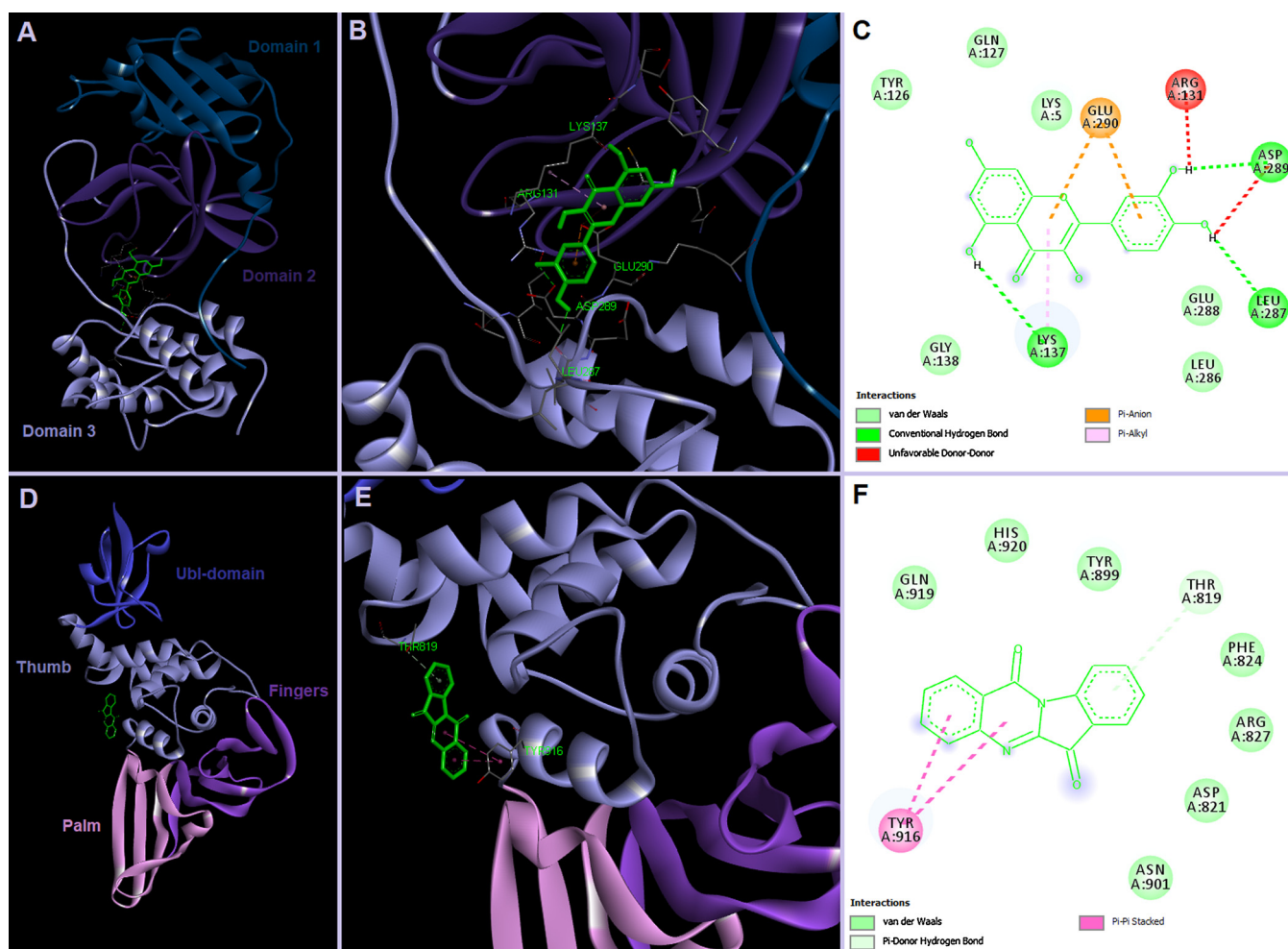
mixed, reversible inhibitors of PLpro through a type I mechanism (i.e. preferentially bind to the free enzyme, rather than the enzyme substrate complex) (Kim et al., 2014).

### 3.2.4. Lectins

Plant lectins, which are proteins that can bind specifically and reversibly to carbohydrate groups (Mitchell et al., 2017), are another group of naturally occurring compounds that may inhibit SARS-CoV. Lectins have shown promise as antiviral agents against viruses such as influenza and herpes simplex virus (Hwang et al., 2020), as well as Ebola (Covés-Datson et al., 2019; Michelow et al., 2011). Remarkably, elevating the plasma levels of recombinant human mannose-binding lectin in mice allowed them to survive otherwise fatal Ebola infections (Michelow et al., 2011). Keyaerts et al. (2007) screened the activity of a broad range of plant lectins (33 in total) against SARS-CoV using a cytopathicity effect (CPE) assay, finding  $EC_{50}$  values as low as  $0.45 \pm 0.08 \mu\text{g/mL}$  for *Lycoris radiata* agglutinin. Although the exact mechanism of action was not determined, activity at the stage of viral attachment or the end of the infectious viral cycle were deemed to be the most likely targets. In clinical trials, other lectins have demonstrated reasonable to good tolerability (Petersen et al., 2006), hence with further testing, they may prove to be one of the more promising classes of naturally derived compound(s) for the treatment of SARS-CoV-2 and other coronavirus infections.

### 3.2.5. Increasing potency via chemical modification

Whilst many natural derived compounds show considerable promise for the inhibition of SARS-CoV and other human coronaviruses, few approach the level of efficacy and/or selectivity required for commercial drugs. For example, the calpain inhibitor MDL28170 has been shown to have an  $IC_{50}$  value of just  $2.5 \text{ nM}$  ( $\sim 1 \text{ ng/mL}$ ) in its activity against the SARS-CoV enzyme cathepsin-L (Simmons et al., 2005). Cathepsin-L is a cysteine protease which is an important lysosomal endopeptidase enzyme involved in the initiation of protein degradation (Sudhan and Siemann, 2015) and mediates entry and infection of SARS-



**Fig. 3.** Possible binding sites of quercetin in SARS-CoV-2 3CLpro and tryptanthrin in PLpro. Docking was performed in AutoDock Vina 1.1.2 (Trott and Olson, 2010) against target proteins generated by SWISS-MODEL (<https://swissmodel.expasy.org/repository/species/2697049>).

CoV in its host cells (Huang et al., 2006). Chemical modification of naturally-derived compounds may be required to increase the potency of their antiviral activity to levels suitable for therapeutic application. For the drug discovery process, beginning with the understanding of the structural conformity (i.e. structure including potential isomers) of the naturally derived compound(s) can reduce timelines and greatly increase the chance of finding effective viral inhibitors.

Working on this principle, Hoever et al. (2005) found that modification of glycyrrhizin, naturally found in liquorice and previously used for the treatment of SARS-CoV (Haiying et al., 2003), could increase its viral inhibition activity. For example, adding 2-acetamido- $\beta$ -D-glucopyranosylamine to the glycoside chain of glycyrrhizin increased its antiviral activity in a CPE assay by 10-fold (decrease in  $EC_{50}$  from  $365 \pm 12 \mu M$  to  $40 \pm 13 \mu M$ ), through increased attraction to the S proteins. Amides and conjugates with amino acid residues and free COOH increased activity of glycyrrhizin by up to 70-fold ( $EC_{50}$  for CPE assay ranging from  $5 \pm 3 \mu M$  to  $139 \pm 20 \mu M$ ), albeit with significantly reduced selectivity. Cho et al. (2013) demonstrated that tomentins A-E (all naturally occurring compounds) showed viral inhibition activity against SARS-CoV greater as compared to their non-geranylated precursor compounds. Similarly, quercetin-7-rhamnoside (Fig. 2) demonstrates over 100 times higher antiviral activity against an animal coronavirus strain compared to its parent compound, quercetin (Choi et al., 2009).

While the examples of modifications presented here vary, it should not be considered that substitutions are added in a random fashion, but rather performed in the context of targeting a specific receptor or

biochemical pathway. For example, the addition of glycosides and specific amino acid residues to glycyrrhizin was performed with the intent of increasing its affinity for the highly glycosylated spike proteins of SARS-CoV. With this in mind, future studies that identify an effective natural inhibitor of coronavirus should also consider the design and testing of potential modifications or synthetic derivatives that could increase its desired activity.

### 3.2.6. The significance of viral strains

A difference in the efficacy of natural therapeutics, mirroring that observed for commercially/synthetically available drugs, has been observed between coronavirus species/strains. For example, Park et al. (2017) found that the compound 3'-(3-methylbut-2-enyl)-3',4,7-trihydroxyflavane, isolated from *Broussonetia papyrifera*, showed good inhibition of PLpro from SARS-CoV ( $IC_{50}$  of  $3.7 \pm 1.6 \mu M$ ), but not against PLpro from MERS-CoV ( $IC_{50}$  of  $112.5 \pm 7.3 \mu M$ ). In contrast, the polyphenolics kazinol F and broussonchalcone A isolated from the same species showed better efficacy against MERS-CoV PLpro. Similarly, O'Keefe et al. (2010) found that the efficacy of the compound griffithsin against SARS-CoV was highest for the Urbani and Tor-II strains ( $EC_{50}$  of 0.61  $\mu g/mL$  using a CPE inhibition assay) and lowest for the Frank strain ( $EC_{50}$  of 1.19  $\mu g/mL$ ). Griffithsin, isolated from the red alga *Griffithsia* sp., is a lectin possessing three large identical carbohydrate-binding domains orientated as an equatorial triangle, which enable multivalent binding (O'Keefe et al., 2010). Although the source of these observed differences in the inhibition activity of griffithsin against different SARS-CoV strains was not explored by O'Keefe et al. (2010), it

may be attributable to genomic differences in the amino acid sequences of the spike proteins between SARS-CoV strains, leading to varying multivalent interactions with the carbohydrates-binding domains and thus differing affinity for binding to the spike proteins. Nevertheless, these studies highlight the importance of conducting *in vitro* tests of potential naturally-derived therapeutics on SARS-CoV-2 prior to animal or clinical trials.

### 3.3. Inhibitors of MERS-CoV

Only a handful of studies have investigated the potential of natural products as therapeutic agents against MERS-CoV. Silvestrol (Fig. 2), a phytochemical from *Aglaia* sp., was found to be a more potent inhibitor of MERS-CoV replication ( $EC_{50}$  of 1.3 nM) (Müller et al., 2018). However, no cell-based antiviral assays were performed. Silvestrol is a specific inhibitor of RNA helicase eIF4A, thus inhibiting viral replication and leading to the inhibition of expression of CoV proteins and preventing the formation of replication/transcription complexes (Müller et al., 2018).

Griffithsin (Fig. 2), a 12.7 kDa lectin found in the *Griffithsia* genus (red algae), is one of the most promising inhibitors of MERS-CoV. It comprises three carbohydrate-binding domains which allow it to bind specifically to glycans on CoV protein spikes and inhibit viral attachment to host cells, with high potency found in *in vitro* trials against MERS-CoV ( $EC_{50}$  of  $\sim 0.125$   $\mu$ M) (O'Keefe et al., 2010) and several HCoV strains ( $EC_{50}$  of 0.0032–0.33  $\mu$ M) (Millet et al., 2016). Griffithsin also appears to have a low systematic toxicity, with its specificity index against HCoV cells (compared to human colorectal adenocarcinoma or fibroblast cell lines) has been estimated at between 30–3100 (O'Keefe et al., 2010), hence showing the potential to be considered as one of the prime candidates for animal and clinical trials against SARS-CoV-2.

### 3.4. Inhibitors of other human coronaviruses

A recent study highlighted the potential anti-HCoV-NL63 activity of the methanolic extract of *Strobilanthes cusia* leaf and its major phytochemicals (Tsai et al., 2020). The *S. cusia* extract effectively reduced virus yield ( $EC_{50}$  = 0.64  $\mu$ g/mL) in cells infected with HCoV-NL63 in a dose-dependent manner. Of the six key compounds isolated, purified and identified via NMR spectroscopy, two exhibited the most potent antiviral activity against HCoV, namely tryptanthrin (Fig. 2), a natural alkaloid containing the basic indoloquinazoline moiety, and indigodole B (5Ar-ethyltryptanthrin), an indole alkaloid derivative. The  $EC_{50}$  values against virus yield from infected cells were 1.52  $\mu$ M and 2.60  $\mu$ M for tryptanthrin and indigodole B, respectively. The increased antiviral activity of tryptanthrin in comparison to indigodole B may result from the double bond at C5a in the former, as compared to the additional ethyl moiety in the latter. This highlights that for compounds based on a tryptanthrin structural conformity, addition of a double bond in the quinazoline ring could significantly increase their antiviral activity. Tryptanthrin has previously been found to have a broad spectrum of biological activities including anticancer, anti-inflammatory, anti-protozoal, antiallergic, antioxidant and antimicrobial action (Kaur et al., 2017). Through manipulation of varying modes of time-of-addition/removal assay, it was found that tryptanthrin prevented the early and late stages of HCoV-NL63 replication, particularly by blocking the viral RNA genome synthesis and papain-like protease 2 activity (Fig. 3), and inhibiting the post-entry stage of HCoV replication (Tsai et al., 2020). Intriguingly, both tryptanthrin ( $EC_{50}$  = 0.06  $\mu$ M) and indigodole B ( $EC_{50}$  = 2.09  $\mu$ M) exhibited strong viricidal activity directly against HCoV-NL63. As the HCoV-NL63 spike protein (S protein) targets the ACE2 receptor, showing a highly conserved sequence and structural similarity to SARS-CoV and SARS-CoV-2 (Letko et al., 2020), tryptanthrin has the potential to be explored as a possible bioactive agent against SARS-CoV-2 and other human coronaviruses.

Another recent study on HCoV-OC43 focussed on the antiviral activities of the bis-benzylisoquinoline alkaloids tetrandrine, fangchinoline, and cepharanthine (CPE assay  $EC_{50}$  values of  $0.33 \pm 0.03$ ,  $1.01 \pm 0.07$  and  $0.83 \pm 0.07$   $\mu$ M, respectively) (Kim et al., 2019). These compounds were the primary bioactive phytochemicals identified in *Stephania tetrandra* and related species. All compounds were found to inhibit virus-induced cell death, through suppressing viral replication, expression of viral S and N proteins (nucleocapsid protein), and the virus-induced host response. However, the effective inhibitory concentrations were not reported for the inhibition of viral protein expression or host response. In addition, tetrandrine activated the p38MAPK pathway in MRC-5 cells, improving the aforementioned host response.

Weng et al. (2019) tested the ethanolic extracts of *Sambucus formosana* (elderberry) stems against the human coronavirus strain HCoV-NL63, finding quite high efficacy ( $EC_{50}$  of  $1.17 \pm 0.75$   $\mu$ g/mL) for viral yield reduction. With further investigation of the phenolic composition of the extracts and antiviral assays performed on the main individual phenolic acids present, caffeic acid (Fig. 2) was identified as the most potent compound present ( $EC_{50}$  of  $3.54 \pm 0.77$   $\mu$ M; or  $\sim 0.64 \pm 0.14$   $\mu$ g/mL). Although no specific binding sites were identified, caffeic acid was found to inhibit the attachment of HCoV to host cells, indicating potential binding to S proteins. Caffeic acid has also been found to inhibit other viruses such as hepatitis B (Wang et al., 2009). Notably, extracts of *Sambucus nigra* (black elderberry), another species in the same genus as *S. formosana*, have been commercialised for the treatment of cold and flu symptoms. Therefore, it is likely that extracts from *S. formosana* would prove similarly nontoxic and suitable for human use, although clinical trials would be required to confirm this. Nevertheless, the bioavailability/delivery mechanism of such extracts would need to be considered in order to reach therapeutic plasma concentrations for viral inhibition (Wittmer et al., 2005).

Cheng et al. (2006) examined the anticoronaviral activity of saikosaponins (A, B<sub>2</sub>, C and D) and their mode of action against HCoV-229E *in vitro*. Saikosaponins represent a group of pentacyclic triterpenoid derivatives, usually present as glycosides, that have been isolated from medicinal plants such as *Bupleurum* spp., *Heteromorpha* spp. and *Scrophularia scorodonia*, previously found to possess efficacy against several viruses, including human immunodeficiency virus (HIV) (Chiang et al., 2003; Ushio and Abe, 1992). All saikosaponins tested showed good to moderate anti-coronavirus activity, with saikosaponin B<sub>2</sub> (Fig. 2) showing the greatest potency ( $EC_{50}$  of  $1.7 \pm 0.1$   $\mu$ M). Subsequent time-of-addition studies indicated that saikosaponin B<sub>2</sub> inhibited viral attachment and penetration. Interestingly, the same compound has been found to inhibit hepatitis C entry (Lin et al., 2015) as well as multidrug resistance-associated drug transporters present on the cell surface (Zhao et al., 2019), indicating that it has potential to display a broad spectrum of bioactivity. This may be advantageous in developing nations where hepatitis C prevalence is high (Pawlotsky, 2014).

Finally, two compounds that have shown efficacy against MERS-CoV (section 3.3) also show promise against HCoV. Müller et al. (2018) found that silvestrol (Fig. 2) inhibited the translation of HCoV-229E proteins, with an  $EC_{50}$  of 3 nM. A follow-up study confirmed that silvestrol can inhibit HCoV-229E in an *ex vivo* bronchial epithelial cell system, with its mechanism of action being the specific inhibition of RNA helicase eIF4A (Müller et al., 2020). Another compound active against MERS-CoV, griffithsin, also has shown great efficacy against several HCoV strains ( $EC_{50}$  of 0.0032–0.33  $\mu$ M) (Millet et al., 2016). This again underscores the prospect of exploring griffithsin for antiviral activity against SAR-CoV-2.

## 4. Animal coronaviruses

Animal coronavirus strains are responsible for severe morbidity events across a wide range of domestic animals and livestock, incurring



major economic demise worldwide (Jackwood et al., 2010; Lelesius et al., 2019; McCutcheon et al., 1995). The genomic diversity, coupled with the ability of coronaviruses to rapidly adapt and mutate, presents unique challenges in the development of novel antiviral agents, hence exploring alternative methods of controlling these viruses could potentially be effective across many or all serotypes. As natural phytochemicals have been shown to have activity across a wide range of viral parsnips (Chen et al., 2014), they form the basis of the studies reviewed here. The majority of plant-derived compounds considered in this review are direct-targeting antivirals, acting through direct inhibition of some part of the virus, such as proteases or spike proteins. For example, silvestrol prevents viral replication occurring through the specific inhibition of RNA helicase eIF4A (Müller et al., 2018), while griffithsin binds directly to the S protein, preventing viral entry to the host cell (O'Keefe et al., 2010). However, host-targeting antivirals form another important group of antiviral compounds. For example, extracts of *Cinnamomi* sp. inhibit the clathrin-dependent endocytosis pathway, thus preventing viral entry to the host cells (Zhuang et al., 2009). Other host-targeting antiviral compounds may stimulate the immune response (Lau et al., 2008).

Table 2 presents a summary of studies reporting anti-viral activity of plant-derived isolates against a range of animal coronavirus strains. Where available, the key compounds responsible for the antiviral activity and their mechanism of action are provided.

#### 4.1. Avian infectious bronchitis virus (IBV)

The antiviral activity of extracts from plant species against the avian IBV viral strains have been extensively studied (Chen et al., 2014; Jackwood et al., 2010; Lelesius et al., 2019; Li et al., 2011; Nguyen et al., 2015; Yang et al., 2010, 2011; Yin et al., 2011). Considering all studies on avian IBV that have established the mechanism of viral inhibition, the main mechanisms of action appear to be through viral envelope disruption or interference with the spike protein (Table 2). A notable exception are the terpenoid compounds 1,8-cineole, (-)- $\alpha$ -pinene and (-)- $\beta$ -pinene, which bind to the IBV nucleocapsid protein (N-protein) (Yang et al., 2010, 2011), inhibiting its interaction with the viral genomic RNA and breaking the IBV replication cycle. These observations were reinforced by conformational models of the terpenoids binding to the active site at the N terminus of the N protein, which indicated that these compounds have the potential to bind strongly to five amino acid residues at this location (TyrA92, ProA134, PheA137, AspA138 and TyrA140) (Yang et al., 2010, 2011). As these residues are highly conserved between IBV strains (Yang et al., 2010), these terpenoids would be expected to show high efficacy against most or all IBV strains, making them a sensible target for further research.

Nevertheless, the envelope protein (E protein) of avian IBV remains the key target for most researchers. The coronavirus E protein is integral to many stages of the viral life cycle, including assembly, budding, envelope formation, and pathogenesis (Schoeman and Fielding, 2019). Interestingly, Chen et al. (2014) reported the inactivation of two distinct enveloped viruses (avian IBV and influenza) following treatment with *Sambucus nigra* extract, suggesting that *S. nigra* extract may have the potential to display a broad spectrum of antiviral effects against many other enveloped viruses. Although the authors raised the possibility of synergistic action of inhibitory compounds within the extract, no fractionation, identification or characterisation of the components was performed. However, flavonoids or lectins were suggested as the likely aetiological agents of the antiviral activity.

#### 4.2. Feline coronavirus

Feline coronavirus is another coronavirus in the Alphacoronavirus group, causing a fatal disease in cats with no effective antiviral treatments available. In one study, *Galanthus nivalis* agglutinin, another lectin, was identified as a potent inhibitor of feline coronavirus (FCoV)

(Hsieh et al., 2010). With an EC<sub>50</sub> of just 0.0088 nM and a high selectivity index (> 218), this carbohydrate-binding protein outperformed all synthetic antiviral agents tested for comparison purposes. For example, the protease inhibitor nelfinavir displayed an EC<sub>50</sub> of 8.19  $\mu$ M and a selectivity index of just 1.4. As with other agglutinins, *Galanthus nivalis* agglutinin binds to the spike and membrane proteins of FCoV, preventing their attachment to the host cells.

McDonagh et al. (2014) conducted a screening study against 19 compounds, focusing on those demonstrated to have previous antiviral effects against coronaviruses or other RNA viruses. Several naturally occurring compounds were included in this study, such as quercetin, curcumin, rutin, glycyrrhizic acid, hesperidin, hesperitin, baicalin and artemisinin. However, none of these compounds reached EC<sub>50</sub> at the concentrations tested. Glycyrrhizic acid was the most promising, causing a 26.7 % reduction in CPE at a concentration of 25  $\mu$ M. As only a single concentration was tested for each compound (ranging from 10 to 50  $\mu$ M, depending on the compound), further investigation of these compounds against FCoV is required.

Theerawatanasirikul et al. (2020) adopted a computer-aided approach to their screening study against feline infectious peritonitis virus (FIPV), a mutated form of the parental enteric form of FCoV. Firstly, compounds from a chemical library were virtually screened for potential binding to the protease 3CL. The 15 most promising compounds were then evaluated *in vitro* using a protease inhibitor assay against FIPV 3CL<sup>pro</sup>. Of the naturally occurring compounds tested, the lowest IC<sub>50</sub> values were shown by 7-methyluteolin (28.5  $\pm$  4.2  $\mu$ M) and stictic acid (29.4  $\pm$  4.6  $\mu$ M). Quercetin 7-rhamnoside (Fig. 2) also showed moderate inhibition (IC<sub>50</sub> of 77.2  $\pm$  13.8  $\mu$ M); however 7-benzyl luteolin and steviol showed no inhibition (IC<sub>50</sub> > 500  $\mu$ M). Subsequent testing of the active compounds using a cytopathic effect (CPE) assay indicated that only stictic acid protected cells from viral-induced death (EC<sub>50</sub> for pre-viral entry treatment of 16.24  $\pm$  1.33  $\mu$ M; selectivity index of 23).

#### 4.3. Porcine endemic diarrhoea virus (PEDV)

Porcine endemic diarrhoea virus is another serious coronavirus from the Alphacoronavirus group. Several studies have investigated the activity of phytochemicals against murine coronaviruses (strain MHV-A59), the most extensive of which were performed by Kim et al. (2008) and Kim et al. (2010). Nevertheless, none of the studies of this viral strain have conclusively managed to determine the specific compounds responsible for viral inhibition, but rather suggested possible compounds or classes of compounds based on their abundance in the extracts tested. The low inhibitory concentrations (< 1  $\mu$ g/mL) of extracts obtained from *Sophorae* sp., *Acanthopanax* sp. and *Torilis* sp. appear to show significant potential. In particular, the high viral specificity (SI = 696) of *Sophorae* sp. root extracts suggests that it could be considered as a prime candidate for future studies on the screening and isolation of compounds from the aforementioned species. The authors suggested that the antiviral activity of these three species are likely to be occurring through inhibition of RNA-dependent RNA polymerase or other protease activity.

Six studies considered natural agents active against porcine endemic diarrhoea virus (PEDV), with the majority of these identifying the compounds with their antiviral activity. In particular, the anti-PEDV activity of griffithsin and quercetin (Fig. 2) and its derivatives deserve particular discussion, given that these compounds have also been found to have antiviral activity against human CoVs (Millet et al., 2016; O'Keefe et al., 2010; Wen et al., 2011). Notably, quercetin was also one of the compounds shortlisted by Zhang et al. (2020) in their virtual screening of inhibitors of SARS-CoV-2 proteases. Against PEDV, quercetin-7-rhamnoside, a disaccharide glucoside, provided 50 % inhibition of viral activity at a concentration of just 0.03  $\mu$ M, approximately 187 times lower than quercetin alone (Choi et al., 2009). Significantly, the specificity of quercetin-7-rhamnoside was exceptionally high (SI =

**Table 2**  
Studies reporting antiviral activity of natural products or isolates against non-human coronavirus strains.

Viral strain	Assay method	Plant species	Plant part/ isolate	EC <sub>50</sub> or IC <sub>50</sub> (μM unless otherwise stated)	SI	Key compounds present (if identified)	Biological action	Reference
Avian IBV	Plaque assay	<i>Alstonia scholaris</i>	Isolated from 50 % ethanol extract	35 55	> 2.8 > 1.8	Alstotide 1 Alstotide 3	As1: interferes with membrane and spike proteins but not nucleocapsid proteins	(Nguyen et al., 2015)
Avian IBV Beaudette strain	CPE and plaque assays	<i>Sambucus nigra</i>	70 % ethanol extract	ND	ND	Possibly flavonols or lectins	Disrupts virion structure and compromises membrane integrity	(Chen et al., 2014)
Avian IBV Beaudette strain	CPE assay <i>In vivo</i> study in chickens	n/a	QR448(a) (emulsion of oleoresins and essential oils)	$\sim 1 \times 10^{-4}$ dilution of extract	ND	Not determined	Possibly disrupt viral membrane or interfere with viral envelope proteins involved in host cell attachment	(Jackwood et al., 2010)
Avian IBV Beaudette strain	CPE (MTT) assay	<i>Mentha piperita</i> <i>Thymus vulgaris</i> <i>Desmodium canadense</i>	40 % ethanol extract	0.004 μg/mL 0.010 μg/mL 0.017 μg/mL	67.5 63.1 17.1	Not determined	Possibly direct inactivation of virus envelope structures	(Lelesius et al., 2019)
Avian IBV Beaudette strain	Plaque assay <i>In ovo</i> and <i>in vivo</i> trials	<i>Houttuynia cordata</i>	<i>H. cordata</i> solution (essential oils and sodium chloride solution)	0.97 mg/mL	> 257	Main component: methyl-nonyl-ketone	Not determined	(Yin et al., 2011)
Avian IBV Gray strain	MTT assay	Found in numerous plants, e.g. eucalypts	Chemical standard used	0.61 ± 0.07 mM	> 16.39	Eucalyptol (1,8-cineole)	Interferes with binding between RNA and IBV nucleocapsid protein	(Yang et al., 2010)
Avian IBV Gray strain	MTT assay	Found in coniferous trees. Produced as by-products of the pulp industry	Chemical standards used	0.98 ± 0.25 mM 1.32 ± 0.11 mM	> 10.20 > 7.58	(-)-α-pinene (-)-β-pinene	May suppress N-protein, hindering binding process between RNA and IBV N-protein	(Yang et al., 2011)
Avian IBV M41	CPE assay and RT-qPCR	Found in: <i>Forsythia suspensa</i>	Chemical standard used	0.64 mM (complete inhibition)	ND	Forsythoside A	Not clear. Appears to affect cell signalling	(Li et al., 2011)
Bovine coronavirus (BCV)	CPE assay	<i>Rosa nulkana</i> <i>Amelanchier alnifolia</i>	Methanol extract (branches)	< 200 μg/mL for both	ND	<i>R. nulkana</i> : ND <i>A. alnifolia</i> : possibly prunasin	Not determined	(McCutcheon et al., 1995)
FCoV NTU156	CPE assay	<i>Galanthus nivalis</i>	Commercial standard	0.0088 nM	> 218	Galanthus nivalis agglutinin Quercetin	Binds to spike and membrane proteins	(Hsieh et al., 2010)
FPV1146 and FECV1683 (FCoVs)	CPE assay	n/a	Commercial standards	> > 10 > > 10 > > 25 > > 25 > > 50 > > 50 > > 10 > > 25	ND	Curcumin Rutin Glycyrrhizic acid Hesperidin Hesperitin Baicalin Artemisinin	Not determined	(McDonagh et al., 2014)
FPV1146 (FCoV)	Virtual screening followed by 3CLpro inhibition assay	Found in: several lichen and plant species	Commercial standards	29.4 ± 4.6 28.5 ± 4.2 77.2 ± 13.8 > 500 > 500	n/a	Stictic acid 7-Methyluteolin Quercetin 7-rhamnoside 7-benzyl luteolin Steviol	Inhibition of 3CLpro	(Theerawatanasirikul et al., 2020)
MHV-A59	Plaque assay	<i>Cimicifuga racemosa</i>  <i>Melia</i> sp. <i>Coptidis</i> sp. <i>Phellodendron</i> sp.  <i>Sophora subprostrata</i>	Methanol extracts	19.4 ± 7.0  13.0 ± 1.4 2.0 ± 0.5 10.4 ± 2.2  27.5 ± 1.1(μg/mL)	12.3  25.6 34.9 13.4  11.1	Ferulic & isoferulic acid? Toosendanin? Berberine? Protuberberine alkaloids? Matrine, oxymatrine, sophoranone & sophocarpine?	Inhibit replication of MHV	(Kim et al., 2008)

(continued on next page)

Table 2 (continued)

Viral strain	Assay method	Plant species	Plant part/ isolate	EC <sub>50</sub> or IC <sub>50</sub> (μM unless otherwise stated)	SI	Key compounds present (if identified)	Biological action	Reference
MHV-A59 Also inhibited MHV-JHM	Plaque assay	<i>Sophora</i> sp. <i>Acanthopanax</i> sp. <i>Sanguisorbæ</i> sp. <i>Tortilis</i> sp.	Methanol extracts	0.8 ± 0.2 0.9 ± 0.1 3.7 ± 1.4 0.8 ± 0.0(μg/mL)	696.0 188.9 105.0 195.6	Not determined	Possibly inhibitors of RNA-dependent RNA polymerase or other protease activity	(Kim et al., 2010)
MHV-A59	Plaque assay	<i>Punica granatum</i>	Pomegranate juice and ethanol/water extract of powder	≥ 200 μg/mL	ND	Possibly polyphenols	May interact with surface glycoproteins spikes	(Sundararajan et al., 2010)
MHV-A59	RT-qPCR	<i>Nigella sativa</i> <i>Anthemis hyaline</i> <i>Citrus sinensis</i>	Ethanol extract	ND	ND	Not determined	Inhibits viral replication via undetermined mechanism	(Ulasli et al., 2014)
PEDV	CPE assay	<i>Ziziphus jujuba</i>	Compounds isolated from methanol extract	13.41 ± 1.13 4.49 ± 0.67 6.17 ± 0.50	> 30.0 47.11 26.75	Jubanine G Jubanine H Nummularine B	Not determined	(Kang et al., 2015)
PEDV CV 777	CPE assay	<i>Ginkgo biloba</i>	Polysaccharides purified from 98 % ethanol extract	1.7 ± 1.3 μg/mL	> 58.8	Mixture of polysaccharides	Dose-dependent inhibition, apparently at viral attachment and entry steps	(Lee et al., 2015)
PEDV CV 777	CPE assay	<i>Houttuynia cordata</i>	Compound isolated from methanol extract	ND	ND	Quercetin 7-rhamnoside	Uncertain. Doesn't obstruct viral mRNA production or interact directly with PEDV	(Song et al., 2011)
PEDV CV 777	CPE assay	<i>Houttuynia cordata</i>	Compounds isolated from methanol extract	~0.03 ± 0.01 ~5.6 ± 2.6 ~0.4 ± 0.4 ~0.7 ± 0.7	7143 215 370 32.7	Quercetin 7-rhamnoside Quercetin Apigenin Luteolin	Not determined	(Choi et al., 2009)
PEDV (KPEDV-9)	CPE assay	<i>Prunus serrulata</i> var. <i>spontanea</i>	80 % methanol extract	1.95 μg/mL	ND	Possibly polyphenols	Not determined	(Yook et al., 2010)
PEDV (NJ-PEDV)	Immunofluorescence assay and RT-qPCR	Found in: <i>Griffithsia</i> sp.	Purified compound used	~0.08	ND	Griffithsin	Prevents viral attachment to host cells	(Li et al., 2019)

BCV = bovine coronavirus; CPE assay = cytopathogenic effect assay; FCoV = feline coronavirus; IBV = (avian) infectious bronchitis virus; MHV = mouse hepatitis virus; n/a = not applicable to this study; ND = no data; PEDV = porcine epidemic diarrhoea virus.



7143), indicating its potential for application in future animal or clinical trials. This study again highlights the importance of considering all possible structural conformities of a compound, including glycosides, in order to identify the most bioactive chemical species. Although the mechanism of action was not determined for quercetin or quercetin-7-rhamnoside against PEDV, previous computer modelling work has indicated that quercetin binds to and inhibits the SARS-CoV proteases PLpro and 3CLpro (Zhang et al., 2020). As previously determined, griffithsin binds to the spike protein of CoV, preventing attachment to host cells (Millet et al., 2016; O'Keefe et al., 2010).

In general, most authors recommend testing multiple coronavirus strains when searching for antiviral activity in naturally occurring compounds. Similarly important is determining their specific mechanism of action. As highlighted by several studies, a selection of compounds that are active against animal coronaviruses are also active against human coronavirus strains (e.g. griffithsin, quercetin). This underscores the potential of utilising compounds with identified activity against animal coronaviruses to guide the discovery of drugs against human coronaviruses, in addition to identifying drugs for the treatment of economically significant animal coronaviruses such as PEDV and avian IBV.

## 5. The significance of chemical polarity

Across both human and animal CoV strains, a clear trend toward the use of chemically polar compounds is evident (Fig. 2). Of the 30 studies that specified the solvent extraction protocols used, ethanol or an ethanol/water combination was the most commonly used (50 % of all studies), followed by methanol or a methanol/water combination (27 %). A further 17 % of relevant literature used a water-based extraction protocol, with only three studies using relatively non-polar solvents (diaminopropane, chloroform and acetone). This is consistent with a wide body of research indicating that relatively polar extracted fractions generally contain a greater level of bioactive and antimicrobial compounds compared to their nonpolar counterparts (Han et al., 2007; Tian et al., 2009; Wigmore et al., 2016). This may also be indicative of increased bioactivity of highly polar glycosylated compounds, similar to the vast difference in antiviral activity observed between quercetin and quercetin-7-rhamnoside (Choi et al., 2009).

In the case of compounds being administered orally, more polar compounds are subject to compartmentalisation within the body, reducing their rate of elimination. However, the potential chemical changes that could occur as a result of highly acidic stomach conditions and the level of absorption in the intestine would need to be considered individually for each compound.

## 6. Conclusions

Naturally occurring phytochemicals provide a valuable and powerful resource of chemical compounds displaying antiviral properties. Further chemical modification of these structures, guided by computer-based docking simulations, may also increase their potency and/or selectivity. Some of the key compounds that show promise for the treatment of coronavirus in humans include scutellarein, silvestrol, tryptanthrin, saikosaponin B<sub>2</sub>, lectins such as griffithsin, lycorine and polyphenolics – including quercetin, myricetin, caffeic acid, psoralidin and isobavachalcone. Needless to mention, these compounds may be toxic at certain levels, and hence *in vitro* and *in vivo* testing is required to determine safe and therapeutic levels for each compound before clinical trials in humans can be performed. Initial studies could focus on compounds which have previously been approved for drug use, or are generally regarded as safe by the FDA or other national organisations, as is the case for some polyphenolic compounds. It is hoped that researchers will be guided by this information presented here in the process of developing safe, effective anti-coronavirus therapeutic agents from naturally derived compounds.

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## Declaration of Competing Interest

The authors declare that no conflict of interest exists.

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