



Structure-based drug designing for potential antiviral activity of selected natural products from Ayurveda against SARS-CoV-2 spike glycoprotein and its cellular receptor

Vimal K. Maurya¹ · Swatantra Kumar¹ · Anil K. Prasad¹ · Madan L. B. Bhatt¹ · Shailendra K. Saxena¹

Received: 17 April 2020 / Accepted: 29 April 2020 / Published online: 24 May 2020
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Abstract The recent outbreak of COVID-19 caused by SARS-CoV-2 brought a great global public health and economic concern. SARS-CoV-2 is an enveloped RNA virus, from the genus *Betacoronavirus*. Although few molecules have been tested and shown some efficacy against SARS-CoV-2 in humans but a safe and cost-effective attachment inhibitors are still required for the treatment of COVID-19. Natural products are gaining attention because of the large therapeutic window and potent antiviral, immunomodulatory, anti-inflammatory, and antioxidant properties. Therefore, this study was planned to screen natural products from Ayurveda that have the potential to modulate host immune system as well as block the virus entry in host cells by interfering its interaction with cellular receptor and may be used to develop an effective and broad-spectrum strategy for the management of COVID-19 as well as other coronavirus infections in coming future. To decipher the antiviral activity of the selected natural products, molecular docking was performed. Further, the drug-likeness, pharmacokinetics and toxicity parameters of the selected natural products were determined. Docking results suggest that curcumin and nimbin exhibits highest interaction with spike glycoprotein (MolDock score – 141.36 and – 148.621 kcal/mole) and ACE2 receptor (MolDock score

– 142.647 and – 140.108 kcal/mole) as compared with other selected natural products/drugs and controls. Also, the pharmacokinetics data illustrated that all selected natural products have better pharmacological properties (low molecular weight; no violation of Lipinski rule of five, good absorption profiles, oral bioavailability, good blood–brain barrier penetration, and low toxicity risk). Our study exhibited that curcumin, nimbin, withaferin A, piperine, mangiferin, thebaine, berberine, and andrographolide have significant binding affinity towards spike glycoprotein of SARS-CoV-2 and ACE2 receptor and may be useful as a therapeutic and/or prophylactic agent for restricting viral attachment to the host cells. However, few other natural products like resveratrol, quercetin, luteolin, naringenin, zingiberene, and gallic acid has the significant binding affinity towards ACE2 receptor only and therefore may be used for ACE2-mediated attachment inhibition of SARS-CoV-2.

Keywords COVID-19 · SARS-CoV-2 · Attachment inhibitor · Natural products · Ayurveda · CAM (Complementary and Alternative Medicine)

Introduction

A recent coronavirus pneumonia pandemic, named COVID-19 by World Health Organization, caused by a novel severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), has been a major threat to global health since December 2019 [[https://www.who.int/emergencies/diseases/novel-coronavirus-2019/technical-guidance/naming-the-coronavirus-disease-\(covid-2019\)-and-the-virus-that-causes-it](https://www.who.int/emergencies/diseases/novel-coronavirus-2019/technical-guidance/naming-the-coronavirus-disease-(covid-2019)-and-the-virus-that-causes-it)]. Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2)/novel coronavirus (2019-nCoV) is a new

Vimal K. Maurya and Shailendra K. Saxena contributed equally to this work.

✉ Shailendra K. Saxena
shailen@kgmcindia.edu

¹ Department of Centre for Advanced Research (CFAR), Faculty of Medicine, King George's Medical University (KGMU), Lucknow 226003, India

strain that has emerged from Wuhan City, Hubei Province of China and spread in more than 195 countries of the world including India [1]. The virus infection leads to over 3.8 million confirmed cases and 0.26 million deaths worldwide [<https://www.who.int/emergencies/diseases/novel-coronavirus-2019>].

SARS-CoV-2 is an enveloped RNA virus, from the genus *Betacoronavirus* and identified as the seventh member of the family *Coronaviridae*, which has infected the human population [2]. Commonly reported symptoms of a COVID-19 patient are fever, headache, vomiting, chills, dyspnea, nausea, sore throat, coughing up blood, shortness of breath, myalgia, diarrhea, and malaise. The severe infection leads to pneumonia, acute respiratory distress syndrome (ARDS) and sometimes multi-organ failures such as kidney failure, and even death [3]. SARS-CoV-2 genome showed higher sequence homology towards SARS-CoV than that of MERS-CoV through the whole genome sequence alignment analysis [4]. The coronavirus genome encodes four structural proteins: spike glycoprotein (S), a small envelope protein (E), matrix glycoprotein (M) and nucleocapsid protein (N) [5]. The spike (S) glycoprotein present in CoVs acts as a viral antigen and responsible for host-receptor binding, virus internalization and induces robust humoral and cell-mediated immune responses in humans during infection. The internalization of SARS-CoV-2 initiates via binding to its cellular receptor angiotensin-converting enzyme 2 (ACE2). The receptor binding and membrane fusion characteristic of spike glycoprotein make its an ideal target for the the attachment inhibitors for the management of COVID-19 [6].

Designing of drugs that can target the host cells or immune system as well as have direct inhibitory action against SARS-CoV-2 may be an effective approach for the treatment of COVID-19 [7]. The innate immune response is critical for controlling the replication and infection of coronavirus. With the advent of new and more efficient screening assays and prediction methods, the efficacy of natural products/drugs that can effectively treat a wide range of viral infections by interfering with different host functions need to be explored during outbreaks [8]. Therefore, the screening of natural products/drugs from Ayurveda that may have the potential to modulate the host immune system as well as block the virus entry into the host cell by interfering with its cellular receptor has been elucidated, which may be an effective and broad-spectrum approach for combating COVID-19. This study may provide the understanding of antiviral activity of natural products which may enable us to establish a potential therapy for patients diagnosed with COVID-19 in a dose-dependent manner with a reduced level of associated toxicity.

Material and methods

Ligand retrieval

A literature search was performed on PubMed and Google to identify natural products that have immunomodulatory properties, shown antiviral activity against RNA viruses, and also used for the management of similar signs and symptoms (clinical manifestations) associated with SARS-CoV-2 infection. The 3 dimensional (3D) structures of; hydroxychloroquine (CID: 3652), nafamostat (CID: 4413), captopril (CID: 44,093), nimbin (CID: 108,058), curcumin (CID: 969,516), withaferin A (CID: 265,237), piperine (CID: 638,024), mangiferin (CID: 5,281,647), thebaine (CID: 5,324,289), berberine (CID: 2353), andrographolide (CID: 5,318,517), quercetin (CID: 5,280,343), luteolin (CID: 5,280,445), resveratrol (CID: 445,154), naringenin (CID: 932), zingiberene (CID: 92,776), β -caryophyllene (CID: 5,281,515), citronellol (CID: 8842), eugenol (CID: 3314), and gallic acid (CID: 370) were retrieved from the NCBI PubChem compound database in SDF format and optimized in Discovery Studio. The chemical structures and their pharmacological properties of selected natural products were given in (Table 1).

Protein retrieval

The 3D crystal structure of spike glycoprotein of SARS-CoV-2 (PDB ID:6VXX) and its receptor Angiotensin-converting enzyme-2 (PDB ID: 1R42) were obtained from the Research Collaboratory for Structural Bioinformatics (RCSB) Protein Data Bank [9, 10].

Molecular docking or structure-based drug design

Molegro Virtual Docker (MVD-3.0.0) has been used for the molecular docking of selected natural products/drugs with spike glycoprotein and angiotensin-converting enzyme-2 [11]. Structure-based drug design approach has been used to identify novel attachment inhibitors against SARS-CoV-2 infection. MVD offers precise predictions of ligand binding modes (87.0%) than other docking software such as (FlexX2: 57.9%, Surflex: 75.3%, GOLD: 78.2%, and Glide: 81.8%) [11]. The potential ligand binding site or cavities was predicted using MVD. Precisely 5 cavities were identified in the protein but only one cavity having maximum surface area and volume was selected as active site based on the active site residues described in the literature [11]. This cavity was utilized in further structure-based docking studies. Nafamostat (spike glycoprotein inhibitor) and captopril (angiotensin-converting enzyme-2 inhibitor) were taken as a control for selected natural

Table 1 List of natural products evaluated for in silico antiviral activity against SARS-CoV-2

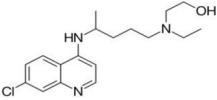
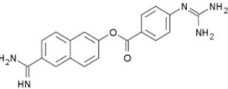
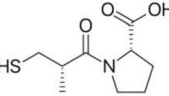
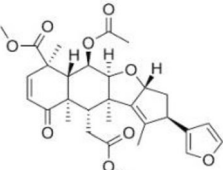
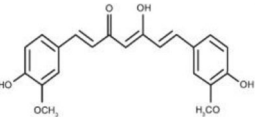
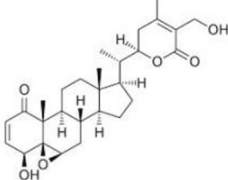
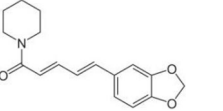
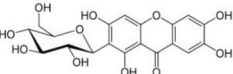
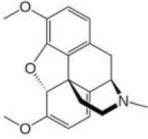
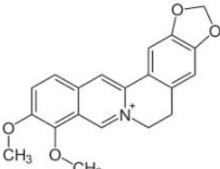
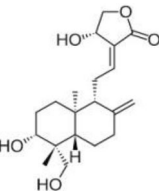
Ligand	Source (<i>Common /Ayurveda or Sanskrit Name</i>)	Structure	Pharmacological functions
Hydroxychloroquine	Synthetic drug		Antimalarial agent, Treatment of COVID-19
Nafamostat	Synthetic drug		Antiviral agent
Captopril	Synthetic drug		Antihypertensive agent
Nimbin	<i>Azadirachta indica</i> (<i>Neem/ Nimbka</i>)		Antiinflammatory, antimicrobial
Curcumin	<i>Curcuma longa</i> (<i>Turmeric/ Haridra</i>)		Antiinflammatory
Withaferin A	<i>Withania somnifera</i> (<i>Indian ginseng/Ashwagandha</i>)		Immunomodulator, antiangiogenic, antitumor agent
Piperine	<i>Piper nigrum</i> (<i>Black pepper/ Maricha</i>)		Antimicrobial, immunomodulator, hepatoprotective, antioxidant
Mangiferin	<i>Mangifera indica</i> (<i>Mango/ Aamra</i>)		Antiviral, andanthelminthic, antiinflammatory, analgesic
Thebaine	<i>Papaver somniferum</i> (<i>Poppy/ Khas-khas</i>)		Analgesic, antitussive
Berberine	<i>Berberis vulgaris</i> (<i>Barberry/ Daruharidra</i>)		Antidiabetic, antihypertensive, antiinflammatory, antioxidant, antidepressant, anticancer, antimicrobial
Andrographolide	<i>Andrographis paniculata</i> (<i>Green chireta/ Kalamegha</i>)		Antiviral, antiinflammatory, antineoplastic

Table 1 continued

Ligand	Source (<i>Common /Ayurveda or Sanskrit Name</i>)	Structure	Pharmacological functions
Quercetin	<i>Citrus aurantium</i> (<i>Bitter orange/Naranga</i>)		Antioxidant, neuroprotective, antiviral, anticancer, antimicrobial, anti-inflammatory
Luteolin	<i>Capsicum annuum</i> (<i>Bell peppers/Mahamarichika</i>)		Antiinflammatory, antioxidant, anticarcinogenic, antihyperalgesic, anxiolytic, antidepressant
Resveratrol	<i>Vaccinium angustifolium</i> (<i>Blueberry/Nila badari</i>)		Antiinflammatory, antioxidative
Naringenin	<i>Citrus paradisi</i> (<i>Grapefruit/Madhukarkati</i>)		Antiinflammatory, antioxidative, antidyslipidemic, antiobesity, antidiabetic, antifibrotic,
Zingiberene	<i>Zingiber officinale</i> (<i>Ginger/Singabera</i>)		Antiinflammatory
β -Caryophyllene	<i>Syzygium aromaticum</i> (<i>Clove/Devakusuma</i>)		Antiinflammatory
Citronellol	<i>Rosa damascena</i> (<i>Damask rose/Satapatri</i>)		Antiinflammatory, antibacterial, antifungal
Eugenol	<i>Ocimum sanctum</i> (<i>Holy basil/Tulsi</i>)		Antioxidant, neuroprotective, antiviral, anticancer, antimicrobial, antiinflammatory
Gallic acid	<i>Emblica officinalis</i> (<i>Indian gooseberry/Amalaki</i>)		Antibacterial, antifungal, antiviral, antiinflammatory, antioxidant, anticancer, antidiabetic

Table 2 Molecular docking results are indicated with maximum binding affinity of selected natural products against SARS-CoV-2 spike glycoprotein and its cellular receptor along with MolDock score and other parameters

Ligand name	MolDock score (kcal/mole)	Interaction (kcal/mole)	H-bond (kcal/mole)	Interacting amino acid residues
<i>Spike glycoprotein of SARS-CoV-2 (6VXX)</i>				
Hydroxychloroquine	- 105.28	- 116.92	- 4.57	Gln314, Ser316, Asn317,Thr761, Asn764, Arg765, Thr768
Nafamostat	- 114.63	- 124.91	- 8.55	Ile312, Gln314, Ser316, Thr761, Arg765
Nimbin	- 148.62	- 142.14	- 15.87	Lys304,Thr315, Ser316, Asn317, Arg765
Curcumin	- 141.36	- 143.70	- 16.39	Thr302, Lys304, Gln314, Thr315, Asn317, Asn764, Arg765, Thr768
Withaferin A	- 108.56	- 136.26	- 7.69	Lys303, Lys304, Tyr313, Gln314, Asn764, Arg765, Thr768
Piperine	- 104.56	- 110.91	- 8.55	Gln314, Ser316, Asn317, Arg765, Thr768
Mangiferin	- 104.26	- 136.54	- 23.51	Gln314,Thr315, Asp737, Thr739, Thr761, Asn764, Arg765,
Thebaine	- 103.58	- 113.19	- 10.77	Tyr313, Gln314, Asn764, Arg765, Thr768
Berberine	- 99.93	- 117.62	- 9.41	Ser316, Asn317, Asn764, Arg765
Andrographolide	- 98.80	- 113.55	- 5.21	Lys303, Tyr313, Gln314, Thr761, Arg765
Quercetin	- 86.22	- 115.16	- 14.48	Gln314, Ser316, Asn317, Asn764, Arg765,Thr768
Luteolin	- 85.01	- 107.21	- 10.32	Ile312, Gln314, Asn764, Arg765,Thr768
Resveratrol	- 83.40	- 90.38	- 6.06	Gln314, Thr761, Asn764, Arg765, Thr768
Naringenin	- 82.10	- 103.13	- 11.73	Lys304, Arg765, Thr768
Zingiberene	- 75.49	- 81.61	0	-
β-caryophyllene	- 72.37	- 71.15	0	-
Citronellol	- 69.44	- 68.55	- 5.59	Gln314, Ser316, Asn317, Arg765
Eugenol	- 66.55	- 73.74	- 6.19	Gln314, Thr315, Asn317, Asn764, Arg765,
Gallic acid	- 63.54	- 72.46	- 16.36	Ser758, Gln957, Lys964, Gln965
<i>Angiotensin-converting enzyme 2 (1R42)</i>				
Hydroxychloroquine	- 105.05	- 113.20	- 3.25	Ala348, Trp349, Asp350, Glu375, His378, Asp382, His401
Captopril	- 73.69	- 81.15	- 13.07	His378, Asn394, Glu398, His401, Glu402
Curcumin	- 142.64	- 139.52	- 9.49	Ala348, His378, Asn394, Tyr385, His401, Glu402
Nimbin	- 140.10	- 125.30	- 12.76	His378, Asn394, His401, Tyr510, Tyr515, Arg518
Piperine	- 112.83	- 120.05	- 1.10	Gly399, His401, Glu402, Arg514, Arg518
Withaferin A	- 112.38	- 139.67	- 11.16	Thr347, Ala348, Glu375, His378, Phe390, Arg393, Phe400, His401
Mangiferin	- 104.15	- 137.74	- 23.11	Asp206, Ala348, Asp382, Asn394, Gly395, Glu398, His401, Arg514
Thebaine	- 100.77	- 108.58	- 6.23	His378, Asn394, His401
Andrographolide	- 99.354	- 116.74	- 14.38	Ala348, His378, Asp382, Tyr385, Asn397, Phe400, His401, Glu402, Arg514
Berberine	- 97.54	- 116.92	- 0.429	Ala348, Glu375, His378, Phe390, His401
Resveratrol	- 96.81	- 103.7	- 5.26	Asp382, Phe390, Arg393, His401
Quercetin	- 92.05	- 112.59	- 7.86	Asp350, Tyr385, Phe390, Arg393
Luteolin	- 89.95	- 108.26	- 22.58	His378, Tyr385, Glu398, His401, Glu402, Arg514
Naringenin	- 83.42	- 102.46	- 5	Ala348, Asp350, His378, Asp382, Tyr385, Arg393, Asn394, His401
Zingiberene	- 82.85	- 88.32	0	His378, His401
Gallic acid	- 75.64	- 84.60	- 16.88	Asp206, Asn394, Gly395, Asn397, His401, Arg514
β-caryophyllene	- 73.42	- 72.20	0	Phe390, Leu391, Arg393, Phe400
Citronellol	- 69.61	- 71.37	- 5	Phe390, Arg393, Phe400, His401
Eugenol	- 67.46	- 77.25	- 2.5	His401, Glu402, Arg514

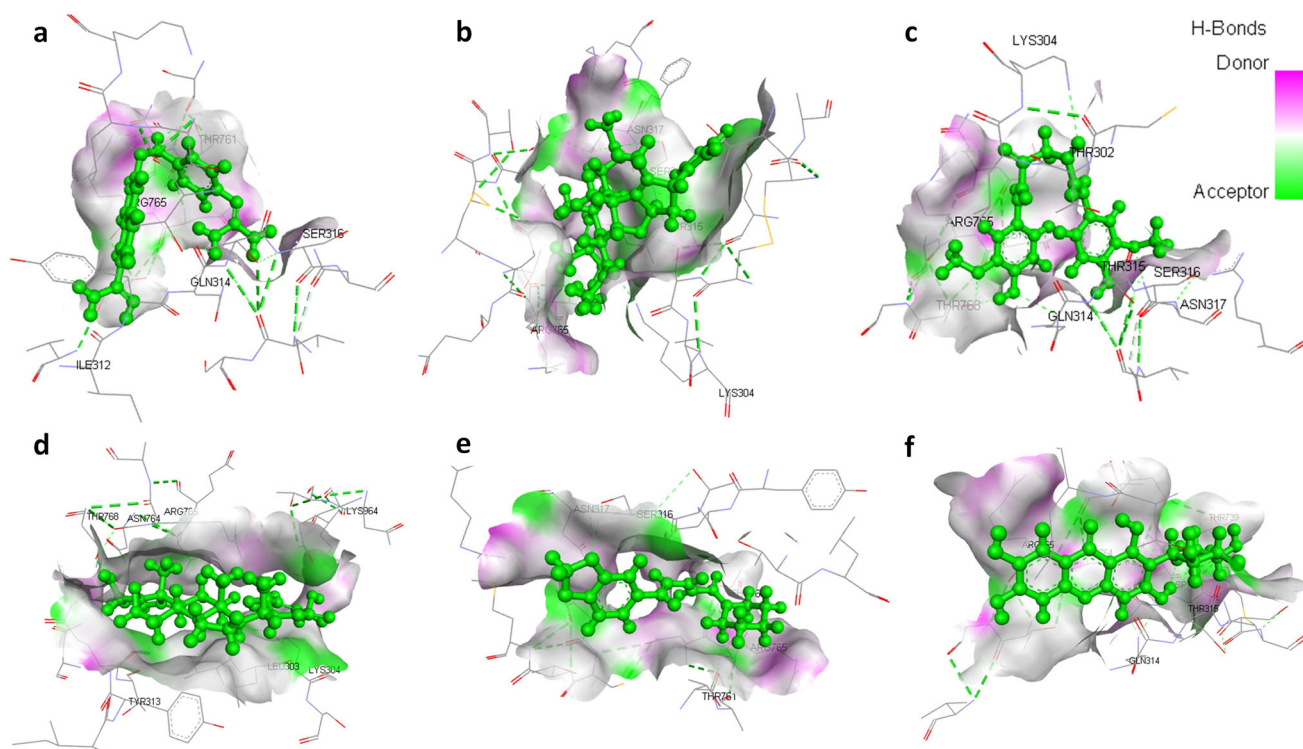


Fig. 1 3D structural views of ligand-binding site. The 3D structures display H-bond interactions (green dashed lines) of ligands **a** nafamostat, **b** nimbin, **c** curcumin, **d** withaferin A, **e** piperine, and **f** mangiferin with the Spike glycoprotein of SARS-CoV-2 (6VXX) (color figure online)

products/drugs [6]. The docking score of selected natural products/drugs was compared with the docking score of nafamostat for spike glycoprotein and docking score of captopril for angiotensin-converting enzyme-2 respectively. The natural products that showed higher docking score than nafamostat and captopril was considered a potent inhibitor of spike glycoprotein and angiotensin-converting enzyme-2 respectively. Considering the suggested use in COVID-19, the hydroxychloroquine was also taken for comparison in this study.

Absorption, distribution, metabolism, excretion, toxicity, and receptor binding analysis

Physico-chemical and pharmacokinetic properties have a vital role in the discovery of novel antiviral agents. The drug-likeness properties of the selected natural products were investigated via swissADME server [12]. ADME-toxicity of the selected natural products was analyzed by the admetSAR server. The absorption, water solubility, human oral bioavailability, blood–brain barrier (BBB) transport, and plasma protein binding were calculated. Additionally, carcinogenicity, hepatotoxicity, acute oral toxicity and receptor binding properties of these compounds were also calculated [13].

Results

Inhibitory effect of natural products against spike glycoprotein of SARS-CoV-2

The binding energies of selected natural products (Table 1) with spike glycoprotein were studied and docking results are given in terms of MolDock score, interaction energy, H-bond energy and interacting amino acid residues present at the active site of protein (Table 2). Docking results revealed that out of 17 selected natural products/drugs; only nimbin and curcumin have shown higher score than nafamostat (MolDock score -114.633 kcal/mole, interaction energy -124.911 kcal/mole and forms five hydrogen bonds with Ile312, Gln314, Ser316, and Arg765 amino acid residues present at the active site of protein) and hydroxychloroquine (MolDock score -105.28 kcal/mole, interaction energy -116.92 kcal/mole and forms two hydrogen bond with Gln314 and Asn317) (Fig. 1). Nimbin showed the highest binding affinity toward spike glycoprotein (MolDock score -148.621 kcal/mole and interaction energy -142.145 kcal/mole) and also forms three hydrogen bond with Lys304, Ser316, and Asn317. Similarly, curcumin has been found to interact with spike glycoprotein with a high binding affinity (MolDock score -141.36 kcal/mole and interaction energy

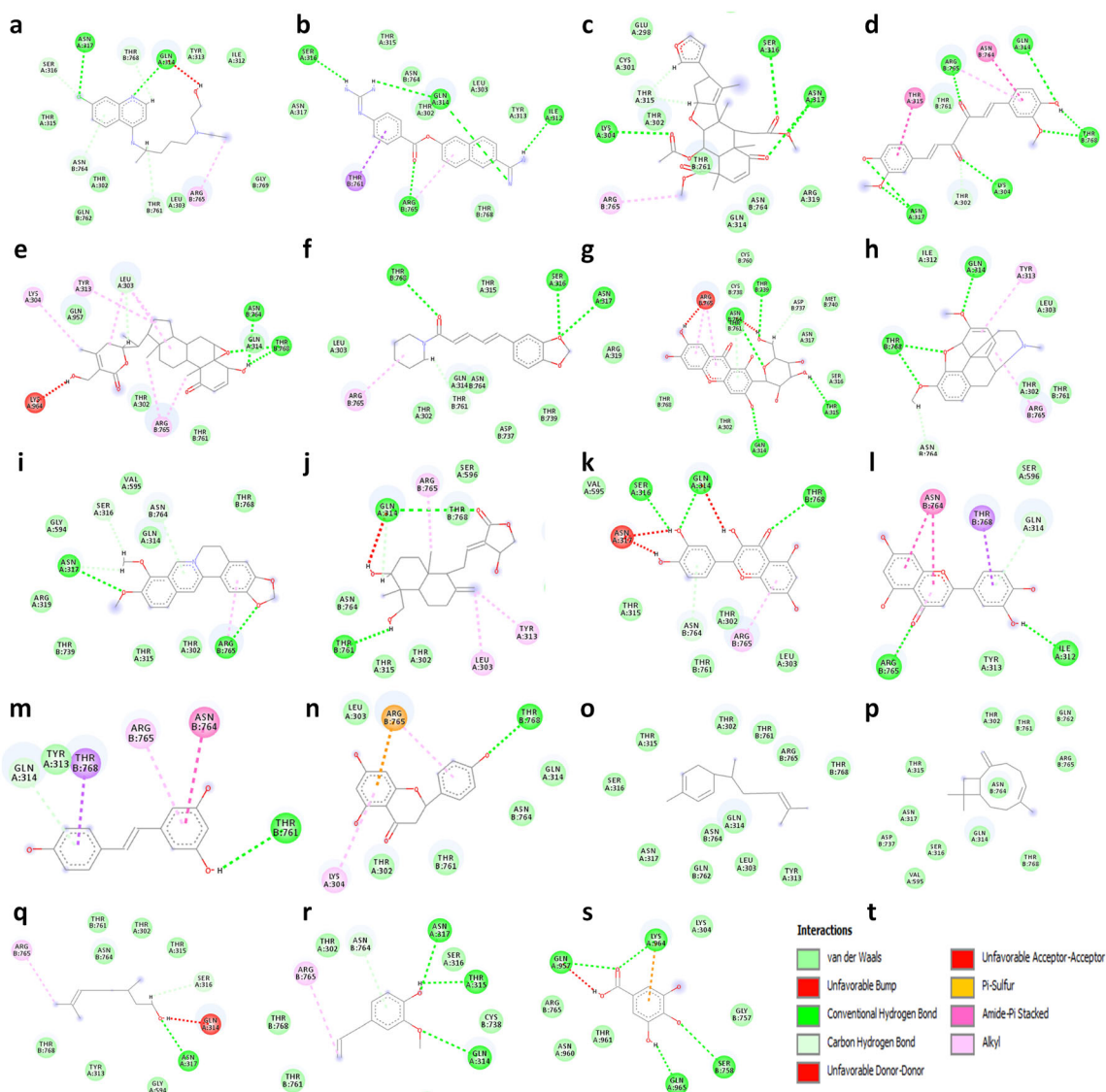


Fig. 2 2D visualization of molecular interaction of Spike glycoprotein of SARS-CoV-2 (6VXX) with **a** hydroxychloroquine, **b** nafamostat, **c** nimbin, **d** curcumin, **e** withaferin A, **f** piperine, **g** mangiferin, **h** thebaine, **i** berberine, **j** andrographolide, **k** quercetin, **l** luteolin, **m** resveratrol, **n** naringenin, **o** zingiberene, **p** β -caryophyllene, **q** citronellol, **r** eugenol, **s** gallic acid and **t** displays the color indication for interactions. H-bonding (dark green circles associated with the green dashed lines); van der Waals forces (medium light

green circles); carbon–oxygen dipole–dipole interaction (light green circles with dashed lines); alkyl-pi interactions (light pink circles with dashed lines); T-shaped pi–pi stacking and (parallel) pi–pi stacking (both indicated with dark violet circles); cation-pi interaction (orange circle). The blue halo surrounding the interacting residues represents the solvent accessible surface that is proportional to its diameter. Images were generated using discovery studio visualizer 4 (color figure online)

– 143.705 kcal/mole) and forms six hydrogen bond with Lys304, Gln314, Asn317, Arg765, and Thr768. In addition, it has been found that the nimbin and curcumin have better binding affinity toward spike glycoprotein as compared to nafamostat and hydroxychloroquine. Apart from nimbin and curcumin, other selected natural products/drugs such as withaferin A, piperine, mangiferin, thebaine, berberine, and andrographolide have also demonstrated a significant binding affinity towards spike glycoprotein of SARS-CoV-2 with significant hydrogen bond interactions with various residues (Fig. 2).

Inhibitory effect of natural products against Angiotensin-converting enzyme 2

Many studies have shown that host ACE2 acts as a specific receptor for the Spike receptor binding domain (RBD) of SARS-CoV-2. Therefore, targeting ACE2 may be considered as host-specific therapy to block SARS-CoV-2 from entering into the host cells and may facilitate the design of the novel attachment inhibitors for the treatment of COVID-19. The docking results revealed that all the selected natural drugs showed higher binding affinity

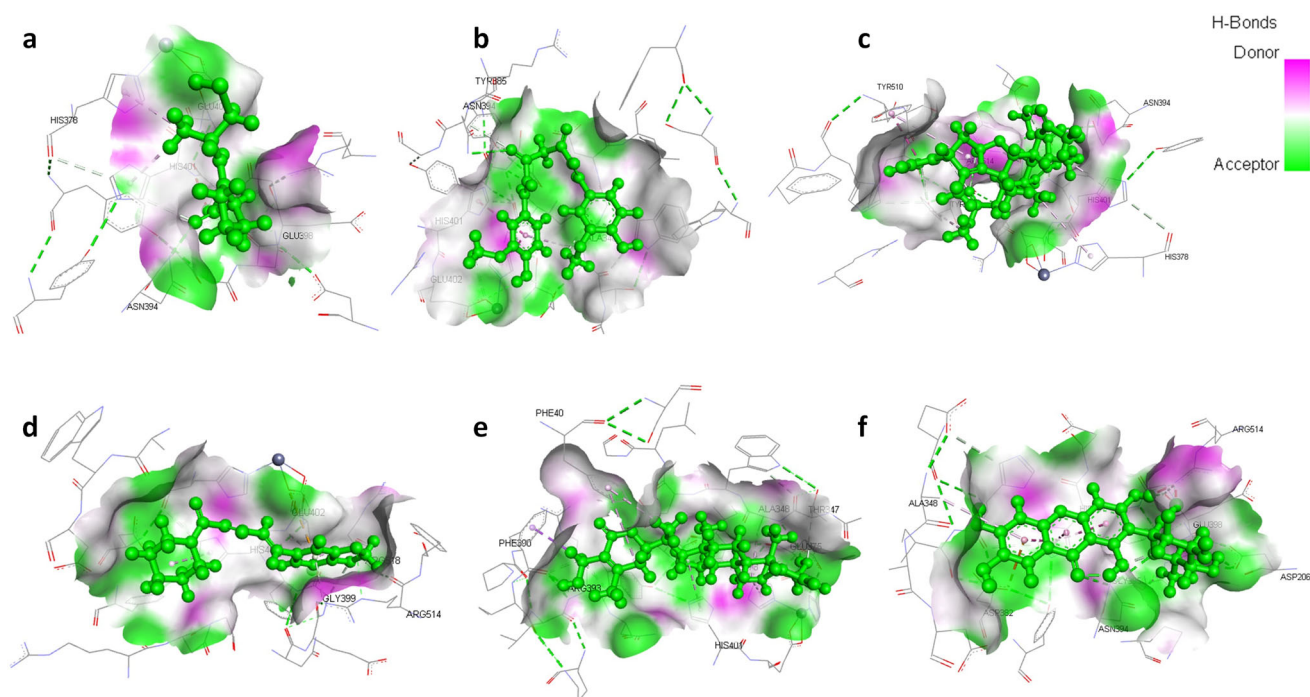


Fig. 3 3D structural views of ligand-binding site. The 3D structures display H-bond interactions (green dashed lines) of ligands **a** captopril, **b** curcumin, **c** nimbin, **d** piperine, **e** withaferin A, and **f** mangiferin with the Angiotensin-converting enzyme 2 (1R42) (color figure online)

towards ACE2 except β -caryophyllene, citronellol, and eugenol as compared to captopril which is a known inhibitor of the enzyme. We have found that captopril interacts with amino acid residues His378, Asn394, Glu398, His401, and Glu402 of ACE2 to generate a binding score (MolDock score of -73.6975 kcal/mole and interaction energy -81.1588 kcal/mole) and forms three hydrogen bonds with Glu402 and Asn394 (Fig. 3). In addition, hydroxychloroquine is also evaluated for ACE2 binding property, which exhibited significant interaction with the protein (MolDock score -105.05 kcal/mole and interaction energy -113.20 kcal/mole) and form hydrogen bond with Asp350 present at the active site of protein. Among the selected natural products, curcumin has the highest binding affinity for ACE2 followed by nimbin, piperine, withaferin A, mangiferin, thebaine, andrographolide, berberine, resveratrol, quercetin, luteolin, naringenin, zingiberene, and gallic acid. Curcumin interacts with Ala348, His378, Asn394, Tyr385, His401, and Glu402 at the active site of the protein and generated MolDock score and interaction energy (-142.647 kcal/mole and -139.525 kcal/mole) and forms two hydrogen bonds with Tyr385 and Asn394. Similarly, nimbin also interacts with His378, Asn394, His401, Tyr510, Arg514, and Tyr515 at the active site and generated moldock score and interaction energy (-140.108 kcal/mole and -125.304 kcal/mole) and forms four hydrogen bonds with Asn394, Arg514, and Tyr515. Unlike spike glycoprotein, curcumin and nimbin

again showed better binding affinity toward ACE2 compared to selected natural drugs, controls, hydroxychloroquine and ranked top two molecules that have inhibitory effect for SARS-CoV-2 spike glycoprotein as well as its cellular receptor ACE2 (Fig. 4).

Absorption, distribution, metabolism, excretion, toxicity and receptor binding analysis:

Drug-likeness properties play a crucial role in antiviral drug development. The calculated drug-likeness parameters of selected natural products using swissADME are given in (Table 3). The absorption profile is directly related to the availability of drug in systemic circulation. The admetSAR results revealed that all the selected natural drugs exhibited good intestinal absorption after the oral uptake in the gastrointestinal tract. In addition, all the selected natural drugs also showed good water solubility and oral bioavailability whereas molecules such as nimbin, piperine, mangiferin, thebaine, berberine, andrographolide, quercetin, luteolin, resveratrol, naringenin, zingiberene, citronellol, and eugenol have shown poor human oral bioavailability. Distribution of drugs to the various tissue and organs guaranteed its better treatment efficiency. After the drug distribution analysis we have found that all the natural drugs have good blood–brain barrier penetration except curcumin, mangiferin, quercetin, luteolin, resveratrol, naringenin, and gallic acid. Furthermore, based on

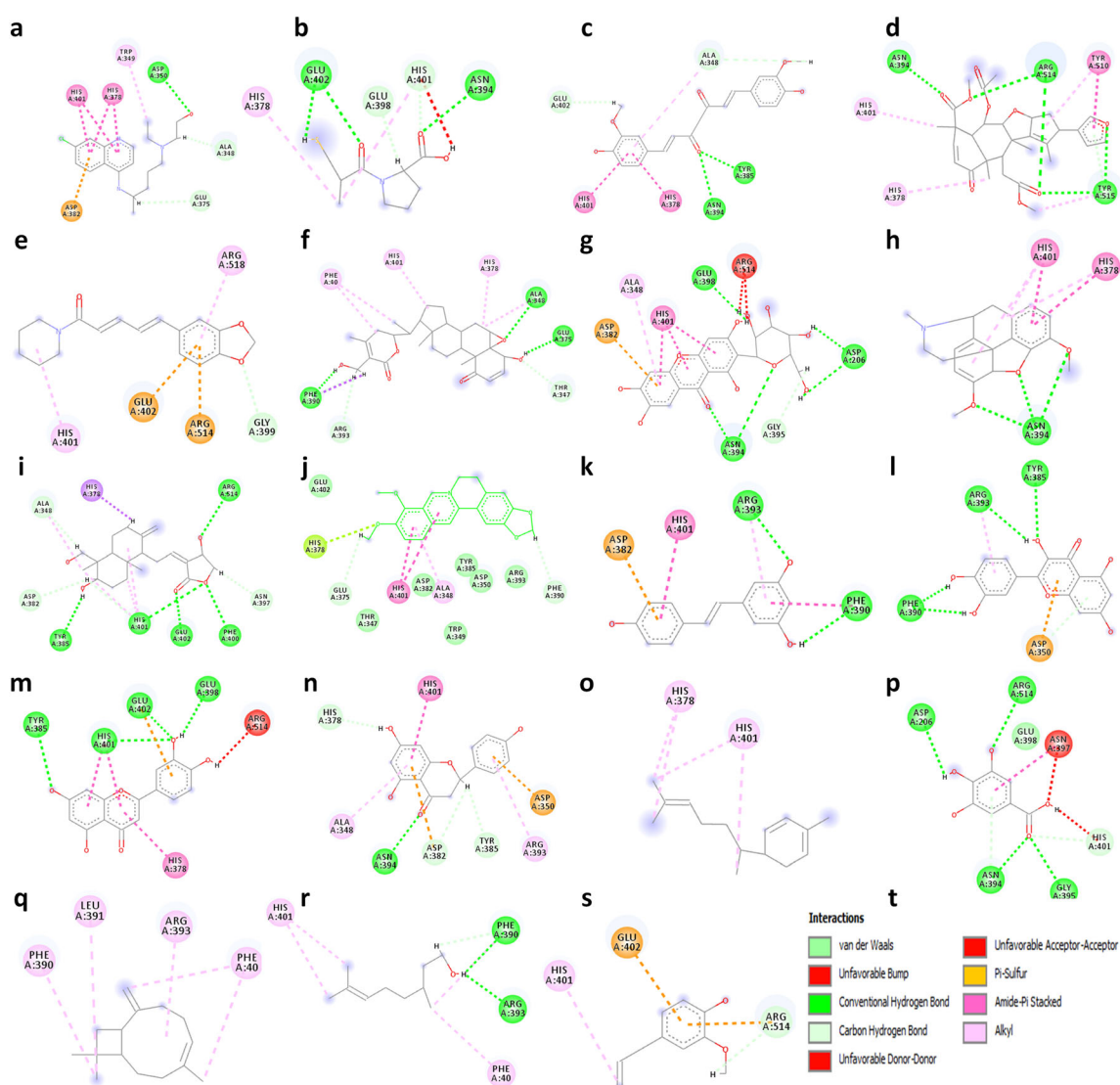


Fig. 4 2D visualization of molecular interaction of Angiotensin-converting enzyme 2 (1R42) with **a** hydroxychloroquine, **b** captopril, **c** curcumin, **d** nimbin, **e** piperine, **f** withaferin A, **g** mangiferin, **h** thebaine, **i** andrographolide, **j** berberine, **k** resveratrol, **l** quercetin, **m** luteolin, **n** naringenin, **o** zingiberene, **p** gallic acid **q** β -caryophyllene **r** citronellol, **s** eugenol and **t** displays the color indication for interactions. H-bonding (dark green circles associated with the green dashed lines); van der Waals forces (medium light green circles);

carbon–oxygen dipole–dipole interaction (light green circles with dashed lines); alkyl-pi interactions (light pink circles with dashed lines); T-shaped pi–pi stacking and (parallel) pi–pi stacking (both indicated with dark violet circles); cation–pi interaction (orange circle). The blue halo surrounding the interacting residues depicts the solvent accessible surface that is proportional to its diameter. Images were generated using Discovery Studio Visualizer 4.5 (color figure online)

distribution profile, non-substrates or non-inhibitors of P-glycoprotein (ABCB1) and renal organic cationic transporter (OCT2) of the drugs were predicted which facilitate efflux of xenobiotics from the cells. The drug metabolism analysis suggested that all the selected natural drugs are predicted as either non-substrates or non-inhibitors of CYP1A2, CYP2D6, CYP2C9, CYP2C19, and CYP3A4 enzymes. Further, the drugs were evaluated for carcinogenicity and hepatotoxicity and no drugs have shown carcinogenic effect whereas nimbin, curcumin, mangiferin,

berberine, quercetin, luteolin, resveratrol, and naringenin were found to be associated with hepatotoxicity. The receptor binding analysis suggested that except the zingiberene, β -caryophyllene, citronellol, eugenol, and gallic acid, all the selected natural drugs have shown significant binding with cellular receptors such as estrogen receptor, androgen receptor, thyroid receptor, glucocorticoid receptor, aromatase and PPAR gamma receptor (Tables 4, 5 and 6).

Table 3 Drug likeness properties of selected natural products

Ligand	Molecular formula	Molecular weight (g/mol)	LogP	Hydrogen bond donor	Hydrogen bond acceptor	Topological polar surface area (Å ²)
Hydroxychloroquine	C ₁₈ H ₂₆ ClN ₃ O	335.9	3.6	2	4	48.4
Nafamostat	C ₁₉ H ₁₇ N ₅ O ₂	347.4	2.0	4	4	141
Captopril	C ₉ H ₁₅ NO ₃ S	217.29	0.3	2	4	58.6
Nimbin	C ₃₀ H ₃₆ O ₉	540.6	2.3	0	9	118
Curcumin	C ₂₁ H ₂₀ O ₆	368.4	3.2	2	6	93.1
Withaferin A	C ₂₈ H ₃₈ O ₆	470.6	3.8	2	6	96.4
Piperine	C ₁₇ H ₁₉ NO ₃	285.34	3.5	0	3	38.8
Mangiferin	C ₁₉ H ₁₈ O ₁₁	422.3	- 0.4	8	11	197
Thebaine	C ₁₉ H ₂₁ NO ₃	311.4	2.2	0	4	30.9
Berberine	C ₂₀ H ₁₈ NO ₄ ⁺	336.4	3.6	0	4	40.8
Andrographolide	C ₂₀ H ₃₀ O ₅	350.4	2.2	3	5	87
Quercetin	C ₁₅ H ₁₀ O ₇	302.23	1.5	5	7	127
Luteolin	C ₁₅ H ₁₀ O ₆	286.24	1.4	4	6	107
Resveratrol	C ₁₄ H ₁₂ O ₃	228.24	3.1	3	3	60.7
Naringenin	C ₁₅ H ₁₂ O ₅	272.25	2.4	3	5	87
Zingiberene	C ₁₅ H ₂₄	204.35	5.2	0	0	0
β-caryophyllene	C ₁₅ H ₂₄	204.35	4.4	0	0	0
Citronellol	C ₁₀ H ₂₀ O	156.26	3.2	1	1	20.2
Eugenol	C ₁₀ H ₁₂ O ₂	164.2	2.0	1	2	29.5
Gallic acid	C ₇ H ₆ O ₅	170.12	0.7	4	5	98

Discussion

Coronavirus is an enveloped RNA virus, from the genus *Betacoronavirus* that is transmitted among birds, humans, and other mammals. The recent outbreak of COVID-19 caused by SARS-CoV-2 is a serious global public health and economic concern that emphasizes the significance of restricting infectious agents at international borders [14]. In the absence of approved antiviral agent for SARS-CoV-2 infection, more than 80 clinical trials of 20 drugs such as traditional Chinese medicines (TCM), human immunoglobulin, interferons, methylprednisolone, chloroquine, arbidol, remdesivir, favipiravir, lopinavir, ritonavir, oseltamivir, bevacizumab, and hydroxychloroquine have been launched for the treatment of COVID-19 [15]. Natural products are gaining interest because of the large therapeutic window and potent antiinflammatory and antioxidant properties. Recently, various drugs from natural origin have shown antiviral activity against the number of viruses such as Influenza virus, Zika virus, Dengue virus, Japanese encephalitis virus, Herpes simplex virus, Hepatitis B and C virus, Respiratory syncytial virus, and Human immunodeficiency virus [16]. The importance of natural products for health and disease treatment have immense importance throughout human evolution. Several natural products derived from plants for thousands of years

have been traditionally used to treat various types of human illnesses including viral infections. Natural plants such as *Azadirachta indica*, *Curcuma longa*, *Withania somnifera*, *Ocimum sanctum*, *Embllica officinalis*, *Piper nigrum*, *Mangifera indica*, *Andrographis paniculata*, and *Zingiber officinale* etc. have been listed in AYUSH system of medicine and extensively used as complementary and alternative medicine for the management of chronic fever, inflammation and auto-immune disorders. In addition, these natural products/drugs from Ayurveda have significant pharmacological activities that regulate various vital cell signaling pathways that cause cytotoxic, genotoxic, and mitogenic reactions leading to various disease pathologies [17].

The development of drugs with broad-spectrum antiviral activities is a long pursued goal in the drug discovery. This research is primarily focused to develop therapeutics targeting the spike glycoprotein of SARS-CoV-2 and its host receptor ACE2 to find potential attachment inhibitors for the treatment of COVID-19. Recently various studies have shown that hydroxychloroquine may have potential while combating the SARS-CoV-2 infection [18–20]. However, we have found that natural drugs such as nimbin, curcumin, withaferin A, piperine and nafamostat (known spike glycoprotein inhibitor) have better binding affinity toward spike glycoprotein and its cellular receptor ACE2

Table 4 Pharmacokinetics, toxicities and receptor binding properties of hydroxychloroquine, nafamostat, captopril, nimbin, curcumin withaferin A and piperine

Parameters	Hydroxychloroquine	Nafamostat	Captopril	Nimbin	Curcumin	Withaferin A	Piperine
Absorption							
Human intestinal absorption	+	+	+	+	+	+	+
Caco-2	+	-	+	-	-	-	+
Human oral bioavailability	+	-	+	-	+	+	-
Water solubility	- 3.566	- 4.074	- 1.798	- 4.455	- 3.364	- 4.203	- 3.398
Distribution							
Subcellular localization	Lysosomes	Mitochondria	Mitochondria	Mitochondria	Mitochondria	Mitochondria	Mitochondria
Blood-brain barrier penetration	+	+	+	+	-	+	+
Plasma protein binding	0.756	0.739	0.367	0.788	0.832	0.944	1.058
P-glycoprotein inhibitor	-	-	-	+	+	+	-
P-glycoprotein substrate	+	-	-	+	-	+	-
Metabolism							
OATP2B1 inhibitor	-	-	-	-	-	-	-
OATP1B1 inhibitor	+	+	+	-	+	+	+
OATP1B3 inhibitor	+	+	+	-	+	+	+
MATE1 inhibitor	-	+	-	-	-	-	-
OCT2 inhibitor	-	-	-	-	-	-	-
BSEP inhibitor	+	+	-	+	+	+	+
CYP3A4 substrate	-	-	-	+	-	+	-
CYP2C9 substrate	+	-	-	-	-	-	-
CYP2D6 substrate	-	-	-	-	-	-	-
CYP3A4 inhibition	+	-	-	+	-	-	+
CYP2C9 inhibition	-	-	-	-	+	-	-
CYP2C19 inhibition	-	-	-	-	+	-	-
CYP2D6 inhibition	-	-	-	-	+	-	+
CYP1A2 inhibition	-	+	-	-	+	-	+
CYP inhibitory promiscuity	-	-	-	+	+	-	+
Toxicity							
Carcinogenicity	-	-	-	-	-	-	-
Human ether-a-go-go inhibition	+	-	-	+	-	+	+
Hepatotoxicity	-	+	-	+	+	-	-
Acute oral toxicity(kg/mol)	2.665	2.33	1.853	3.955	1.992	3.276	2.201
Receptor binding							
Estrogen receptor binding	-	+	-	+	+	+	+
Androgen receptor binding	-	+	-	+	+	+	+
Thyroid receptor binding	+	+	-	+	+	+	+
Glucocorticoid receptor binding	-	+	-	+	+	+	-
Aromatase binding	+	+	-	+	+	+	+
PPAR gamma	+	+	-	+	+	+	+

OATP2: organic anion-transporting polypeptide; OCT2: organic cation transporter 2; BSEP: bile salt export pump; CYP: cytochrome P450; Human ether-a-go-go-related Gene (hERG); (-): absence of activity; (+): presence of activity

Table 5 Pharmacokinetics, toxicities and receptor binding properties of mangiferin, thebaine, berberine, andrographolide, quercetin and luteolin

Parameters	Mangiferin	Thebaine	Berberine	Andrographolide	Quercetin	Luteolin
Absorption						
Human intestinal absorption	+	+	+	+	+	+
Caco-2	-	+	+	+	-	+
Human oral bioavailability	-	-	-	-	-	-
Water solubility	- 2.398	- 2.979	- 2.974	- 2.853	- 2.999	- 2.999
Distribution						
Subcellular localization	Mitochondria	Mitochondria	Mitochondria	Mitochondria	Mitochondria	Mitochondria
Blood-brain barrier penetration	-	+	+	+	-	-
Plasma protein binding	0.97	0.718	0.834	0.536	1.175	1.043
P-glycoprotein inhibitor	-	-	+	-	-	-
P-glycoprotein substrate	-	+	-	-	-	-
Metabolism						
OATP2B1 inhibitor	+	-	-	-	+	+
OATP1B1 inhibitor	+	+	+	+	+	+
OATP1B3 inhibitor	+	+	+	+	+	+
MATE1 inhibitor	-	-	-	-	+	+
OCT2 inhibitor	-	-	-	-	-	-
BSEP inhibitor	-	-	+	+	-	-
CYP3A4 substrate	+	+	+	+	+	-
CYP2C9 substrate	-	-	-	-	-	-
CYP2D6 substrate	-	+	-	-	-	-
CYP3A4 inhibition	-	-	-	-	+	+
CYP2C9 inhibition	-	-	-	-	-	-
CYP2C19 inhibition	-	-	-	-	-	-
CYP2D6 inhibition	-	+	+	-	-	-
CYP1A2 inhibition	-	-	+	-	+	+
CYP inhibitory promiscuity	-	-	+	-	+	+
Toxicity						
Carcinogenicity	-	-	-	-	-	-
Human ether-a-go-go inhibition	-	-	-	-	-	-
Hepatotoxicity	+	-	+	-	+	+
Acute oral toxicity (kg/mol)	2.979	3.597	1.545	2.795	2.559	2.525
Receptor binding						
Estrogen receptor binding	+	+	+	+	+	+
Androgen receptor binding	+	-	+	+	+	+
Thyroid receptor binding	-	+	+	+	+	+
Glucocorticoid receptor binding	+	+	+	+	+	+
Aromatase binding	-	-	-	+	+	+
PPAR gamma	+	+	+	-	+	+

OATP2: organic anion-transporting polypeptide; OCT2: organic cation transporter 2; BSEP: bile salt export pump; CYP: cytochrome P450; Human ether-a-go-go-related Gene (hERG); (-): absence of activity; (+): presence of activity

compared to hydroxychloroquine. It has been reported that patients receiving hydroxychloroquine may experience several severe adverse effects such as visual disturbances, ventricular tachycardia, cardiovascular collapse, convulsions, hypokalemia, rhythm and conduction disorders

including QT prolongation, and ventricular fibrillation. Therefore, development of potential attachment inhibitor from natural products is a safe and effective strategy for the treatment of COVID-19. Our docking results demonstrated that curcumin and nimbin are the best-ranked drugs among

Table 6 Pharmacokinetics, toxicities and receptor binding properties of resveratrol, naringenin, zingiberene, β -caryophyllene, citronellol, eugenol, and gallic acid

Parameters	Resveratrol	Naringenin	Zingiberene	β -caryophyllene	Citronellol	Eugenol	Gallic acid
Absorption							
Human intestinal absorption	+	+	+	+	+	+	+
Caco-2	+	+	+	+	+	+	-
Human oral bioavailability	-	-	-	+	-	-	+
Water solubility	- 2.778	- 3.19	- 4.937	- 4.687	- 2.78	- 1.918	- 1.097
Distribution							
Subcellular localization	Mitochondria	Mitochondria	Lysosomes	Lysosomes	Lysosomes	Mitochondria	Mitochondria
Blood Brain Barrier penetration	-	-	+	+	+	+	-
Plasma protein binding	0.677	1.001	1.038	0.832	0.769	0.94	0.668
P-glycoprotein inhibitor	-	-	-	-	-	-	-
P-glycoprotein substrate	-	-	-	-	-	-	-
Metabolism							
OATP2B1 inhibitor	-	-	-	-	-	-	-
OATP1B1 inhibitor	+	+	+	+	+	+	+
OATP1B3 inhibitor	+	+	+	+	+	+	+
MATE1 inhibitor	-	-	-	-	-	-	-
OCT2 inhibitor	-	-	-	-	-	-	-
BSEP inhibitor	-	-	-	-	-	-	-
CYP3A4 substrate	-	-	-	+	-	-	-
CYP2C9 substrate	-	-	-	-	-	-	-
CYP2D6 substrate	-	-	-	-	-	+	-
CYP3A4 inhibition	+	+	-	-	-	-	-
CYP2C9 inhibition	+	+	-	-	-	-	-
CYP2C19 inhibition	+	+	-	-	-	-	-
CYP2D6 inhibition	-	-	-	-	-	-	-
CYP1A2 inhibition	+	+	-	-	-	-	-
CYP inhibitory promiscuity	+	+	-	-	-	-	-
Toxicity							
Carcinogenicity	-	-	-	-	-	-	-
Human ether-à-go-go inhibition	-	-	+	+	-	-	-
Hepatotoxicity	+	+	-	-	-	-	-
Acute Oral Toxicity (kg/mol)	2.644	1.87	1.827	2.366	2.077	1.645	1.552
Receptor binding							
Estrogen receptor binding	+	+	-	-	-	-	-
Androgen receptor binding	+	+	-	-	-	-	+
Thyroid receptor binding	+	+	-	-	-	-	-
Glucocorticoid receptor binding	+	+	-	-	-	-	-
Aromatase binding	+	+	-	-	-	-	-
PPAR gamma	+	+	-	-	-	-	-

OATP2: organic anion-transporting polypeptide; OCT2: organic cation transporter 2; BSEP: bile salt export pump; CYP: cytochrome P450; Human ether-a-go-go-related Gene (hERG); (-): absence of activity; (+): presence of activity

the selected natural products showing an inhibitory effect for spike glycoprotein and ACE2. Both the drugs have been traditionally used in AYUSH system of medicine for the management of various disease conditions such as inflammation, fever, and treatment of infections,

autoimmune disorders like arthritis and neurological disorders [21, 22]. Most of these natural products/drugs have been recognized as safe for human usage. Besides, these natural products/drugs are also well-known regulator of various interleukins, chemokines, and cytokines. Recent

studies have shown that the level of IL-6, TNF- α , and IFNs were found to be elevated in patients with SARS-CoV-2 [23]. Therefore, these natural drugs can be used as alternative therapy for the management of various sign and symptoms associated SARS-CoV-2 infection as well as other coronaviruses in the absence of specific antiviral drug.

We further analyzed the drug-likeness and ADME toxicity and receptor binding properties of selected natural products by using the swissADME and admetSAR server. According to In Lipinski rule for drug-likeness, any drug molecule having MW > 500 g/mol, hydrogen-bond-donating atoms > 5, Hydrogen-bond-accepting atoms > 10, or ClogP > 5 is not considered as a good pharmaceutical agent in terms of oral activity. Even though they may have strong docking energies, compounds violating more than 2 Lipinski's rule of 5 were not considered for drug-likeness. Molecules violating more than one of these rules may have problems with oral bioavailability [24]. Similarly, the knowledge of pharmacokinetics and toxicity profile of drugs is essential in order to measure the safety and efficacy of in biological system. Therefore, designing an attachment inhibitor from natural products may provide a safe, effective and broad-spectrum approach while combating viral infections. The findings of the present study would facilitate us to understand the antiviral mechanism of natural products which enables us to establish a potential therapy for patients diagnosed with COVID-19 in a dose-dependent manner with a reduced level of associated toxicity.

Collectively, for the first time our data demonstrated the efficacy of selected natural products against spike glycoprotein of SARS-CoV-2 and its cellular receptor ACE2. We have found that curcumin, nimbin, withaferin A, piperine, mangiferin, thebaine, berberine, and andrographolide may effectively inhibit the interaction of spike glycoprotein with its receptor compared to other selected natural drugs. Also, the pharmacokinetics data illustrated that all selected natural products have better pharmacological properties (low molecular weight; no violation of Lipinski rule of five, good absorption profiles, oral bioavailability, good blood–brain barrier penetration, and low toxicity risk). Therefore, based on our results it is suggested that these natural products/drugs from Ayurveda may be useful as a therapeutic or prophylactic agent during COVID-19 for restricting viral attachment to the host cells via inhibition of spike glycoprotein or its cellular receptor. Consequently, resveratrol, quercetin, luteolin, naringenin, zingiberene, and gallic acid could be a useful starting material for the development of potent antiviral agents for the prevention and treatment of SARS-CoV-2 and other coronaviruses if tested further.

Acknowledgements The authors are grateful to the Vice Chancellor, King George's Medical University (KGMU), Lucknow, India for the encouragement for this work. The authors have no other relevant affiliations or financial involvement with any organization or entity with a financial interest in or financial conflict with the subject matter or materials discussed in the manuscript apart from those disclosed.

Funding None.

Compliance with ethical standards

Conflict of interest The authors declare no conflict of interest.

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