ORIGINAL ARTICLE

Implementing in vitro and in silico approaches to evaluate anti‑infuenza virus activity of diferent Bangladeshi plant extracts

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Abstract

Emergence of antiviral drug resistance in infuenza virus remains a major public health concern worldwide. Nowadays, different herbs receive renewed attention because of their enormous antiviral potential. In this study, we investigated the antiviral activity of *Camellia sinensis, Persicaria hydropiper, Persicaria orientale*, *Persicaria lapathifolia, Persicaria stagnina, Mucuna pruriens* and *Chenopodium album* against diferent infuenza strains using both in vitro and in silico approaches. Antiviral efect of plant extracts was evaluated by cytopathic efect (CPE) inhibition assay on infuenza infected MDCK (Madin Darby Canine Kidney) cell line. Later, the herb demonstrating antiviral activity was virtually screened for their available bioactive compounds and multiple in silico tools were performed to prioritize and establish these compounds as potential inhibitor. The methanol, but not the n-hexane and ethyl acetate extracts of *C. sinensis, P. hydropiper, M. pruriens* and *C. album* exhibited anti-influenza effect with EC_{50} values within 32–46 μ g/ml. Importantly, the extracts remained effective against both amantadine-resistant and -sensitive infuenza isolates. The molecular docking analysis showed that favonoids, steroid and derivatives had strong binding affinity to the target proteins which may remain responsible for the anti-influenza characteristics of plant extracts. Pharmacokinetic properties, bioavailability and drug-likeness score revealed that ferulic acid, sinapic acid, campesterol, cryptomeridiol, eupatin and genistein could be attractive leads as potential infuenza inhibitors. Taken together, the botanical ingredients of these herbs could be used as valuable candidates for developing novel therapeutics to control infuenza related illnesses.

Keywords Infuenza · Plant extract · Antiviral · Docking · Natural products · Bioactive molecules

Introduction

Infuenza (or 'fu') caused by infuenza virus is a highly contagious respiratory illness that may cause mild to severe infections and may end to death occasionally, posing severe impact on public health and economy (Peteranderl et al.

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[2016;](#page-13-0) Moghadami [2017\)](#page-13-1). Based on the antigenic specifcity of the envelope proteins, infuenza viruses are classifed into infuenza A, B and C, of which, the predominant type A infects both humans and other animals such as birds, horses and dogs. (Pulendran and Maddur [2014](#page-13-2)). The virus spreads rapidly from person to person by airborne droplets that attack the upper respiratory tract cells, causing typical fu symptoms like fatigue, fever, cough and body aches which may make immunocompromised and elderly people more vulnerable to potentially life-threatening secondary infections (McCullers et al. [1999;](#page-13-3) Dawood et al. [2012](#page-12-0)). Globally, more than 1 billion people become infected with infuenza of which, 3 to 5 million sufer from severe illness and 300,000 to 500,000 people died each year (Naeem et al. [2020\)](#page-13-4). In 2010, the estimated incidence rate of infuenza associated respiratory illnesses was 6.5 and 1.3 per 1000 persons aged \textdegree 5 and \geq 5 years, respectively, resulted in an estimated economic loss of US\$ 169 million in Bangladesh (Azziz-Baumgartner et al. [2012](#page-12-1)). Moreover, the estimated infuenza associated mortality rate was 6 and 41 per 100 000 children $<$ 5 years and persons > 60 years in 2010–2011, respectively which became 13 and 88 in 2011–2012 (Ahmed et al. [2018](#page-12-2)). Previously, we reported the prevalence of infuenza 25.8% among children living in the slums of Dhaka city that was signifcantly associated with age, poor economic status, malnutrition and poor hygiene practices (Rahman et al. [2016](#page-13-5)).

Multiple approaches such as vaccines as well as antiviral drugs have been developed to minimize the burden of the infuenza illnesses (Yamayoshi and Kawaoka [2019\)](#page-13-6). Currently, three types of infuenza vaccines: inactivated, live attenuated and recombinant are used in various countries that have several advantages as well as drawbacks (Rajão and Pérez [2018](#page-13-7)). Although World Health Organization (WHO) recommends to change the vaccine strain composition every year, antigenic mismatches between the vaccine viruses and the circulating strains often reduce the efectiveness of vaccination (Carrat and Flahault [2007](#page-12-3); Chan et al. [2018](#page-12-4)). In addition, two classes of infuenza antivirals: M2 channel inhibitor like amantadine and rimantadine, and neuraminidase inhibitor like zanamivir and oseltamivir are available at present to treat infuenza infection (Betakova [2007\)](#page-12-5). Among these, zanamivir and oseltamivir perform poorly against infuenza in adults, while oseltamivir exhibit less efectiveness in reducing mortality among 2009A/H1N1 infected patients (Heneghan et al. [2016](#page-12-6)). Furthermore, emergence of antiviral- resistant infuenza A subtypes H1N1, H1N2, H3N2, H4N2, H5N1, H5N2, H6N1, H6N6, H6N8, H9N2 and H11N3 from diferent regions of the world have been reported in many studies (Bright et al. [2005;](#page-12-7) Deyde et al. [2007;](#page-12-8) Cheng et al. [2009](#page-12-9)). A rapid increase of M2 inhibitors-resistant infuenza A/H3N2 strain was reported between 2002 and 2007 (Bright et al. [2005](#page-12-7); Deyde et al. [2007\)](#page-12-8). Again, an elevated level of amantadine-resistant infuenza was observed in South East Asia since 2007 (Barr et al. [2008\)](#page-12-10). Recently, emergence of new infuenza A variant have been reported that bore mutations associated with reduced susceptibility to Baloxavir marboxil, a new replication inhibiting drug against infuenza (Imai et al. [2020](#page-12-11)). Thus, regular screening for new drugs against these drugresistant infuenza's is essential.

Plant materials are the vital sources of lead compounds in modern drug development (Rajasekaran et al. [2013\)](#page-13-8). For example, the extract of *C. sinensis* contains higher number of polyphenols that exhibit signifcant health protecting activity (Manzocco et al. [1998](#page-13-9)). Catechin derivatives are the major polyphenols in green tea that show antiviral efects against hepatitis C virus, human papilloma virus, rotavirus and enterovirus, infuenza virus (Mukoyama et al. [1991](#page-13-10); Song and Lee [2005;](#page-13-11) Gross et al. [2007;](#page-12-12) Chen et al. [2012](#page-12-13)). *P. hydropiper* is useful to mitigate infammatory responses

in treating viral infections (Ren et al. [2020](#page-13-12)). All parts of *M. pruriens,* widely known as "velvet bean", possess antidiabetic, antineoplastic and antimicrobial properties (Sathiyanarayanan & Arulmozhi, 2007). *C. album* contains two antiviral proteins CAP-I and CAP-II that inhibited two plant viruses, Tobacco mosaic virus and Sunnhemp rosette virus (Dutt et al. [2003](#page-12-14)). However, the anti-influenza effect of these herbs yet remains vague.

In this study, we determined the antiviral activity of the diferent extracts of seven locally available herbs *C. sinensis, P. hydropiper, P. orientale*, *P. lapathifolia, P. stagnina, M. pruriens* and *C. album* against diferent infuenza strains circulating in Bangladesh.

Materials and methods

The entire workflow is shown in Fig. [1.](#page-2-0)

Preparation of plant extracts

The plants were collected from their natural habitats from diferent parts of Bangladesh by an experienced plant taxonomist. The extracts of *P. orientale, P. lapathifolia, P. stagnina, and P. hydropiper* were prepared from whole plant, whereas leaves were used for *C. sinensis*, *M. pruriens* and *C. album.* The fresh plant materials were cut into small pieces, washed with tap water and allowed for sun drying in plastic boxes. The dried plant parts were crushed to prepare fne powder with mortar and pestle and kept in tightly closed containers at room temperature and away from light. For extraction, the dry powder (100 g) was soaked in 500 ml of solvent (hexane, ethyl acetate or methanol) for 2 days at room temperature. The extracts were then fltered through cotton and Whatman flter paper, transferred into a vacuum fask and dried at 45 °C under reduced pressure on a rotary evaporator. The resulting powder was stored in a tightly sealed falcon tube at room temperature. It was then weighed and dissolved in 100% DMSO to prepare stock solution and stored at 4 °C until use. To avoid the toxic efect of DMSO on cell line, the fnal concentration of DMSO was maintained at about 1% v/v during in vitro test.

Virus propagation

Eleven infuenza strains (9 Infuenza A and 2 Infuenza B virus), isolated earlier from nasal and throat swab of different slums dwellers in Dhaka city (Rahman et al. [2016](#page-13-5)), were propagated on MDCK cell line as described previously (Rahman et al. [2017\)](#page-13-13). Virus titration was performed by infecting MDCK cells. Viral suspension was diluted in serum free DMEM media and concentration that yielded 100% CPE by 48 h post-inoculation observed under

Fig. 1 Workfow of the entire study

microscope and confrmed by MTT assay was selected as optimum for antiviral assay.

Screening of anti‑infuenza efect

The anti-influenza effect of n-hexane, ethyl acetate and methanol extracts of herbs was determined by CPE inhibition assay against 6 influenza A/H1N1, 3 influenza A/H3N2 and 2 infuenza B/Yamagata isolates. In brief, 125 µl of cell suspension $(15 \times 10^4 \text{ MDCK}$ cells/ml in 10% FBS containing DMEM media) was seeded in each well of 96 well micro plate. When cell growth became confuent, media was removed, washed twice with PBS and 25 μl of virus inoculum was added into each well. After 1 h of incubation at 37 °C, the plate was washed twice with PBS to remove unbound viruses. Finally, 0, 10.0, 50.0, 90.0 and 120.0 μg/ml of extracts solution with DMEM containing 2% TPCK treated trypsin were added in each infected well. 25 μl of serum free DMEM media was used as control instead of virus inoculum in mock infected wells. Anti-influenza effect was determined based on survival of MDCK cells after 48 h of incubation.

MTT assay

MTT (3-(4,5- dimethyl thiazol-2yl)-2,5-diphenyl tetrazolium bromide) assay was performed to determine the EC_{50} value of extracts as described previously (Rahman et al. [2017](#page-13-13)). After observing CPE under light microscope, 20 μl of 5 mg/ ml MTT solution (Sigma-Aldrich) was added for cell staining. After 2 h incubation at 37 °C, SDS (Promega, USA) solution was used to dissolve formazan crystals. Finally, absorbance was measured at 570 nm to determine the percentage of survivability of MDCK cells and plotted against extract concentration to determine EC_{50} values of extracts.

Ligand preparation

The phytochemical constituents of four extracts that were found to be efective in inhibition of infuenza through in vitro experiments were retrieved from Indian Medicinal Plants, Phytochemistry and Therapeutics (IMPPAT) database (Mohanraj et al. [2018](#page-13-14)). The 3D structure of compounds was downloaded and converted into pdb (Protein Data Bank) format using OpenBabel tool integrated in PyRx (Dallakyan and Olson [2015\)](#page-12-15). The Universal force feld (UFF) was applied for energy

minimization of ligands followed by conversion into pdbqt format for docking.

Protein preparation

The molecular structure of three target proteins was obtained from RCSB protein data bank. The PDB ID of the neuraminidase, transmembrane M2 protein and haemagglutinin protein of influenza virus were 2HU4, 6BKK and 6CF7, respectively. All the water molecules and heteroatoms including ions and bound molecules were removed from protein structure and polar hydrogen was added using Discovery Studio software v2020. The energy minimization of the cleaned structure was performed by GROMOS 43B1 forcefield in SWISS PDB viewer (Kaplan and Littlejohn [2001](#page-12-16)) and later saved in pdb format for further analysis. The active site of the target protein was predicted by Discovery Studio software v2020 and PDBsum (Laskowski [2001\)](#page-12-17).

Molecular docking

The binding energy of the protein–ligand complex was calculated using PyRx Autodock Vina (Dallakyan and Olson [2015\)](#page-12-15). The loaded protein structure was converted into pdbqt format and the grid box parameter was set manually to cover the active site of receptor with an exhaustiveness of 8. The maximum binding affinity was identified by observing the highest negative binding energy. Moreover, the docking procedure was validated by docking the active site of optimized protein with extracted ligand from retrieved protein structure following the same protocol (Shivanika et al. [2020](#page-13-15)). The protein ligand complex was visualized and their binding interactions were observed using PyMOL, Discovery Studio software v2020 and LigPlot +(Laskowski and Swindells [2011](#page-12-18)).

Exploration of drug‑likeness properties

Compounds that had favorable docking score $(< - 7.0 \text{ kcal/mol})$ were selected to study drug-likeness and ADMET (Absorption, Distribution, Metabolism, Excretion and Toxicity) properties. The Lipinski rule is regarded as one of the most important factors for predicting a drug's oral drug-likeliness (Lipinski [2004](#page-13-16); Benet et al. [2016](#page-12-19)). Pharmacokinetic properties were predicted by admetSAR (<http://lmmd.ecust.edu.cn/admetsar2>) and SwissADME (Daina et al. [2017](#page-12-20)). Drug-Likeness Tool (DruLiTo) was used for Quantitative estimation of Druglikeness (QED) (Bickerton et al. [2012\)](#page-12-21).

Results

Screening of in vitro anti‑infuenza activity of herbs

The methanol, n-hexane and ethylacetate extracts of seven available herbs: *C. sinensis, P. hydropiper, P. orientale*, *P. lapathifolia, P. stagnina, M. pruriens* and *C. album* were examined against a total of 11 influenza isolates. All the extracts were tested at concentrations ranging from 10.0, 50.0, 90.0, 120.0 μg/ml.

Among these, only methanol extracts of *C. sinensis, P. hydropiper, M. pruriens* and *C. album* exhibited antiviral efect with average EC_{50} of 41.36, 46.0, 41.68 and 32.22 µg/ml, respectively. All the extracts were also tested on uninfected cells (control). But no CPE was evident even in highest concentration. So, the extracts had no cytotoxicity in MDCK cell up to 120 µg/ml.

The average EC_{50} values of the methanol extract of *C. sinensis* against 6 infuenza A/H1N1, 3 infuenza A/H3N2 and 2 infuenza B/Yamagata isolates were 40.91, 39 and 46.25 µg/ ml, respectively. The average survivability of cells at 10.0, 50.0, 90.0 and 120.0 μg/ml of extracts were 29, 56, 77 and 90%, respectively for infuenza A virus (Table [1\)](#page-4-0).

The percentage of survivability at diferent concentrations of the methanol extract of *P. hydropiper* varied between isolates. The average EC_{50} values against influenza A and infuenza B/Yamagata isolates were 45.1 and 50.25 µg/ml, respectively.

When six infuenza A/H1N1, three infuenza A/H3N2 and two infuenza B/Yamagata viruses were challenged with methanolic extract of M . pruriens, the EC_{50} values ranged between 34 and 46.5 µg/ml (Table [1](#page-4-0)). The cell survivability pattern was similar to *C. sinensis* for all isolates.

Nearly 92% cell survivability was observed at 120 µg/ml concentration of the methanolic extract of *C. album* for infuenza A/H1N1, influenza A/H3N2 and Influenza B. The EC_{50} values for all isolates were between 26.5 and 37 µg/ml.

The EC₅₀ values of the methanolic extract of *C. sinensis*, *P. hydropiper, M. pruriens* and *C. album* against amantadineresistant infuenza strain A/ 105/ H1N1 were 42, 45, 43 and 33 µg/ml, respectively. The amount of these extracts needed to achieve the half of the maximum efect against Infuenza B subtypes were quite higher than the same against Infuenza A.

Though methanol, n-hexane and ethyl acetate extracts of *P. orientale*, *P. lapathifolia, P. stagnina* could not exhibit antiinfuenza activity even at highest concentration, they were found to be non-toxic to MDCK cells.

Findings of the potential bioactive candidates

As the methanol extract of *C. sinensis, P. hydropiper, M. pruriens and C. album* were effective against the influenza virus, the phytochemicals of these antiviral herbs were

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Protein	Name of the target with attached molecule in PDB	Center grid values $(x \times y \times z)$ A	Dimension $(x \times y \times z)$ Å
2HU4	N1 neuraminidase in complex with osel tamivir 2	$-1.53 \times 80.90 \times 109.57$	$30.42 \times 25.15 \times 29.78$
6BKK	Influenza A M2 transmembrane domain bound to amantadine	$66.11 \times 10.03 \times 46.02$	$21.03 \times 21.88 \times 24.80$
6CF7	Crystal structure of the A/Solomon Islands/3/2006(H1N1) influenza virus hemagglutinin in complex with small molecule JNJ4796	$18.09 \times 45.08 \times 29.66$	$36.24 \times 36.40 \times 31.27$

Table 2 Name of target protein and Grid box parameters in PyRx

virtually screened and retrieved their chemical structure with relevant information. Thereafter, docking protocols (Table [2\)](#page-5-0) were utilized to observe the binding affinity as well as active interacting residues between the targeted viral receptors and phytochemicals of herbs. The 3D and 2D interactions between protein–ligand complex are shown in Figs. [2](#page-6-0) and [3](#page-8-0), respectively.

Among 98 phytochemicals of *C. sinensis*, half of the them showed favorable binding affinity to three different target receptors ranged from -5.3 to -10.4 kcal/mol. Epicatechin gallate, delphinidin and theafavin were identifed as the strongest binder to 2HU4 (− 9.4 kcal/mol), 6BKK (-10.4 kcal/mol) and 6CF7 (-8.9 kcal/mol) , respectively. The binding energy of quercetin, fsetin and afzelechin to transmembrane domain of influenza virus, 6BKK were − 10.1, − 9.9 and − 9.1, respectively. ARG118, ASP151, ARG292, ASN294, ARG371, TYR347 of 2HU4; HIS37(A), HIS37(B), HIS37(C), VAL27(D) of 6BKK; and THR37(A), HIS38(A), THR49(B) of 6CF7 were found to be involved in forming hydrogen bond with epicatechin gallate, delphinidin and theafavin, respectively (*Fig. [3](#page-8-0)*).

In case of 38 phytochemicals present in *P. hydropiper*, the highest binding score of miquelianin-2HU4, quercetin-6BKK, daucosterol-6CF7 complex were -9.2 , -10.1 , − 7.2, respectively. The amino acids that involved in hydrogen bond with ligands were ARG118, GLU119, ASN221, GLU227, GLY244, GLU277, ARG292, TYR347, TYR406 of 2HU4; HIS37(A), HIS37(B), HIS37(C), GLY34(B) of 6BKK and HIS38(A), GLY20(B) of 6CF7. Quercetin, kaempferol, confertiforin were able to bind to all analyzed targets. Molecules that bound at least two targets were isoquercetin $(-8.7 \text{ kcal/mol}$ to $2\text{HU}4, -6.8 \text{ kcal/mol}$ to 6CF7), astragalin (-7.6 kcal/mol) to $2HU4$, -6.8 kcal/mol to 6CF7), ellagic acid (-8.4 kcal/mol to 2HU4, -9.6 kcal/ mol to 6BKK), 6-hydroxyluteolin (− 7.9 kcal/mol to 2HU4, − 6.8 kcal/mol to 6CF7) (Fig. [3](#page-8-0)). The average binding energy of all molecules to 2HU4, 6BKK and 6CF7 were − 6.5, − 6.14 and − 5.6 kcal/mol, respectively.

Cardenolide-B3 of *M. pruriens* showed highest binding activity to the active site of $2HU4$ ($- 8.6$ kcal/mol) and $6CF7$ ($- 7.6$ kcal/mol) that formed hydrogen bond with ARG118, GLU119, ASP151, ASN221, ARG224, GLU277, ASN294, TYR347, ARG371 of 2HU4 and HIS18(A), ALA19(A), ASN20(A), HIS38(A), THR15(B) of 6CF7 (Fig. [3](#page-8-0)). 2'-hydroxygenistein had highest binding affinity to $6BKK$ ($- 8.8$ kcal/mol) followed by genistein $(- 8.7 \text{ kcal/mol})$ and glutathione $(- 7.7 \text{ kcal/mol})$. Compounds that were observed to interact with all target proteins were 2'-hydroxygenistein, ambroxol hydrochloride and genistein. Eupatin, coumarin, tryptamine, levodopa bound to at least two targets. The average binding affinity of all molecules to 2HU4, 6BKK and 6CF7 were − 6.08, − 6.16 and − 5.5 kcal/mol, respectively.

CID 508,204 of *C. album* strongly bound to 2HU4 (-8.9 kcal/mol) and 6CF7 (-7.7 kcal/mol) whereas cryptomeridiol demonstrated the same to 6BKK. TYR347, TYR406 of 2HU4; HIS37(A), HIS37(D) of 6BKK and ASN50(B), SER113(B) of 6CF7 were involved in hydrogen bond with cid 508,204, cryptomeridiol and cid 508,204, respectively (Fig. [3](#page-8-0)). Metamitron, astragalin and cryptomeridiol were able to bind to all targets. The mean binding affinity of all molecules to 2HU4, 6BKK and 6CF7 were $-6.3, -5.9$ and -5.6 kcal/mol, respectively.

The redocked structure of the extracted ligands oseltamivir, amantadine and jnj4796 bound to the target proteins 2HU4, 6BKK and 6CF7 with rmsd of 0.00, 0.042 and 0.035 Å, respectively. The lower rmsd value signifed the accuracy and validation of docking protocol.

Analysis of pharmacokinetic properties

Most of the compounds belong to favonoids class. Other classes include steroid and derivatives, purine nucleotides, carboxylic acid and derivatives, cinnamic acid and derivatives etc. Diverse patterns of absorption (human intestinal absorption, blood brain barrier, Caco-2 permeability), metabolism and toxicity (AMES toxicity, carcinogenicity) were observed. No molecules were found to be carcinogenic. All molecules except isoquercetin, astragalin, glutathione, ambroxol hydrochloride, levodopa were nontoxic in AMES test (Table [3](#page-9-0)). The bioavailability score of maximum molecules was 0.55. The drug-likeness of sinapic acid, lariciresinol, campesterol, cryptomeridiol, ferulic acid, genistein, eupatin and kaempherol were 0.797, 0.772, 0.769, 0.742, 0.748, 0.719, 0.71 and 0.618, respectively (*Fig. [4](#page-11-0)*).

Fig. 2 3D interaction between ligands (phytochemicals) and target protein. 3D images of target protein–ligand (phytochemicals) interactions. A1) 2HU4- CID 508,204 (*C. album*), A2) 6BKK- Cryptomeridiol (*C. album*), A3) 6CF7- CID 508,204(*C. album*), B1) 2HU4- Epicatechin gallate (*C. sinensis*), B2) 6BKK- Delphinidin (*C. sinensis*),

B3) 6CF7- Theafavin (*C. sinensis*), C1) 2HU4- Cardenolide B-3 (*M. pruriens*), C2) 6BKK- Hydroxygenistein (*M. pruriens*), C3) 6CF7- Cardenolide B-3 (*M. pruriens*), D1) 2HU4- Miquelianin (*P. hydropiper*), D2) 6BKK- Quercitin (*P. hydropiper*), D3) 6CF7- Daucosterol (*P. hydropiper*)

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Fig. 3 2D interaction between ligands (phytochemicals) and target ◂protein. 2D images of target protein–ligand (phytochemicals) interactions. A1) 2HU4- CID 508,204 (*C. album*), A2) 6BKK- Cryptomeridiol (*C. album*), A3) 6CF7- CID 508,204(*C. album*), B1) 2HU4- Epicatechin gallate (*C. sinensis*), B2) 6BKK- Delphinidin (*C. sinensis*), B3) 6CF7- Theafavin (*C. sinensis*), C1) 2HU4- Cardenolide B-3 (*M. pruriens*), C2) 6BKK- Hydroxygenistein (*M. pruriens*), C3) 6CF7- Cardenolide B-3 (*M. pruriens*), D1) 2HU4- Miquelianin (*P. hydropiper*), D2) 6BKK- Quercitin (*P. hydropiper*), D3) 6CF7- Daucosterol (*P. hydropiper*)

Discussion

Medicinal plants have long been used for the treatment of infuenza infections and now become increasingly popular as alternatives to synthetic drugs (Amić et al. [2003](#page-12-22); Rajasekaran et al. [2013](#page-13-8)). Several studies reported the antiviral efect of diferent Asian medicinal plants against infuenza virus through in vitro experiments (Enkhtaivan et al. [2015;](#page-12-23) Shoji et al. [2017](#page-13-17); Liu et al. [2018](#page-13-18)). Here, we determined the anti-infuenza activity of a number of traditional plant extracts that could serve as sources of new antivirals to combat drug-resistant viral infections. A signifcant criterion for an antiviral treatment is safety and it is critical to evaluate potential adverse efects when searching for novel medications. Herein, no cytotoxicity was observed (up to the highest concentration used) in the tested extracts. These nontoxic characteristics indicated the potential of extracts in the development of safer and less harmful medications. Generally, natural compounds with ethnomedical background are regarded as safer and more efective compared to substances missing this framework (Grienke et al. [2012;](#page-12-24) Rajasekaran et al. [2013](#page-13-8)).

The methanol extract of *C. sinensis* showed a potent antiviral efect that is in agreement with previous studies, describing the antiviral characteristics of tea against infuenza A and B viruses (Nakayama et al. [1990](#page-13-19); Zu et al. [2012](#page-13-20)). Aqueous extract of green tea contains diferent polyphenolic compounds such as catechins, theafavins that are known to exert antioxidant, antiviral, antifungal and antibacterial effects (Friedman [2007](#page-12-25)). A study on antiviral potential of catechin in green tea reported the EC_{50} (the 50% effective inhibition concentration) of EGCG (epigallocatechin gallate), ECG (epicatechin gallate), and EGC (epigallocatechin) against infuenza A virus were 22–28, 22–40 and 309–318 μ M, respectively (Song and Lee [2005](#page-13-11)). The methanol extract of *P. hydropiper* was known to have antiinfammatory properties (Yang et al. [2012\)](#page-13-21). Here, the average EC_{50} value of the same extract was 46 µg/ml that was in line with another study that found the EC₅₀ of *P. chinese* against infuenza virus ranged between 38.4 to 55.5 µg/ml (Tran et al. [2017](#page-13-22)). Such anti-infuenza efect might be attributed to their pharmacological and biological efects and the known phytochemicals of potential therapeutic importance that have been isolated so far (Fan et al. [2011](#page-12-26)). Phytochemical analysis of *M. pruriens*, an important medicinal plant, revealed the presence of a wide variety of chemicals such as alkaloids, β-sitosterol, glutathione, steroids, favonoids, coumarins, cardenolides, tryptamine, alkylamines, oleic acid, linoleic acid, and palmitic acid having antiviral, antiparkinson, antioxidant, antimicrobial and antiprotozoal efects (Gupta et al. [1997](#page-12-27); Adebowale et al. [2005;](#page-12-28) Rajeshwar et al. [2005;](#page-13-23) Sivaraman et al. [2010\)](#page-13-24). We observed the moderate antiviral characteristics of *M. pruriens*, in which the average EC_{50} was 41.6 µg/ml. *C. album* is used in diet as a source of minerals, fber, vitamins and essential fatty acids that has been traditionally used as a blood purifer, sedative, antiscorbutic laxative and anthelmintic (Poonia and Upadhayay [2015](#page-13-25)). In this study, the methanol extract of *C. album* was found to be most effective (EC_{50} =32 µg/ml) in preventing infuenza virus compared to other tested herbs. Such lower EC_{50} indicated the presence of higher amount of anti-influenza molecules that might be active at low concentration. Medicinal herbs have many secondary metabolites. Sequential method of extraction using n-hexane, ethyl acetate and methanol is mostly recommended if active metabolites of herbs are uncharacterized and thus it is possible to extract non polar, mid polar and polar compounds orderly (Poonia and Upadhayay [2015](#page-13-25)). As only methanolic extracts of diferent herbs showed anti-infuenza activity, it can be interpreted that the active compounds of these herbs are polar in nature. Similarly, the polar extract of *Jatropha multifda* strongly inhibited infuenza virus and thus promote the survivability of infected MDCK cells (Shoji et al. [2017\)](#page-13-17). However, no anti-infuenza activity was observed in n-hexane and ethyl acetate extracts of plants described herein which might be due to the chemical nature of solvents that did not facilitate the extraction of higher amount of polar compounds (da Costa Cordeiro et al. [2018](#page-12-29)).

Plant extracts contain many active components that can efectively prevent virus inhibition, playing a major role in exerting antiviral activity (Rajasekaran et al. [2013\)](#page-13-8). The used extracts contain a lot of polyphenols such as catechins, isoquercetin, delphinidin, ellagic acid etc. and their antiviral activity had been demonstrated both experimentally and through molecular docking studies in numerous literatures (Kim et al. [2010;](#page-12-30) Liu et al. [2015;](#page-13-26) Sadati et al. [2019](#page-13-27)). In this study, catechin derivatives in tea exhibited noticeable interactions with target proteins. The binding affinity between *C. sinensis* phytochemicals and target protein revealed the catechin derivatives as potential inhibitors of infuenza virus. Apart from the catechin derivatives, favonoid compounds such as quercetin, delphinidin, cyanidin, fisetin and luteolin also showed strong binding affinity to neuraminidase, haemagglutinin and M2 protein. The derivatives namely miquelianin, isoquercetin, quercetin, confertiforin, kaempferol, daucosterol from *P. hydropiper* exhibited diferential binding

Table 3 Drug likeness and Pharmacokinetic properties of representative ligands

Fig. 4 Graphical representation of the drug-likeness and bioavailability of selected molecules

afnity to receptor molecules. The binding of cardenolide from *M. pruriens* exerted strong affinity to neuraminidase and haemagglutinin of infuenza which is similar to a study in which cardenolide was proven as an antiviral inhibitor of Influenza A virus (Boff et al. [2020\)](#page-12-31). According to the in silico studies, the molecules abundant in *C. album* that were found to be responsible for exerting anti-infuenza activities were cid508204, isoquercetin, metamitron, cryptomeridiol, astragalin, sinapic acid etc. The maximum nine hydrogen bond was observed in 2HU4-miquelianin interactions followed by eight in cardenolide-B3-2HU4 complex and six in 2HU4-epicatechin gallate complex. Importantly, based on higher docking score, better pharmacokinetic properties, higher drug-likeness and bioavailability score, it can be deduced that molecules such as sinapic acid, ferulic acid, lariciresinol, cryptomeridiol, genistein, eupatin, kaempherol, fsetin, scutellarein could have potential as drug candidate to control infuenza.

Infuenza virus has been a great threat to public health and health care system. Vaccines and a few anti-viral drugs are available as antiviral therapy for the treatment of patients. However, as evidenced by the appearance of the new 2009 H1N1 type pandemic virus of swine origin in april 2009, vaccinations are not always available in time (Pleschka et al. [2009](#page-13-28)). The acquisition of resistance to M2 channel inhibitors such as amantadine, is also identifed as a potential health risk, globally. Therefore, alternative therapeutic approaches to overcome such resistance are urgently needed to minimize fu symptoms. To meet the growing need for new antiviral agents to overcome the increasing problem of antiviral drug resistance, plants can be suitable candidates due to their ability to produce a large number of phytochemical substances and have a long history of safe and successful use as traditional medications against infectious diseases (Guo et al. [2006\)](#page-12-32). From our results, we speculated that these preparations can be promising choice in the prevention and treatment of infuenza virus infections because of their antiviral characteristics, natural abundance and non-toxic properties.

Conclusions

The present study focused on the investigation of the efficacy of diferent herbs on isolated infuenza viruses. The study reported the inhibitory efects of the methanol extract of *C. sinensis, P. hydropiper, M. pruriens, C. album* on diferent subtypes of infuenza virus strains and later identifed their compounds as the potential drug candidates by employing extensive in silico methods. However, further animal studies must be needed to confrm their efectiveness against the crucial targets of infuenza virus.

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Declarations

Ethical statement This article does not contain any studies involving animals performed by any of the authors. This article does not

contain any studies involving human participants performed by any of the authors.

Conflict of interest Md Abu Sayem Khan has no confict of interest. Rifat Parveen has no confict of interest. Sheikh Ariful Hoque has no confict of interest. Md Firoz Ahmed has no confict of interest. Abu Shara Shamsur Rouf has no confict of interest. Sabita Rezwana Rahman has no confict of interest.

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