Potential natural compounds for preventing 2019-nCoV infection

Hansen Chen¹ *, Qiaohui Du²*

1. Department of neurosurgery, School of Medicine, Stanford University
2. School of Chinese Medicine, The University of Hong Kong

*These two authors contributed equally to this study

Corresponding:

Dr. Hansen Chen: chenhhs@stanford.edu
300 Pasteur Dr, Palo Alto, CA 94304

Or Qiaohui Du: elvisdu@hku.hk

10 Sassoon Road, Pokfulam, Hong Kong
Abstract

2019-nCoV, a novel coronavirus, caused the pneumonia outbreak in China and continue to expand. 2019-nCov share the same host receptor with SARS-Cov, the Angiotensin-converting enzyme 2 (ACE2). Therefore, targeting ACE2 holds the promise for preventing 2019-nCov infection. Chinese Medicine herbs could be a valuable pool for identifying active compounds for treating infection of 2019-nCov. In this study, we summarize several active compounds including baicalin, Scutellarin, Hesperetin, Nicotianamine and glycyrrhizin that could have potential anti-2019-nCov effects, and we conduct molecular docking to predict their capacity for binding ACE2, subsequently preventing the 2019-nCov infection. We propose that these selected compounds worth further investigation for preventing 2019-nCov.

Keywords: 2019-nCov, Baicalin, Scutellarin, Hesperetin, Nicotianamine, Glycyrrhizin
Introduction

2019-nCoV, a novel coronavirus, caused the pneumonia outbreak in Wuhan city, Hubei Province, China and subsequently expands. The original pneumonia cases were linked to a large seafood and animal market in Wuhan. This is an emerging, rapidly evolving situation. One genome sequence (WH-Human_1) of the Wuhan CoV was first released on Jan 10, 2020, and subsequently, five additional Wuhan CoV genome sequences were released (Zhang, 2020; Shu and McCauley, 2017). By comparing to the genomes of SARS-CoV and MERS-CoV, the WH-human_1 genome has a better sequence homology toward the genomes of SARS-CoV than that of MERS-CoV. By using structural modeling of its S-protein, scientists suggest the strong interaction of human ACE2 molecules with 2019-nCoV [https://doi.org/10.1007/s11427-020-1637-5]. ACE2 is a type I integral membrane protein, with its active site domain exposed to the extracellular surface of cells. ACE2 has been demonstrated to be a functional receptor for the SARS-coronavirus (CoV) (Kuhn et al., 2006). Michael Letko et al. showed that the 2019-nCoV receptor-binding domain (RBD) was capable of entering cells expressing human ACE2, but not any of the other receptors, further confirming that human ACE2 is the receptor for the recently emerging 2019-nCoV [https://doi.org/10.1101/2020.01.22.915660]. As the host cell receptor is critical
for the virus entry, targeting ACE2 holds the promise for preventing infection of 2019-nCov infection.

The development of drugs for targeting 2019-ncov could be time-consuming, and the safety of newly-developed drugs could be a major concern, which needs time for testing. It seems unrealistic to synthesize new drugs and tests for safety and toxicity within such limited time. Traditional Chinese Medicine has been practiced in China for thousands of years, and Chinese medicine licorice was suggested to be promising for treating SARS (Pilcher, 2003). Considering the low toxicity and availability, screening active compounds from Chinese herbal medicine for targeting the ACE2 receptor could be a potential strategy for treating 2019-nCov. In this mini-review, we summarize the potential natural compounds that could target ACE2 for the potential treatment of 2019-nCov. By using molecular docking, we proposed that baicalin, Scutellarin, Hesperetin, glycyrrhizin and Nicotianamine are potential compounds that target the ACE2 receptor and exert anti-virus effects for preventing 2019-nCov infection.
Natural compounds candidates for 2019-nCov treatment

In the following session, we will summarize these natural compounds that may have therapeutic effects against 2019-nCov infection. To generate putative binding poses, we used the AutoDock Vina software package with the default scoring function (Trott O, Olson AJ, 2010). In the AutoDock Vina configuration files, the parameter num_modes was set to 1000 and exhaustiveness to 100. We identified the receptor binding pocket based on the structures of ACE2 proteins. We chose all the rotatable bonds in ligands to be flexible during the docking procedure, and we kept all the protein residues inside the binding pockets rigid. We assigned the Gasteiger atomic partial charges and converted all receptors and ligands to the PDBQT format using the AutoDockTools package. We did not use explicit hydrogens either for the receptors or for the ligands.

Baicalin

Baicalin is extracted and purified from the Chinese medicinal plant Scutellaria baicalensis Georgi (Chinese name: Huang Qin). Baicalin has broad therapeutic effects, including anti-oxidative stress, anti-inflammation, anti-apoptosis (Chen et al., 2017; Ishfaq et al., 2019). Baicalin has been shown its antiviral activities for SARS coronavirus using the foetal rhesus kidney-4 (fRhK-4) cell line, with a EC50
12.5µg/ml at 48 hours, and selectivity index more 4 to 8 (Chen et al., 2004). The plaque reduction assay showed that baicalin has a EC 50 of 11µg/ml (Chen et al., 2004). Those results suggest that baicalin has anti-SARS effects. Since the 2019-nCOV shared similarity with the SARS virus, we suspect that baicalin may also show anti-virus effects on 2019-nCov. In addition, a study showed that baicalin can inhibit ACE, with an IC50 value of 2.24 mM in vitro (Deng et al., 2012). However, whether baicalin can bind to ACE2 is not yet studied. Therefore, we use the molecular docking to test whether baicalin could bind to the ACE2 receptor, and subsequently may block the entry of 2019-nCov. The docking result shows that baicalin may have strong binding to the ACE2 enzyme (Figure 1), with an estimated ΔG (kcal/mol) -8.46, and the potential binding site at ASN-149, ARG-273, HIS-505. Given the low toxicity of baicalin, its efficacy on anti-2019nCov worth further investigation.
Scutellarin is another active compound from Chinese Medicine Erigeron breviscapus (Vant.) Hand Mazz, which showed broad pharmacological effects, including anti-oxidant, anti-inflammation, vascular relaxation, anti-platelet, anti-coagulation (Wang and Ma, 2018). A study showed that scutellarin treatment could reduce the expression and activity of ACE in brain tissue \textit{in vivo} (Wang et al., 2016). The IC50 value of scutellarin against ACE was $48.13 \pm 4.98 \mu M$ (Wang et al., 2016). However, whether scutellarin could inhibit ACE2 is not yet reported. Here we conduct a molecular docking and find that scutellarin may bind to ACE2, with estimated $\Delta G$ (kcal/mol) $-14.9$, with binding site GLU-495, UNK-957, ARG-482.

\textbf{Figure 1}, molecular docking result of baicalin to ACE2 enzyme.
Therefore, it’s worthwhile to test whether scutellarin could inhibit ACE2 and block the infection of 2019-nCov.

**Figure 2**, molecular docking result of Scutellarin to ACE2 enzyme.

**Hesperetin**

Hesperetin is a bioflavonoid compound abundant in Chinese Medicine citrus aurantium and Citri Reticulatae Pericarpium. Hesperetin dose-dependently inhibited cleavage activity of the 3C-like protease (3CLpro) of SARS-coronavirus in cell-free and cell-based assays, with an IC50 8.3 uM (Lin et al., 2005). Whether hesperetin could inhibit 2019-nCov replication is not yet investigated. To
understand whether hesperetin has the potential to inhibit ACE2, we conduct the molecular docking of Hesperetin to the ACE2 enzyme, and the results showed that hesperetin has the potential biding to ACE2 with an estimated ΔG (kcal/mol) -8.3, with binding cites TYR-613, SER-611, ARG-482, GLU-479 (Figure 3). This result suggests that hesperetin may bind to ACE2, therefore, block the infection of 2019-nCOV.

Figure 3 molecular docking result of Hesperetin to ACE2 enzyme.

Nicotianamine

Nicotianamine is rich in soybean (Takenaka, 2009). Toshihiro et al. reported that nicotianamine is a potent inhibitor of ACE2, with an IC₅₀ value of 84 nM. The authors screened ACE2 inhibitors from various foodstuffs and found that soybean
contained vigorous ACE2 inhibitory activity. They isolated the active compound “soybean ACE2 inhibitor” (ACE2iSB), which was identical to nicotianamine by direct comparison with a standard compound. Since ACE2 is critical for the 2019-nCov infection, we hypothesize that nicotianamine may block the infection of 2019-nCov through inhibiting ACE2, which needs further investigation.

**Glycyrrhizin**

Glycyrrhizin, another plant product isolated from Chinese Medicine herb licorice root (Glycyrrhiza radix), a herb that is promising for SARS treatment (Pilcher, 2003). Glycyrrhizin is used for treating chronic hepatitis and is relatively non-toxic. *In vitro* study showed that glycyrrhizin has anti-SARS-CoV effects. It inhibited viral adsorption and penetration and was most effective when administered both during and after the viral adsorption period (Cinatl et al., 2003). The introduction of specific chemical modifications increase the antiviral potency of glycyrrhizin, but also increased the cytotoxicity, thus the selectivity index was reduced as compared with that of glycyrrhizin (selectivity index: ≥65) (Hoever et al., 2005). Whether glycyrrhizin has anti-2019-nCov effects need further investigation. Our docking
results showed that glycyrrhizin has the potential biding to ACE2 with an estimated \( \Delta G \) (kcal/mol) - 9, with binding cites ARG-559, GLN-388, ARG-393, ASP-30 (Figure 4)

Figure 4 molecular docking result of glycyrrhizin to ACE2 enzyme.

Summary

Drug development for treating 2019-nCov is timely important due to its rapid expansion. Vaccine development could take a long time to complete, and its safety needs to be verified. Synthesized agents for blocking ACE2 also needs to test its toxicity. Chinese Medicine is applied for anti-virus treatment for a long time, and active compounds from Chinese Medicine may be applied for 2019-nCov virus. Due to the low toxicity and availability of some active compounds from Chinese Medicine, it is worthwhile to select potential candidates for 2019-nCov treatment.
Since 2019-nCov share some common sequence with SARS-Cov, and used the same host receptor ACE2, we review the potential active compounds for anti-SARS-Cov, and at the same time predict the binding affinity of those compounds to bind ACE2. In this study, by using the molecular docking and reviewing the literature, we report for the first time that baicalin, Scutellarin, Hesperetin, Nicotianamine, glycyrrhizin has the potential to bind to ACE2 and block the entry of 2019-nCov. Further studies are needed to verify our results and test the anti-2019n-Cov effects of these compounds.

Acknowledgment

The authors did not receive grants for this study

Conflict of interest

None
Reference


