



A narrative review of the scientific natures of the prevention and treatment of COVID-19 with traditional Chinese medicine

Qiangzhong Pi¹, Wanying Tan², Zhenghuai Tan²

¹Department of Respiratory and Critical Care Medicine, Southwest Hospital, Chongqing, China; ²Institute of Traditional Chinese Medicine Pharmacology and Toxicology, Sichuan academy of Chinese Medicine Sciences, Chengdu, China

Contributions: (I) Conception and design: Q Pi; (II) Administrative support: None; (III) Provision of study materials or patients: None; (IV) Collection and assembly of data: W Tan; (V) Data analysis and interpretation: None; (VI) Manuscript writing: All authors; (VII) Final approval of manuscript: All authors.

Correspondence to: Zhenghuai Tan. Institute of Traditional Chinese Medicine Pharmacology and Toxicology, Sichuan Academy of Chinese Medicine Sciences, 51, section 4, Renmin South Road, Chengdu 610041, China. Email: tanzhh616@sohu.com.

Abstract: Corona virus disease 2019 (COVID-19) has shown a pandemic around the world and it has caused more than a million cases in more than 130 countries or regions. While it has taken thousands of lives, there are no vaccines, antibodies nor anti-viral drugs at present. Confronted by massive difficulties, researchers have still gained great success in the prevention and treatment of COVID-19 via vigorously using the integrated therapy of traditional Chinese medicine (TCM) and western-style medicine. To describe and discuss the cognitions of SARS-CoV-2, pathogenesis of COVID-19, mainly around ACE2, and application TCM in the prevention and treatment of COVID-19 to provide clinical references of this disease. An overview of pathogenesis of SARS and COVID-19 and TCM may be used in the treatment of SARS and COVID-19 is presented. Narrative overview of the literature synthesizing the findings of literature retrieved from searches of computerized databases. Characteristics of TCM are the organic conception of the human body and discriminate and dialectical demonstration. It is worth noting the application of TCM in the prevention and treatment of COVID-19 to provide clinical references of this disease. Through block of key links of the disease and the combined treatment, we would finally heal SARS and COVID-19 patients.

Keywords: SARS-COV-2; corona virus disease 2019 (COVID-19); SARS CoV; traditional Chinese medicine (TCM)

Received: 20 December 2020; Accepted: 22 March 2021; Published: 30 March 2021.

doi: 10.21037/lcm-20-58

View this article at: <http://dx.doi.org/10.21037/lcm-20-58>

Introduction

Corona virus disease 2019 (COVID-19) has been worldwide detected in more than 216 countries (1). Chinese scientists identified structure of this virus at first time. The novel coronavirus was finally named as SARS-CoV-2 by Coronavirus Study Group (CSG) of International Committee on Taxonomy of Viruses (ICTV) because the virus is thought as a sister species of severe acute respiratory syndrome associated coronavirus (SARS-CoVs) (2). This paper intends to summarize the previous researches in PubMed of SARS-CoV and COVID-19 by combining

with the current comprehending treatment with therapy of TCM.

We present the following article in accordance with the Narrative Review checklist (available at <http://dx.doi.org/10.21037/lcm-20-58>).

Methods

Narrative overview of the literature synthesizing the findings of literature retrieved from searches of computerized databases of PubMed.

Structural and biological characteristics of SARS-CoV and SARS-CoV-2

SARS CoV spike virus is isolated from Himalayan palm civet gained from Guangdong wild animal market. Compared with human SARS CoV, animal SARS CoV has 29 additional nucleotide sequences. In the additional 29 nucleotide sequences of animal SARS CoV 10 and 11 open reading frame (ORF) combine as a novel ORF, encoding 122 putative proteins which biological significance is unknown (3). It is not clear how the precursor virus adapts itself in human to achieve effective transmission in populations. But it showed that replication of SARS CoV is attenuated by a 29-nucleotide deletion in SARS-coronavirus acquired during the early stages of human-to-human transmission. Without this mutation, the SARS epidemic in 2003 would have gone through a more serious result (4). It is known that SARS-CoV is a virus with positive-sense, non-segmented, single-stranded RNA genomes and belongs to the genus Betacoronavirus of Coronaviridae (5). Its first two thirds of RNA sequences encode the replicase genes, which are translated into two large polyproteins that are processed into 15 or 16 non-structural proteins (nsp) via proteolytic cleavage (5). The other one third of the RNA sequences are ORFs for the structural proteins, namely the spike (S), envelope (E), membrane (M), and nucleocapsid (N) proteins and accessory proteins of SARS-CoV, namely ORFs 3a, 3b, 6, 7a, 7b, 8a, 8b, and 9b (6). SARS-CoV-2 also belongs to the genus Betacoronavirus and has a 30-kb genome which 80% are homologous genome compared with SARS-CoV (7). The protein structures of SARS-CoV-2 and SARS CoV are very similar, mainly are spike (S), membrane (M), and nucleocapsid (N) proteins (7), and also contains SARS-CoV-2 accessory proteins namely orf1a, orf1b, S, ORF3a, ORF6, ORF7a, ORF7b, ORF8, ORF10 (8).

Study indicated that both SARS-CoV and SARS-CoV-2 entire human cells through binding with receptors which is a metalloproteinase, namely angiotensin-converting enzyme 2, or ACE2 receptor (9). And it is clearly that SARS-CoV infects target cells mainly via bindings spike proteins with ACE2. ACE2 has already been proved to be a functional receptor of SARS CoV (10) and its binding domain of SARS CoV to ACE2 is known as the 270-510 amino acid sequence of S protein (11). S protein is also thought as specific to bind to ACE2 with strong affinity via biochemistry and crystal structure analysis (12,13). The N-terminal of S protein (S1) of SARS CoV is receptor binding region, while the C-terminal (S2) is a membrane anchored membrane

fusion subunit, which contains two heptapeptide repeat regions (HR1 and HR2) (14). S1 subunit contains receptor binding domain, which can bind to the peptidase domain of ACE2, and S2 is responsible for membrane fusion function (14). When S1 region binds to the receptor of ACE2, the cleavage site on S2 is exposed and cleaved by the protease of host, which is crucial for virus infection (15).

After binding to ACE2 of the target cell, the transmembrane protein changes its conformation through the binding to HR1 and HR2, forming a six helix-oligomer complex, which leads to the fusion of the virus and the target cell membrane (16). While cells expressing ACE2 mediate SARS CoV infection, its role in initiating virus replication is not clear.

After binding with ACE2, S protein was destroyed by a way of acid dependent hydrolysis through protease (17). SARS CoV can infect cells from lung, intestine, liver, distal convoluted renal tubule, sweat gland, parathyroid gland, pituitary gland, pancreas, adrenal gland and brain, defecate, lymph node, spleen, heart and skeletal muscle (18,19).

Clinical characteristics of SARS and COVID-19 (Table 1)

The main clinical manifestations SARS and COVID-19 are shown in *Table 1*.

Pathogenesis of SARS and COVID-19

The damage effect of SARS CoV and SARS CoV-2 induced by combination with ACE2

Extensive lung injury in SARS patients seems to be related to a high initial virus load (29), which may relate with two damage direct effects: the first is to combine with ACE2, the other is to cause a series of immune reactions SARS-CoV.

The binding of S protein of SARS CoV with ACE2 on target cells can cause the down-regulation of ACE2 and the increasing of local of Ang II (30) and finally leads to acute lung injury (31), which can be improved by blocking the renin-angiotensin pathway. Low ACE2 can transform Ang II into Ang 1-7 that has effects of vasodilation and can increase pulmonary vascular permeability, by activating AT1R and leads to further lung tissue damage (30). SARS-CoV-2 mainly infects the lower respiratory tract and infect alveolar epithelial cells via binding to ACE2 (32). But studies have shown that ACE2 gene polymorphism is not related to the susceptibility and prognosis of SARS, suggesting that the reduction of ACE2 caused by SARS

Table 1 Clinical characteristics of SARS and COVID-19

Clinical characteristics	SARS	COVID-19
Fever	Fever was the main clinical feature of SARS (19)	99% cases had symptom of fever, in some cases, there was no sign of fever before the onset (20)
Gastrointestinal symptoms	Watery diarrhea was the most common extrapulmonary symptom, 40–70% of SARS patients will have diarrhea symptoms within first week (19)	10.1% cases had symptoms of diarrhea or nausea (20)
Liver function	Most cases have liver dysfunction and the liver enzyme level is obviously increased (21)	Some show a high liver enzyme level (22)
Renal function	Acute renal dysfunction is not common in SARS, but lead a high mortality (23)	29% cases have acute renal injury (24)
Respiratory symptoms	ARDS mainly infected airway and alveolar epithelial cells (19)	Dry cough, dyspnea and acute respiratory distress syndrome (20)
Cardiovascular symptoms	Tachycardia was the most common cardiovascular complication. Other complications include hypotension shock, atypical chest pain, bradycardia, cardiac hypertrophy, and atrial fibrillation in very few patients (25,26)	Increased troponin, palpitation, and cardiac failure often occurs in patients with potential cardiovascular disease (27,28)

Overview of the main clinical manifestations SARS and COVID-19. COVID-19, corona virus disease 2019; SARS, severe acute respiratory syndrome.

CoV has a relatively limited role in the occurrence and development of the disease (33).

Damages of SARS-CoV, SARS-CoV-2 caused through immune mechanism

SARS and COVID-19 patients present with various underlying medical conditions, so far, no specific cytokine markers have been found, which also indicates that the immune response caused by SARS-CoV and SARS-CoV-2 may involve in multiple ways. Lymphopenia with rapid decrease of CD4 and CD8 T cells is common in acute phase of SARS (34), which mechanism is unknown and may be related with lymphocyte apoptosis induced by virus without the involvement of ACE2. There is no ACE2 expression in T, B lymphocytes or macrophages in all hemolymph organs (35).

Most SARS-CoV-2 patients (82.1%) were found to have peripheral lymphocytopenia, indicating that lymphocytes may infiltrate in the lung and/or cell damage pathway through apoptosis or heat shock which leads to the release of a large number of proinflammatory factors, which can lead to the release of a large number of proinflammatory factor (17). The imbalance of inflammatory pathway of endothelial cells is a key factor that leads to coagulation

disorder and the increased vascular permeability, which also can promote sepsis and cause organ failures. Endothelial cells also participate in many physiological processes, such as regulation of vascular tension, selective permeability of blood vessels, and provision of coagulant and anticoagulant surfaces (36,37). Strong pro-inflammatory cascade may be induced by LPS which eventually leads to acute injury of capillaries and alveolar epithelial cells (38-40) and stimulates the production of proinflammatory mediators which indirectly activate endothelial cells and causes endothelial dysfunction. LPS promotes the combination of eNOS and NADPH oxidase 2 (NOX2) and the superoxide produced by NOX2 leads to the uncoupling of eNOS and the dysfunction of endothelial barrier and influences adhesion of leukocytes to the vascular wall (41,42). LPS can also down regulate the expression of protein and mRNA of eNOS (43).

Possible mechanism of prevention and treatment of COVID-19 by TCM

TCM is used for treatment of COVID-19 and is effective in prevention of process which moderate stage trans to severe stage. And progress of COVID-19 or mortality of COVID-19 are greatly reduced by TCM using.

Protecting the function of respiratory barrier is helpful to the prevention and treatment of early stage COVID-19

Shegan Mahuang Decoction is known as an expectorant, which can warm the lung and disperse the cold and is used to purify phlegm. Study indicated that Shegan Mahuang Decoction can reduce the lung infiltration of CD3+ and CD4+ T cells in mice (44). Maxingshigan decoction (Decoction of Ephedra, Apricot Kernel, Gypsum and Licorice) is a compound Chinese medicine preparation composed of ephedra, almond, gypsum and licorice. It is reported that Maxingshigan decoction can significantly improve the lung injury in rats and protect the alveolar capillary barrier by regulating HMGB1/TLR4/NF- κ B pathway and inflammatory response (45).

Mucous of respiratory tract is not only the main site of virus attack, but also is the defense system against virus infection. The innate immune system first senses virus invasion and the starts nonspecific clearance of virus. But if the virus escapes the early defense, it will be cleared by adaptive immunity. The main adaptive immune mechanisms include: (I) the participation of specific SIgA antibody and CD8+ CLTS in the recovery period of the initial virus infection; (II) when the virus-ig complex cleans up the virus, the pre-formation of specific SIgA antibody and IgG play an important role. And in fact, many single traditional Chinese medicine (TCM) or compound prescription, such as Astragalus, Sanqi, Siwu Decoction and some jiebiao prescriptions, regulate mucosal immunity by increasing the secretion of S-IgA (46). Studies showed that Astragalus, Yupingfeng powder and other TCM have certain regulatory effect on respiratory symbiotic bacteria in the treatment of influenza (46). The anti-influenza effect of volatile oil and cinnamaldehyde in cassia twig was also promoted by activating TLR7 pathway in mucosal immunity, inducing the expression of i-rak-4 protein and promoting the high expression of IFN- β (46), which suggested that TCM can prevent or reduce the invasion of SARS-CoV-2 and the direct damage of it by improving the resistance of respiratory mucosa to virus.

Inhibition or elimination of coinfecting virus and reduction of further invasion of SARS-CoV-2

Many TCMs have the antiviral effect via direct or indirect ways, which is the material base for the clinical application of one or two TCMs to treat cold. For example, Emodin is one of the main effective components of rhubarb, Polygonum cuspidatum and other TCMs (47). Emodin can

also inhibit the interaction between S-protein and ACE2 protein and inhibit the infection of S-protein retrovirus in a dose-dependent manner (48).

TCM with bacteriostatic that treat the deterioration of COVID-19 disease

Many TCMs for clearing away heat and detoxifying have antibacterial and virus killing properties, among which Coptis, phellodendron, Houttuynia cordata, Ligusticum chuanxiong and prickly ash are the most famous.

Pudilan Xiaoyan oral liquid, which is composed of dandelion, isatis root, kuditin and scutellaria, has the special effects of bacteriostasis and antiviral and clear away heat and detoxification via improving pharynx symptoms, detumescence, effectively blocking the replication and reproduction of the virus *in vivo* (49).

Houttuynia cordata is also a traditional herbal medicine used to clear away heat and detoxify. Its polysaccharide (CHCP) can significantly reduce the acute lung injury caused by hemorrhagic shock and LPS infusion in rats and inhibit the fever, the deposition of complement activation products in the lungs and increase of leukocyte (50). It can also restore the level of serum complement, reduce the pulmonary edema and protein exudation in BALF. The total flavonoids extracted from the leaves of crotalaria mucronat also showed bacteriostatic effect on Escherichia coli, Bacillus subtilis, Proteus vulgaris, Staphylococcus aureus and Pseudomonas aeruginosa (51).

Berberine is an alkaloid of broad-spectrum antibacterial drug, which can inhibit and eliminate a variety of G+ bacteria, G-bacteria, fungi and molds. It has a good inhibitory effect on Staphylococcus aureus and methicillin-resistant Staphylococcus aureus (51). Cinnamon oil also has obvious antibacterial activity against Escherichia coli, Staphylococcus aureus and resistant Staphylococcus aureus (51).

Tea polyphenols can destroy the cell membrane structure of Staphylococcus aureus and Pseudomonas aeruginosa, especially a higher concentration of tea polyphenols would destroy the cell membrane structure in a short period of time, resulting in the leakage of electrolyte inside the cell, and the gradual leakage of carbohydrate outside the cell, thus affecting the stability of cell structure, and finally making the cell gradually aureus (51).

Regulation of the endothelial function and reversion of the process of COVID-19

Statistical analysis showed that there were significant gender differences in incidence rate and mortality rate of SARS and

COVID-19, that is, male patients and mortality rates were significantly more than females, suggesting that estrogen plays an important role in the occurrence and development of SARS and COVID-19.

It was reported that estrogen receptors, including ER α , ER β and G protein coupled estrogen receptors (GPERs), also expressed in endothelial cells (52).

Many TCM and its effective components have the function of activating estrogen receptor, and play a protective role on the structure and function of vascular endothelial cells. Flavonoids is thought to be the main active components that have estrogen like effect (53).

Salvia miltiorrhiza (Danshen) can significantly increase the expression of ER α and ER β protein and their mRNA, upregulates serum estradiol (E2) but decrease follicle stimulating hormone and luteinizing hormone and luteinizing hormone, which is very similar to the effect of estrogen (54). But compared with estrogen, Salvia miltiorrhiza can play the role of estrogen without adverse reactions, and improve the expression of ER in target tissues (54). Both Daidzein and Genistein can produce estrogen like effects via activation of the transcription of estrogen response primitives reporter gene through ER α and ER β , two subtypes of ER, which can be blocked by ER antagonist ICI 182780 (54).

Discussion

An overview of pathogenesis of SARS and COVID-19 and TCM may be used in the treatment of SARS and COVID-19 is presented. LPS is a key factor of moderate and severe COVID-19. ACE2 is also involving in the dysfunction of vascular endothelial cells, the exploration of pathogenesis of COVID-19 may involving the interaction between LPS and ACE2.

TCM is a cultural and medical treasure left by the Chinese ancestors which is even a treasure to the world. In history, it has been made great contributions in the Chinese people's victory over disease and plagues. In modern times, it also showed a strong ability to cure and save patients presented with SARS and COVID-19. From the previous TCM we talked, it can be seen that TCM can block the progress of SARS and COVID-19 and make it good in a good prognosis.

But Studies included in article are non-RCTs, which leads to lack of reliability in the conclusions should be considered in the treatment of TCMs. It is worth noting that th TCM may be inspiration of future research in SARS

and COVID-19.

Conclusions

It is worth noting that the characteristics of TCM are the organic conception of the human body and discriminate and dialectical demonstration. Through block of key links of the disease and the combined treatment, we would finally heal SARS and COVID-19 patients.

Acknowledgments

Funding: None.

Footnote

Reporting Checklist: The authors have completed the Narrative Review reporting checklist. Available at <http://dx.doi.org/10.21037/lcm-20-58>

Conflicts of Interest: All authors have completed the ICMJE uniform disclosure form (available at <http://dx.doi.org/10.21037/lcm-20-58>). The authors have no conflicts of interest to declare.

Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved.

Open Access Statement: This is an Open Access article distributed in accordance with the Creative Commons Attribution-NonCommercial-NoDerivs 4.0 International License (CC BY-NC-ND 4.0), which permits the non-commercial replication and distribution of the article with the strict proviso that no changes or edits are made and the original work is properly cited (including links to both the formal publication through the relevant DOI and the license). See: <https://creativecommons.org/licenses/by-nc-nd/4.0/>.

References

1. WHO. Coronavirus disease (COVID-19) outbreak situation. Available online: <https://www.who.int/emergencies/diseases/novel-coronavirus-2019>
2. WHO. Naming the coronavirus disease (COVID-19) and the virus that causes it. Available online: <https://www.who.int/emergencies/diseases/novel-coronavirus-2019/>

- technical-guidance/naming-the-coronavirus-disease-(covid-2019)-and-the-virus-that-causes-it
3. Lau YL, Peiris JS. Pathogenesis of severe acute respiratory syndrome. *Curr Opin Immunol* 2005;17:404-10.
 4. Muth D, Caormn V, Roth H, et al. Attenuation of replication by a 29 nucleotide deletion in SARS-coronavirus acquired during the early stages of human-to-human transmission. *Sci Rep* 2018;8:15177.
 5. Thiel V, Ivanov K, Putics Á, et al. Mechanisms and enzymes involved in SARS coronavirus genome expression. *J Gen Virol* 2003;84:2305-15.
 6. Liu D X, Fung T S, Chong K L, et al. Accessory proteins of SARS-CoV and other coronaviruses. *Antiviral Res* 2014;109:97-109.
 7. Lu R, Zhao X, Li J, et al. Genomic characterisation and epidemiology of 2019 novel coronavirus: implications for virus origins and receptor binding. *Lancet* 2020;395:565-74.
 8. Ahmed SF, Quadeer AA, McKay MR. Preliminary Identification of Potential Vaccine Targets for the COVID-19 Coronavirus (SARS-CoV-2) Based on SARS-CoV Immunological Studies. *Viruses* 2020;12:254.
 9. Zhou P, Yang XL, Wang XG, et al. A pneumonia outbreak associated with a new coronavirus of probable bat origin. *Nature* 2020;579:270-3.
 10. Li W, Moore MJ, Vasilieva N, et al. Angiotensin-converting enzyme 2 is a functional receptor for the SARS coronavirus. *Nature* 2003;426:450-4.
 11. Babcock GJ, Eshaki DJ, Thomas WD, et al. Amino Acids 270 to 510 of the Severe Acute Respiratory Syndrome Coronavirus Spike Protein Are Required for Interaction with Receptor. *J Virol* 2004;78:4552-60.
 12. Walls AC, Park YJ, Tortorici MA, et al. Structure, function and antigenicity of the SARS-CoV-2 spike glycoprotein. *Cell* 2020;181:281-292.e6.
 13. Li F, Goff SP. Receptor recognition mechanisms of coronaviruses: a decade of structural studies. *J Virol* 2015;89:1954-64.
 14. Celigoy J, McReynolds S, Caffrey M. The SARS-CoV heptad repeat 2 exhibits pH-induced helix formation. *Biochem Biophys Res Commun* 2011;412:483-6.
 15. Millet JK, Whittaker GR. Host cell proteases: Critical determinants of coronavirus tropism and pathogenesis. *Virus Res* 2015;202:120-34.
 16. Supekhar VM, Bruckmann C, Ingallinella P, et al. Structure of a proteolytically resistant core from the severe acute respiratory syndrome coronavirus S2 fusion protein. *Proc Natl Acad Sci U S A* 2004;101:17958-63.
 17. Fu Y, Cheng Y, Wu Y. Understanding SARS-CoV-2-Mediated Inflammatory Responses: From Mechanisms to Potential Therapeutic Tools. *Virology* 2020;35:266-71.
 18. Farcas GA, Poutanen SM, Tony M, et al. Fatal severe acute respiratory syndrome is associated with multiorgan involvement by coronavirus. *J Infect Dis* 2005;191:193-7.
 19. Hui DSC, Zumla A. Severe Acute Respiratory Syndrome: Historical, Epidemiologic, and Clinical Features. *Infect Dis Clin North Am* 2019;33:869-89.
 20. Wang D, Hu B, Hu C, et al. Clinical Characteristics of 138 Hospitalized Patients With 2019 Novel Coronavirus-Infected Pneumonia in Wuhan, China. *JAMA* 2020;323:1061-9.
 21. Chan HL, Leung WK, To KF, et al. Retrospective analysis of liver function derangement in severe acute respiratory syndrome. *Am J Med* 2004;116:566-7.
 22. Jin YH, Cai L, Cheng ZS, et al. A rapid advice guideline for the diagnosis and treatment of 2019 novel coronavirus (2019-nCoV) infected pneumonia (standard version). *Mil Med Res* 2020;7:4.
 23. Chu KH, Tsang WK, Tang CS, et al. Acute renal impairment in coronavirus-associated severe acute respiratory syndrome. *Kidney Int* 2005;67:698-705.
 24. Yang X, Yu Y, Xu J, et al. Clinical course and outcomes of critically ill patients with SARS-CoV-2 pneumonia in Wuhan, China: a single-centered, retrospective, observational study. *Lancet Respir Med* 2020;8:475-81.
 25. Gu J. Multiple organ infection and the pathogenesis of SARS. *J Exp Med* 2005;202:415-24.
 26. Yu CM. Cardiovascular complications of severe acute respiratory syndrome. *Postgrad Med J* 2006;82:140-4.
 27. Huang C, Wang Y, Li X, et al. Clinical features of patients infected with 2019 novel coronavirus in Wuhan, China. *Lancet* 2020;395:497-506.
 28. Liu K, Fang YY, Deng Y, et al. Clinical characteristics of novel coronavirus cases in tertiary hospitals in Hubei Province. *Chin Med J (Engl)* 2020;133:1025-31.
 29. Peiris JSM, Chu CM, Cheng VCC, et al. Clinical progression and viral load in a community outbreak of coronavirus-associated SARS pneumonia: a prospective study. *Lancet* 2003;361:1767-72.
 30. Glowacka I, Bertram S, Herzog P, et al. Differential downregulation of ACE2 by the spike proteins of severe acute respiratory syndrome coronavirus and human coronavirus NL63. *J Virol* 2010;84:1198-205.
 31. Imai Y, Kuba K, Rao S, et al. Angiotensin-converting enzyme 2 protects from severe acute lung failure. *Nature* 2005;436:112-6.
 32. Yang M. Cell Pyroptosis, a Potential Pathogenic

- Mechanism of 2019-nCoV Infection. SSRN Electronic Journal 2020. doi: 0.2139ssrn.3527420.
33. Chiu RW, Tang NL, Hui DS, et al. ACE2 gene polymorphisms do not affect outcome of severe acute respiratory syndrome. *Clin Chem* 2004;50:1683-6.
 34. Wong RSM, Wu A, To KF, et al. Haematological manifestations in patients with severe acute respiratory syndrome: Retrospective analysis. *BMJ* 2003;326:1358-62.
 35. Hamming I, Timens W, Bulthuis MLC, et al. Tissue distribution of ACE2 protein, the functional receptor for SARS coronavirus. A first step in understanding SARS pathogenesis. *J Pathol* 2004;203:631-7.
 36. Grandel U, Grimminger F. Endothelial Responses to Bacterial Toxins in Sepsis. *Crit Rev Immunol* 2003;23:267-99.
 37. Dauphinee S M, Karsan A. Lipopolysaccharide signaling in endothelial cells. *Laboratory Investigation* 2006;86:9.
 38. Goddard LM, Iruela-Arispe ML. Cellular and molecular regulation of vascular permeability. *Thromb Haemost* 2013;109:407-15.
 39. Chen H, Bai C, Wang X. The value of the lipopolysaccharide-induced acute lung injury model in respiratory medicine. *Expert Rev Respir Med* 2010;4:773-83.
 40. Okada H, Takemura G, Suzuki K, et al. Three-dimensional ultrastructure of capillary endothelial glycocalyx under normal and experimental endotoxemic conditions. *Crit Care* 2017;21:261.
 41. Wu F, Szczepaniak W, Shiva S, et al. Nox2-dependent glutathionylation of endothelial NOS leads to uncoupled superoxide production and endothelial barrier dysfunction in acute lung injury. *Am J Physiol Lung Cell Mol Physiol* 2014;307:L987-997.
 42. Förstermann U. Oxidative stress in vascular disease: Causes, defense mechanisms and potential therapies. *Nat Clin Pract Cardiovasc Med* 2008;5:338-49.
 43. Bernardini C, Greco F, Zannoni A, et al. Differential expression of nitric oxide synthases in porcine aortic endothelial cells during LPS-induced apoptosis. *J Inflamm (Lond)* 2012;9:47.
 44. Lin CC, Wang YY, Chen SM, et al. Shegan-Mahuang Decoction ameliorates asthmatic airway hyperresponsiveness by downregulating Th2/Th17 cells but upregulating CD4+FoxP3+ Tregs. *J Ethnopharmacol* 2020;253:112656.
 45. Fei YX, Zhao B, Yin QY, et al. Ma Xing Shi Gan Decoction Attenuates PM2.5 Induced Lung Injury via Inhibiting HMGB1/TLR4/NFκB Signal Pathway in Rat. *Front Pharmacol* 2019;10:1361.
 46. Yang JH, Zheng QG, Zhang XC, et al. Research progress of traditional Chinese medicine in immune response of respiratory mucosa against influenza virus infection. *Chinese Journal of Immunology* 2020;36:375-81.
 47. Dong X, Fu J, Yin X, et al. Emodin: A Review of its Pharmacology, Toxicity and Pharmacokinetics. *Phytother Res* 2016;30:1207-18.
 48. Yang L, Liu J, Bian Y, et al. The possibility of using emodin from Chinese herbal medicine as treatment for novel coronavirus pneumonia. *Pharmacy Today* 2020;4:1-19.
 49. Lu H, Chen L, Wan S, et al. Systematic Evaluation on Pudilan Xiaoyan Oral Liquid for Treating Pediatric Herpangina. *China Pharmaceuticals* 2013;22:24-7.
 50. Lu Y, Jiang Y, Ling L, et al. Beneficial effects of *Houttuynia cordata* polysaccharides on "two-hit" acute lung injury and endotoxic fever in rats associated with anti-complementary activities. *Acta Pharmaceutica Sinica B* 2018;8:218-27.
 51. Liu YN, Li XF, Ban XX, et al. The review on active antibacterial ingredients of Chinese medicine and the antibacterial mechanism. *Global Traditional Chinese Medicine* 2015;8:1012-7.
 52. Caulin-Glaser T, García-Cardeña G, Sarrel P, et al. 17β-Estradiol Regulation of Human Endothelial Cell Basal Nitric Oxide Release, Independent of Cytosolic Ca²⁺ Mobilization. *Circ Res* 1997;81:885-92.
 53. Vaya J, Tamir S. The relation between the chemical structure of flavonoids and their estrogen-like activities. *Curr Med Chem* 2004;11:1333-43.
 54. Zhao Y, Zheng HX, Ying XU, et al. Research progress in phytoestrogens of traditional Chinese medicine. *China Journal of Chinese Materia Medica* 2017;42:3474-87.

doi: 10.21037/lcm-20-58

Cite this article as: Pi Q, Tan W, Tan Z. A narrative review of the scientific natures of the prevention and treatment of COVID-19 with traditional Chinese medicine. *Longhua Chin Med* 2021;4:1.