Re-understanding anti-influenza strategy: attach equal importance to antiviral and anti-inflammatory therapies

Zhengtu Li^{1#}, Li Li^{2#}, Shuai Zhao^{3#}, Jing Li¹, Hongxia Zhou¹, Yunhui Zhang⁴, Zifeng Yang^{1,5}, Bing Yuan^{1,4}

¹State Key Laboratory of Respiratory Disease, National Clinical Research Center for Respiratory Disease, Guangzhou Institute of Respiratory Health, First Affiliated Hospital of Guangzhou Medical University, (Guangzhou Medical University), Guangzhou 510120, China; ²Department of Respiration, The First Hospital of Yulin, Yulin 719000, China; ³Department of Emergency, The Second Clinical College of Guangzhou University of Chinese Medicine, Guangzhou 510120, China; ⁴Department of Respiration, First People's Hospital of Yunnan Province, Yunnan 650032, China; ⁵Department of Faculty of Chinese Medicine, Macau University of Science and Technology, Macau 519020, China

Contributions: (I) Conception and design: Z Li, Y Zhang, B Yuan, Z Yang; (II) Administrative support: L Li, S Zhao, H Zhou, J Li, B Yuan; (III) Provision of study materials or patients: L Li, S Zhao; (IV) Collection and assembly of data: H Zhou, J Li; (V) Data analysis and interpretation: Z Li, L Li, S Zhao, B Yuan; (VI) Manuscript writing: All authors; (VII) Final approval of the manuscript: All authors.

[#]These authors contributed equally to this work.

Correspondence to: Zifeng Yang. State Key Laboratory of Respiratory Disease, National Clinical Research Center for Respiratory Disease, Guangzhou Institute of Respiratory Health, The First Hospital of Guangzhou Medical University, No. 151, Rd. Yanjiang, Guangzhou 510000, China. Email: jeffyah@163.com; Bing Yuan. Department of Respiration, First People's Hospital of Yunnan Province, No. 157, Jinbi Road, Kunming 650032, China. Email: 506638016@qq.com.

Abstract: The direct replication of influenza virus is not the only cause of harm to human health; influenza infection leading to a hyper-inflammatory immune response can also result in serious conditions. So, the treatment strategy for influenza needs to keep balance between antivirus and anti-inflammation. Herein, we review the treatment strategies of anti-influenza drugs and traditional Chinese medicines.

Keywords: Influenza; antivirus; anti-inflammation; chemical drug; traditional Chinese medicine

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Introduction

Influenza is an acute, contagious infectious disease caused by influenza virus; the disease can be harmful to human health, even leading to death. Great progress has been made in the treatment of influenza, especially for patients with severe influenza infection. The most effective approaches to prevention and control measures against influenza disease are vaccination and antiviral therapy. However, owing to antigenic variation of influenza viruses and individual differences in host immunity, the influenza vaccine can be less than effective. Thus, timely and regular antiviral treatment is especially important for confirmed influenzapositive patients, which can greatly reduce the patient's symptoms and complications and shorten the course of the disease (1,2). In recent years, together with new understanding of the pathogenesis of influenza, it has been recognized that influenza viruses can directly affect the host and can also cause indirect effects via severe immunological disorders. Influenza virus infection can promote chemokine and cytokine secretion, which help to clear influenza viruses from the host. However, in some cases, excessive or uncontrolled release of inflammatory cytokines occurs after influenza infection, resulting in a so-called "cytokine storm" that can lead to multiple organ damage and even death (3-6). Methods to alleviate the damage caused by a cytokine storm in response to influenza virus infection have become an important part of anti-influenza strategies.

We now have greater understanding of influenza viruses, including the direct harm caused by the virus itself as well

as the cytokine storm induced by influenza viruses. These can act together and cause multiple organ dysfunction, such as acute respiratory distress syndrome (ARDS). In this review, we discuss reliable and effective antiviral and antiinflammatory therapies, which must both be considered to guide the treatment of influenza patients.

Etiology, epidemiology, and impact of influenza viruses

Influenza viruses belong to the family Orthomyxoviridae and are single-stranded, negative-sense, segmented RNA viruses. According to nucleocapsid protein (NP) and matrix protein, influenza viruses are classified into influenza A, B, C, and D. Influenza A and B viruses contain eight different RNA segments, whereas influenza C viruses only have seven different RNA segments, lacking a fragment encoding the neuraminidase (NA) protein. The influenza virus genome is segmented, thus re-assorting among different virus strains readily occurs. In addition, influenza virus RNA often does not function correctly during replication; consequently, influenza has a higher mutation frequency than other viruses. Therefore, new influenza subtypes can easily emerge, leading to influenza epidemics. Influenza viruses have different species specificity. Influenza A virus can infect humans as well as pigs, horses, seals, whales, and ferrets, among other species. Influenza B virus has thus far been isolated in only humans, and influenza C virus has been isolated in only humans and pigs (7). Influenza D virus, recently reported in 2016, has not been implicated in human infection (8).

Influenza A and B viruses contain eight different negative sense (anti-mRNA sense) RNA segments that encode at least 11 viral proteins, including hemagglutinin (HA), NA, NP, matrix proteins (M1, M2), nonstructural proteins (NS1, NS2), and polymerase (PA, PB1, PB2, PB1-F2). Influenza C viruses only have seven different RNA segments, lacking a fragment encoding NA. The current study confirms that some of these proteins have an important role in the infection of the host by influenza virus. HA is responsible for virus attachment to host respiratory epithelial cells and facilitates virus entry into cells. NA can help the virus to pass through mucous membranes, which helps to promote the release and spread of newly synthesized virions from infected cells (9). The PB1, PB2, and PA subunits of influenza virus can form a heterotrimer that associates with NP-complexed viral RNA (vRNA) to form virion ribonucleoproteins

(vRNPs) (10). M2 is a multifunctional protein whose transmembrane domain plays a critical role in ion channel activity and whose cytoplasmic domain is required for vRNP incorporation and virion morphogenesis (11). Detailed information about influenza virus replication and the function of each protein have been well summarized in other recent reviews (12). These proteins are also the key targets for anti-influenza therapies. Currently, influenza virus infection can be effectively treated with anti-influenza antiviral agents, which are directed against these influenza virus proteins. These agents either inhibit viral replication by blocking the ion channel at the stage of virus entry into cells or they inhibit the function of virus-coded RNA polymerases or prevent the release of progeny virions from infected cells, among other functions (13,14).

About 20-30% of children and 5-10% of adults in the general population are affected by influenza annually. Approximately 3-5 million people develop severe influenza infection, leading to about 1 million deaths per year (15). Four influenza A pandemics have occurred during the 20th century: the 1918 Spanish influenza pandemic (H1N1), the 1957 Asian influenza pandemic (H2N2), the 1968 Hong Kong influenza pandemic (H3N2), and the 1977 Russian influenza pandemic (H1N1). In the present century, the threat of influenza A virus pandemics remains. In 2009, a new H1N1 influenza virus pandemic swept through 74 countries in 2 months, causing nearly 20,000 deaths (16). Furthermore, research has been focusing more on avian influenza A viruses, which are transmitted among birds and can cause cross-species transmission to infect humans; these viruses include H5N1 (17), H7N9 (18,19), and H5N6 (20). Human infection with avian influenza virus can cause acute lung injury, leading to severe ARDS and multiple organ dysfunction syndrome (MODS); the disease severity and mortality rates are higher than those owing to severe acute respiratory syndrome (SARS) coronavirus. Data published by the China Centers for Disease Control (CDC) show that the mortality rate of highly pathogenic avian influenza A (H5N1) virus is >60%, and influenza A (H7N9) virus has a mortality rate of about 40%. Thus, influenza virus continues to be a major threat to human health and public safety, especially avian influenza.

Pathogenesis of influenza virus infections

Influenza can lead to different degrees of inflammation. Initial symptoms are usually mild, and patients typically develop fever, cough, expectoration, and other symptoms of upper respiratory tract infection. However, some patients quickly develop severe symptoms and pneumonia, ARDS, and other organ dysfunction (21). It is generally believed that the pathogenesis of influenza virus infection includes two main aspects: direct damage caused by virus replication and indirect damage caused by an influenza-induced cytokine storm.

Direct damage caused by influenza virus

The surface hemagglutinin protein (HA) of influenza virus is a glycoprotein that binds to sialic acid receptors on host cells in the respiratory tract; HA mediates viral adsorption and fusion. Influenza viruses then enter host cells via endocytosis, the viral genome is transcribed and replicated in the nucleus, and viral nucleoprotein synthesized in the cytoplasm enters the nucleus and combines with viral RNA to form a nucleocapsid that is exported to the cytoplasm. The nucleocapsid is tightly wrapped in a cell membrane modified by virus-specific membrane proteins, and viral particles are subsequently released through a budding process to infect other normal cells (22). Depending on the viral receptors present, seasonal influenza viruses will generally adhere to and multiply in epithelial cells of the upper respiratory tract. However, avian influenza A viruses, such as H7N9 and H5N1, can invade epithelial cells of the lower respiratory tract and alveolar type II pneumocytes where it can efficiently replicate, which may cause deteriorating lung function (23-25).

Adhesion, multiplication, and release of influenza virus in epithelial cells can damage the barrier function of the respiratory tract mucosa, which can cause further damage to the host. First, this process damages the integrity of the normal epithelial cell layer and enhances bacterial colonization, allowing strongly adhesive bacteria [such as Streptococcus pneumoniae (S. pneumoniae)] to colonize (26,27). Second, viral replication exposes the basement membrane, which is rich in fibers that connect proteins, causing respiratory pathogens (such as S. pneumoniae) to have high-affinity binding to proteins. Third, the NA protein of influenza virus can promote viral release from infected cells and help the virus spread in the respiratory tract while simultaneously exposing bacterial receptors in respiratory tissue, which can increase secondary bacterial infection by cleaving sialic acid receptors of epithelial cells (28-30). Finally, influenza virus-infected cells may have reduced expression of antimicrobial peptides, thereby affecting the natural defense function of the host epithelium (31).

Indirect damage caused by influenza-induced excessive inflammatory response

In addition to direct damage, influenza virus also causes indirect damage by inducing an excessive inflammatory response. Influenza virus is highly infectious in human respiratory cells, including endothelial cells, plasma cell-like dendritic cells, epithelial cells, and macrophages. Influenza virus infection causes these cells to release chemokines and cytokines, leading to aggregation of inflammatory cells such as neutrophils, lymphocytes, and monocytes in inflammatory regions (22). For example, airway epithelial cells, macrophages, and dendritic cells of the lung that are infected by influenza virus, viral RNA within an infected cell, and/ or virus debris taken up by phagocytic cells lead to NLRP3 signaling activation, interleukin (IL)-1ß production, and recruitment of neutrophils and monocytes to the infection site, where they further secrete inflammatory cytokines (32). Dysfunction of the chemokine and cytokine response is associated with severe avian influenza virus infection (17,33), which may contribute to disease severity (34). Additional evidence has showed that patients with H5N1 infection have higher serum levels of macrophage and neutrophil chemoattractant chemokines (CXCL10, CXCL2, IL-8) and pro- and anti-inflammatory cytokines [IL-6, IL-10, interferon (IFN)- γ], particularly those patients who died (17,35). Furthermore, a study on the relationship between viral load and cytokines in the second wave of H7N9 patients in Guangdong Province, China during 2013-2014 showed that there was a positive correlation between viral titers in different clinical specimens and cytokine levels in plasma and bronchoalveolar fluid. Specifically, upregulation of proinflammatory cytokines IP-10, MCP-1, MIG, MIP-1 α/β , IL-1 β , and IL-8 was found in plasma and bronchoalveolar fluid (19). Increased levels of IP-10, MIG, MIP-1β, MCP-1, IL-6, IL-8, and IFN-α were detected in sera from acute patients with H7N9 infection, which were significantly higher than those in healthy study participants. Some low pathogenic avian influenza viruses, such as avian influenza A (H9N2) virus, can also promote the expression of proinflammatory cytokines and chemokines in human airway epithelial cells. H9N2 virus can stimulate production of proinflammatory cytokines (IL-1 β , IL-6) and chemokines (IL-8) (36). Similar findings have been observed in patients with severe influenza A (H1N1) pdm09 and seasonal influenza (37).

As described above, proinflammatory cytokine levels increase after influenza virus infection, which can help

clear the virus from host cells. However, this increase may lead to an imbalance in the host regulatory network, leading to a cytokine storm, which can cause more serious conditions such as ARDS and MODS (3-5). There is evidence that cytokine dysregulation contributes to the pathogenesis of H5N1 and H7N9 viruses, which cause severe respiratory disease (17,25). Additionally, cytokine levels have been positively correlated with prolonged hospitalization and C-reactive protein levels and negatively correlated with oxygenation index (38). Another study suggested IP-10 and IL-6 levels as biomarkers for severe cases of H7N9 disease (39).

Treatment strategy for influenza

As described above, damage caused by influenza virus infection can be classified as direct, such as that caused by virus replication, and indirect, such as that caused by a cytokine storm. Therefore, the clinical treatment of influenza should be focused on both antiviral and antiinflammatory therapies.

Importance of antiviral therapy

More and more evidence has confirmed that antiviral inhibitors are important for influenza treatment. The viral load in influenza patients correlates with disease severity and prolonged viral shedding (40), particularly in infants and young children, who are at high risk of influenza-related hospitalization and adverse disease outcomes. Therefore, antiviral agents are necessary in influenza treatment. One major class of antiviral drug is neuraminidase inhibitors (NAI), which can inhibit NA activities such as glycoside hydrolase activity, which cleaves glycosidic linkages of neuraminic acids to reduce viral spread and secondround infection. Such drugs include older NAIs, such as oseltamivir and zanamivir, and newer NAIs like laninamivir and peramivir. The other major class of antiviral drug is M2 ion channel protein inhibitors (M2 inhibitors), which target the influenza virus M2 protein and can inhibit the replication of influenza virus by interfering with the M2 protein ion channel activity. M2 inhibitors include amantadine and rimantadine. Other antivirals include hemagglutinin inhibitors, such as EB peptide and arbidol; viral RNA polymerase inhibitors, such as ribavirin; and traditional Chinese medicine, which contains anti-influenza activity.

NAI and M2 inhibitors have been approved by the

United States Food and Drug Administration for the clinical treatment of influenza. However, because of high resistance (41-43), M2 inhibitors have been withdrawn from the market; currently, NAIs are mainly used. NAIs, such as oseltamivir, remain the mainstay of influenza therapy as they have good tolerance, low toxicity, lower resistance rate, and can effectively inhibit viral replication and transmission of various influenza virus strains (44,45). Thus, NAIs are currently included in national and international guidelines for the use of influenza drugs to guide clinical treatment (45,46).

An accumulation of data has shown that anti-influenza drugs, such as oseltamivir, can significantly reduce the median duration of symptoms in patients with influenza infection and viral shedding (47). Clinical analysis of antiviral treatment has revealed that the median length of hospitalization of participants who received antiviral therapy was significantly shorter than that of participants who did not (2,48,49). Findings of a double-blind, randomized, controlled trial showed that median duration of symptoms and virus isolation were lower in patients who received oseltamivir treatment within 48 hours of symptom onset compared with those given a placebo (50,51); this finding suggests that early antiviral treatment is efficacious. However, further clinical trials and basic research studies suggest that delayed treatment with NAIs (>48 hours since illness onset) remains effective against influenza infection (52,53).

Although antiviral monotherapy has been shown to be effective, there are some limitations to its use. Primarily, viral resistance has been shown to reduce the activity of antivirals such as amantadine and rimantadine, which have been rendered largely ineffective owing to the high prevalence of resistant viruses in nature (54). Initially, NAI and M2 inhibitors were highly successful in inhibiting and preventing influenza infection, and nearly no resistance to influenza virus was detected (55,56). However, in 2003 and 2004, the frequency of adamantane-resistant A (H3N2) viruses in China increased substantially to 57.5% and 73.8%, respectively (57). Our previous study suggested that clinical isolates of H1N1 and H3N2 developed amantadine resistance at 93.1% and 100%, respectively, in Guangzhou, China, 2009 (43). Dong et al. found that 45.2% of influenza A (H1-H17) viruses circulating globally in 2013 were resistant to adamantanes (58). Up until 2007, influenza viruses resistant to the NA inhibitor oseltamivir had been isolated at a low level (13); however, since the beginning of 2007, the resistance of influenza A (H1N1) viruses

to oseltamivir has increased significantly, by 7.1% (59). Moreover, resistance to zanamivir has been reported among immunodeficient patients (60). Recently, according to antiviral susceptibility monitoring of the World Health Organization Global Influenza Surveillance and Response System, virtually all influenza A viruses circulating in humans have been deemed resistant to one category of antiviral drugs, namely, M2 inhibitors (amantadine and rimantadine). However, as of October 2017, the frequency of resistance to the NA inhibitor oseltamivir remains low (1–2%), (http:// www.who.int/mediacentre/factsheets/fs194/en/).

Thus, clinicians should consider combination therapy using two or three antiviral drugs to produce synergistic effects, increase anti-influenza virus activity, reduce the emergence of drug-resistant virus strains, and lower influenza mortality rates (61,62). However, combination therapy can reduce the use of individual antivirals and may have dose-related drug toxicity and adverse effects (13). One such example is oseltamivir-ribavirin combination therapy, which significantly inhibits influenza A virus replication in experimental mouse models, controls the spread of influenza viruses in the respiratory tract, and improves parameters of lung function, particularly mean breath volume (63,64). In addition, oseltamivir-zanamivir combination therapy has been found to be less effective than oseltamivir or zanamivir monotherapy for seasonal influenza A (H3N2) virus and H1N1 virus (65). Some combination therapies may exhibit antagonistic effects. One group noted that oseltamivir and peramivir both act on N1 NA enzymes; however, peramivir is more likely to bind to N1 NA enzymes than oseltamivir, which may lead to antagonistic interactions (66). Moreover, a retrospective study of adults infected with H7N9 virus showed that oseltamivir-peramivir combination therapy was not superior to oseltamivir monotherapy (67). Thus, antiviral drugs included in combination therapies should have different antiviral mechanisms (68). If the antivirals are in the same class, then an antagonistic effect between the two drugs is likely to occur because they compete for the same binding site, which ultimately affects the antiviral activity of the drug.

Owing to the rapid and extensive emergence of drug resistance among the available drugs, as well as the increasing need for combination therapy, new antiviral strategies with novel mechanisms of action and reduced drug resistance potential are urgently required. Recent years have seen the development of inhibitors that block virus-cell interactions occurring at different stages of influenza virus replication, such as attachment, entry, viral genome transcription and replication, nuclear export of viral products, and viral particles release. The new antiviral strategies target HA (69,70), RNA polymerase (71-73), NP (74), and NS1 (75); however, to date, no anti-influenza drugs targeting NS1 are under clinical development. These have been well summarized in other recent reviews (12,76). There are new developments in NA inhibitors, such as laninamivir, which holds great promise for its long-acting inhibitory activity (77). The characteristics of these anti-influenza virus drugs have been summarized in *Table 1*.

Here, we only review anti-influenza drugs according to the important functional proteins of influenza virus. However, other anti-influenza therapies should also be adopted, including those that regulate cell metabolism, host signaling pathways, and host antiviral response. Detailed information has been provided in other recent reviews (12,80).

Importance of anti-inflammatory therapy

In some cases, the cytokine storm induced by influenza virus infection is an important cause of ARDS, MODS and even patient death (6). Therefore, in addition to conventional antivirus therapies, the use of inflammation inhibitors in patients, especially those infected by highly pathogenic avian influenza viruses, is another possible therapeutic approach. Currently, corticosteroids are the most widely used and effective adjuvant therapy. Glucocorticoids can reduce alveolar effusion, capillary permeability, and the inflammatory response of the lung parenchyma and interstitium, to shorten the course of severe viral pneumonia. The use of glucocorticoids in the treatment of severe viral pneumonia has come from the experience of SARS treatment in Guangzhou, China. SARS coronavirus induces a IFN-y-related cytokine storm in patients, which might be involved in the immunopathological damage observed in patients with SARS (81). A retrospective study of Guangzhou SARS cases showed that corticosteroid use reduced mortality and shortened hospitalization (82). In an experimental mouse model of H1N1 infection, corticosteroid treatment ameliorated acute lung injury induced by 2009 H1N1 virus (83). However, other reports have suggested that corticosteroid therapy does not significantly improve the survival of patients with influenza A virus infection (84) and may even be associated with higher mortality in patients with severe influenza H1N1 infection (85). Among H1N1positive patients with ARDS in particular, the early use of

Table 1 Inhibitors of influenza virus	and their characteristics
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Subtype	Viral target	Classical drugs	Advantage	Disadvantage	Reference
M2 inhibitors	A/M2 ion channel	Amantadine and rimantadine	The earlier used; prevention and treatment of influenza	Have no activity again IBV and often associated with serious side effects; have high resistance	(43,57,78)
NA inhibitors	Neuraminidase	Oseltamivir, zanamivir, peramivir and laninamivir	Have activity again IAV and IBV; the long-acting inhibitory activity drug has been developed	Drug resistance gradually increases	(44,45,59,60,77)
HA inhibitors	Hemagglutinin		Have the broad antiviral effects; more and more this target drugs have been developed	The main limit is the hyper-variability of the globular head of HA	(69,70,79)
RdRP inhibitors	PB1, PB2 and PA	Favipiravir, Benzbromarone	Has the development of drugs with broad efficacy	Lack the clinical data	(71-73)
NP inhibitors	Nucleoprotein	Developmental medications; lack data to present.			(74)
NS1 inhibitors	Nonstructural protein 1	Developmental medica	tions; lack data to present.		(75)

IBV, influenza B virus; IAV, influenza A virus.

corticosteroids may be harmful (86). Therefore, there is no clear consensus on corticosteroid use in the treatment of severe viral pneumonia or on the timing, dosage, and course of treatment.

Passive immunotherapy may be a good option for the treatment of influenza A (H5N1 or H1N1) infection. Treatment with convalescent plasma in patients with H1N1 infection can effectively reduce respiratory tract viral load, serum cytokine response, and mortality (87,88). Furthermore, some groups have applied monoclonal antibodies to combat the abnormal secretion of cytokines under pathological conditions or have used specific recombinant protein to regulate the cytokine storm, which can also alleviate immune damage to some extent. For example, some studies have found that IP-10 and IL-17 were significantly elevated in the serum of patients with H1N1-induced ARDS. In an experimental mouse model of acute lung injury caused by influenza A (H1N1) virus infection, anti-IL-17 and anti-IP-10 monoclonal antibodies significantly reduced the expression of related cytokines and reduced lung injury (89,90).

Anti-inflammatory therapy has become an important component of anti-influenza strategies, and equal importance should be placed on antiviral and antiinflammatory therapies to effectively treat severe influenza virus infection. However, the optimal anti-inflammatory drug, duration of treatment, and timing of treatment remain unclear and require further investigation.

Influenza treatment with traditional Chinese medicine

There are some disadvantages to using antiviral and antiinflammatory therapies; mainly, most drugs do not contain both antiviral and anti-inflammatory activities. It has been postulated that novel antiviral agents could control influenza, thus attenuating excessive proinflammatory responses and limiting symptoms and tissue damage associated with infection. An increasing number of antiviral drugs influence host immune responses (91), with the most successful drugs possessing both antiviral and antiinflammatory functions (92). Based on this concept, traditional Chinese herbal medicine (TCM) has an innate advantage in that it can target the virus and the host simultaneously, offering more effective inhibition of host inflammatory responses induced by influenza virus infection. Thus, the development of novel therapeutic agents from TCM is a promising approach for anti-influenza strategies.

Various TCMs have been shown to have anti-influenza effects, such as the herb Radix isatidis (R. isatidis), honeysuckle, forsythia, and compound formulas Lianhua-Qingwen and MaxinShigan Tang. Lianhua-Qingwen capsules are an important TCM that has been confirmed to have anti-influenza A activity in a double-blind, randomized, positive-controlled clinical trial (93,94). Furthermore, we have found that Lianhua-Qingwen capsules have strong efficacy in inhibiting virus-induced excessive inflammation immune responses (95), which has also been observed with R. isatidis (96). Thus, in addition to directly inhibiting viral replication, TCM regulates influenza virus-induced excessive inflammatory immune responses; the latter effect is has a beneficial role in fighting influenza infection.

R. isatidis, also known as Ban-Lan-Gen in Chinese, has traditionally been used for the treatment of influenza, viral pneumonia, mumps, pharyngitis, and hepatitis (97). Clinical studies have also demonstrated high efficacy of R. isatidis in acute pharyngitis (50% caused by a virus) and influenza virus-induced acute upper respiratory infections and pneumonia, including cough, nasal congestion, runny nose, sneezing, and other inflammatory catarrhal symptoms (98,99). Another study showed that R. isatidis also plays a significant role in the treatment of viral myocarditis and other viral-mediated inflammatory diseases (100). These clinical observations suggest that R. isatidis has therapeutic effects for mild to moderate influenza-like symptoms and virus-induced inflammatory catarrhal symptoms.

Pharmacological studies have also demonstrated that R. isatidis components serve two main functions: antiviral and anti-inflammatory. Various types of compounds have been isolated from R. isatidis including polysaccharides, indirubin, clemastanin B, alkaloids, lignans, and flavonoids. Our research and previous studies have shown that R. isatidis polysaccharides inhibit in vitro replication of influenza virus (101); the underlying mechanism may include direct competitive binding to the HA of the virus, thereby inhibiting viral invasion and replication (102). We further elucidated that polysaccharides have a strong antiinflammatory effect on influenza virus via suppression of host toll-like receptor 3 (TLR3) signaling in vitro (96). Moreover, Kido et al. reported that one of the antiviral components of R. isatidis, namely, isatis root polysaccharide, induces the production of protective IgG antibodies, which may serve as an antiviral vaccine adjuvant (103).

It is unclear whether indirubin, an indole alkaloid, has a direct effect on influenza virus *in vitro* (104). However, it is clear that indirubin and its derivatives can significantly inhibit influenza virus (H5N1)-induced overproduction of inflammatory cytokines, such as IP-10 and MIG, in airway epithelial cells and macrophages (105). Moreover, indirubin has potent anti-influenza viral activity via inhibition of RANTES (also known as CCL5) expression (106). Concerning lignans, our preliminary study demonstrated that clemastanin B, a lignan, can slightly inhibit replication of influenza virus *in vitro* by inhibiting the expression and nuclear export of early influenza virus NP (97). In recent studies, we demonstrated that lariciresinol-4-O- β -dglucopyrano-side, another lignan, exhibited strong antiinfluenza virus A activity and decreased proinflammatory cytokine expression via inhibition of the NF- κ B pathway in influenza A virus-infected alveolar epithelial (A549) cells (107). Thus, as described above, R. isatidis contains numerous chemical substances with a broad spectrum of anti-viral and anti-inflammatory activities that may serve as potential therapeutic targets.

In-depth study of the clinical and basic pharmacology of R. isatidis has revealed its broad anti-influenza effects: it is not only a direct antiviral but, more significantly, R. isatidis also inhibits virus-induced inflammation by regulating excessive host immune responses. These pharmacodynamic characteristics are not restricted to R. isatidis; other anti-influenza medicines also exhibit these activities. Therefore, a deeper understanding of new anti-influenza strategies using combined antiviral and anti-inflammatory therapies is required. Particular emphasis should be placed on the pharmacodynamic characteristics of TCM in inhibiting influenza virus-induced host inflammatory and immune responses.

In addition, based on the superior anti-inflammatory efficacy of TCM, some researchers have investigated the treatment of influenza with oseltamivir plus TCMs. Using this approach, the median duration of fever after treatment was significantly shorter in patients who received Mao-to (a Japanese traditional herbal medicine) plus oseltamivir compared with those who received oseltamivir only. Clinical trials have indicated that combination therapy that includes Mao-to is effective in controlling fever owing to type A influenza infection in children (108). Another clinical trial also found that maxingshigan-yingiaosan (a TCM) in combination with oseltamivir obviously reduced the time to fever resolution in patients with H1N1 influenza virus infection, as compared with treatment using oseltamivir alone (94). These clinical trials indicate that TCMs are a good support in oseltamivir treatment that can reduce the duration time of fever. There is also evidence showing that TCM (Lianhua-Qingwen capsules) significantly reduces the severity of illness and duration of symptoms including cough, sore throat, and fatigue (93). However, data are lacking regarding whether treatment with TCMs plus oseltamivir is effective for inhibiting viral shedding or reducing the outbreak rate of viral resistance. These topics warrant further research in the future. Nevertheless, drug interactions must be seriously considered when using TCMs plus oseltamivir.

Conclusions

Influenza viruses, especially avian influenza viruses, still cause significant human disease and can lead to direct or indirect harm via viral replication or hyper-inflammatory immune responses, respectively. Antiviral therapy is particularly beneficial for the treatment of mild influenza or influenza-like pneumonia in adults and children. Although early antiviral use is critical for controlling influenza, delayed use of some NAIs (oseltamivir or peramivir) may also be effective. However, anti-inflammatory treatment of influenza must not be omitted, especially for avian influenza. Therefore, we stress the equal importance of antiviral and anti-inflammatory therapies when developing anti-influenza strategies. Additionally, the use of TCMs for influenza treatment has gained recent attention as TCMs have both antiviral and anti-inflammatory activities and may represent potent new anti-influenza agents.

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Footnote

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