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Review Article



Molecular mechanisms and signaling pathways in-volved in immunopathological events of COVID-19

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ABSTRACT

Introduction: COVID-19, a novel coronavirus that causes severe acute respiratory syndrome (SARS-CoV-2), is currently regarded as the most serious viral disease. During corona infection, viruses bind to host proteins and employ a variety of cellular pathways for their own purposes. Cell signaling is important for the regulation of cellular function. SARS-CoV-2 infection alters multiple signal transduction pathways that are critical for cell survival. The virus causes a severe and prolonged period of hypercytokinemia with misusing of these signaling cascades. Hyperactivation of the host immune system after infection with SARS-CoV-2 is the main cause of death in COVID-19 patients. Thus, to develop effective therapeutic approaches, it is necessary to first understand the problem and the underlying molecular pathways implicated in host immunological function/dysfunction. A number of intracellular signaling cascades have been implicated in infected cell pathways, including MAPK pathway, NF- κ B pathway, JAK–STAT signaling pathway, PI3K/AKT/mTOR pathway and TLRI signaling cascades. Here, we have presented the molecular insights on the potential mechanisms involved in immunopathological events of COVID-19.

Introduction

Infectious diseases such as influenza, acquired immunodeficiency syndrome (AIDS), malaria and meningitis remain the leading causes of death in human populations worldwide (Morens et al., 2004). Humans are infected with a new coronavirus that causes serious pneumonia, which was recognized on 11 2020 by the WHO as coronavirus disease 2019 (COVID-19) (Lai et al., 2020). COVID-19 cause epidemic in all countries and rapidly increasing pandemics move (Gössling et al., 2020). It is not the first outbreak of severe respiratory disease from coronavirus. Coronaviruses have caused three infectious diseases in only the past two decades, namely Middle East respiratory syndrome (MERS), severe acute respiratory syndrome (SARS) and COVID-19 (Rockx et al., 2020; Mahase, 2020).

Severe acute respiratory coronavirus syndrome 2 (SARS-CoV-2) is genetically related to SARS-CoV, the first pandemic threat of a new and fatal coronavirus that appeared at the end of 2002 and triggered an

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SARS-CoV-2 Infection Signal transduction pathways Cytokine storm epidemic of SARS. SARS-CoV was extremely lethal but disappeared due to strong public health control (Petersen et al., 2020). According to a recent report, SARS-CoV-2 and SARS-CoV overlap about 80% of their genes (Gralinski and Menachery, 2020; Xu et al., 2020). Another analysis found a 96% similar sequence between SARS-CoV2 and the isolated CoV from Rhinolophus affinis, suggesting bats as a virus source (Xu et al., 2020). COVID-19 symptoms involve cough, fever, headache and experiencing shortness of breath. Furthermore, most COVID19 patients developed lymphopenia, with markedly elevated concentration of cytokines such as interleukin (IL)-1b and IL-6 (Prompetchara et al., 2020). COVID-19 uses the angiotensin-converting enzyme II (ACE2) as an entry receptor to infect lung alveolar epithelial cells (Velavan and Meyer, 2020). COVID-19 has the capacity to induce symptoms that range from common cold to acute respiratory distress syndrome (ARDS) (Liu et al., 2020b; Zimmermann and Curtis, 2020). In older COVID19 patients with one or more co-morbidities such as hypertension, diabetes





FIGURE 1. Signaling pathways involved in COVID-19 pathophysiology. SARS-CoV-2 down-regulates ACE2 expression and result in production of proinflammatory cytokines and inflammatory mediators. Angiotensin-I is transformed into angiotensin-II by the action of ACE. ACE2 catalyzes Ang-II conversion to Ang (1-7), which promotes anti-inflammatory effects (Gheblawi et al., 2020). Since action of ACE2 is impaired during viral infection, Ang II cause chronic stimulation of AT1R. AT1R signaling induces p38 MAPK activation and inflammatory mediator's production such as TNF and IL-6. The cytokines IL-6 and TNF bind to specific receptors and promote further NF-kB nuclear translocation and phosphorylation of p38 MAPK, which will result in cytokines storm (Feng et al., 2019; Grimes and Grimes, 2020). IL-6 activates JAK/STAT-3 pathway. Toll-like receptors identify SARS-CoV-2 RNA and trigger the inflammatory response through expression of interferon gene and NF-kB pathway (Battagello et al., 2020). PI3K activation lead to Akt phosphorylation and subsequent activation of mTOR. PI3K-Akt-mTOR pathway was also found to regulate cytokine production in COVID-19 (Ramaiah, 2020).

ACE2, Angiotensin-converting enzyme II; Ang, Angiotensin; AT1R, Angiotensin II type 1 receptor; MAPK, Mitogen activated protein kinase; TNF, Tumor necrosis factor; IL, Interleukin; NF-κB, Nuclear factor kappa-B; JAK, Janus kinase; STAT, Signal transducer and activator of transcription; SARS, Severe acute respiratory syndrome; COVID-19, Coronavirus disease 2019; PI3K, Phosphatidylinositol-3-kinase; mTOR, Mammalian target of rapamycin.

Virus	Mechanism of modulation	Implications for therapy	Ref
SARS- CoV-2	MAPK Pathway	Silmitasertib (CK2 inhibitor) Ralimetinib (p38 inhibitor) ARRY-797 (p38 inhibitor)	(Bouhaddou et al., 2020)
		Losmapimod (p38 inhibitor) Dilmapimod (p38 inhibitor)	(Grimes and Grimes, 2020a)
	NF-kB pathway	Artesunate (Inhibitor of NF-kB downregulation) Aspirin (Inhibition of ATP-binding to IKKβ)	(Uzun and Toptas, 2020)
		Sulfasalazine (Inhibitor of the NF-kB activation)	(Elkhodary, 2020)
	JAK-STAT signaling	Ruxolitinib (JAK/STAT pathway inhibitor)	(Bagca and Avci, 2020)
		Baricitinib (a selective JAK1 and JAK2 inhibitor)	(Cingolani et al., 2020b)
	Toll-like Receptor Signaling Pathway	Tocilizumab (IL6 inhibitor) Anakinra (IL1 inhibitor)	(Birra et al., 2020)
	PI3K/AKT/mTOR pathway	Inhalation of buformin or phenformin	(Lehrer, 2020)

TABLE 1: Potential treatment against COVID-19 disease.

Research strategy

The search for scientific papers was performed by researchers in the electronic databases, including Web of Science, Medline (PubMed) and Scopus. The initial search was carried out in the PubMed database based on the combinations of the following words: SARS-CoV-2, MAPK pathway, NF-κB pathway, JAK–STAT signaling, PI3K/AKT/mTOR pathway and TLRI signaling cascades, cytokine storm and immune defenses.

Mitogen activated protein kinase (MAPK) pathway

In response to certain environmental stimuli, MAPK signaling pathways are responsible for controlling several cell functions such as proliferation, differentiation and apoptosis (Cowan and Storey, 2003). The three main MAPK pathways in mammals are MAPK/extracellular signal-regulated kinase (ERK), Jun amino-terminal kinases/stress-activated protein kinases (JNK/SAPK) and p38 MAPK. Pro-inflammatory substances and environmental stimuli primarily activate the p38

MAPK pathway, which has a significant effect on a subset of physiological events such as immune response and inflammatory processes (Deak et al., 1998). Activation of the p38 pathway is required to increase the levels of pro-inflammatory cytokines such as IL-6, tumor necrosis factor (TNF) and IL-1, which appear to play critical roles in the cytokine storm induced by SARS-CoV-2 infection (Catanzaro et al., 2020). Indeed, the excessive immune reaction to COVID-19 infection may be triggered by overly up-regulated p38 activity, as two mechanisms have clarified. Activation of p38 MAPK has been involved in the ACE2 endocytosis (Xiao et al., 2013; Koka et al., 2008; Deshotels et al., 2014). First, the ACE2 activity is lost during SARS-CoV-2 viral entry. ACE2 inhibits related ACE activity by decreasing angiotensin-II and increasing angiotensin 1-7. The stimulation of angiotensin II type 1 receptor (AT1R) by angiotensin II leads to the activation of p38 MAPK and phosphorylation of A disintegrin and metalloprotease 17 (ADAM-17) (Xu and Derynck, 2010; Scott et al., 2011). Phosphorylation increases ADAM17's catalytic

activity, resulting in increased ACE2 shedding and reduced conversion of angiotensin II into angiotensin 1–7, culminating in renin-angiotensin system (RAS)-mediated adverse consequences in a positive feedback cycle (Patel et al., 2014; Xu et al., 2017). Angiotensin 1-7 is critical for suppressing MAPK cascades and reducing inflammation (Zhang et al., 2014).

Angiotensin II promotes proinflammatory, pro-vasoconstrictive and pro-thrombotic activity through p38 MAPK activation, which is reversed by angiotensin 1-7 down-regulation of p38 activity. When ACE2 is lost during viral infection, it may shift the balance towards harmful p38 signaling via angiotensin II (Grimes and Grimes, 2020). ACE2 activity was found in both the lung and the heart. SARS-CoV-2 binds in the respiratory epithelium and lung alveoli to the same ACE2 receptor (Chen et al., 2020; Li et al., 2020). When a virus gets inside a cell, induce ACE2 shedding. Deficiency of ACE2 is associated with alveoli damage and increases permeability of the pulmonary vascular by angiotensin II (Li and De Clercq, 2020). Angiotensin II levels were directly linked with degree of lung injury and viral load in a study of COVID-19 patients, indicating RAS imbalance in COVID-19 etiology (Liu et al., 2020c).

Second, it has been previously shown that SARS-CoV directly up-regulates p38 activity through a viral protein, identical to several other RNA respiratory viruses that can hijack p38 activation to facilitate reproduction. Because SARS-CoV and SARS-CoV-2 are so similar, the latter may use a similar mechanism. As a result, SARS-CoV-2 might cause widespread inflammation by directly activating p38 and down-regulating a crucial inhibitory pathway, all while exploiting p38 activity to reproduce (Grimes and Grimes, 2020).

According to a report, permissive cell SARS- CoV-infection triggered the p38 MAPK signaling pathway. Up-regulation of the p38 MAPK pathway triggers the activation of IL-6, TNF- α and IL-1 pro-inflammatory cytokines (Zarubin and Jiahuai, 2005). MAPK-activated protein kinase-2 (one of the downstream effectors of p38 MAPK) is triggered in response to SARS-CoV-infection in Vero E6 cells (Foltz et al., 1997; Mizutani et al., 2004). Diffuse alveolar damage, which includes significant infection and viral load in type II pneumocytes, and also pulmonary edema, is the most common finding in COVID-19 postmortem tissue from all vital organs (Bradley et al., 2020; Carsana et al., 2020). CT- scans with several ground glass opacities are common and have diagnostic value (Parekh et al., 2020). Angiotensin II levels are particularly high in ACE/angiotensin II receptor blocker naive COVID-19 cases and elevated concentrations are related to greater intensity (Liu et al., 2020a). Immune effector cells release massive amounts of pro-inflammatory cytokines and chemokines, lead to lethal uncontrolled systemic inflammation (Cameron et al., 2008; Channappanavar and Perlman, 2017; Huang et al., 2020a). Furthermore, because some COVID-19 patients have endothelial cell apoptosis, these biological effects could be linked to increased MAPK signaling activity (Grimes and Grimes, 2020; Zhou et al., 2020). The over-activation of p38 MAPK in infected cardiomyocytes, which has been demonstrated to cause apoptosis, impair contractility and promote fibrosis, could be part of the reason for cardiac dysfunction in COVID-19 patients (Grimes and Grimes, 2020). Another pathway involved in SARS-CoV infection is the c-jun NH2-terminal kinase (JNK) pathway, which may result in a rise in proinflammatory factors, as well as increased lung harm (Mizutani, 2010; Liu et al., 2014). JNK signaling pathway could be a target for SARS-CoV-2 as it includes proteins which are similar in both viruses. This mechanism induces ACE2 receptor binding and opened the way for COVID-19 virus internalization into the respiratory tract's alveolar epithelial cells. JNK signaling is implicated in the extrinsic and intrinsic apoptotic pathway, tissue cytokine production, inflammation and metabolism (Vellingiri et al., 2020).

Although the role of RAS in the pathophysiology of SARS-CoV-2 is still being explored, a recent study indicated that blocking RAS with ACE inhibitors or angiotensin receptor blockers may reduce overall mortality in COVID-19 patients (de Abajo et al., 2021). The specialized viral entrance mechanism of SARS-CoV-2 deactivates a key counterbalancing mechanism that the cell employs to reduce p38 signaling through ACE2 activation, which causes inflammation while also extending the viral lifespan. As a result, SARS-CoV-2 may cause excessive inflammation by directly activating p38 and downregulating a key inhibitory pathway, while also exploiting p38 activity to proliferate. COVID-19 infection could be reduced if p38 is suppressed medically. Losmapimod is the most researched p38 inhibitor in clinical trials and it has a good efficacy. Therefore, p38 MAPK inhibitor could be beneficial in patients with serious COVID-19 health problems (Grimes and Grimes, 2020).

Nuclear factor kappa-B (NF-κB) signaling pathway

The NF-kB signaling pathway regulates a variety of essential genes in the innate and adaptive immune systems (Hoesel and Schmid, 2013; Liu et al., 2017). NF-kB signaling pathway plays a key role in gene expression involving cytokine/chemokine encoding and anti-apoptotic genes (Tak and Firestein, 2001; Gupta, 2003). NF-kB and its inhibitor (the inhibitory kappa B kinases, IkB) are present as a complex. Release from this complex requires IkB kinase (IKK) activation. The kinase complex of the IKK is the central element of the cascade of the NF-κB. Essentially, it consists of two kinases (IKKα and IKKB) and a regulatory sub-unit, NEMO (NF-kB essential modulator) /IKKy (Bonizzi and Karin, 2004). In most unstimulated cells, NF-kB dimers are kept inactively in the cytosol by interacting with IkB proteins (Oeckinghaus and Ghosh, 2009). After activation, the IKK complex will induce the phosphorylation of the IkB proteins leading to their degradation (Viatour et al., 2005). The degradation of these inhibitors by the IKK complex upon their phosphorylation resulting in the nuclear translocation of NF-kB and the induction of target gene transcription (Magnani et al., 2000). In other types of cell, including mature B cells, macrophages as well as a significant number of tumor cells, NF-kB may also be recognized as a nuclear protein which is constitutively active (Oeckinghaus and Ghosh, 2009).

During a virus infection, the NF-kB signaling pathway is activated and the gene expression of interferon beta (IFN- β)/ TNF α / IL8 are increased (Pfeffer, 2011; Liu et al., 2017), suggesting that IKK-mediated NF-kB signaling is necessary for the host's innate immune response (Banoth et al., 2015). In the Vero E6 cells, fulllength N protein considerably enhances NF-kB activity. In addition, T helper cells develop proinflammatory cytokines by NF-κB signaling (Liao et al., 2005). NF-κB activation is a characteristic of most infections, including those caused by viruses, which lead to defensive and pathological reactions. Following mice infection with rSARS-CoV-MA15, increased expression of inflammatory cytokine TNF, C-C motif (CC) chemokines [CC chemokine ligand (CCL) 2, CCL5], C-X-C motif (CXC) chemokines [CXC chemokine ligand CXCL1, CXCL2, and CXCL10], and IL-6 were found in neutrophils and infected lungs. Elevated levels of IL-6 and chemokines such CCL2 and CXCL10 have also been found in human lungs with fatal SARS (Jiang et al., 2005; Tang et al., 2005; Cameron et al., 2007). Researchers recently investigated the regulatory relation between the protein SARS-CoV-2 mediated pro-inflammatory cytokine/chemokine response and the NFκB signaling pathway (Huang et al., 2020; Ingraham et al., 2020; Islam and Fischer, 2020; Neufeldt et al., 2020; Rian et al., 2021). Huang et al. (2020b) showed

flammatory cytokine/chemokine response and the NFκB signaling pathway (Huang et al., 2020b; Ingraham et al., 2020; Islam and Fischer, 2020; Neufeldt et al., 2020; Rian et al., 2021). Huang et al. (2020b) showed that a significant transcriptomic transition in infected cells, characterized by a change to an inflammatory phenotype with activation of NF-kB signaling and NF-kB target genes by day 1 post-infection, leads to the loss of the mature alveolar program in a human in vitro model that simulates the initial apical infection of alveolar epithelium with SARS-CoV-2, leads to a loss of the mature alveolar program. Differentially expressed genes are enriched for components of pathways related to NF- κ B, TNF- α and IL-17 signaling in bronchial epithelial cells infected with SARS-CoV-2 (Enes and Pir, 2020). Elements in the ACE2 gene regulate pirin, a negative regulator of the NF-kB subunit RELA (p65). Pirin expression is thought to be reduced when SARS-CoV-2 disrupts ACE2 (Fadason et al., 2020). Furthermore, in human bronchial epithelial cells, SARS-CoV-2 spike protein subunit 1 (CoV2-S1) caused high rates of NFκB activation, the development of pro-inflammatory cytokines and chemokines including IL-1, TNF, IL-6 and CCL2, as well as mild epithelial damage. S1 interaction with the human ACE2 receptor, as well as early activation of the endoplasmic reticulum stress, subsequent unfolded protein response and MAP kinase signaling pathways, were all necessary for CoV2-S1-induced NFкВ activation. CoV-2-S1 had a higher NF-кВ activation than CoV-S1, which may be attributed to CoV-2-S1's higher affinity for the ACE2 receptor (Hsu et al., 2020). Previous research has shown that an elevated cytokine/ chemokine response during extreme SARS infection indicates a dysregulated immune response. In vivo, IL-6 is the primary stimulator of signal transducers and activators of transcription (STAT-3), and STAT3 is needed for complete NF-kB pathway activation, particularly during inflammation (Hirano and Murakami, 2020; Murakami et al., 2019). Both NF-kB and STAT-3 are triggered as a result of SARS-CoV-2 infection in the respiratory system, resulting in activation of the IL-6 amplifier, a mechanism for STAT-3 hyperactivation of NF- κ B, leading to a variety of inflammatory and autoimmune diseases (Murakami et al., 2019). Moreover, previous study reported that thalidomide as an immunomodulatory agent modulates the NF- κ B activities in combination with celecoxib (the cyclooxygenase-2 inhibitor) which can restrict the symptoms of inflammation if used to treat severe pneumonia (Hada, 2020). Since immunomodulatory drugs can affect the cytokine storm, these drugs may be effective in treating COVID-19. Immunomodulation of NF- κ B activity and inhibitors of NF- κ B (I κ B) degradation, in combination with TNF- α inhibition may reduce the cytokine storm and lessen the severity of COVID-19. Inhibition of NF- κ B pathway may be useful in treating COVID-19 in its most severe form.

Many of the drugs appear to have binds to the NF- κ B cascade of immune regulation in COVID-19. Dexamethasone is one of two glucocorticoids (the other being prednisolone) that has an inhibitory effect on the NF- κ B pathway (Ye et al., 2020; D'Acquisto et al., 2002). Remdesivir (GS-5734) is a nucleotide analogue that inhibits the RNA dependent RNA polymerase, causing viral replication to be disrupted. It decreases the cytokine storm and severe illness by lowering dsRNA-related activation of the NF- κ B pathway. Remdesivir patients had a faster time to recover in the Adaptive COVID-19 Treatment Trial, which compared to a placebo (Beigel et al., 2020). TNF- α , TNF-1 β , IgG and IFN- γ are all reduced by hydroxychloroquine, which suppresses the NF- κ B pathway (Liang et al., 2018).

Janus kinase (JAK)–STAT pathway

The JAK-STAT pathway signaling mechanism, may be a valuable marker of a strong immune response to COVID-19 infections (Bouwman et al., 2020). According to one study, SARS-CoV-2 triggers the biochemical mechanisms mediated by JAK–STAT in the lungs, leading in viral cell proliferation and transmitting (Singh et al., 2020). In another study, inhibiting the JAK-STAT pathway reduced hyperinflammatory conditions while having no effect on viral clearance (Rojas and Sarmiento, 2021). The JAK-STAT pathway is also activated by IL-6 (Billing et al., 2019). The finding demonstrates that induction of the JAK-STAT pathway, particularly through cytokines such IL-6, is associated with the inflammatory response to COVID-19 (Luo et al., 2020b). Angiotensin II binds to the AT1R and activates the JAK- STAT pathway, leading to the production of IL-6 (Ni et al., 2020). The SARS-CoV-2 S protein inhibits ACE2, causing an increase in angiotensin II expression and, as a result, enhanced IL-6 production. Anti-inflammatory drugs, in particular JAK-STAT inhibitors may be useful against increased cytokine levels and may be effective to prevent viral infection. Ruxolitinib is a JAK1 and JAK2 inhibitor that suppresses STAT activation and nuclear translocation by blocking JAK kinase activity. Ruxolitinib also suppresses the IL6/JAK-STAT3 pathway, decreasing IL-6 levels in the blood (Caocci and La Nasa, 2020; Kusoglu et al., 2020). The role of baricitinib (a specific JAK1 and JAK2 inhibitor) in the treatment of COVID-19 has been proposed, despite its true safety profile has yet to be determined (Cingolani et al., 2020a).

Toll-like receptor (TLR) signaling pathway

The TLRs are important in the innate immunity by detecting microbes to invade pathogens. TLR signaling pathways are the recruitment of different adaptor molecules resulting in the activation of NF-kB and the IFN regulatory factor transcription factors dictating the outcome of TLR's innate immune responses (Barton and Medzhitov, 2003). While the immune system's effective functioning protects the body from infections, the cytokine storm associated with extreme COVID-19 manifestations is mainly caused by the adaptive immune system's over-expression and exhaustion, rather than an innate immune response (Coperchini et al., 2020). The virus's spread is limited by the host immune response during infection or mild COVID-19 disease, but the innate immune response may also trigger immune-related dysfunction, resulting in extreme pneumonia in cases of high viral load (Soraya and Urmia, 2020). In viral diseases, TLR activators have both defensive and therapeutic effects. The study also discovered that the SARS-CoV-2 spike protein binds to TLR1, 4 and 6 with a higher affinity for TLR4 than the others (Khadke et al., 2020). A recent study offers that TLRs may be involved in both the initial viral clearance failure and the subsequent production of the deadly clinical manifestations of severe COVID-19 primarily ARDS. Lung macrophages can play a critical role in massive release of IL-6 and other cytokines such as IL-1 β , IL-10, IL-12 and TNF-α via activation of TLRs in patients with severe COVID-19 (Onofrio et al., 2020).

In addition to the development of proinflammato-

ry cytokines, TLRs' interaction with virus particles has immunopathological effects that lead to death in COVID19 patients (Patra et al., 2020). TLR4's pathologic role in patients with an excessive inflammatory response has been documented in other SARS-CoV-2 studies. COVID-19 patients had substantially higher levels of CCL 2, CCL7, CCL8, CCL24, CCL20, CCL13, CCL3, CXCL 2, CXCL10 and IL-1b, and its down-stream inflammatory signaling molecules (IL1R1, Myeloid differentiation primary response [MYD88], interleukin 1 receptor associated kinase 1 [1IRAK1], TNF receptor associated factor [TRAF6], NF-KBIA, NF-KB1, RELA). TLR4 and related/down-stream signaling molecules (CD14, MYD88, IRAK1, TRAF6, TIRAP, TICAM) as well as most NF-κB signaling pathway genes (NF-KBIA, NF-KB1, RELA, NF-KB2) were also highly up-regulated, implying that activation of the NFκB signaling pathway by TLR4 is thought to be responsible for the up-regulation of inflammatory responses in COVID-19 infection patients (Sohn et al., 2020). Furthermore, COVID19 patients have a higher level of neutrophil myeloperoxidase, which triggers oxidized phospholipids and TLR4 pathway activation causes oxidative injury during the pulmonary process of infection (Khadke et al., 2020; Onofrio et al., 2020). Tocilizumab, an anti-IL-6 monoclonal antibody is used to treat rheumatoid arthritis, may be useful in the treatment of critically ill patients with COVID-19 (Kaly and Rosner, 2012). Findings support the use of therapeutic approaches such as dexamethasone that inhibits TLR4-mediated inflammatory signaling through molecular checkpoints (Sohn et al., 2020).

Phosphatidylinositol-3-kinase (PI3K)/ AKT/ mammalian target of rapamycin (mTOR) pathway

The PI3K/AKT/ mTOR signaling pathways is an important intracellular signaling pathway in the regulation of the cell cycle and cell growth. Therefore, it is specifically associated with cellular proliferation, quiescence and survival. The plasma membrane protein AKT is phosphorylated and activated when PI3K is activated (King et al., 2015). Insulin-like growth factor, epidermal growth factor, sonic hedgehog signaling molecule insulin and CaM can enhance the PI3K / AKT pathway (Man et al., 2003; Peltier et al., 2007; Ojeda et al., 2011; Rafalski and Brunet, 2011). The mTOR signaling pathway modulates protein synthesis in response to stress, hor-

mones and genetic factors. Rapamycin inhibits mTOR by interfering with the PI3K/AKT/mTOR pathway and activating AMP-activated protein kinase (Huang, 2013). MTOR signaling is required for influenza development and regulates the antibody response, resulting in cross-protective immunity against lethal influenza virus infections. Treatment of serious pneumonia caused by H1N1 influenza with rapamycin and steroids has been shown to enhance reporting outcomes in human studies (Chuang et al., 2014; Wang et al., 2014; Lehrer, 2020). The PI3K/AKT/mTOR signaling responses have a key role in MERS-CoV infection which making it a target for therapeutic intervention. Buformin or phenformin (mTOR inhibitor) inhalation may be an effective novel treatment for coronavirus (Lehrer, 2020). Cytokine storms are the main reason of COVID-19-related serious illness and death. The most significant cause of cytokine storms can be the antibody-dependent enhancement. mTOR inhibitors may suppress antibody-dependent enhancement and decrease the severity of COVID19 by selectively inhibiting memory-B cell activation (Zheng et al., 2020).

The mTOR-PI3K-AKT pathway was identified as a key signaling pathway in SARSCoV2 infection in a recent report. The in vitro testing of three mTOR inhibitors showed that they significantly inhibited SARSCoV2 (Garcia Jr et al., 2020). Regarding to recent reports, activation of the PI3K/ AKT/ mTOR pathway appears to be important to promote replication of different viruses and drugs that inhibit PI3K/ AKT/ mTOR signaling pathways may be recommended for SARS-CoV-2 infection. In order to identify potential drug targets, a human protein-protein interaction map for SARSCoV2 was recently developed. The proposed drugs included the mTOR inhibitors rapamycin and sapanisertib, as well as the mTORC1 protein complex modulator metformin. Metformin-treated COVID19 patients have been shown to have a lower mortality rate (Bramante et al., 2020; Cariou et al., 2020; Luo et al., 2020a).

Inflammatory cytokines can be a double-edged sword when it comes to viral infection and disease pathogenesis. To battle viral infection and avoid a cytokine storm, the innate immune system must be fine-tuned (Säemann et al., 2009). As a result, clinical trials should include early and short-term intervention with mTOR inhibitors to reduce the undesirable immunosuppressive effect. Furthermore, IL-6 may play a crucial role in the cytokine storm's substantial negative consequences and IL-6 inhibition has been used to treat severe COVID19 disease with respiratory distress (Zheng et al., 2020). In addition to mTOR inhibitors, combination therapy with an anti-IL6 antibody could be included in the clinical trial for patients suffering SARS-CoV2 pneumonia (Zheng et al., 2020).

Conclusion

Infection with SARS-CoV-2 changes multiple signal transduction pathways, which contribute to important physiological functions of the cell. The balance of signaling pathway activities is important for cell death, or cell survival determination. The virus takes over mechanisms from the host cell to utilize it for its own benefit. SARS-CoV-2 involved MAPK signaling pathway, NF-kB pathway, PI3K/ AKT/ mTOR pathway, JAK-STAT pathway and toll-like receptors cascades through different mechanisms. In certain infected individuals, SARS-CoV-2 induces excessive and prolonged cytokine/chemokine responses. ARDS, or multi-organ dysfunction, is caused by the cytokine storm and it leads to physiological deterioration and death. The virus manipulates these signaling pathways for inhibiting cytokine antiviral effects.

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Conflict of interest

The authors declare no conflict of interest.

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