Cellular Physiology and Biochemistry Published online: 20 December 2022

Cell Physiol Biochem 2022;56:707-729 DOI: 10.33594/000000595

Accepted: 30 October 2022

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Review

Possible Therapeutic Targets from Derivatives of Natural Marine Products Based on PI3K/AKT Dependent Inhibitors in Viral Infection COVID-19

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Key Words

Natural marine products • Viral infection • COVID-19 • PI3K/AKT • ACE2 • Medicinal purposes

Abstract

Natural resources have long played a prominent part in conventional treatments as a parental source due to their multifaceted functions and lesser side effects. The diversity of marine products is a significant source of possible bioactive chemical compounds with a wide range of potential medicinal applications. Marine organisms produce natural compounds and new drugs with unique properties are produced from these compounds. A lot of bioactive compounds with medicinal properties are extracted from marine invertebrates, including Peptides, Alkaloids, Terpenoids, Steroids. Thus, it can be concluded that marine ecosystems are endowed with natural resources that have a wide range of medicinal properties, and it is important to examine the therapeutic and pharmacological capabilities of these molecules. So, finding particular inhibitors of the COVID-19 in natural compounds will be extremely important. Natural ingredients, in this light, could be a valuable resource in the progression of 707

Cellular Physiology and Biochemistry Published online: 20 December 2022 Cell Physiol Biochem Press GmbH&Co. KG

Cell Physiol Biochem 2022;56:707-729

DOI: 10.33594/000000595 © 2022 The Author(s). Published by

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COVID-19 therapeutic options. Controlling the immunological response in COVID-19 patients may be possible by addressing the PI3K/Akt pathway and regulating T cell responses. T cell effector activity can be improved by preventing anti-viral exhaustion by suppressing PI3K and Akt during the early anti-viral response. The diversity of marine life is a significant supply of potentially bioactive chemical compounds with a broad range of medicinal uses. In this study, some biologically active compounds from marine organisms capable of inhibiting PI3K/AKT and the possible therapeutic targets from these compounds in viral infection COVID-19 have been addressed.

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Introduction

Due to the accelerating spread of the COVID-19 pandemic and the considerable time required to develop vaccines and effective drugs, a pressing need exists to develop strategies for preventing and treating COVID-19 using the already existing natural products [1-3]. Numerous natural products have been widely studied and used as potential options for the treatment of COVID-19 [4-6]. It seems that the endosomal-autophagic vesicles networks play a key role in the infections caused by Coronaviruses (CoVs), including SARS-CoVs [7, 8]. Autophagy plays an important role in pulmonary infections, improving the host's immune defenses against bacterial and viral infections of the respiratory tract. Considering the various roles of autophagy in viral infection, it was postulated that three groups of autophagy modulators might inhibit viral replication clinically related to COVID-19 [9, 10]. The first group is the drugs with lysosomotropic activity, which inhibit cathepsin activity, preventing infections caused by CoVs by neutralizing the acidic pH in the endosomes-lysosomes [8]. The second group of drugs includes protease inhibitors, which might inhibit the S protein's proteolytic cleavage and limit viral cell entry. The third group of drugs includes PI3K/AKT/ mTOR regulators, which might mean autophagic machinery's CoVs-mediated appropriation, despite being considered autophagy regulators [11]. In the next subsections, several welltolerated clinically approved autophagy-modulating compounds will be discussed, which might be investigated as potential COVID-19 modulators and replication used in COVID-19 management. Currently, there is not any acceptable antiviral drug to treat SARS-CoV-2 infections. Hence, preventive and supportive cares of complications are essential strategies to minimize the harms [11, 12]. Suggested strategies for the treatment of SARS-CoV-2 might be divided into four different categories: SARS-CoV-2 RNA synthesis prevention, through vital proteins and enzymes inhibition, virus-cell receptor binding impedition or virus selfassembly suppletion, the host's immunity stimulation by producing virulence factors, and virus entry blockage into host cells by acting on the host's enzymes or through cellular entry receptors [13-16]. Alkaloids, macrolides, polypeptides, and terpenoids are just a few of the molecular classes represented by marine-derived natural products and contain a variety of unique and fascinating structures that contribute significantly to biological activity and clinical therapeutic uses. In terms of kinase research, several marine-derived kinase inhibitors have been developed from a variety of various sources and have been shown to inhibit a wide range of protein kinases. Recently, several biologically active kinase inhibitors have been introduced in different ocean life forms, such as soft corals, algae, fungi, bacteria, cyanobacteria, sponges, and animals. Basic bioactivity tests are usually included in the initial reports of these natural agents, and subsequent researches should go on to evaluated the pharmacological activity of each in greater depth [17-20].

Lipid Kinase Inhibitors

Phosphoinositide 3-Kinase (PI3K) Inhibitors

PI3K, belonging to a large family of lipid kinases, has been proven to play a significant role in cancer, aging, and diabetes. PI3Ks have a "PI3K characteristic motif", composed of

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a C2 domain (which may bind to the membrane), a catalytic kinase domain, and a helical domain [21, 22]. There are three types of PI3K (I, II, III) and four isoforms (α , β , γ , δ). Class I (I PI3K) has been implicated in various cancers [22, 23].

Class I PI3Ks. Class I PI3K is a heterogeneous mer, including coordination ($p50\alpha$, $p55\alpha$, p55y, p85 α , p85 β , or p101) subunits with catalysts (p110 α , p110 β , p110 δ , or p110 γ). The known carcinogenic PI3K/mTOR signaling has been explored elsewhere [24]. In outline, this pathway is actuated by cell surface receptors, such as oncogenes like RAS, G protein-coupled receptors, and receptor tyrosine kinases. Furthermore, the activated p110 subunit can catalyze converting phosphatidylinositol 4,5-bisphosphate (PIP2) to phosphatidylinositol 3,4,5-trisphosphate (PIP3), activating protein kinase B/AKT. AKT usually sends signals to downstream effectors, such as glycogen synthase kinase 3, forkhead transcription factor, mTOR complex (mTORC) 1, and cell death antagonist BCL2 to regulate different cellular processes. mTORC2 also promotes AKT activation through the serine 473 phosphorylation. Contrasting to this, the role of phosphotensin homolog (PTEN) is antagonizing the PI3K function and dephosphorylating PIP3 to PIP2 [24, 25]. Many growth factor pathways are controlled by activated RTK or GPCR. These RTKs or GPCRs recruit the p85p110 complex to the plasma membrane to alleviate the inhibition of p85, and p110 converts PtdIns (4.5) P2 to PtdIns (3,4,5) P3 to trigger a signal conduction response [25, 26]. In particular, PtdIns (3,4,5) P3 recruits Akt kinase, which controls the activation of FOXO, mTORC1, etc. to regulate metabolism, autophagy, survival, cell growth, and proliferation [27, 28]. Restricted class I PI3K movement additionally assumes a part in cortical F-actin elements, which causes chemotaxis and phagocytosis of particles [29-32]. For instance, at the neutrophil driving edge, p110γ-incited PtdIns (3,4,5) P3 development brings about the enrollment of Rac GTPase, which advances F-actin polymerization, lamellipodia arrangement, and cell relocation [33]. 3phosphatase PTEN downregulates PtdIns (3,4,5) [3-5] P3 and class I PI3K activation pathways [34]. Type 4 I catalyst isomorphs are nested but share distinct functions. While p110 γ and p110 δ isomorphs are primarily restricted to functions expressed in immune cells, p110 α and p110 β also exhibit isomorphic-specific cell types and context-dependent requirements. Most Class I PI3K functions are associated with catalytic properties. However, there is increasing evidence of the role of kinase-independent scaffolding of $p110\gamma$ and p110β [35-37].

Class II PI3Ks. Class II PI3K includes Class I and Class III and other high molecular weight monomers with long N- and C-terminal domains [38]. The magnificence II enzymes lack structural records that could assist in recognizing those mechanisms. While the magnificence II enzymes include a PI3K center and a Ras-binding area (RBD) like different individuals of the PI3K family, all magnificence II enzymes include C-terminal domain names, along with a phox-homology (PX) area observed with the aid of using a C2 area. A prolonged N-terminal area is particular to magnificence II. Unlike their magnificence I and III spouse and children, which can be focused on their web website online of movement through special regulatory subunits, magnificence II enzymes lack solid affiliation with regulatory subunits. Instead, those particular domain names at each terminus are idea to play an essential position inside the law and localization of those kinases with the aid of using mediating particular protein and lipid interactions [39]. In mammals, 3 totally different category II members are identified: the ubiquitously expressed PI3K-C2 α , the liver-specific PI3K-C2 γ , and PI3KC2 β [38, 40]. Contrasting to class I, class II PI3Ks are primarily recognized as substrates PtdIns and PtdIns (4)P, which are involved in the regulation of follicular transport, resulting in PtdIns (3) P and PtdIns (3,4) P2, respectively. Several stimuli, including cytokines, chemokines, growth factors, and hormones can activate class II PI3Ks through various membrane receptors, such as GPCRs and RTKs (EGFR and PDGFR) [38-41]. The best studied in class IIPI3K is PI3KC2a. At the cellular level, it is demonstrated that PI3KC2 α regulates the complete translocation of the glucose transporter GLUT4 to the plasma membrane of muscle cells, an important event in the regulation of glucose homeostasis [41].

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Class III PI3Ks. PI3K is a lipid kinase that can phosphorylate phosphatidylinositol (PI) and its phosphorylated derivatives phosphatidylinositol 4,5 diphosphate (PIP2) and phosphatidylinositol 4 The D3 hydroxyl group of monophositol (PI4 ring). PI3K accepts entry from activated receptor tyrosine kinases and heterotrimeric guanine nucleotide (G protein) -binding protein-coupled receptors (GPCRs), and affects cell proliferation, growth, metabolism, motility, and second messenger transport of intracellular lipids [42]. In recent years, mammalian class III PI3K (phosphatidylinositol 3 kinase) complex has become the key to several basic cellular processes through the aftereffect of its catalytic product phosphatidylinositol 3-phosphate (PtdIns (3) P) Modifier. One of these processes is the ligand-dependent down-regulation of growth factor receptors through membrane transport events. Specifically, PtdIns (3) P mediates the load classification from early endosomes to multivesicular bodies and finally degraded lysosomes [43]. Inhibition of mTOR results in the activation of the important molecule Unc51like autophagy effective kinase 1 (ULK1). which is potentiated into the endoplasmic reticulum (ER) for the initiation of autophagy. ULK1 is a crescent-shaped bilayer structure known for omega cotton that emerged from the ER by recruiting additional III-type four-spoy city Ted 3 kinase (PI3K) complexes composed of Beclin1, VPS34, and ATG14. Induces formation and macrophage nucleation. The class III PI3K complex mediates the accumulation of phosphatidylinositol triphosphate (PIP3) on the surface of omega cotton to recruit LC3, and phosphatidylethanolamine (PE) is conjugated to LC3I to form LC3II, which forms a phagocytic membrane. Phagocytes eventually seal, form self-predatory bodies, and then fuse with lysosomes to break down the packaged contents. Apart from recycling useful substances, self-predation is also an important mechanism for removing intracellular pathogens, including viruses, which is called virophage [43-45].

Immune system and activation of PI3K

Unique PI3K lineage members are active in the immune system, depending on the type of receptor and/or cell [41, 46-48]. For instance, cytokines like interleukin (IL)-2, IL-3, IL-6, IL-7, and IL-15, interferons (IFNs), oncostatin M, erythropoietin, granulocyte colonystimulating factor activate class IA PI3Ks in a number of immune cells, such as dendritic cells (DCs) and T cells, by activating Janus kinase and tyrosine phosphorylation of different proteins. Cytokine receptors with intrinsic PTK activity can also activate class IA PI3Ks. These enzymes are further activated on antigen recognition through natural killer (NK) stimulatory receptors, the B cell receptor (BCR), T cell receptor (TCR), and Fc receptors, the same a IgG ($Fc\gamma RI$) and high-affinity IgE ($Fc\epsilon RI$) receptors [49-52]. Co-stimulatory receptors such as CD28 and CD19 on T cells and B cells, respectively, and cell adhesion molecules also activate class IAPI3Ks. The cytoplasmic region of CD28 and CD19 binds to the p85 subunit, so binding of both antigen receptors and costimulatory receptors potently activates PI3K [52-54]. Various adapter proteins including BCAP and linkers for T cell (LAT) activation that mediate protein-protein interactions are involved in the activation of IAPI3K, a downstream class of antigen and costimulatory receptors, possibly recruited to the plasma membrane of tyrosine phosphorylation [53-55]. In particular, the tumor necrosis factor receptor family, such as the IL1 receptor Toll-like receptor (TLR) and CD40, activate class IAPI3Ks in many cell types, including macrophages and DCs, despite no apparent association with PTK [41, 46, 53, 54]. It is not yet clear how signal transduction through these receptors enables downstream PI3K. Unlike class IA PI3Ks, class IB PI3Ks are mostly activated via GPCRs like chemokine receptors [56-59]. Class II and class III PI3Ks are likely to play a role in the immune system, but the mechanisms supporting their activation remain undetermined. Giant predation, the process by which long-lived proteins and intracellular components such as organelles are broken down and recycled, is at least partially regulated by the PI3KAKT path [58-60]. I It can be additionally induced by activating kinase Ulk1 throu AMP-activated protein kinase (AMPK). Activating the PI3KAKT mTORC1 path suppresses self-predation through suppression of the interaction between AMPK and Ulk1 by mTORC1 [61]. Auto-

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purging with respect to viral infections has been well-studied, but it is not necessarily clear whether the effectiveness of the route is pro-virus or anti-virus [62, 63]. The advantages of autophagy appear to be distinct for distinct viruses, depending on the time after infection. Some viruses, inclusive of rotavirus and ZIKV, require the manner early in infection [27, 61-63], while that is unfavorable for others. Conversely, a few viruses, including influenza A virus result in autophagy overdue after contamination to boom replication. Finally, there are viruses wherein autophagy does now no longer seem to steer replication. Autophagy has been studied withinside the context of CHIKV, SFV, and SINV infection, aleven though maximum of those research did now no longer check out the direct position of PI3K-AKT activation. These alphaviruses seem to have special consequences on autophagy, aleven though there additionally is probably version because of special mobileular sorts used and special experimental conditions [63-65].

Akt

Akt, a serine/threonine kinase, which was previously known as protein kinase B (PKB), is composed of Akt1, Akt2, and Akt3 that regulate glucose metabolism, cell cycle progression and cell size. Akt plays an important role in key cellular functions, including neovascularization, protein synthesis, genome, stability, and transcription. Akt facilitates cell survival by blocking apoptosis through the ineffectiveness of pro-apoptosis proteins, which mediates cell growth factors [66-70]. Akt/PKB is similar to supermolecule kinase A (PKA) and kinases C (PKC), further on the retroviral oncoprotein microorganism akt (v akt) [71-74]. Akt consists of a central, an amino terminus (N terminus), and a carboxyl terminus fragment (C terminus). The purecustorin homology (PH) domain (N-terminal domain), consisting of 100 amino acids, is similar to other domains seen in 3-hoshoinoshichido binding molecules that interact with membrane lipid products such as phosphatidylinositol triphosphate (PIP3) and phosphatidylinositol 4,5-diphosphate (PIP2) [68-70]. The kinase domain, being similar to the AGC protein kinase, shares a regulatory residue named Thr308, which its phosphorylation activates Akt. The C-terminal home is composed of forty amino acids that form a hydrophobic region containing the regulatory serine residue Ser473 [71, 72].

PI3K/AKT (Structure and function)

Akt kinase hobby is precipitated following PI3K activation in numerous boom thing receptor-mediated signaling cascades [75]. PI3K phosphorylates hoshoinoschido at the 300H position of the inositol ring. The secondary messenger products of the kinase response in growth factor-stimulated animal cells are phosphatidylinositol 3,4 phosphate (PI3, 4P2) and phosphatidylinositol 3,4,5 phosphate (PIP3) [76]. The discovery of phosphoinositidespecific phospholipid phosphatases, such as PTEN, further enhances our knowledge due to the regulation of secondary messenger signals by phospholipids and cellular phosphoinositide metabolism. Activation of Akt by PI3K is initiated through binding specific 30 phosphorylated Force Spoi City Ted in the Akt PH domain [77]. The Akt PH area binds to each PIP3, and PI-3,4-P2 and suggests a fairly better affinity for the binding of PI-3,4-P2. The binding specificities of various PH domain names for one of a kind phosphoinositides shown with the aid of using structural studies, and the specificity of the Akt PH area for each PI-3,4-P2 and PIP3 can be applicable for accomplishing a prolonged sign of Akt pastime in cells [76]. The initial binding result of plasma-bound Force Spoiled Ted Akt is the reorganization of the cellular Akt protein of the plasma transduction complex. The PH domain is important for mediating these reorganizations, consistent with its ability to bind to phospholipid second messenger molecules. The discovery of increased activity of the N-terminal deleted Akt mutation was interpreted as an indication of intramolecular inhibition, but only recent evidence suggested an interaction of the N- and C-terminal regions of the Akt molecule [78]. Oncogenic mutations in Akt, which would result in constitutive factor-independent activation, also account for activity in the vakt configuration by involving the plasma membrane binding

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to the N-terminal myristoyl, thereby circumventing the initial translocation step [79, 80]. Phosphorylation with the aid of using upstream kinases is needed for complete activation of Akt and is essential for protection of its activity. Therefore, it isn't always sudden that a few protein phosphatases are capable of inactivate Akt with the aid of using dephosphorylation. Nonspecific inhibition of endogenous phosphatase hobby turns on Akt efficiently [81]. Furthermore, phosphatase activity is concerned in mediating the results of extracellular stresses on Akt sign transduction [80]. Akt dephosphorylation was also observed after increasing ceramide levels, possibly due to inhibition of PI3K or other upstream signaling molecules [82]. Information on the importance of phosphatases, which dephosphorylated the phosphoinositide products of PI3K. Among these, the homology of the src2 (SH2) domain containing the inositol SHIP1/2 phosphatases and the PTEN phosphatase is important mediators in the determination of Akt activity. SHIP1 dephosphorylates inositides and phosphoinositides at position 50 and regulates B cell / myeloid function [82-84].

PI3K signaling activation

In normal situation, the catalytic subunit (p110) is synthetized through dimerization with the regulatory subunit (p85) and different extracellular stimuli, including cytokines, hormones, and growth factors activate PI3K [85]. When activated, PI3K phosphorylates PtdIns (4,5) P2(PIP2) to make PtdIns (3,4,5) P3(PIP3) and recruits a subset of pleckstrinhomology (PH), such as Phox (PX), FYVE, C1, C2 to the cell membrane. Several signaling proteins, including kinases PDK1 and AKT, may localize to the cell membrane by binding to the PI3K lipid products to activate cell growth and survival [86]. The pathway is normally regulated through dephosphorylating PIP3 to PIP2 and preventing the downstream kinases' activation [87].

PI3K and SARS-CoV-2

PI3K/AKT signaling plays a crucial role in migration, invasion, cell proliferation, growth, and survival, and can promote angiogenesis and subset cell apoptosis. Abnormal PI3K/AKT signaling pathway has been shown to cause health problems and diseases such as cancer. With life science advancements, targeted therapy such as PI3K/AKT inhibition has become a popular method for the treatment of malignant tumors. AKT and PI3K are antitumor drug targets, and their antitumor therapies have shown attractive prospects. A potent panAKT kinase inhibitor, Capivasertib, which inhibitor, Capivasertib is used in clinical trials for the treatment of drug-resistant breast cancers [88, 89]. The kinase inhibitor capivasertib is also an anticancer drug that targets AKT preventing SARSCoV2 from entering the cell. There are a lot of antitumor drugs that target the PI3K/AKT signaling pathway. This might be used in treating critical COVID19 cancer patients in the era of the pandemic [89-91].

(PI3K)/AKT and development of immune responses

The development of immune responses involves entering the virus into the cell and immune response development. The relationship between virus entery and this pathway should be examined. The main receptors involved in virus entery are and CD147 and ACE2 [91-93]. Furin and TMPRSS2 are two other non-endosomal pathways for viral entry [93]. Furin and CD147activation can induce the PI3K/AKT signaling pathway [93, 94]. Endocytosis occurs when the virus binds to ACE2. The occurrence of SARS-CoV-2 endocytosis is regulated by the PI3K/AKT signaling pathway and is mediated through a clathrin pathway [93-95].

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Suppression of this pathway can inhibit the entry of any clathrin-mediated endocytosis. ACE2 reduction at the cell surface enhances the serum levels of Ang II, which is seen in COVID-19 patients [96-99]. Ang II elevation in serum can enhance the inflammatory cytokines such as TNF- α and IL-6, as seen in COVID-19 patients. Ang II also induces the fibrosis of various organs, which can be attributed to the PI3K/AKT signaling pathway [97, 99, 100]. The binding of Ang II to ang II receptor type 1 (AT1R) can activate this pathway [101]. Activation of PI3K/AKT signaling pathway induces lung tissue fibrosis, which is frequently seen in COVID-19 patients [91, 97, 102]. The activation of some factors including nuclear factor kappa B (NF- κ B) and activated protein-1 (AP-1) has a crucial role in induction of inflammation [103]. If the PI3K/AKT signaling pathway is suppressed, AP-1 and NF- κ B is inhibited reducing the inflammatory cytokines such as TNF- α and IL-6 [91, 104]. Hence, PI3K/AKT inhibitors can suppress inflammation, and their uses along with antiviral drugs help combat COVID-19 [105, 106].

PI3K/AKT signaling in human viral infections

The PI3K/AKT signaling pathway is known to be modulated by many viruses. Most viruses activate the pathway through different measures; therefore, they prevent cell apoptosis, an effective and primitive host defense mechanism [107, 108]. As an essential step to maintain reproduction and dissemination, interruption in programmed cell death enables viruses to enhance replication and even cause persistent infections. Adenoviruses activate PI3K/AKT pathway in corneal and lung epithelial cells, respectively.

Hepatitis B virus (HBV) is known to be an activator of PI3K/AKT cascade [107, 109-112]. On the other hand, HBV DNA replication is downregulated by the active PI3K/AKT signaling pathway, making the latter an equalizer of acute infection and hepatocarcinogenesis [113, 114]. Interestingly some viruses affect the course of other viral infections by the means of PI3K/AKT pathway. Hepatitis C virus (HCV) NS5A protein reduces HBV DNA replication by activating PI3K/AKT cascade [113]. Another example is human immunodeficiency virus (HIV) Tat protein, an activator of PI3K/AKT pathway, which prevents apoptosis. In HIV infected patients Kaposi's sarcoma-associated herpes virus (KSHV) causes a more aggressive form of Kaposi's sarcoma, which is believed to be associated with the anti-apoptotic features of Tat protein. It is worth mentioning that KSHV itself also upregulates PI3K/AKT signaling and inhibits cell apoptosis [115, 116]. A number of studies have found that different viral activators of PI3K/AKT pathway are used by Epstein-Barr virus (EBV) to infect the host cells latently and trigger carcinogenesis [107, 117, 118]. Inhibition of apoptosis and changes in cellular morphology are proposed to be the responsible measures taken by the virus PI3K/ AKT pathway, which plays a role not only in coordination of apoptosis but also in cytoskeletal regulation of cell polarity and mobility. Viruses control penetration and intracellular transport of viral products via this pathway [107, 119]. Previous studies have identified viral activators of PI3K/AKT pathway in other viruses such as human papilloma virus type 16 (HPV16), paramyxoviruses like respiratory syncytial virus (RSV), poliovirus and rhinovirus, cardiovirus, encephalomyocarditis virus (EMCV), coxsackie virus, dengue virus, and Japanese encephalitis virus [120]. Some viruses such as influenza A virus, herpes simplex virus 1 (HSV-1), HIV, and HCV initiate the process as soon as entering the host cell. It is concluded that the early activation of PI3K/AKT pathway promotes viral entrance besides preparing the host environment for infection; for example, activated PI3K/AKT augments HCV entry and genome replication [121, 122]. In contrast, some viruses downregulate the activity of PI3K/AKI. Foot and mouth disease virus (FMDV) and measles virus are some examples of the aforementioned viruses. Some viral products of the latter facilitate PI3K/AKT activity. Interestingly, it has been shown that certain viral activators of PI3K/AKT also work as facilitators of cell apoptosis. A possible explanation for these blind spots is their function in different stages of viral life cycle. Over time, an extensive literature has been developed on the mechanisms of PI3K/AKT modulation by viruses; however, all the components are not fully

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understood [120]. Among coronaviruses, severe acute respiratory syndrome coronavirus (SARS CoV) has been shown to promote apoptosis by downregulating PI3K/AKT activity [123, 124]. At the same time, the virus has the ability to activate PI3K/AKT components by phosphorylation, prevent apoptosis, and establish a state of persistent infection *in vitro* [125, 126]. This is a phenomenon speculated to be responsible for prolonged presence of SARS-CoV particles in human feces in some patients. According to research, the activity level of PI3K/AKT pathway is not high enough to avoid apoptosis in a specific cell line [127, 128]. Meanwhile, some of the cells escaped apoptosis and were infected persistently [125]. A decade after SARS-CoV, a new coronavirus emerged and was named Middle East respiratory syndrome coronavirus (MERS-CoV) [129]. Kindrachuk and coworkers demonstrated the modulation of PI3K/AKT pathway during the infection and reduction in viral replication by adding PI3K/AKT inhibitors to host cells in vitro [130]. Also, MERS-CoV promotes its replication by blocking autophagy [27, 131]. Furthermore, an indirect inhibitor of PI3K/AKT pathway, Saracatinib, has been identified to show antiviral effects against MERS-CoV [132]. Severe acute respiratory syndrome related coronavirus 2 (SARS-CoV-2) activates PI3K/AKT pathway as well [133-137].

Communication between the PI3K-Akt pathway and ACE2 activation in COVID-19 patients

The phosphatidylinositol 3-kinase (PI3K)/Akt pathway mediates different and various types of cell functions, including metabolism, growth, proliferation, and cell survival [138]. It was demonstrated that the virus enters the cell via two pathways; the non-endosomal pathway and endosomal pathway. In the non-endosomal entry, the virus genome is directly transmitted to the host cell's cytosol through some molecules on the surface of the cell, including TMPRSS2 and furin. In the endosomal pathway, SARS-CoV-2 S protein can bind to CD147 and ACE2, letting the virus to enter the host cells and cause infection [93, 139]. In addition, it has been indicated that binding the virus spike proteins to furin and CD147 mediated with some signaling pathway, including PI3K/Akt signaling [140]. Also, it has been demonstrated that endocytosis of SARS-CoV-2 into the cell by receptor-mediated entry follows the clathrin-mediated pathway, regulated by PI3K/Akt signaling, which typically forms an early endosome and becomes a mature endosome before releasing contents in the cytosol [141]. Furthermore, it has been shown that the blocking of the PI3K/Akt signaling pathway has a suppression effect on the viruses that are utilizing clathrin-mediated pathway to enter the host cells [96]. ACE2 is expressed primarily by the lung vascular endothelial cells. However, it has been placed on the surface of some other extrapulmonary cells and tissues, such as the heart, nervous system, intestine, kidney, blood vessels, and muscle [142, 143]. The presence of this enzyme on the surface of various organs in the body may clarify the multiorgan failure and dysfunction in COVID-19 patients. As we know, the ACE2 acts as one of the SARS-CoV-2 receptors, and the reduction of ACE2 on the cells' surface causes the improvement in producing angiotensin II (Ang II), which has been found in COVID-19 patients. Increasing the level of angiotensin II leads to increasing some inflammatory cytokines like interleukin (IL)-6 and TNF- α that consequently shows us the proinflammatory effect properties of Ang II. Increasing the level of Ang II stimulates fibrosis in different organs [97]. In addition, it is believed that angiotensin II (Ang II) type 1 receptor (AT1R) regulates most of the functions of the Ang II in the body. The AT1R exists in different organs and tissues, such as vascular smooth muscle, endothelium, heart, brain, kidney, adrenal gland, and adipose tissue, and assists most of the physiological functions stimulated by Ang II [144]. These different effects can be relevant to PI3K/AKT signaling pathway; by binding the Ang II to its receptor, AT1R, the PI3K/AKT signaling pathway is activated, and it may be responsible for those consequences [101] (Fig. 1).

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Targeting PI3K/AKT in SARS-CoV-2: mechanisms

Since its emergence, SARS-CoV-2 has caused nearly 200 million diseased people and over 4 million deaths. Scientists are trying to find new therapeutic agents against the virus and lower the morbidity and mortality of the disease. SARS-CoV-2 has been shown to be an activator of ATR [136, 145], which serves as a part of PI3K/AKT pathway by detecting DNA damage, ceasing cellular proliferation, and upholding DNA integrity. It has been proposed that berzosertib acts as a protein kinase inhibitor of ATR and a potential antiviral agent against SARS-CoV-2. Berzosertib decreased viral production in vitro with minimal side effects compared with other protein kinase inhibitors. It also reduced infection associated programmed cell death [136, 146]. A strong relationship between PI3K/AKT activation and pulmonary fibrosis has been observed in SARS-CoV-2 induced pneumonia. D-Limonene an ingredient of citrus oil has been proposed as a potential antifibrotic agent in SARS-CoV-2 pulmonary fibrosis. D-Limonene not only suppresses PI3K/AKT expression but also inhibits its phosphorylation. Another possible explanation of D-Limonene induced inactivation of PI3K/AKT is reduction in reactive oxygen species (ROS) levels and TGF-B, which are known as the up-regulators of the pathway. Taken together D-Limonene improved chemical induced pulmonary fibrosis in rats and might be of use in SARS-CoV-2 induced pulmonary fibrosis, vet further research needs to be done [137]. Biguanides are other well-known PI3K/AKT inhibitors and are widely used as antidiabetic agents. An observational study showed lower incidence and morbidity of influenza among patients treated with biguanides compared with patients taking other antidiabetic agents. Another research on animals demonstrated lower pulmonary inflammation and mortality with better general condition in the biguanide treated group. Researchers assumed inhaled biguanides with lower systematic side effects may be useful for SARS-CoV-2 treatment [147]. It is worth mentioning that metformin, a biguanide antidiabetic medication-first used as anti-influenza drug has been demonstrated to be associated with lower mortality in COVID-19 patients However, there are some researches against its efficacy [148-150]. An increasing number of studies have found that dysregulated immune response and hyperinflammation are the reasons behind more severe SARS-CoV-2 infection and pulmonary fibrosis Mammalian target of rapamycin (mTOR), a kinase that functions in PI3K/AKT pathway and regulates inflammatory response by modulating different kinds of T cells and cytokine release. Inhibition of mTOR suppresses the cytokine storm, which maybe the cause of hyperinflammatory phase of COVID-19 [133, 151]. Inhibition of PI3K/AKT pathway by blocking mTOR has been claimed to be beneficial in SARS-CoV-2 infection [130, 152]. Sirolimus also known as rapamycin, an mTOR inhibitor used as an immunosuppressant against transplant rejection, was found to be advantageous in MERS-CoV. Its use might be beneficial in the hyper-inflammatory phase of SARS-CoV-2 infection [153-156]. Until 6 January 2021, 8 clinical trials have been investigating sirolimus use in COVID-19. Terrazzano and coworkers claimed everolimus, a rapamycin derivate, to be a potential antiviral agent against SARS-CoV-2 by blocking mTOR in PI3K/AKT pathway. Everolimus may decrease SARS-CoV-2 replication as it does in some other viral infections [152]. Azithromycin, a macrolide antibiotic has been used for its anti-inflammatory and immunomodulatory effects in some respiratory diseases. It inhibits PI3K/AKT pathway and thus reduces inflammatory cytokines level and modulates T cells. Along with that, Azithromycin may directly decrease viral entrance and replication via PI3K/AKT pathway modulation [157, 158] (Fig. 1, Table 1).

Cell Physiol Biochem 2022;56:707-729 DOI: 10.33594/00000595 Published online: 20 December 2022 Cell Physiol Biochem Press GmbH&Co. KG Aleebrahim-Dehkordi et al.: Natural Marine Drugs to Target COVID-19: Based on PI3K/

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Fig. 1. The virus SARS-CoV-2 enters the cell via the non-endosomal pathway and/or endosomal pathway. In the Non-endosomal entry, the virus genome is directly transmitted to the host cell's cytosol through some molecules on the surface of the cell, including TMPRSS2 and furin. In the endosomal pathway, SARS-CoV-2 S protein can bind to angiotensin-converting enzyme 2 receptor (ACE2) and CD147 that let the virus enter the host cell and active PI3K/AKT/mTOR pathway, resulting in lung fibrosis. Ascidian, Sponges, Corals, Algaes, and Bryozoa are natural PI3K/AKT/mTOR inhibitors suppressing pulmonary fibrosis and cytokine storm.

Classification	Composition name	Source	Kinase	Ref.	
Sponges	Macrolide	Latrunculia magnifica Latrunculia corticata Spongia mycofijiensis Hyatella sp.	РІЗК	[202-204]	
	Bicyclic carboxylic acid	Plakortis sp.	PI3K		
	Sterol sulfates	Spheciospongia sp.	PI3K		
	Bis-steroids	Lissodendoryx (Acanthodoryx) fibrosa	PI3K		
	Liphagal	Aka coralliphaga Fascaplysinopsis sp.	PI3K –Akt	[205-218]	
	Macrocyclic oxaquinolizidine alkaloid	Xestospongia species	PI3K –Akt		
	Triterpenoid	Jaspis stellifera	PI3K –Akt		
	12-epi-scalaradial	Cacospongia sp.	Akt/PKBPI3K		
Corals	Simple sulfone alcohol	Cladiella australis	Akt	[205-218]	
Ascidian	Benzo[d] [1, 2, 3] Trithiole	Ascidium Lissoclinum sp.	PI3K	[205-218]	
Algaes	Phenol	Red alga Rhodomela confervoides	PI3K		
	Fucoidan	Fucos vesiculosus	PI3K –Akt		
	Phlorotannin	Eisenia bicyclis and Undaria pinnatifida	p-Akt	Akt [205-218] t t	
	Carotenoid	Laminaria japonica	Akt		
	Bromo-honaucin	Leptolyngbya crossbyana	Akt		
Bryozoa	Triterpenoids	Conopeum seuratum	PI3K	[205-218]	
sacoglossan (sea slug)	C75 cyclic tridecapeptide	Elysia rufescens	PI3K	[205-218]	

 Table 1. Biologically active compounds from marine organisms capable of inhibiting PI3K/AKT

Cellular Physiology and Biochemistry Published online: 20 December 2022

Cell Physiol Biochem 2022;5	6:707-729

DOI: 10.33594/000000595 2022 The Author(s). Published by Cell Physiol Biochem Press GmbH&Co. KG

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Sponges

Marine sponges (Phylum porifera) are some of the earliest multicellular invertebrate organisms. They are also a great source of valuable therapeutic natural products, some of which are thought to be important lead components for new therapy development. The majority of them are secondary compounds synthesized by sponges [159, 160], that might be used to protect them from harmful bacteria, algae, fungi, and other possible predators; a mechanism sponges have evolved over thousands of years of development. Marine sponges have a soft body, are immobile, and filter feeders, accumulating microscopic bits of food from the seawater that rises across their bodies. Bioactive compounds that come from sponges and other marine microorganisms have been found to contain anti-inflammatory, muscle relaxants, immunosuppressive, antiviral, antifungal, antibacterial anthelminthic, and antimalarial, properties. Sponge substances have exhibited a surprising range of chemical properties. In addition to rare nucleosides, marine sponges can synthesize other kinds of amino acid derivatives such as peroxides, alkaloids, sterols, cyclic peptides, fatty acids, terpenes, and so on. Antiviral compounds found in sponges are among the numerous innovative ways being used to put these compounds into medicinal applications [161-165].

A lot of molecules and anti-infective strategies have been developed for the protection of patients against virus and microbe attacks. These marine compounds can inhibit RNA and DNA viruses such as coronaviruses. They are in various structural classes such as peptides, terpenoids, alkaloids, polysaccharides, and steroids [165-168].

Macrolide

Macrolides are a class of antibiotics distinguished by the presence of a macrolide ring to which one or more deoxy sugars can be linked. It is common for lactone rings to have 14, 15, or 16 parts. Macrolides, which aggregate within leukocytes and are delivered to the site of infection, are used to treat Gram-positive bacterial infections of the respiratory and soft tissues, including S. pneumoniae and H. influenzae. Macrolides are known to inhibit the inflammatory process, minimize excessive production of cytokines in virus infection, and may lessen virus-related aggravation. Additionally, macrolides may affect phagocyte activity by altering a variety of processes such as chemotaxis, phagocytosis, oxidative burst, bacterial death, and cytokine production in addition to other effects [169-171] (Fig. 2).



Cell Physiol Biochem 2022;56:707-729

and Biochemistry Published online: 20 December 2022 Cell Physiol Biochem Press GmbH&Co. KG

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Fucoidan

Fucoidan is a sulfated anionic polysaccharide found in brown marine algae. A unique feature of this sulfated polysaccharide class is its fucose-rich skeleton, which is made up of a variety of sugars that vary considerably between species (e.g., mannose, galactose, glucose, xylose). Due to their low cytotoxicity and antiviral activity, sulfated polysaccharides have recently been shown to have antiviral activity both *in vitro* and *in vivo*. This raises the possibility of their usage in drug and gene delivery systems, wound healing formulations, and other applications. Fucoidans isolated from *Adenocytis utricularis* have inhibitory effects against the reproduction of a variety of enveloped viruses, such as the human immunodeficiency virus and the human cytomegalovirus. Activation of human neutrophils and natural killer cells (NK), as well as the generation of pro-inflammatory cytokines (IL-6, IL-8, and TNF- α), were all enhanced by purified fucoidan from the brown seaweed Undaria pinnatifida, which also inhibited the cells' spontaneous apoptosis. Dendritic cell maturation, cytotoxic T cell stimulation, Th1 immunological responses, antibody production in response to antigen exposure, and formation of memory T cells were all improved by fucoidan produced from *Fucus vesiculosus* [172] (Fig. 3, Fig. 4).

Association between idiopathic pulmonary fibrosis and COVID-19 severity: role of the PI3K/Akt route

Pulmonary fibrosis (PF) is initiated by a subgroup of macrophages named M2. M2 macrophages release transforming growth factor- β 1 (TGF- β 1), which activates fibroblasts and triggers fibrogenesis [173]. Furthermore, TGF- β 1 activates PI3K/Akt pathway, by which autophagy and apoptosis of activated fibroblasts are suppressed and fibrotic remodeling is boosted [174-176]. SARS-CoV-2 may cause pneumonia followed by PF [177-180], which shares similar molecular basis with idiopathic pulmonary fibrosis (IPF) along with comparable role of PI3K/Akt pathway [181]. For instance, interleukin-6 (IL-6) and IL-8, proinflamatory cytokines upregulated by PI3K/Akt pathway [182-185], have been shown to be elevated in IPF and worsening its prognosis, with fibrogenic contribution to the disease [186, 187]. In the same way IL-6 and IL-8 are elevated during COVID-19 infection with significantly higher levels in severe disease [188-190]. Another example is significantly lower levels of Interferon- γ (IFN- γ) in patients with COVID-19 induced PF in comparison





Fig. 4. Potential of fucoidan as a natural marine product that covers a wide range of antiviral activities, anti-bacterial, antioxidant, immunomodulatory, protective effect against oxidative stress and anti-inflammatory effects.



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Cell Physiol Biochem 2022;56:707-729

Published online: 20 December 2022 Cell Physiol Biochem Press GmbH&Co. KG

DOI: 10.33594/000000595

with COVID-19 patients without PF [191]. PI3k inhibitors enhance Interferon- γ (IFN- γ) levels *in vivo* [133]. Given the fact that SARS-CoV-2 is an activator of PI3K/Akt pathway [134-137, 192], lower level of IFN- γ in severe form of the disease (marked with PF [193] is understandable. PI3K/Akt cascades, specially some of its isoforms, are over expressed in IPF [194, 195]. Altogether, with the use of IFN- γ as a therapeutic agent in IPF [187-189], enhances our understanding of pathogenesis of both diseases. Moreover, positive genetic correlation between IPF and severe SARS-CoV-2 has been investigated [196-198]. Hence, there is no surprise that idiopathic pulmonary fibrosis (IPF) patients are more susceptible to get infected by COVID-19, with more severe disease and higher mortality rates [199-201].

Conclusion

The PI3K/AKT signaling pathway is known to be modulated by many viruses. Most viruses activate the pathway through different measures; and therefore, prevent cell apoptosis an effective and primitive host defense mechanism. Marine-derived natural products contain a variety of unique and fascinating structures that contribute significantly to biological activity and clinical therapeutic uses. Because of their bioactive and structure variety, natural compounds from marine organisms are regarded as an outstanding source of novel treatments with structural and chemical properties not present in the terrestrial environment. Several marine-derived kinase inhibitors have been developed from a variety

Cell Physiol Biochem 2022;56:707-729 DOI: 10.33594/000000595 Published online: 20 December 2022 Cell Physiol Biochem Press GmbH&Co. KG Aleebrahim-Dehkordi et al.: Natural Marine Drugs to Target COVID-19: Based on PI3K/

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of various sources and have been shown to inhibit a wide range of protein kinases. A huge spectrum of marine substances exhibits chemical structures with promising biological activity, indicating that they could be exploited to develop drugs for a variety of human diseases caused by viruses, notably COVID-19. Additionally, marine products have the benefit of being drug-like in nature with high levels of bioavailability, allowing them to be used as effective medications against viral illnesses in a very short period of time. Therefore, targeting the PI3K/Akt pathway can be useful as part of a COVID-19 response strategy. If this route is targeted early during the first phases of the disease, it has the potential to block the virus's entry and replication and reduce the viral load, resulting in improved patient outcomes.

Acknowledgements

Author Contributions

Koochaki P, Samami E, Ghoshouni H, Ghadimi D.J, Shadnoush F, Saberianpour Sh: Investigation; writing and preparation of the initial draft. Aleebrahim-Dehkordi E: Study design; Idea and the initiator of the figures; Project administration; writing-final draft and editing. Rafieian-Kopaei M: Supervised the project; editing and critically appraised the manuscript.

Disclosure Statement

The authors declare that no conflict of interests exists.

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