Review

Flavonoids against the SARS-CoV-2 induced inflammatory storm

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ABSTRACT

The disease severity of COVID-19, especially in the elderly and patients with co-morbidities, is characterized by hypercytokinemia, an exaggerated immune response associated with an uncontrolled and excessive release of proinflammatory cytokine mediators (cytokine storm). Flavonoids, important secondary metabolites of plants, have long been studied as therapeutic interventions in inflammatory diseases due to their cytokine-modulatory effects. In this review, we discuss the potential role of flavonoids in the modulation of signaling pathways that are crucial for COVID-19 disease, particularly those related to inflammation and immunity. The immunomodulatory ability of flavonoids, carried out by the regulation of inflammatory mediators, the inhibition of endothelial activation, NLRP3 inflammasome, toll-like receptors (TLRs) or bromodomain containing protein 4 (BRD4), and the activation of the nuclear factor erythroid-derived 2-related factor 2 (Nrf2), might be beneficial in regulating the cytokine storm during SARS-CoV-2 infection. Moreover, the ability of flavonoids to inhibit dipeptidyl peptidase 4 (DPP4), neutralize 3-chymotrypsin-like protease (3CL^{pro}) or to affect gut microbiota to maintain immune response, and the dual action of angiotensin-converting enzyme 2 (ACE-2) may potentially also be applied to the exaggerated inflammatory responses induced by SARS-CoV-2. Based on the previously proven effects of flavonoids in other diseases or on the basis of newly published studies associated with COVID-19 (bioinformatics, molecular docking), it is reasonable to assume positive effects of flavonoids on inflammatory changes associated with COVID-19. This review highlights the current state of knowledge of the utility of

flavonoids in the management of COVID-19 and also points to the multiple biological effects of flavonoids on signaling pathways associated with the inflammation processes that are deregulated in the pathology induced by SARS-CoV-2. The identification of agents, including naturally occurring substances such as flavonoids, represents great approach potentially utilizable in the management of COVID-19. Although not clinically investigated yet, the applicability of flavonoids against COVID-19 could be a promising strategy due to a broad spectrum of their biological activities.

Nomenclature

3CL^{pro}	3-Chymotrypsin-like protease
ACE-2	Angiotensin-converting enzyme 2
Ang1-7	Angiotensin1-7
AngII	AngiotensinII
APC	Antigen-presenting cells
ARDS	Acute respiratory distress syndrome
AT1R	Angiotensin II receptor type 1
BET	Bromodomain and extra terminal domain
BRD4	Bromodomain containing protein 4
CCL2	Monocyte chemo-attractant protein-1
CRP	C-reactive protein
CXCL10	C-X-C motif chemokine ligand 10
DPP4	Dipeptidyl peptidase 4
EGCG	Epigallocatechin-3-gallate
IL	Interleukin
JAK	Janus kinase
KEAP1	Kelch-like ECH-associated protein 1
Mas	Mitochondrial assembly receptor
MCP-1	Monocyte chemoattractant protein 1
MERS-CoV	Middle East respiratory syndrome coronavirus
NF-κB	Nuclear factor kappa B
NK	Natural killer
Nrf2	Nuclear factor erythroid-derived 2-related factor 2
RAAS	Renin-angiotensin-aldosterone system
RNA	Ribonucleic acid
SARS-CoV	Severe acute respiratory syndrome coronavirus
SARS-CoV-2	Severe acute respiratory syndrome coronavirus 2
STAT-2	Signal transducer and activator of transcription 2
TCM	Traditional Chinese medicine
TLRs	Toll-like receptors
TMPRSS	Transmembrane serine protease
TNF-α	Tumor necrosis factor-alpha
VEGF	Vascular endothelial growth factor

1. Introduction

The ongoing COVID-19 pandemic, caused by the severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), is currently one of the most discussed research topics worldwide [1]. The clinical manifestation of COVID-19 occurs after an incubation period of approximately 5–7 days and usually includes cough, fever, fatigue, headache, diarrhea, dyspnoea, and/or lymphopenia [2]. Most patients are asymptomatic or experience only mild to moderate symptoms. However, some patients may develop respiratory failure, acute respiratory distress syndrome (ARDS), or multiple organ failure [1,3]. Advanced age and co-morbidities such as diabetes, cardiovascular disease, hypertension, chronic kidney disease, and chronic obstructive pulmonary disease are highly associated with increased risk of severe course of COVID-19. Complications of severe COVID-19 includes ARDS, septic shock, coagulation dysfunction, metabolic acidosis, cardiac arrhythmia, kidney damage, liver dysfunction, heart failure, or secondary infections [2]. There is ample evidence that lung damage and multiple organ failure in COVID-19 result from systemic hyper-inflammation [1]. Similar to patients with SARS-CoV and MERS-CoV, the subgroup of COVID-19 patients with severe disease have a high level of serum ferritin, D-dimer, fibrinogen, C-reactive protein, interleukin 6 (IL-6) and procalcitonin and are in increased risk of thrombosis and disseminated intravascular coagulation. These clinical and laboratory traits are linked to hyper-inflammation and are associated with macrophage activation syndrome [3]. Therefore, the identification of relevant anti-inflammatory agents applicable to COVID-19 patients could support strategies to overcome the current pandemic [1]. Reducing uncontrolled inflammation should be a therapeutic strategy, especially in patients with exaggerated immune responses demonstrated by over-production of proinflammatory mediators also known as a cytokine storm. Cytokine hyper-production is frequently followed by the development or worsening of acute respiratory failure, acute cardiac damage, or multiple organ failure [2]. Recently, the therapeutic use of naturally

occurring phytochemicals has been revived, building on the traditional use of herbs for medicinal purposes [4,5]. Flavonoids are secondary plant metabolites with a polyphenolic structure [6] that exhibit a wide range of biological effects, including anti-viral, anti-inflammatory, and immunomodulatory activities [7–12]. In addition, flavonoids are relatively cheap and environmentally-friendly substances with minimal or no side effects [7,8]. During the pandemic that limits people's lives worldwide, it is necessary to investigate all potential treatment options for COVID-19. Flavonoids, which are among the most important plant substances found in nature, might advance the treatment of excessive inflammation in COVID-19.

2. Pathogenesis of SARS-CoV-2 infection

The primary immunogenic components of coronaviruses are spike glycoproteins that bind to host cells' receptors to enter them. The human receptor of SARS-CoV-2, angiotensin-converting enzyme-2 (ACE-2), is intensely expressed on the surface of alveolar epithelial type II, renal, cardiac, intestinal, endothelial, and brain cells, which is in a consistency with target organs of COVID-19 [3,13]. ACE-2 is crucial for the entry and replication of SARS-CoV-2; nevertheless, once attached, viral translocation and replication is followed by depletion of membrane ACE-2 [14]. However, the role of ACE-2 is to cleave the angiotensinII (AngII) into angiotensin1-7 (Ang1-7) [15]. Therefore, the binding of SARS-CoV-2 prevents the production of anti-inflammatory Ang1-7 and leads to the accumulation of proinflammatory AngII. This mechanism is believed to be crucial in the pathogenesis of acute lung injury (ALI) in COVID-19 [16]. The protective effects of ACE-2 on ALI were observed in animal models; after infection by SARS-CoV, these injuries were more severe in mice with inactivated ACE-2 [17]. Dipeptidyl peptidase 4 (DPP4) has also been suggested as a co-receptor for SARS-CoV-2 [18, 19]. However, the evidence of a potential role of DPP4 in COVID-19 is still unclear. While Pitocco et al. have not so far supported the inhibition of DPP4 as a credible approach to mitigate COVID-19 [20], other authors discuss the potential role of DPP4 as a target of therapeutic strategies in the pathology of SARS-CoV-2 [19,21]. As is shown in Fig. 1 [18,

19,22], the primary route of SARS-CoV-2 transmission includes direct contact through droplets of saliva or discharge from the respiratory tract when a person sneezes or coughs. After the binding of spike glycoprotein to ACE-2 receptor on the cell surface, SARS-CoV-2 enters the cell, releases RNA genome, and replicates [3]. When the cell is invaded by many viral particles and the viral load is high, the whole protein synthesis apparatus is dedicated to the replication of the virus. The new viral particles are assembled and released by exocytosis and the cell is up to the death by apoptosis or as a result of energetic-metabolic chaos. The released copies of the virus spread and infect other cells and organs in a chain expansion. Eventually, an inflammatory reaction develops in tissues with many dead cells, initially in the lungs and then systematically (lymph, blood, immune system, coagulation, liver, kidney) while the reaction can be clinically severe, especially in patients with co-morbidities [13].

After the recognition of viral antigens by the immune system, antigen-presenting cells present viral antigens to natural killer (NK) cells and CD8-positive cytotoxic T-cells, which activate innate and adaptive immunity leading to the production of proinflammatory cytokines and chemokines [3]. Although the appropriate cytokine release is crucial for the defense against viral infection, an aberrant immune response can lead to organ injury [1]. Cytokines are secreted proteins with specific effects on the development, differentiation, and regulation of immune cells [23]. The immune activation by SARS-CoV-2 infection [24], especially in the elderly or co-morbid patients [25], can become so intense that it may cause deregulated inflammatory processes and uncontrolled systemic inflammatory responses, also known as a cytokine storm [24]. The cytokine storm, which has also been observed in other viral diseases (such as influenza, MERS, and SARS) is defined as an exaggerated immune response demonstrated by the overproduction of proinflammatory cytokines such as IL-6, tumor necrosis factor-alpha (TNF-α), and IL-1β. Currently, there is a clear correlation between elevated pro-inflammatory cytokines as a surrogate marker of cytokine storm and the severity of COVID-19 as well as unfavorable outcomes in hospitalized patients. Elevated IL-6 is recognized as one of the most important marker of unfavorable outcomes [1,23]. The cytokine storm could lead to thrombotic events, ARDS, multiple organ failure, and death [3,26]. The direct cause of death from acute COVID-19 includes cytokine storm-induced damage predominantly to the lungs. However, other organs, including heart, kidney, and liver are also affected. ARDS is characterized by the cytokine storm as one of its main features [24]. Therefore, the over-stimulated immune response may cause more damage to host cells than the foreign invader (SARS-CoV-2) [23]. Other immunologic features of COVID-19 include a lower level of
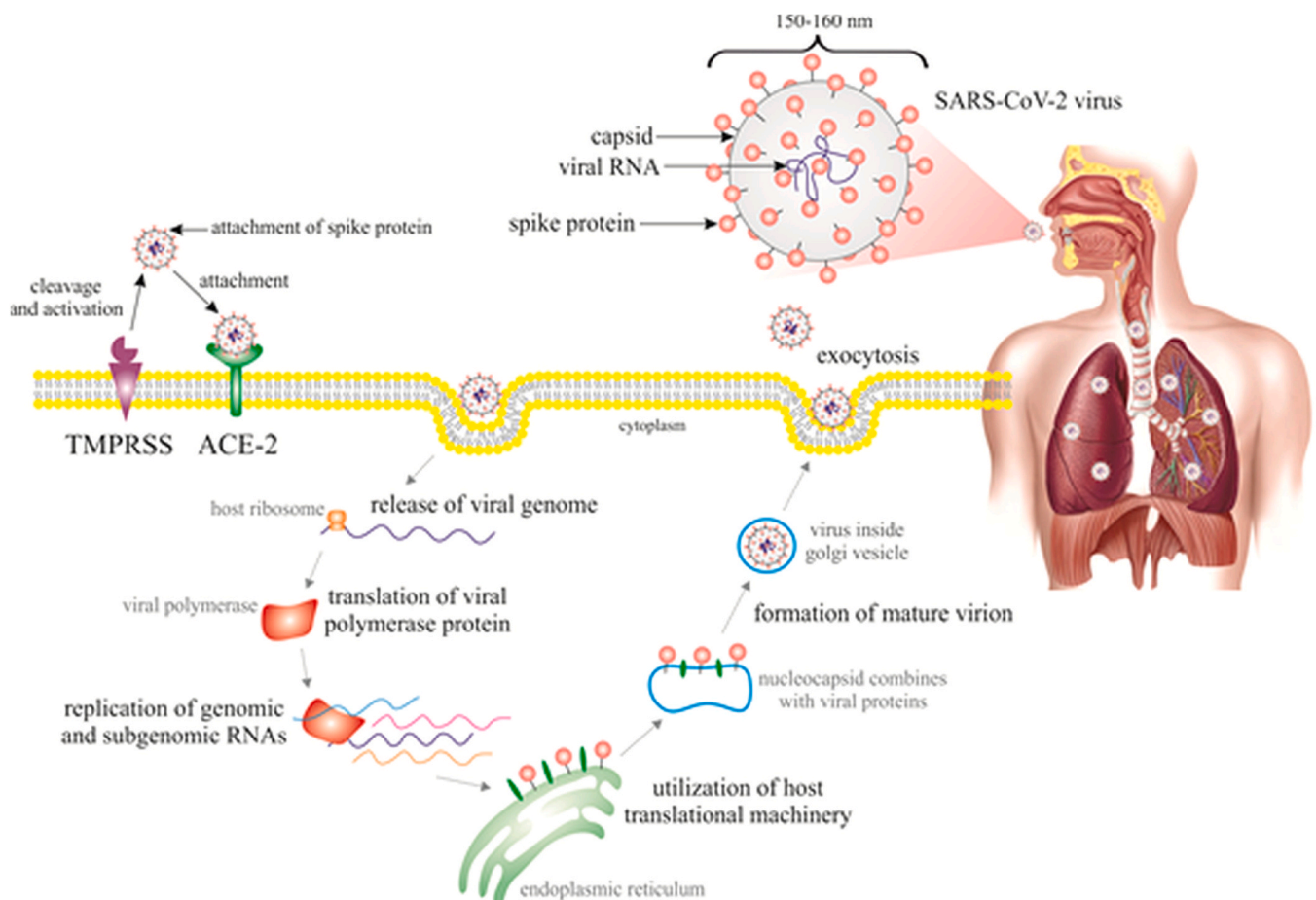

Fig. 1. SARS-CoV-2 infection of host cells. *Abbreviations:* ACE-2, angiotensin-converting enzyme-2; TMPRSS, transmembrane serine protease. *Explanatory notes:* The spike glycoprotein of SARS-CoV-2 is composed of two subunits: S1 mediates the binding of the virus to the ACE-2 receptor and S2 drives host cell membrane fusion allowing viral entry. After the binding of S1 region of the virus to the receptor (ACE-2), the S protein is cleaved by host proteases such as TMPRSS (more specifically TMPRSS2) to be functional and to activate fusogenicity. Then, the fusion of the viral envelope and host plasma membrane and acidified endosomes results in the release of viral genome into the cytoplasm. The next process is facilitated by low pH of endosomes and S2 functional subunit of spike protein. SARS-CoV-2 takes advantage of host endoplasmatic reticulum to form numerous double-membrane vesicles that shield the viral genome and enable replication through the replication-transcription complex. The viral genome is translated into viral polyproteins by the protein translation machinery of the host cell that split by viral proteases into structural and non-structural viral proteins. The assembly of viral particles takes place in the endoplasmatic reticulum/Golgi compartment, and then the assembled virions are carried to the cell surface and are discharged from the cell via exocytosis [18,19,22].

T-lymphocytes, NK cells, and regulatory T-cells in patients with severe manifestations of the disease. On the contrary, an increased level of monocytes and macrophages in COVID-19 patients may explain the elevation of proinflammatory cytokines associated with the cytokine storm [3]. The above-discussed immunologic features of COVID-19 are illustrated in Fig. 2 [1,3,23,24]. Also, the normal gut microbiota can maintain immune response to a viral infection and improve respiratory symptoms [18].

2.1. Inflammatory pathways associated with SARS-CoV-2

Elevated levels of cytokines and inflammatory markers including IL-6, IL-10, IL-1β, and D-dimer have been observed in severely ill COVID-19 patients compared to those with moderate symptoms [1]. Also, the examination of bronchoalveolar lavage fluid revealed the excessive release of chemokines, including C-X-C motif chemokine ligand 10 (CXCL10) and monocyte chemo-attractant protein-1 (CCL2), due to SARS-CoV-2 infection [26]. Table 1 [27–31] provides a detailed overview of the
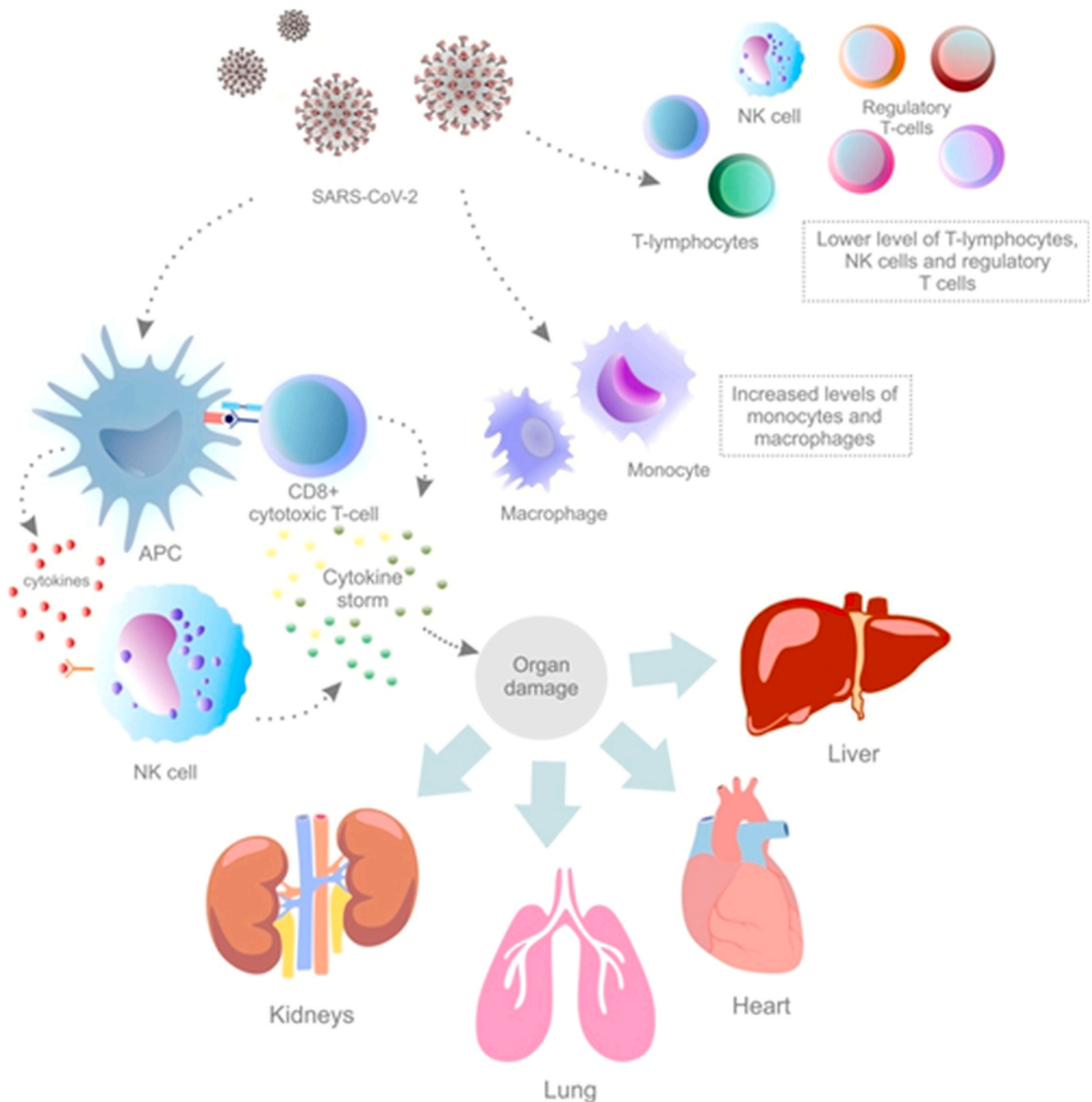
Fig. 2. Immunologic features of SARS-CoV-2-associated pathology. *Abbreviations:* NK cells, natural killer cells; APC, antigen-presenting cells. *Explanatory notes*: APC presents viral antigens to NK cells and CD8-positive cytotoxic cells to activate innate and adaptive immunity and to produce proinflammatory mediators (cytokines) [3]. The immune activation might become so intense that it can lead to exaggerated immune response (cytokine storm) [1,23]. The cytokine storm can result in the damage of lungs, kidneys, heart, and/or liver [24]. The immunologic features of COVID-19 include also lower levels of T-lymphocytes, NK cells, and regulatory T-cells in patients with severe disease progression. An increased level of monocytes and macrophages in COVID-19 patients can also explain the elevation of proinflammatory cytokines [3].

Table 1

Excessive levels of cytokines and chemokines in COVID-19 patients.

The level of inflammation-associated molecules in COVID-19 patients	Reference
Increased IL-1B, IL-1RA, IL-7, IL-8, IL-9, IL-10, FGF, GM-CSF, IFNγ, G-CSF, IP10, MCP1, MIP1A, PDGF, TNFα, VEGF in COVID-19 patients (among which IL-2, IL-7, IL-10, G-CSF, IP10, MCP1, MIP1A, TNFα higher in severe patients)	[27]
Increased plasmatic concentration of IL-2, IL-7, IFN-γ, GCSF, IP-10, MCP1, MIP, and TNF-α in severe COVID-19 patients	[28]
Increased IL-6 in patients with ARDS who died in comparison with patients with ARDS who survived	[29]
Increased IL-6 in patients with pneumonia	[30]
Increased IL-6 associated with death	[31]

Abbreviations: ARDS, acute respiratory distress syndrome; FGF, fibroblast growth factor; G-CSF, granulocyte-colony stimulating factor; GM-CSF, granulocyte-macrophage colony-stimulating factor; IFNγ, interferon-gamma; IL, interleukin; IP10, interferon-γ-inducible protein 10; MCP1, monocyte chemo-attractant protein 1; MIP, macrophage inflammatory proteins; MIP1A, macrophage inflammatory protein 1 alpha; PDGF, platelet-derived growth factor; TNFα, tumor necrosis factor-alpha; VEGF, vascular endothelial growth factor.

clinical features of COVID-19 patients associated with the inflammatory chaos.

Inflammatory pathways associated with SARS-CoV-2 during innate immune response include toll-like receptors (TLRs), which represent a subfamily of pattern recognition receptors (PRR). PRR are capable to recognize viruses including SARS-CoV-2 in the extracellular milieu or endosomes and mediate the inflammatory signaling leading to the activation and production of inflammatory cytokines [23]. Another important inflammatory mediator of the SARS-CoV-2-related exaggerated immune response might be the inflammasome triggered by NLRP3 that also belong s to PRR family. It is expressed in various cell types, including innate immune, endothelial, lung epithelial, hematopoietic, kidney, or cardiac cells [32]. Another mediator of innate immune reaction to SARS-CoV-2 is the bromodomain-containing protein 4 (BRD4). It is an epigenetic reader of acetylated lysines belonging to the bromodomain and extra terminal domain (BET) family that play an important role in epithelial-driven and nuclear factor kappa B (NF-κB)-dependent innate inflammation during viral infection [1]. Furthermore, respiratory viral infections are associated with inflammatory processes and oxidative stress of the epithelium lining cells that activate the transcription factor nuclear factor erythroid-derived 2-related factor 2 (Nrf2), which protects cells from oxidative damage and inflammation. The severity of COVID-19 disease is related to pre-existing conditions, including impaired immune responses, obesity, or age. These conditions are associated with decreased Nrf2 levels [25].

2.2. Targeting inflammation in COVID-19 patients

Based on the known mechanisms of SARS-CoV-2 infection, substances with potentially beneficial effects may act at various stages such as preventing the binding of the virus to the receptors or inhibiting the function of the receptor, suppressing viral replication, helping cells to resist viral attack via inhibition of cytotoxicity processes (potentially slowed down by natural antioxidants), and blocking the virus spread in the body [13]. Moreover, due to the association between over-inflammation or exaggerated immune response and SARS-CoV-2 infection, the cytokine is necessary to be early identified and appropriate anti-inflammatory treatment must be applied. Corticosteroids are currently anti-inflammatory agents with most solid evidence of benefit in the treatment of COVID-19 [1].

However, their applicability is still associated with many unknowns [1]. Notably, as SARS-CoV-2 mainly increases IL-6, and therefore the administration of corticosteroids acting against a wide range of cytokines might be excessive [26]. Therefore, in addition to the global targeting of inflammation, the neutralization of a single key inflammatory

mediator is a currently evaluated therapeutical approach in COVID-19 management [33]. Anti-cytokine therapies such as IL-6, TNFα, and IL-1 cytokine antagonists are suggested to alleviate the hyper-inflammation associated with COVID-19 [23]. Potential treatment strategies targeting the cytokine storm in COVID-19 include the recombinant humanized anti-human IL-6 receptor monoclonal antibody tocilizumab. Nevertheless, the efficacy and safety of tocilizumab in COVID-19 patients needs to be evaluated in large samples and high quality studies and its benefits are yet to be established [26]. In addition, anti-TNF antibodies (infliximab, adalimumab) are suggested to modulate hyper-inflammation and cause a rapid decrease in proinflammatory cytokines in COVID-19 patients. As mentioned above, NLRP3 inflammasome plays a crucial role in the activation of IL-1β. Therefore, drugs targeting NLRP3 inflammasome and IL-1β are promising agents to mitigate hyper-inflammation and cytokine storm induced by NLRP3 inflammasome in severely ill COVID-19 patients [23]. Other potential therapeutic strategies against COVID-19 include cytokine-adsorption device [26], JAK inhibitors, chloroquine, hydroxychloroquine [27], or interferon [34]. Despite the lack of any reported cases of COVID-19 patients treated with BRD4 inhibitors, these drugs can be also considered as potential candidates to ameliorate the excessive inflammation and cytokine storm associated with COVID-19 [1]. Nevertheless, before implementing anti-inflammatory strategies in COVID-19 patients, it is highly important to balance the risks and benefits of specific therapeutic modalities due to the impairment of host immune system which might lead to secondary infections [27].

3. Flavonoids in COVID-19

Flavonoids are defined as important naturally occurring compounds with a phenolic structure [7]. Chemically, flavonoids are composed of fifteen-carbon skeleton that consists of two benzene rings connected through a pyrane ring [35]. Flavonoids are classified by their chemical structure, level of oxidation, and the pattern of the substitution of their heterocyclic pyrane ring also known as C ring. The classification of individual compounds within each class is based on the substitution of benzene rings (A and B rings) [7]. Table 2 provides an overview of the classification and food sources of flavonoids [7–9,36–41].

Flavonoids are secondary plant metabolites responsible for their color, flavor, and are also related to plants' pharmacological activities. Flavonoids possess significant anti-bacterial, anti-oxidant, anti-cancer,

Table 2

Classification and food sources of flavonoids [7–9,36–41].

Flavonoid classification	Members of the flavonoids subgroup	Food sources
Flavanones	Hesperidin, naringenin, naringin, taxifolin, eriodictyol, naringenin	Oranges, lemons, oregano, grapes, medicinal plants
Flavonols	Kaempferol, quercetin, fisetin, myricetin, morin, rutin	Onion, apples, tomatoes, kale, grapes, berries, lettuce, tea, red wine, olive oil, medicinal plants
Flavanols	Catechin, epicatechin, epigallocatechin-3-gallate	Green tea, apples, bananas, blueberries, cacao beans, peaches, pears, medicinal plants
Flavones	Apigenin, luteolin, hispidulin, wogonin, oroxylin, scutellarin, rhamnocitrin baicalein, chrysin, morusin, tangeretin, pectolinarigenin, scutellarin	Chamomile, mint, celery, parsley, Ginkgo biloba, tomatoes, fruit skin, red wine, medicinal plants
Isoflavonoids	Genistein, glycitein, daidzein	Soya, medicinal plants
Chalcones	Phloretin, xanthohumol, isoliquiritigenin, velutone F	Strawberries, apples, medicinal plants
Anthocyanidins	Cyanidin, delphinidin, apigenidin, malvidin	Black/cran/rasp/straw/blue-berries, grapes, cherries, blackcurrants, nuts, medicinal plants

anti-inflammatory, and immunomodulatory abilities [7,8,42]. In addition, flavonoids exert a strong anti-viral capacity in numerous pathologies [43–46]. More importantly, flavonoids demonstrated anti-viral and immunomodulatory activities against coronaviruses [47]. Therefore, the anti-viral properties of flavonoids might be applicable also in the current COVID-19 pandemic. The potentially beneficial role of flavonoids or flavonoid-rich whole plants in COVID-19 pandemic is currently a widely discussed topic [48–51]. One of the suggested targets of SARS-CoV-2 therapies is the ACE-2 receptor [52]. The applicability of flavonoids is associated with the SARS-CoV-2 spike-protein that engages ACE-2 receptors to entry into host cells as was demonstrated by bioinformatics and molecular docking in case of tea flavonoids, especially epigallocatechin-3-gallate (EGCG) and theaflavin gallate [53], fisetin, quercetin, kaempferol [54], quercetin, luteolin, or naringenin [55]. Moreover, the biological activity of flavonoids predetermines them to be effective also in terms of the modulation of inflammatory and immune pathways of SARS-CoV-2-associated pathology.

3.1. Flavonoids as potential inflammatory modulators in COVID-19

The anti-inflammatory and immunomodulatory properties of flavonoids are well-described [7–12]. Thus, flavonoids could potentially be useful in the modulation of COVID-19-related inflammatory processes and immune responses. Due to the excessive immune responses that trigger cytokine release and can result in the overproduction of proinflammatory cytokines, the modulation of systemic immune responses and reversion of hyper-inflammation are suggested to possess a potential role in the management of COVID-19 patients [23]. Generally speaking, flavonoids can modulate the production of inflammatory mediators. Luteolin inhibited IL-1β-induced inflammation in rat chondrocytes [56]. Similarly, apigetrin, a glucoside conjugate of apigenin [57], reduced inflammatory factors including IL-lβ, TNF-α, IL-6, and VEGF in mice with acute otitis media [58]. Also, *Smilax campestris* aqueous extract, which contains catechin and glycosylated derivatives of quercetin (quercetin-3-O-glucoside, quercetin-3-O-galactoside, rutin, and quercetin-3-rhamnoside) as its main constituents, reduced the production of proinflammatory cytokines TNF-α, IL-1β, IL-6, IL-8, and MCP-1 in lipopolysaccharide (LPS)-activated macrophages derived from the monocytic cell line THP-1 [59]. In addition, apigenin alleviated inflammation demonstrated through reduced plasma levels of IL-6, TNF-α, and interferon-γ (IFN-γ) in vivo [60]. Furthermore, flavonoids can modulate the switch of macrophages from proinflammatory to anti-inflammatory phenotype [61]. As was demonstrated by et al, flavonoids (quercetin, naringenin and naringin) induced metabolic variations opposite to proinflammatory metabolic reprogramming elicited by LPS and IFN-γ stimulation in in vitro cultured human macrophages [62]. Also, flavonoids regulate the immune cell functions through the enhancement of the activity of NK cells and cytotoxic T lymphocytes and also through the macrophage functions via modulation of lysosomal activity and the release of nitric oxide [63]. Moreover, potent modulatory properties of hesperidin on systemic immunity (demonstrated by enhanced NK cell cytotoxicity and proportion of phagocytic monocytes, amelioration of the secretion of cytokines stimulated by macrophages, or increased T helper cells) were observed in rats following an intensive training and exhausting exercise [64]. Therefore, a significant benefit of flavonoids is associated with their potent immunomodulatory properties [11,62,65].

Based on the immunomodulatory properties of flavonoids discussed above, we can assume their significant effects on the production of proinflammatory cytokines also within COVID-19. Despite the high affinity to the spike protein, helicase and protease sites on ACE-2 demonstrated in *in silico* studies, the flavonoid-based phytomedicine caflanone also exhibited the ability to inhibit the production of cytokines including IL-1β, IL-6, IL-8, Mip-1α, TNF-α [47]. Moreover, Bellavite and Donzelli [13] recently discussed the nutraceutical properties of citrus fruits, primarily focusing on its flavonoid component hesperidin

as a potential substance against SARS-CoV-2 due to its anti-viral, anti-oxidant, and inflammation-modulatory activities [13]. Hesperidin ameliorated altered level of inflammatory mediators in ischemia/reperfusion-induced kidney injury in rats [66] and also triggered anti-inflammatory responses resulting in a decreased level of IL-33 and TNF-α in mice co-treated with hesperidin and LPS [67]. Nevertheless, the hypothetical role of citrus flavonoid hesperidin against COVID-19 requires corroboration in further pre-clinical, epidemiological, and clinical studies [13].

Moreover, vascular endothelial activation has a crucial role in the excessive cytokine production leading to the cytokine storm and severe pathologies in infectious diseases such as SARS or COVID-19. Therefore, rhamnocitrin, a flavonoid extracted from *Nervilia fordii*, which has been identified as a potent inhibitor of endothelial activation, might be suggested as a potential modulator of cytokine storm in the management of these diseases [68].

Notably, the traditional Chinese medicine (TCM) with its main active ingredients including flavonoids (such as quercetin, kaempferol, luteolin, baicalein, naringenin, and wogonin) could also exert beneficial effects in the management of COVID-19 via the inhibition of viral adsorption and replication as well as the regulation of inflammatory mediators, anti-inflammatory, and immune-regulatory effects to prevent cytokine storm and to protect the target organs [69]. Indeed, Niu et al. [70] recently evaluated chemical constituents of three TCM formulas that were proven to be effective in COVID-19. Eventually, the network pharmacology research revealed decreased IL-6 through several TCM compounds, including but not restricted to quercetin, luteolin, and rutin. These observations suggest the positive association between TCM efficiency in the prevention and rehabilitation of at-risk COVID-19 [70].

Table 3

Effects of TCM (mostly flavonoids among core compounds) in COVID-19 evaluated through network pharmacology and molecular docking.

TCM	Potential effects in COVID-19	Reference
TCM prescription Dayuanyin	Suppression of the inflammatory storm and regulation of immune function. Observed affinity between the core compounds of Dayuanyin (kaempferol, quercetin, 7-Methoxy-2-methyl isoflavone, naringenin, formononetin) and its target genes such as IL-6, IL1β, and CCL2.	[72]
Maxingyigan decoction	Recognized and verified gene targets (including IL-6) and three components of Maxingyigan (quercetin, formononetin, luteolin). The potential role of Maxingyigan in the prevention and treatment of COVID-19 could be based on its anti-inflammatory and immunity-based actions including the activation of T-cells, lymphocytes, leukocytes, cytokine-cytokine-receptor, and chemokine signaling pathways.	[73]
Toujie Quwen granule	The potential role of Toujie Quwen granule and its key active ingredients (including quercetin, kaempferol, luteolin, and oroxylin A, among others) in the treatment of COVID-19 associated with the mechanisms that elevate immunity, suppress inflammatory stress, and regulate inflammatory responses among others.	[74]
Qing-Fei-Pai-Du decoction	Observed immuno-regulatory, anti-inflammatory and multi-organ protective abilities (attributed to four compounds including also flavonoids baicalin and hesperidin and its targets) that could be applicable in COVID-19 management (thrombin and TLR signaling suggested as essential pathways of its anti-inflammatory effects).	[75]

Abbreviations: CCL2, monocyte chemo-attractant protein-1; IL, interleukin; TCM, Traditional Chinese medicine; TLR, Toll-like receptor.

Similarly, other TCM formula, which includes quercetin among others, could inhibit COVID-19 through ACE2 downregulation [71]. Moreover, Table 3 [72–75] briefly summarizes the applicability of other TCM formulas with defined core compounds (represented mostly by flavonoids among others) in COVID-19 through the modulation of inflammatory/immune pathways, based on studies of network pharmacology and molecular docking.

Thus, flavonoids are currently a widely discussed source of agents potentially applicable in the management of COVID-19, as demonstrated by network pharmacology and molecular docking experiments.

3.2. The potential of flavonoids to modulate specific inflammatory pathways deregulated in SARS-CoV-2 infection

Apart from the above-discussed effects of flavonoids against the SARS-CoV-2, research databases offer a wide range of results that could indirectly point to flavonoids' role in targeting inflammatory mediators that are altered in COVID-19. Based on the above-described inflammatory processes associated with COVID-19 and with diverse biological effects of the flavonoids taken into account, it is possible to hypothesize their significant effects on SARS-CoV-2-associated pathways such as the modulation of NLRP3 inflammasome, TLRs, or BRD4, and activation of Nrf2, or the effects on ACE-2 (Fig. 3) [1,16,23,32,76–78].
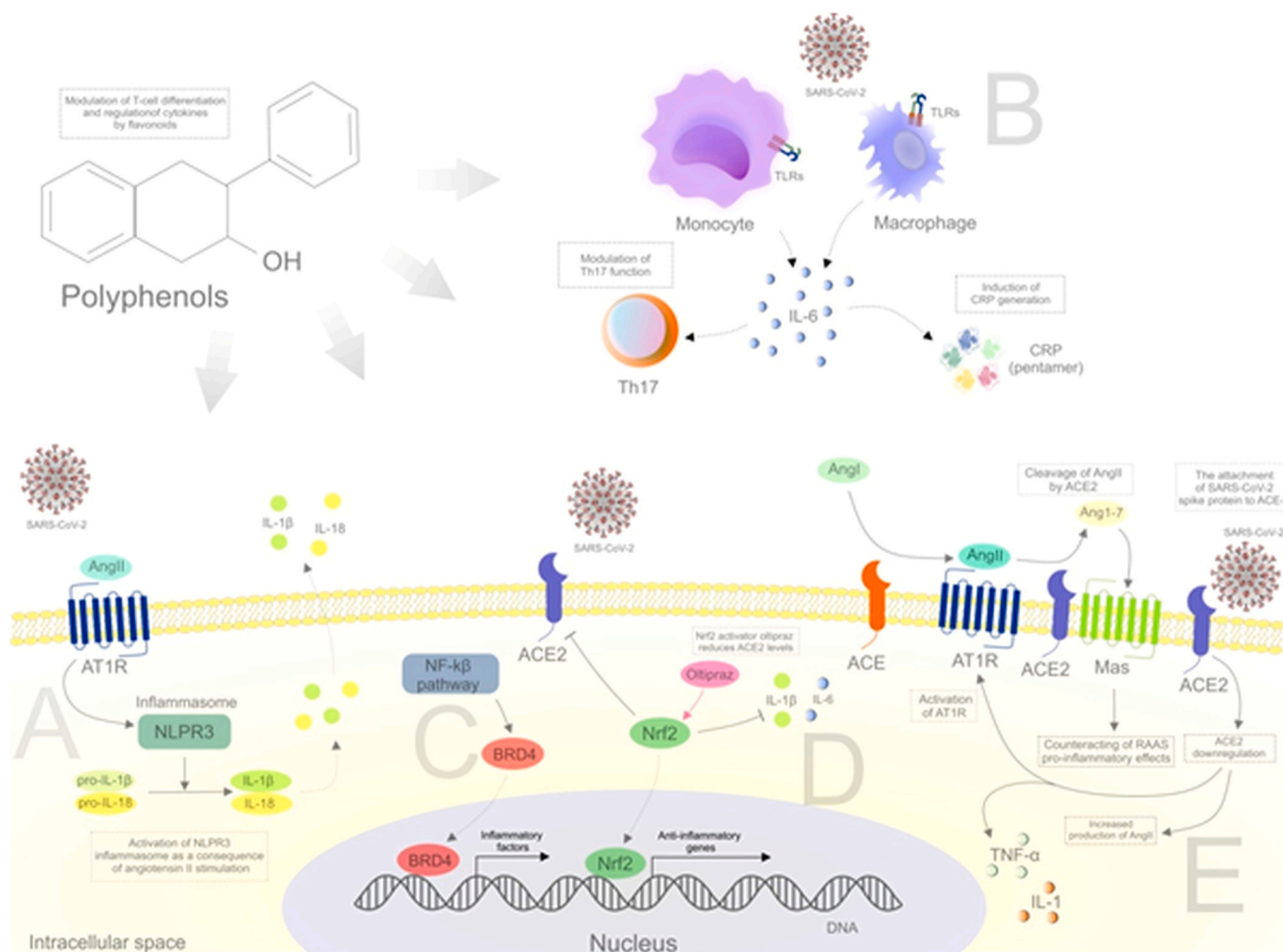

Fig. 3. Inflammatory pathways associated with SARS-CoV-2 that can be potentially targeted by flavonoids. *Abbreviations:* AngII, angiotensin II; ACE, angiotensin-converting enzyme; ACE-2, angiotensin-converting enzyme 2; Ang1-7, angiotensin 1-7; AngII, angiotensin II; AT1R, angiotensin II receptor type 1; BRD4, bromodomain-containing protein 4; CRP, c-reactive protein; IL, interleukin; Mas, mitochondrial assembly receptor; NF-κB, nuclear factor kappa B; Nrf2, nuclear factor erythroid 2-related factor 2; RAAS, renin-angiotensin-aldosterone system; TLRs, toll-like receptors. *Explanatory notes:* (A) An essential determinant of the inflammatory response is the cleavage and secretion of pro-IL-1β and pro-IL-18 into bioactive cytokines activated by the NLRP3 inflammasome [23]. The NLRP3 inflammasome is activated in response to AngII stimulation [32]. (B) TLR activation followed by viral infection can induce the production of IL-6 by macrophages and monocytes. TLRs, TNFα, and IL-1β are considered as the most important stimulators of IL-6. IL-6 is the main regulator of T-cells and can modulate the function of Th17 cells to serve as proinflammatory self-reactive T-cells. IL-6 can also induce the production of acute phase proteins such as CRP [23]. (C) The recruitment of BRD4 by NF-κB leads to the activation of NF-κB-mediated proinflammatory signaling while BRD4 inhibitors decrease the recruitment of macrophages and infiltration of T-cells. The transmembrane E protein of SARS-CoV-2 has been recently demonstrated to bind to BRD4 [1]. (D) The activity of Nrf2 is associated with the modulation of execution and resolution of inflammation through the repression of proinflammatory signals such as IL-6 or IL-1β [76]. (E) Despite the crucial role for viral entry, ACE-2 paradoxically exerts protective effects via conversing AngII to Ang1-7 [77]. SARS-CoV-2 spike protein attachment to ACE-2 leads to ACE-2 downregulation (increase in the level of AngII and augmentation of AngII/AT1R axis activation that are associated with proinflammatory responses). RAAS activation can promote proinflammatory responses through AT1R in kidney and vascular system [16]. The ACE-2-cleaved protein Ang1-7 bind to Mas that is followed by a decrease in proinflammatory cytokine production (TNF-α, IL-6) [78]. Therefore, the binding of SARS-CoV-2 to ACE-2 prevents the production of anti-inflammatory Ang1-7 and leads to the accumulation of proinflammatory AngII [16].

3.2.1. Flavonoids potentially targeting NLRP3 inflammasomes and TLRs in COVID-19

Flavonoids demonstrated reasonable effects in the modulation of inflammatory mediators or signaling cascades including TLRs and NLRP3 inflammasomes (Table 4) [41,79–87]. Altogether, these results highlight flavonoids' capacity to target inflammatory processes associated with TLRs and NLRP3 inflammasome, the deregulation of which is discussed also in terms of SARS-CoV-2 pathology. Therefore, it can be assumed that flavonoids exert significant anti-viral and immunomodulatory effects mediated through TLRs or NLRP3 inflammasomes in COVID-19 patients while these effects need to be precisely evaluated in well-defined pre-clinical and clinical studies.

3.2.2. Flavonoids potentially targeting BRD4 in COVID-19

As mentioned above, the transmembrane E protein of SARS-CoV-2 has been recently demonstrated to bind to BRD4 while its recruitment by NF-κB activates proinflammatory signaling [1]. Importantly, fisetin and amentoflavone are putative ligands of BRD4 and amentoflavone can establish contacts with non-canonical residues for BET inhibition [88]. Moreover, Yokoyama [89] recently discussed the structural and thermodynamic characteristics of isoliquiritigenin interactions with BRD4 and suggested it as a novel template for BDR inhibitors [89]. Hence, BRD4 inhibition represents another approach potentially applicable to overcome COVID-19 associated inflammatory chaos.

3.2.3. Flavonoids potentially targeting Nrf2 in COVID-19

The Nrf2-response has been recently demonstrated to be suppressed in COVID-19 patient biopsies while the Nrf2 agonists 4-octyl-itaconate and the clinically approved dimethyl fumarate inhibited SARS-CoV-2 replication across cell lines in vitro [90]. The poor reproducibility of COVID-19 in animal models limits the effectiveness of the development of therapies against SARS-CoV-2. Nevertheless, genetic or pharmacological activation of Nrf2 demonstrated anti-inflammatory and anti-viral effects in other pathologies while the most relevant mechanisms of its action are associated with targeting specific cysteine receptors within KEAP1 [76]. Although the evaluation of Nrf2 inducers for the reduction of oxidative stress and inflammation in SARS-CoV-2 infections has not been yet performed, a wide range of compounds including flavonoids can activate Nrf2 within other pathologies (7-O-methylbiochanin A, biochanin A, flavonoids of *Abelmoschus esculentus* L. flowers, cyanidin chloride) [91–95]. Moreover, xanthohumol protected LPS-induced acute lung injury against inflammatory damage and oxidative stress via induction of the AMPK/GSK3β-Nrf2 signaling axis in vivo and in vitro [96]. Also, EGCG was observed to protect endothelial cells against inflammation induced by environmental pollutants while the mechanisms of action included the induction of Nrf2-regulated genes [97]. Moreover, *Crateva nurvala* Buch. Ham extracts containing flavonoids as the major class of bioactive phytochemicals activated Nrf2 and decreased proinflammatory TNF-α, NF-κβ, and IL-6 in vivo [98]. Besides, flavonoids from *Apios americana* Medikus leaves reduced inflammatory cytokines and activated Nrf2-KEAP1 pathways in RAW264.7 cells that are, when induced by LPS, accepted as a classic inflammatory model [99]. Furthermore, a study evaluating flavonoids' role in the inhibition of Influenza A viral replication demonstrated that 6-demethoxy-4′-O-methylcapillarisin, a flavonoid derivative of *Artemisia rupestris* L., activated Nrf2/heme oxygenase pathway [100]. It can therefore be hypothesized that flavonoids could decrease the severity of SARS-CoV-2 via the activation of Nrf2 and subsequent modulation of inflammatory and immune processes [25].

3.2.4. Flavonoids potentially targeting ACE-associated pathways in COVID-19

The dual impact of ACE-2 in COVID-19 is associated with its ability to convert AngII to Ang1-7, thus counteracting the inflammatory action of AngII [77]. The attachment of SARS-CoV-2 spike protein to ACE-2 is followed by down-regulation of ACE-2 through its intracellular binding site and then an increase in the level of AngII and augmentation of AngII/AT1R axis activation. Increased production of AngII and

Table 4

Effects of flavonoids on inflammatory cascades TLRs and NLRP3 inflammasomes.

Target of inflammatory pathway	Flavonoid	Aim of the study	Effects	Reference
TLR	Epigallocatechin-3-gallate	BALB/C mice (lipopolysaccharide-induced acute lung injury)	Ameliorated lipopolysaccharide-induced acute lung injury by suppression of TLR4/NF-κB signaling. Decreased proinflammatory cytokines TNF-α, IL-1β, and IL-6 in lung, serum, and bronchoalveolar lavage fluid.	[79]
	Luteolin	C57BL/6J mice (inflammation-mediated metabolic diseases)	TLR signaling modulation. Reduction of macrophage infiltration and modulation of the inflammatory response.	[80]
	Nobiletin	Prostate cancer cells (anti-inflammatory activities)	Anti-inflammatory effects (inhibition of TLR4 and TL9-dependent signaling).	[81]
	Pycnogenol® (extract of French maritime pine bark rich in flavonoids)	TLR-dependent immunomodulatory activities	TLRs inhibition (after gastrointestinal metabolization).	[82]
	Flavonoids from *Houttuynia cordata*	Effects and mechanism of flavonoid glycosides from *H. cordata* on influenza A virus-induced acute lung injury in mice	Attenuation of H1N1-induced acute lung injury (inhibition of TLR signaling).	[83]
NLRP3 inflammasomes	Apigenin	Effects on NLRP3 inflammasome pathways – measurement of active IL-1β (differentiated THP-1 cells)	Inhibition of IL-1β.	[84]
	Scutellarin	Effects on NLRP4 inflammasome activation (macrophages)	Suppression of NLRP3 inflammasome activation in macrophages.	[85]
	Myricetin	Effects on NLRP3-driven inflammatory diseases	Inhibition of NLRP3 inflammasome assembly.	[86]
	Baicalin	Effects on neuroinflammation (amyloid beta precursor protein/presenilin-1 mice)	Protection of neurons from microglia-mediated neuroinflammation via suppression of NLRP3 inflammasomes and the TLR4/NF-κB signaling pathway.	[87]
	Flavonoids isolated from *Millettiavelutina* (velutone F)	Effects on NLRP3 inflammasome activation (THP1 cells)	Suppression of NLRP3 inflammasome activation and serum IL-1β release.	[41]

Abbreviations: IL, interleukin; NF-κB, nuclear factor kappa B; TLRs, Toll-like receptors; TNFα, tumor necrosis factor.

activation of angiotensin II receptor type 1 (AT1R) are processes associated with proinflammatory response. The activation of NF-κB by AngII also leads to the production of TNF-α, IL-6, IL-1β. AT1R activation is followed by the regulation of mitogen-activated protein kinase (MAPK) by AngII, which also affects the release of cytokines (TNF-α, IL-1). Furthermore, the activation of the renin-angiotensin-aldosterone system (RAAS) can be associated with proinflammatory responses through AT1R in the kidney and vascular system. In fact, increased circulatory levels of AngII have been observed in COVID-19 patients with the association of its plasma level and lung injury [16]. However, the ACE-2-cleaved protein Ang1-7 is hypothesized to possess beneficial effects on immune regulation and its low expression during SARS-CoV-2 infection can be associated with COVID-19 severity. The antagonist effects of AngII-derived pathway is associated with the binding of Ang1-7 to the mitochondrial assembly (Mas) receptor and consequent decrease in proinflammatory cytokine production (TNF-α, IL-6) [78]. Therefore, the binding of SARS-CoV-2 to ACE-2 decreases its anti-inflammatory Ang1-7 production and promotes the accumulation of proinflammatory AngII [16].

In 1982, Agarwal demonstrated anti-inflammatory effects of a flavonoid nepitrin that could be mediated through its anti-angiotensin action [101]. More recently, in hypertensive rats models, high levels of AngII were attenuated by hesperidin [102] and RAS activation was inhibited by tangeretin [103]. Similarly, kaempferol exerted inflammation-inhibitory effects mediated by a decrease in AngII-induced collagen accumulation in cardiac fibroblasts [104]. A precise analysis of the effects of ACE-2 on the overall course of SARS-CoV-2 pathogenesis may contribute to the identification of key agents, potentially even among flavonoids, targeting the discussed ACE-2 and related signaling mechanisms.

3.2.5. Other potential mechanisms of flavonoids targeting inflammation as a strategy against COVID-19

Other mechanisms that could potentially be modulated by flavonoids in COVID-19 include the inhibition of DPP4, the neutralization of 3CL^{pro}, or the effects on gut microbiota.

3.2.5.1. Dipeptidyl peptidase 4.
In addition to ACE-2, recently suggested interaction between SARS-CoV-2 and dipeptidyl peptidase 4 (DPP4) as a co-receptor could lead to the development of novel therapeutic COVID-19 strategies. DPP4 is a ubiquitous membrane-bound aminopeptidase with multiple roles in metabolism, nutrition, and the endocrine and immune system [105]. A large interface was observed between the SARS-CoV-2 spike glycoprotein and DPP4 using a docked complex model. The capacity of DPP4 to cleave numerous substrates, such as chemokines or growth factors, is associated with its ability to regulate numerous physiological processes [18] and diseases of the immune system. DPP4 is expressed on epithelia and endothelia of the systemic vasculature, lung, small intestine, kidney, and heart. Accordingly, DPP4 distribution may contribute to the virus's entry through the respiratory tract and may also facilitate the development of cytokine storm and immune-pathologies associated with fatal COVID-19 consequences [19]. Therefore, some investigators suggest the inhibition of DPP4 as a therapeutic strategy to slow the progression of COVID-19 or to hamper cytokine storm and inflammation [18,19]. DPP-4 inhibitors, generally known as anti-diabetic drugs, may possess immunomodulatory functions and could be beneficial in inflammatory diseases [106,107] and could act beneficially also in COVID-19 patients through the reduction of inflammation [108]. Natural compounds, including flavonoids such as EGCG also target DPP4 [109–112]. Indeed, citrus flavonoids [113], epicatechin [114], and chrysin [115] were demonstrated to be potent DPP4 inhibitors. Overall, flavonoids may represent a source of DPP4 inhibitors with potential efficacy against COVID-19.

3.2.5.2. 3-Chymotrypsin-like protease.
3-Chymotrypsin-like protease (3CL^{pro}) is a non-structural protein of coronaviruses. The 3CL^{pro} of SARS-CoV-2 shares 96.1% of its sequence with other SARS-CoV family members such as SARS-CoV or MERS-CoV. 3CL^{pro} cleaves polyproteins into viral replication-related proteins, a process essential for viral replication and maturation. Another crucial function of 3CL^{pro} is to cleave host proteins related to innate immune responsiveness such as the signal transducer and activator of transcription 2 (STAT-2) and NF-κB transcription factor. Therefore, the neutralization of 3CL^{pro} can avert viral maturation and restore the natural immune response [18]. As in the case of SARS-CoV [116,117] and MERS-CoV [118], the promising role of flavonoids has been recently suggested also in SARS-CoV-2 pathology. Interestingly, quercetin has been demonstrated as a potent inhibitor of SARS-CoV-2 3CL^{pro} in vitro and can be considered a proper candidate for further optimization and development [119]. Similarly, the docking study revealed quercetin, scutellarin, and myricetin as potent candidates to target 3CL^{pro} [120]. Despite quercetin's promising efficiency in COVID-19 targeting 3CL^{pro}, its anti-viral properties may be challenged by its poor oral bioavailability. Therefore, Di Pierro et al. have recently demonstrated an increased bioavailability of quercetin's phospholipid complex (Quercetin Phytosome®) in humans [121]. Furthermore, a 3CL^{pro} inhibitory activity of quercetin-3-O-rutinoside (rutin), kaempferol-3-O-rutinoside (nicotiflorin), and their human metabolites has been demonstrated using molecular docking approach [122]. Thus, the neutralization of 3CL^{pro} by flavonoids to restore immune response represents another potential strategy against COVID-19.

3.2.5.3. Gut microbiota.
Despite that SARS-CoV-2 is associated with acute respiratory syndrome, there are also gastrointestinal manifestations that may precede respiratory events. Therefore, due to the capacity of normal gut microbiota restoration to maintain immune response to viral diseases and improve respiratory symptoms, the prebiotic properties of polyphenolic compounds may represent a therapeutic strategy for COVID-19, primarily due to the possibility to affect the gut microbiota of patients [18]. Flavonoids can modulate intestinal immune function through the modulation of T-cell differentiation, alteration of gut microbiota as well as the regulation of cytokines [123,124]. Also, Estruel-Amades [125] demonstrated the immunomodulatory activity of hesperidin on gut-associated lymphoid tissue and reinforced its prebiotic role [125]. Hence, the restoration of the immune responses through gut microbiota may potentially support the organism to overcome COVID-19.

Above all, flavonoids modulate the synthesis or activity of a plethora of inflammatory mediators and immunomodulatory signals [51,74,113, 114,118,121,123,126,127]. However, pre-clinical evidence might be limited by the utilization of non-physiological concentrations in in vitro models while flavonoids are extensively metabolized in vivo [126]. Low bioavailability of flavonoids can limit or hinder their activity. Generally speaking, the bioavailability of flavonoids is affected by several factors including molecular weight, glycosylation, or metabolic conversion [128]. The metabolism of flavonoids occurs in small and large intestine, and liver. Also, the absorption, distribution, and metabolism of flavonoids and their circulating concentrations, elimination, and tissue exposure are also affected by age, sex, genotype, habitual diet, prescribed medicine, and gut microbiome [7]. Gut microbiota plays an essential role in the metabolism of flavonoids [129]. However, some metabolites were observed to paradoxically exert more robust physiological functions than their precursors [7]. Nevertheless, the increase in flavonoids bioavailability as well as the safety evaluation of their application are the goals of ongoing research [7,9,130–132]. Furthermore, human studies are necessary to clarify the anti-inflammatory properties of flavonoids [126].

4. Conclusion

The well-known capacity of flavonoids to regulate anti-viral, anti-

inflammatory, and immunomodulatory responses underscores their potential importance also in the treatment of COVID-19. Although the results of studies utilizing flavonoids against COVID-19 published so far are promising, the body of literature on this topic as a whole remains partial and does not provide sufficient evidence for its applicability in COVID-19 patients. Nevertheless, current intensive research suggests the great potential of flavonoids as natural substances promoting the prevention or overcoming the SARS-CoV-2 infection due to the wide range of their biological effects including the modulation of inflammatory processes and immune responses. Due to the extensive biological effects of flavonoids demonstrated in various pathologies, promising results are hypothesized also in the association of their usefulness in the management of COVID-19. Based on the findings discussed in this review, the beneficial effects of flavonoids in the modulation of inflammatory and immune processes in COVID-19 can be expected. Nevertheless, the precise and detailed evaluation of flavonoid effects in well-defined research are necessary to analyze the exact mechanisms of their action, to specify the population that could benefit from such treatment, and to evaluate their bioavailability or the potentialities to improve their efficacy as a single compound or in combination with other SARS-CoV-2-targeted agents. COVID-19 as a new pathological process represents a unique challenge in the search for and identification of therapeutic substances enabling the overcoming of this pandemic, which is a severe current problem paralyzing the whole society.

Conflict of interest statement

All authors declare that they have no conflict of interest.

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