CD147 as a Target for COVID-19 Treatment: Suggested Effects of Azithromycin and Stem Cell Engagement

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Abstract
The expressive number of deaths and confirmed cases of SARS-CoV-2 call for an urgent demand of effective and available drugs for COVID-19 treatment. CD147, a receptor on host cells, is a novel route for SARS-CoV-2 invasion. Thus, drugs that interfere in the spike protein/CD147 interaction or CD147 expression may inhibit viral invasion and dissemination among other cells, including in progenitor/stem cells. Studies suggest beneficial effects of azithromycin in reducing viral load of hospitalized patients, possibly interfering with ligand/CD147 receptor interactions; however, its possible effects on SARS-CoV-2 invasion has not yet been evaluated. In addition to the possible effect in invasion, azithromycin decreases the expression of some metalloproteinases (downstream to CD147), induces anti-viral responses in primary human bronchial epithelial infected with rhinovirus, decreasing viral replication and release. Moreover, resident lung progenitor/stem are extensively differentiated into myofibroblasts during pulmonary fibrosis, a complication observed in COVID-19 patients. This process, and the possible direct viral invasion of progenitor/stem cells via CD147 or ACE2, could result in the decline of these cellular stocks and failing lung repair. Clinical tests with allogeneic MSCs from healthy individuals are underway to enhance endogenous lung repair and suppress inflammation.

Keywords SARS-CoV-2 · Lung stem cells · Coronavirus · Cellular therapy · Asthma · Diabetes mellitus · Invasion · Anti-viral responses · Basigin · EMMPRIN

Introduction
Currently the world is going through an outbreak of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) that can trigger the coronavirus disease 2019 (COVID-19). On March 31, 2020, over 826,000 coronavirus cases, including 40,708 deaths, have already been confirmed around the world, more specifically in 179 countries [1]. These expressive numbers of confirmed cases and death rates call for an urgent demand of effective and available drugs for COVID-19 treatment.

The open-label non-randomized clinical trial of Gautret and co-workers [2] suggested hydroxychloroquine and azithromycin for COVID-19 therapy. In this trial, twenty-six cases were treated daily with 600 mg of hydroxychloroquine (six patients were lost during the monitoring) and presented a significant decrease in viral load after six days compared to untreated controls. Among hydroxychloroquine-treated patients, six also received azithromycin (500 mg on the first day followed by 250 mg daily) in order to prevent bacterial infection. The most interesting finding of this study was that all patients treated with hydroxychloroquine and azithromycin had negative PCR results in nasopharyngeal samples, i.e., 100% of patients treated with both drugs were virologically cured compared to 57.1% of subjects treated with hydroxychloroquine only, and 12.5% of untreated individuals. Hydroxychloroquine and chloroquine have direct antiviral activity, as demonstrated in vitro [3]. Nine studies have used such drugs for the treatment of patients with COVID-19, including one phase 4 clinical trial (ClinicalTrial.gov Identifier: NCT04316377). Moreover, different action mechanisms of hydroxychloroquine and chloroquine on the COVID-19 have been proposed based on the known action of SARS-CoV-2 and its supposed similarity with the SARS-CoV replication cycle (reviewed by [4]). However, to the best of our
knowledge, except the clinical trial of Gautret and co-workers [2], there are no other studies and clinical trials investigating the effects of azithromycin in COVID-19. A possible reason for this fact may be that azithromycin is a classical antibiotic with very little effects reported against virus [5–7] and so far unknown effects on SARS-CoV-2 infection.

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SARS-CoV-2 invades host cells via two receptors: angiotensin-converting enzyme 2 (ACE2) and CD147 (also known as Basigin or EMMPRIN) [8, 9] (Fig. 1). Spike protein (SP) from virus binds to ACE2 or CD147 on the host cell, mediating viral invasion and dissemination of virus among other cells. The structure of SARS-CoV-2 SP is similar to that of SARS-CoV SP, and both bind to ACE2, invading host cells [8, 9]. In addition to ACE2, Wang and co-worker [8] recently demonstrated that SARS-CoV-2 SP also binds to CD147. They used Meplazumab, an anti-CD147 humanized antibody, co-immunoprecipitation, ELISA, and immuno-electron microscope to demonstrate the new CD147-SP route of viral invasion. This discovery provides a key target for the development and administration of specific anti-SARS-CoV-2 drugs. A clinical trial in phase II entitled “Clinical Study of Anti-CD147 Humanized Meplazumab for Injection to Treat With 2019-nCoV Pneumonia” (ClinicalTrials.gov Identifier: NCT04275245) is currently underway in China aiming to block CD147 protein by monoclonal antibodies, preventing SARS-CoV-2 SP binding and subsequent infection. This trial is monitored by the virologic clearance rate using Real-Time PCR of respiratory tract samples to evaluate the effects of Meplazumab. This drug is a highly specific molecule directed against CD147, but such a specificity does not exclude that other drugs affecting CD147 expression may also have beneficial effects on COVID-19 treatment.

CD147, known as Basigin or extracellular matrix metalloproteinase inducer (EMMPRIN), was also identified as a red blood cell (RBC) receptor for the parasite Plasmodium falciparum - protozoan that causes Malaria in humans [10]. The complex PrfH5 of the parasite membrane binds CD147 on RBC, and this interaction was essential for invasion by all tested strains *in vitro* [10]. Experiments with recombinant chimeric antibody (Ab-1) against CD147 also demonstrated that the inhibition of the PrfH5-CD147 interaction potently blocked RBC invasion and cured established infections in a humanized mouse model [11]. Interestingly, azithromycin, which destroys asexual blood-stage parasites by blocking the ribosomal 50S subunit, acts also as an inhibitor of RBC invasion [11–13]. Other macrolide antibiotics were tested and derivatives of azithromycin and erythromycin presented different invasion inhibitory activity [12]. Invasion inhibition assays and experiments of live video microscopy showed rapid azithromycin action to prevent the essential step of tight junction formation, which occurs very early in the invasion process [12]. Removal of two glycan groups of azithromycin induced a remarkable loss of invasion inhibition [12]. In fact, there are some possible mechanisms, by which azithromycin might prevent invasion. A fundamental step in this invasion process is the ligand/receptor interaction [13]. Thus, the interference of azithromycin in PrfH5/CD147 interaction should be considered. If confirmed, this drug might provide a novel therapeutic strategy against other pathogens that invade cells through binding to CD147, such as SARS-CoV-2.

CD147 is a highly glycosylated transmembrane protein of the immunoglobulin super family that acts as the main upstream stimulator of matrix metalloproteinases (MMPs) and may be upregulated in its expression levels during asthmatic and diabetic complications. Expression levels of CD147 and MMPs are often increased in inflammatory processes and tumors and are associated with cancer progression [14]. Cancer stem cells characterized by high cell surface CD147 expression produce more hyaluronan, resulting in stabilization of lipid rafts, as well as reveal enhanced MMP and ABC drug transporter expression, resistance and survival [15]. Recent studies have shown that patients with severe asthma have high levels of MMP-9 in sputum [16], and influenza A virus infection increases CD147 expression in cells from asthmatics [17]. In addition, another study showed that CD147 expression is induced by high glucose (25 mM) concentration in monocytes [18]. High glucose *in vitro* also increased expression of MMP-1 and MMP-9, and this increase was reverted by suppression of CD147 expression by small interfering RNA and activity inhibition by an anti-CD147 antibody. It is well known that MMPs function in monocyte migration. Thus, inhibition of their regulator CD147 may have beneficial effects in the prevention of diabetic complications involving inflammatory processes [18], including severe acute respiratory syndrome triggers by coronavirus 2. However, specific studies are needed for investigating possible correlations between CD147 and diabetes mellitus in clinical complications due to SARS-CoV-2 infection.

Interestingly, doxycycline, an antibiotic analogue of tetracycline, reduced CD147 levels in a gallbladder carcinoma cell line [19] and gingival crevicular fluid (soluble form) in patients with chronic periodontitis [20]. In addition, monocytes and peripheral blood mononuclear cells treated with azithromycin expressed less MMP-9 [21, 22], as well as airways of lung transplant patients did upon treatment with azithromycin [23]. Furthermore, MMP-3 and MMP-1 expression was reduced in respiratory epithelial cells *in vitro* following azithromycin
In fact, several studies have demonstrated that azithromycin decreases MMP expression and action [21–24]. Despite CD147 and MMPs are closely related molecules, further studies are necessary to evaluate whether azithromycin directly changes expression levels of the CD147 receptor.

Patients with severe COVID-19 present high levels of cytokines in the plasma, specially IL-6, a biomarker for inflammation and robust immune response. In this direction, clinical trials using antibodies anti-IL6 receptors (Sarilumab and Tocilizumab) are already recruited for testing such therapeutic strategy in hospitalized patients with COVID-19. Besides invasion inhibitory activity and antibiotic properties, azithromycin also has immunomodulatory effects, and for this reason has been administrated for the therapy of chronic inflammatory conditions, such as bronchiolitis and rosacea. Although the mechanisms of this anti-inflammatory effect are still little known, some studies presented a reduction of IL-6 levels after azithromycin treatment [25, 26]. Moreover, a relevant study from Gielen and co-workers [27] demonstrated that azithromycin induces anti-viral responses in primary human bronchial epithelial cells, increasing rhinovirus-induced interferons and interferon-stimulated mRNA and protein expression as well as decreasing rhinovirus replication and release [27].

Stem cells might also be a target for viral infection and interesting therapy for combating COVID-19 lung disease. In a previous seminal work, Ling et al. [28] studied the cellular tropism of SARS-CoV in mouse neonatal Oct-4 positive, SSEA-1, Sca-1, cytokeratin-7 and ACE2 expressing lung cell epithelial colonies surrounded by a stroma of neonatal lung tissue. These residing lung stem cells expressed ACE2, one of the entry receptors of SARS-CoV-2 (Fig. 2). The authors observed Oct-4+, long-term BrdU label-retaining cells in the broncho-alveolar junction of the neonatal lung, being in line with the in vitro results. Cultures of Oct-4 lung stem cells could be infected with SARS-CoV with subsequent virus replication, suggesting that failing lung repair of SARS-CoV-infected involves the loss of mobilization and differentiation of lung stem and progenitor cells. On the other hand, CD147, the second entry receptor for SARS-CoV-2, has been proposed to be a marker protein of undifferentiated embryonic stem cells, and the monoclonal antibody against this protein also stained lung cancer cells [29]. It is possible that CD147 is also expressed by untransformed lung stem and progenitor cells; however, to the best of our knowledge, there is no experimental evidence for such affirmation. CD147 is expressed...
by mesenchymal stem cells of human cord blood and bone-marrow origin, suggesting that the CD147 protein is expressed by tissue-specific stem cells [30]. Thus, besides the loss of airway epithelial cells by viral infection and replication, a second component responsible for missing cellular regeneration is proposed, since stem and regenerating cells can be directly infected and lost.

In addition to direct viral invasion and loss of lung stem cells, stocks of these cells also can be affected during SARS-CoV-2 infection by extensive differentiation into myofibroblasts. These cell deposit extracellular matrix proteins and are the main cells in fibrotic diseases [31], including pulmonary fibrosis and fibrous stripes due to COVID-19. In fact, current clinical evidences suggest that pulmonary fibrosis or fibrous stripes are important complications due to COVID-19 [32]. Pan and co-workers [33] observed that 17% of hospitalized COVID-19 patients presented fibrous stripes. Myofibroblasts seem to be originated from several types of cells, such as resident fibroblasts [34], pericytes [35, 36], type II pneumocytes via epithelial-to-mesenchymal transition [37], and resident stem cells/MSC-like cells [38, 39]. Recent evidence using lineage tracing showed that resident mesenchymal cells are the main sources of myofibroblasts in pulmonary fibrosis, rather than epithelial or bone marrow-derived cells [40]. It was observed that resident stem cells/MSC-like cells (side population (SP) and GLI1+ cells) [39, 41] play an important role in pulmonary fibrosis, and their numbers decline in mice and patients with this complication compared to healthy controls [38, 39, 41].

ACE2, present in multiple cellular types in lung, including resident stem cells [28], belongs the renin angiotensin system that is involved in pulmonary fibrosis [42, 43]. It was observed that angiotensin-II (Ang-II) is necessary and sufficient to promote fibrosis in mice [42]. On the other hand, ACE2 activity cleaving Ang-II inhibits experimental lung fibrosis, and patients with this clinical complication presented...
downregulated levels of this receptor [42]. CD147, the second entry receptor for SARS-CoV-2, is also present in multiple cellular types in lung and highly expressed in type II pneumocytes and macrophages at the edges of the fibrotic zones [44]. Normal human lung fibroblasts were transiently transfected to overexpress CD147 and presented a significant increase in TGF-β1-induced proliferation and expression of α-smooth muscle actin, a marker of myofibroblasts. On the other hand, anti-CD147 antibodies inhibited TGF-β1-induced proliferation and differentiation of fibroblasts into myofibroblasts [44]. Therefore, the blockade of CD147 could also play a beneficial role in pulmonary fibrosis due to COVID-19.

Proliferating fibroblasts invade the intra-alveolar zone and origin the so called “fibroblast foci”. This process is considered the first step in pathogenesis of pulmonary fibrosis [31]. Lung sections from two patients with early-phase COVID-19 pneumonia presented characteristics similar to this initiation step of fibrosis [45]. Fibroblast foci were observed in the airspaces, in addition to edema, type II pneumocyte hyperplasia with inflammatory cellular infiltration and multinucleated giant cells. Some regions of reactive epithelial hyperplasia had also abundant alveolar macrophages [45]. In fact, the continuing injury to the alveolar zone caused by SARS-CoV-2 infections possibly origins a pro-inflammatory microenvironment triggering this aberrant response with partial replacement of normal tissue by fibrous tissue.

Since both early-phase and severe COVID-19 stages show strong involvement of inflammatory components and possibly loss of resident stem cell stocks, a considerable amount of research activities are engaged in investigating the role of cellular therapy using immune response-suppressing MSCs for COVID-19 therapy. In this direction, several clinical trials are planned or already recruited for testing such cellular therapeutic strategy [46]. These trials shall study possible reduction of inflammation and promotion of tissue regeneration following allogeneic MSC transplantation.

Conclusions

Thus, we have presented here enough evidence to raise some questions: How can the effects of azithromycin in COVID-19 observed by Gautret and co-workers [2] (virologically curing patients in combination with hydroxychloroquine) be explained? Can azithromycin have any invasion inhibitory activity (as already reported elsewhere [11–13]) for the case of COVID-19? Does azithromycin treatment decrease the expression of MMPs and possibly CD147, reducing SARS-CoV-2 infection? Both Plasmodium falciparum causing Malaria and SARS-CoV-2 causing COVID-19 are susceptible to hydroxychloroquine treatment; however, are both pathogens also susceptible to azithromycin because they need of the CD147 protein for cell invasion? Can resident progenitor/stem cells be targets of SARS-CoV-2 infection and/or affected in their differentiation capabilities? Can effects of immunomodulatory and anti-viral response of MCSs and azithromycin, respectively, be therapeutically exploited in COVID-19 treatment? Can patients with asthma or diabetes mellitus have high expression levels of CD147 receptor, and are they for this reason more prone to complications?

We have addressed these issues, highlighting CD147 as a target for COVID-19 treatment and suggesting this protein as a possible pathway for azithromycin-induced effects. Azithromycin has been worldwide used for the treatment of infectious diseases with few side effects and can be rapidly produced in large scale, unlike monoclonal antibodies that generally have a slower production process. Nevertheless, further studies need to be carried out to confirm the possible role and mechanism of the azithromycin in COVID-19. Moreover, progenitor/stem cells can be extensively recruited in pulmonary fibrosis of COVID-19 patients and possibly suffer direct invasion via CD147 or ACE. Thus, besides the loss of airway epithelial cells, a second component of missing cellular regeneration is proposed. Certainly, more basic and clinical investigations into this issue are necessary for effective therapeutic approaches. Combinations of pharmacological and MSC therapy for successful therapy of COVID-19 and other viral diseases might be available in the future.

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Compliance with Ethical Standards

Conflict of Interest The authors declare that there are no conflicts of interest regarding the publication of this paper.

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