

# 1                                   **Macrophage Responses Associated with COVID-19:**

## 2   **A Pharmacological Perspective**

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31 **Abstract**

32 COVID-19 has caused worldwide death and economic destruction. The pandemic is the result of  
33 the severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), which has demonstrated  
34 high rates of infectivity leading to great morbidity and mortality in vulnerable populations. At  
35 present, scientists are exploring various approaches to curb this pandemic and alleviate its health  
36 consequences, while racing to develop a vaccine. A particularly insidious aspect of COVID-19 is  
37 the delayed overactivation of the body's immune system that is manifested as the cytokine storm.  
38 This unbridled production of pro-inflammatory cytokines and chemokines can directly or  
39 indirectly cause massive organ damage and failure. Systemic vascular endothelial inflammation  
40 and thrombocytopenia are potential consequences as well. In the case of COVID-19, the  
41 cytokine storm often fits the pattern of the macrophage activation syndrome with  
42 lymphocytopenia. The basis for the imbalance between the innate and adaptive immune systems  
43 is not clearly defined, but highlights the effect of SARS-CoV-2 on macrophages. Here we  
44 discuss the potential underlying basis for the impact of SARS-CoV-2 on macrophages, both  
45 direct and indirect, and potential therapeutic targets. These include granulocyte-macrophage  
46 colony-stimulating factor (GM-CSF), interleukin 6 (IL-6), interferons, and CXCL10 (IP-10).  
47 Various biopharmaceuticals are being repurposed to target the cytokine storm in COVID-19  
48 patients. In addition, we discuss the rationale for activating the macrophage alpha 7 nicotinic  
49 receptors as a therapeutic target. A better understanding of the molecular consequences of  
50 SARS-CoV-2 infection of macrophages could lead to novel and more effective treatments for  
51 COVID-19.

52 **Key words:** Immunopharmacology, immunomodulation therapy, pandemic, cytokine storm,  
53 macrophage activation syndrome, biologicals.

54 **1. Introduction**

55 Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) emerged in Wuhan,  
56 China in December of 2019 and quickly wreaked havoc around the world in the form of the  
57 pandemic COVID-19, causing death, undermining economies, overwhelming medical  
58 professionals, and challenging the scientific community (Liu et al., 2020b). This positive-sense  
59 single-stranded RNA virus has proven to be highly contagious, being spread by symptomatic and  
60 likely asymptomatic individuals (Furukawa et al., 2020; Huff and Singh, 2020; Oran and Topol,  
61 2020). As the name suggests, the primary target of SARS-CoV-2 is the lungs, but other organs  
62 such as blood vessels, heart, and brain are susceptible as well. All age groups are vulnerable to  
63 infection, but generally exhibit different degrees or classes of symptoms, with those over 60,  
64 male, and with underlying medical conditions more likely to exhibit severe symptoms and  
65 succumb to viral toxicity (Conti and Younes, 2020; Team, 2020). Some 81% exhibit mild,  
66 moderate, or no symptoms; 14% show severe symptoms; and 5% experience critical disease with  
67 high mortality (Wu and McGoogan, 2020). An especially alarming complication of COVID-19 is  
68 the cytokine storm that develops after a week or two of delay in severely infected individuals.

69 SARS-CoV-2 has 4 structural proteins, namely the E (envelope), S (spike), M  
70 (membrane), and N (nucleocapsid) proteins (Guo et al., 2020). The N protein holds the RNA  
71 genome, while the S, E, and M proteins form the viral envelope. The virus primarily gains entry  
72 into a human cell by binding to the exopeptidase angiotensin converting enzyme 2 (ACE2). This  
73 protein is located on the membrane surface of several cell types including alveolar type II and  
74 endothelial cells. Proteins other than ACE2 may function as receptors for entry as well (Guo et  
75 al., 2020). Cell entry is facilitated by cleavage of the spike protein by the serine protease  
76 TMPRSS2 or a furin-like proprotein convertase, thereby exposing the fusion peptide. Besides

77 inducing cell death, viral infection can initiate an inflammatory response, which with SARS-  
78 CoV-2 is thought to manifest among other things as widespread vascular endothelial dysfunction  
79 (Teuwen et al., 2020). Beyond this, however, increasing evidence supports the conclusion that  
80 SARS-CoV-2 may exert some of its lethal effects by insidiously compromising the body's  
81 immune response. Here we summarize evidence for macrophages as targets of SARS-CoV-2 and  
82 the implication that has for immunomodulatory treatments of COVID-19 (Fig. 1).

83

## 84 **2. Cytokine storm**

85 Progression of COVID-19 in more severe cases is marked by the delayed occurrence of a  
86 cytokine storm or cytokine release syndrome, due to overactivation of the immune system.

87 Although not definitively established, this phenomenon is thought to contribute to the acute  
88 respiratory distress syndrome (ARDS) and widespread organ damage that foretells death. Nor is  
89 it clear what relationship there is between the cytokine storm and thrombocytopenia, which is  
90 common in patients with COVID-19 and may ultimately contribute to adverse outcome, although  
91 both enhanced platelet activation/consumption and destruction are likely outcomes of the  
92 cytokine storm. Multi-organ (micro-) thrombosis seems to characterize severe COVID-19 cases  
93 (McFadyen et al., 2020; Prieto-Pérez et al., 2020), and likely reflects in part the production of  
94 pro-inflammatory cytokines, such as IL-1 $\beta$  and TNF- $\alpha$ , by macrophages (Conti et al., 2020a).

95 Notably, excessive activation or proliferation of macrophages is a contributing factor to  
96 hemophagocytic histiocytosis (HH) also known as secondary hemophagocytic  
97 lymphohistiocytosis (Xu et al., 2020). HH has been identified as a deregulation of the immune  
98 system, characterized by hemophagocytosis by macrophages, overactivation of cytotoxic T cells,  
99 and pro-inflammatory cytokine massive release (Ramos-Casals et al., 2014). HH is the

100 histological counterpart of the macrophage activation syndrome. A clinical study performed on  
101 post-mortem bone marrow samples taken from patients who died from COVID- 19 showed  
102 findings highly consistent with the diagnosis of HH (Prieto-Pérez et al., 2020). Elevated blood  
103 ferritin has also been shown to be associated with poor outcome in a retrospective study of 150  
104 COVID-19 patients (Mehta et al., 2020a).

105         From multiple observations, both CD4<sup>+</sup> and especially CD8<sup>+</sup> (or cytotoxic) T-cells appear  
106 to be over-activated early-on in COVID-19 resulting in the excessive production of granulocyte-  
107 macrophage colony-stimulating factor (GM-CSF), which in turn stimulates  
108 monocytes/macrophages to produce interleukin-6 (IL-6) and other inflammatory factors. With  
109 time, there is a significant decrease in peripheral CD4<sup>+</sup> and CD8<sup>+</sup> T lymphocytes, as well as  
110 natural killer (NK cells) in COVID-19 patients, perhaps secondarily to their sustained activation  
111 by macrophage-derived interferon gamma-induced protein 10 (IP-10), also known as CXCL10.  
112 With disease progression, neutrophilia may occur, especially in those with severe critical  
113 pulmonary conditions (Liu et al., 2020a).

114

### 115 **3. Macrophage (monocytes)**

#### 116 **3.1 Inflammatory signature**

117         Human monocytes and macrophages express ACE2, as well as TMPRSS2 and furin, and  
118 would seem to be a widespread target for SARS-CoV-2 infection (Abassi et al., 2020; Wang et  
119 al., 2020b). Evidence was reported in COVID-19 patients for the infection of macrophages of the  
120 spleen and lymph nodes with SARS-CoV-2, which was associated with severe lymphocyte  
121 apoptosis (Wang et al., 2020b). Moreover, infected macrophages were shown to produce IL-6, a  
122 pro-inflammatory cytokine that directly promotes lymphocyte necrosis and would explain in part

123 the common characteristic of lymphocytopenia in COVID-19 patients. Based on their  
124 morphology and ability to produce IL-6, TNF- $\alpha$ , and IL-10, as well as surface expression of  
125 CD11b, CD14, CD16, CD68, CD80, CD163, and CD206, circulating monocytes have an  
126 activated or pro-inflammatory phenotype. The expression of CD163 and CD206 suggests a bias  
127 towards the intermediate or regulatory phenotype, with CD163 expression being a feature of  
128 activated monocytes/macrophages in hemophagocytic lymphosistocytosis syndrome (Wang et  
129 al., 2020b). An increase in the pool size of the intermediate subtype of monocytes may be  
130 characteristic of severe COVID-19 (Merad and Martin, 2020b). The activated plasma blood  
131 monocyte phenotype and lymphocytopenia would seem to persist into the recovery stage as well  
132 (Wen et al., 2020).

133 Multiple studies have demonstrated that the lungs are a target of macrophages in COVID-  
134 19 (Chua et al., 2020; Wang et al., 2020b). Inflammatory macrophages are increased with  
135 increased levels of nonresident macrophages, which in the upper respiratory tract have a highly  
136 inflammatory phenotype with the expression of a number of chemokines and pro-inflammatory  
137 cytokines IL-1B, IL-8, IL-18, and TNF- $\alpha$  (Chua et al., 2020; Liao et al., 2020). Macrophages in  
138 the lower airways were found to have an even stronger inflammatory signature and overall there  
139 was a strong correlation between activation status of non-resident macrophages and COVID-19  
140 disease severity (Chua et al., 2020). Other immune cells, such as mast cells, likely act  
141 synergistically with macrophages to cause lung damage (Kritas et al., 2020).

142

### 143 **3.2 Interferon suppression**

144 Although CXCL10, as well as CCL2, are interferon (IFN)-induced genes, there is  
145 evidence for impaired or delayed Type 1 IFN signaling in SARS-CoV-2-infected cells. One ex

146 vivo experiment with lung tissue showed that SARS-CoV-2 induced less IFNs and pro-  
147 inflammatory mediators than SARS-CoV (Chu et al., 2020). Single-cell RNA sequencing  
148 analysis of bronchoalveolar lavage samples from severe and mild COVID-19 patients revealed  
149 that SARS-CoV-2 mainly infects the epithelial and recruited inflammatory macrophage subsets  
150 (Bost et al., 2020). In the latter, a disease severity-associated downregulation of type I IFN genes  
151 was noted. Notably, IFN is known to exhibit multiple biological functions such as antiviral,  
152 antiproliferative, and immunomodulatory effects (Nile et al., 2020; Wang et al., 2019). How  
153 SARS-CoV-2 thwarts intrinsic innate immune responses in monocyte-macrophages is not  
154 defined, although in monocyte-derived dendritic cells (but not macrophages) viral antagonism of  
155 STAT1 phosphorylation was reported (Yang et al., 2020). In contrast, work in Vero cells,  
156 indicates that SARS-CoV-2-infected cells are still responsive to type I IFN treatment unlike  
157 SARS-CoV-infected cells (Lokugamage et al., 2020). Of note, ACE2 was shown to be an  
158 interferon-stimulated gene in human lung cells, which is also upregulated by smoking and viral  
159 infections (Smith et al., 2020). A discussion of possible means by which SARS-CoV-2 attenuates  
160 the interferon response can be found elsewhere (Paces et al., 2020). Recently, it was reported  
161 that the SARS-CoV-2 viral ORF6, ORF8 and N proteins were potential inhibitors of the type I  
162 interferon signaling pathway (Li et al., 2020).

163 In light of these observations and urgent need to identify new therapies to control  
164 COVID-19 severity, IFN approved drugs have emerged as a potential treatment for COVID-19  
165 patients. For instance, it has been demonstrated that the administration of recombinant IFNs to  
166 SARS-CoV and SARS-CoV-2 patients decreased viral protein synthesis and replication  
167 (Falzarano et al., 2013; Li et al., 2019; Zumla et al., 2016). In agreement, a recent published  
168 study on MERS-CoV patients reported that a combination of remdisevir and IFN beta showed a

169 superior antiviral effect when compared with lopinavir/ritonavir combination (Sheahan et al.,  
170 2020). Therefore, testing the efficacy and safety of recombinant IFNs may be a worthwhile  
171 promising approach in the setting of COVID-19. Triple antiviral therapy with lopinavir-ritonavir,  
172 ribavirin and interferon beta-1b was reported to be safe and superior to lopinavir-alone in  
173 improving symptoms and reducing viral shedding and hospitalization in those with mild to  
174 moderate COVID-19 (Hung et al., 2020). On the other hand, there is evidence that IFN might be  
175 playing an important role in COVID-19 hyper-inflammation, suggesting that timing is a  
176 consideration (Conti et al., 2020c; Lee et al., 2020). Analysis of monocytes by single-cell RNA-  
177 seq from patients with severe COVID-19 exhibited signs of a type I IFN response along with  
178 TNF/IL-1 $\beta$ -driven inflammation.

179

### 180 **3.3 Possible contribution of nicotine and nicotinic acetylcholine receptors**

181 Although multiple investigations report a detrimental impact of nicotine on COVID-19  
182 patients through up-regulating ACE2 receptors in the lungs (Farsalinos et al., 2020a; Leung et  
183 al., 2020; Russo et al., 2020), recently published epidemiological studies reveal that smokers are  
184 either asymptomatic or show less severe respiratory symptoms compared with non-smokers  
185 (Covid et al., 2020; Farsalinos et al., 2020b; Kloc et al., 2020; Miyara et al., 2020; Petrilli et al.,  
186 2020). A disruption of the cholinergic anti-inflammatory pathway in COVID-19 patients has  
187 been noted (Farsalinos et al., 2020a; Farsalinos et al., 2020c). It has been reported that over-  
188 responsiveness of the immune system, otherwise known as the cytokine storm, highly correlates  
189 with enhanced severity of COVID-19 infection, substantially increasing the mortality rate (Wang  
190 et al., 2020a; Ye et al., 2020). In the human lungs, the inflammatory response is mainly  
191 mediated by lung macrophages with two main types: the alveolar and interstitial macrophages



192 (Kloc et al., 2020). Under physiological conditions, the alveolar macrophages exhibit anti-  
193 inflammatory characteristics by dampening the adaptive immune response and suppressing pro-  
194 inflammatory cytokines release (Kloc et al., 2020). Following a viral infection such as COVID-  
195 19, the alveolar macrophages switch from the anti- to pro-inflammatory phenotype, initiating  
196 consequently an inflammatory response, then switch back during the resolution phase to the anti-  
197 inflammatory phenotype, promoting thereafter tissue repair in the site of injury (Hu and  
198 CHRISTMAN, 2019; Hussell and Bell, 2014). In the context of COVID-19 infection, an  
199 accumulation of macrophages in the lungs of COVID-19 patients has been observed (Wang et  
200 al., 2020a). Besides resident macrophages, monocyte-derived and non-resident macrophages  
201 have been described in COVID-19 patients (Chua et al., 2020); however, a better understanding  
202 of their interrelationship is needed.

203         Of note, lung macrophages have been shown to express ACE2 receptors, facilitating  
204 therefore the entry of SARS-CoV-2 to host cells (Tsaytler et al., 2011; Verdecchia et al., 2020).  
205 Besides ACE2 receptors, lung macrophages express alpha 7 nicotinic receptors (nAChRs  $\alpha 7$ )  
206 (Abrial et al., 2012). nAChRs  $\alpha 7$  are potentially implicated in attenuating the cytokine storm  
207 through decreasing pro-inflammatory cytokine release (Kalamida et al., 2007; Tracey, 2002). For  
208 instance, it has been indicated that activation of nAChRs  $\alpha 7$  located on lung macrophages by  
209 acetylcholine and/or nicotine mitigates the hyper-inflammatory response mediated disease  
210 severity (Lu et al., 2014; Tindle et al., 2020). Strong evidence reveals that the cholinergic anti-  
211 inflammatory pathway mediated by nAChRs  $\alpha 7$  inhibits the translocation of the pro-  
212 inflammatory marker NF- $\kappa$ B to the nucleus and activates the JAK2-STAT3 pathway,  
213 consequently suppressing the inflammatory response and decreasing the cytokine storm in the  
214 lungs (Báez-Pagán et al., 2015; Changeux et al., 2020; Lu et al., 2014). Given the observed lower

215 number of hospitalized COVID-19 patients among smokers, the potential role of medicinal  
216 nicotine to alleviate COVID-19 progression and development should be rapidly studied and  
217 clearly distinguished from conventional smoking that has no therapeutic effects.

218

### 219 **3.4 Chemokine profile: a possible role for CXCL10 (IP-10)**

220 Longitudinal profiling of 71 COVID-19 patients identified early expression of inhibitory  
221 mediators IL-10 and IL-1RA, along with the chemokine CCL5 (aka RANTES), in those with  
222 mild but not severe disease (Zhao et al., 2020). CCL5 is chemotactic for T cells, as well as  
223 eosinophils and basophil. On the other hand, the majority of cytokines associated with the  
224 cytokine storm in viral infections, including IL-6 and IFN- $\gamma$ , were only increased at a late stage  
225 in severe illness, with TNF and GM-CSF not showing a difference between mild and severe  
226 cases.

227 Multiple studies have documented the upregulation of not only inflammatory cytokines  
228 but also chemokines in COVID-19 patients. Chemokines are low molecular weight proteins that  
229 act largely as chemoattractants for immune cell recruitment during inflammation, as well as  
230 modulators of immune cell homeostasis and angiogenesis (Coperchini et al., 2020). Compared to  
231 non-ICU patients, COVID-19 patients admitted to the ICU, exhibited higher plasma levels of  
232 IL2, IL7, IL10, GSCF, CXCL10 (IP-10), CCL2 (MCP1), CCL3 (MIP1A), and TNF $\alpha$ , indicating  
233 activation of T-helper 1 (Th1) cell function (Huang et al., 2020), although increased circulating  
234 levels of Th2-immune related cytokines IL-4 and IL-10 implicated in inflammation suppression  
235 are noted as well (Han et al., 2020). Transcriptomic analysis of bronchoalveolar lavage fluid of  
236 COVID-19 patients revealed an upregulation of CXCL1, CXCL2, CXCL6, CXCL8 (IL8),  
237 CXCL10 (IP-10), CCL2 (MCP-1), CCL3 (MIP-1A), and CCL4 (MIP1B) (Xiong et al., 2020).

238 CXCL10 (IP-10) is a chemoattractant for monocytes/macrophages, dendritic cells, NK cells, and  
239 T cells; CCL2 (MCP-1) is a chemoattractant for monocytes, dendritic cells, and memory T cells.  
240 CXCL2 and CXCL8, which are secreted by monocytes/macrophage, serve as potent  
241 chemoattractants for neutrophils. Single cell RNA sequencing of nasopharyngeal and bronchial  
242 samples from COVID-19 patients identified increased inflammatory macrophages that express  
243 CCL2, CCL3 (MIP-1A), CCL20, CXCL1, CXCL3, CXCL10 (IP-10), CXCL8 (IL8), IL1B and  
244 TNF- $\alpha$  (Chua et al., 2020). Levels correlated with disease severity. CXCL10 (IP-10) levels were  
245 previously associated with the severe acute respiratory syndrome (SARS) disease progression  
246 and resolution due to the SARS-CoV virus (Altara et al., 2016; Jiang et al., 2005), and  
247 development of ARDS in preclinical models (Coperchini et al., 2020). The elevated  
248 nasopharyngeal levels of CXCL10 with COVID-19 may permit this chemokine to be used in  
249 widespread immunoassay testing for early detection of SARS-CoV-2-infection (Cheemarla et al.,  
250 2020).

251

### 252 **3.5 Possible contribution of GM-CSF**

253 Mounting evidence suggests that immunomodulatory agents, including GM-CSF, could  
254 be a promising therapy for COVID-19 (Lang et al., 2020; Mehta et al., 2020b). GM-CSF is  
255 known to be implicated in the production of granulocytes, monocytes, macrophages, and  
256 dendritic cells from progenitor cells, a process known as myelopoiesis (Egea et al., 2010;  
257 Fleetwood et al., 2007). It has been demonstrated that GM-CSF is secreted by different cell types  
258 including alveolar type II epithelial cells, playing therefore a key role in the integrity of alveolar  
259 barriers and maturation of alveolar macrophages (Cakarova et al., 2009; Rösler and Herold,  
260 2016). Multiple investigations have considered GM-CSF as a pivotal cytokine that activates both

261 the innate and adaptive immune response. For instance, GM-CSF can polarize myeloid cells into  
262 a pro-inflammatory phenotype, releasing subsequently reactive oxygen species and pro-  
263 inflammatory cytokines such as IL-1 $\beta$ , IL-6, TNF- $\alpha$ , and chemokines including CCL17, CCL2,  
264 and IL8, which can attract lymphocytes, monocytes, and neutrophils to the site of inflammation  
265 (Hamilton, 2020). It has also been reported that GM-CSF can prime dendritic cells to activate T  
266 cells, boosting thereafter the immune response by enhancing the recruitment of myeloid cells to  
267 the site of injury (Cao et al., 2015; Komuczki et al., 2019; Zhang et al., 2013). Since the goal of  
268 enhancing lung tissues integrity and dampening hyper-active immune response may lead to a  
269 drastic decrease in morbidity and mortality rate in COVID-19 patients, administration of GM-  
270 CSF as a promising therapy is being clinically investigated (Lang et al., 2020). Pre-clinical  
271 investigations revealed that overexpression of GM-CSF decreased apoptosis in alveolar wall  
272 cells, consequently preventing hyperoxia-induced lung damage (Baleeiro et al., 2006; Paine III et  
273 al., 2003). A clinical study performed by Matute-Bello et al. reported that in patients ARDS,  
274 increased GM-CSF in bronchoalveolar lavage fluid was associated with decreased mortality rate  
275 through potentially improved alveolar macrophage survival (Matute-Bello et al., 2000). This  
276 observation was further strengthened with a clinical study completed by Herold et al. showing  
277 that administration of inhaled GM-CSF to patients with pneumonia-associated ARDS enhanced  
278 oxygenation and lung compliance (Herold et al., 2014). Currently, a clinical study is assessing  
279 the potential beneficial effect of using inhaled and intravenous GM-CSF agonist in respiratory  
280 failure COVID-19 patients (Movers et al.).

281 The potential benefits of administering GM-CSF agonist in the context of COVID-19  
282 patients, however, should be carefully studied, particularly in the late stage of COVID-19 where  
283 lung injury is thought to be driven by the cytokine storm rather than viral overload (Siddiqi and

284 Mehra, 2020). Paradoxically, considerable interest in administrating anti-GM-CSF is gaining  
285 interest in the setting of COVID-19, given that a marked increase in GM-CSF expressing natural  
286 killer, B cells, and CD<sup>+</sup> 4 and CD<sup>+</sup> 8 T cells was observed in COVID-19 ICU patients when  
287 compared to mild cases (Zhou et al., 2020). However, given the role of GM-CSF in boosting the  
288 immune response to remove pathogen and enhancing lung repair, it is important to consider that  
289 the observed increase could be a result of exacerbated COVID-19 severity and related  
290 comorbidities. The rationale is that during COVID-19 infection, over-activation of myeloid cells  
291 could be a critical mediator of enhanced cytokine storm, consequently aggravating tissue  
292 damage. Therefore, anti-GM-CSF therapy may decrease the detrimental immune response, and  
293 thus exert beneficial effects (Barnes et al., 2020; Mehta et al., 2020a; Merad and Martin, 2020a),  
294 a hypothesis that was supported by a preclinical study of SARS-CoV infection animal model,  
295 showing that GM-CSF mediated the infiltration of inflammatory monocytes/ macrophages into  
296 the lungs (Channappanavar et al., 2016). Taking together, these findings suggest that GM-CSF is  
297 a key player in regulating myeloid cell induced hyper-inflammation in many tissues including  
298 the lungs. Anti-GM-CSF approach in patients with COVID-19, however, should be well  
299 monitored, given the critical contribution of GM-CSF in alveolar macrophage function and  
300 pathogen clearance.

301 As of the start of May 2020, there were some 49 clinical trials underway targeting the  
302 cytokine storm in COVID-19 patients (Wang et al., 2020b). The vast majority involve  
303 biologicals. Besides those involving GM-CSF, prominent among them are a number of studies  
304 involving anti-IL-6 strategies. In addition, antagonistic antibodies directed against TNF, IL-1,  
305 IL-1R, and IL-8 are being investigated for attenuating excessive immune activation and the  
306 cytokine storm (Conti et al., 2020b). The rationale behind those targeting the actions of GM-CSF

307 latter in COVID-19 is that this cytokine constitutes an autocrine/paracrine positive feedback loop  
308 that helps drive the cytokine storm (Mehta et al., 2020c). In a preliminary study, dexamethasone  
309 showed promise in reducing mortality of hospitalized COVID-19 patients if they were receiving  
310 respiratory support (mechanical ventilation or oxygen) (Group et al., 2020), but targeting the  
311 cytokine storm via broad-spectrum immunosuppression does raise a number of concerns  
312 (Theoharides and Conti, 2020).

313

### 314 **3.6 Possible contribution of the renin angiotensin system**

315 SRS-CoV-2 can gain entry into monocytes/macrophages via ACE2, although the virus is  
316 not thought to replicate in these cells. In this way, macrophages may act as a sort of “Trojan  
317 horse”, allowing for the delivery of the virus to lung and other tissue parenchyma (Abassi et al.,  
318 2020). ACE2 is a protease that forms part of the beneficial counterpoint to the renin-angiotensin  
319 system (Forrester et al., 2018). By removing the carboxy-terminus amino acid, it converts the  
320 vasoconstrictive and pro-inflammatory octapeptide angiotensin II (Ang II) to Ang (1-7), which  
321 has beneficial effects including vasodilation and anti-inflammation actions via the Mas receptor.

322 An additional consequence of virus-mediated ACE2 loss might be increased Ang II  
323 inflammatory effects via the Ang II type 1 (AT1) receptor or diminished protective signaling via  
324 the Mas receptor (Abassi et al., 2020). Although multiple studies reported increased ACE2  
325 expression in COVID-19 patients who are on angiotensin converting enzyme inhibitors (ACEIs)  
326 and angiotensin II receptor blockers (ARBs) (Ferrario et al., 2005; Igase et al., 2008), recent  
327 emerging investigations suggested that ACEIs and ARBs could exert protective effects through  
328 up-regulating ACE2, modulating negatively therefore the severity of COVID-19 (Kuba et al.,  
329 2005) and reversing the marked increase in Ang II levels, decreasing consequently its deleterious

330 effects on the cardiopulmonary system (Danser et al., 2020; Sommerstein et al., 2020; Zheng et  
331 al., 2020). A study done by Kuba et al. showed that the administration of exogenous ACE2 to  
332 ARDS animal model substantially decreased inflammation and enhanced oxygenation (Kuba et  
333 al., 2005). Similarly, epidemiological studies revealed that ACEIs and ARBs decreased the risk  
334 of pneumonia in general population (Liu et al., 2013; Shinohara and Origasa, 2012). Therefore,  
335 investigation aimed at testing the potential beneficial or detrimental effects of ACEIs and ARBs  
336 in the context of COVID-19 is being undertaken (Buckley et al., 2020).

337

#### 338 **4. Conclusions**

339 Substantial evidence indicates that pro-inflammatory macrophages play a critical role in  
340 the pathological consequences of COVID-19. Additional evidence is needed concerning the  
341 presence phenotype of these cells. Nor is it clear what the relationship is between SARS-CoV-2  
342 infection and monocyte/macrophage activation status, namely whether these immune cells are  
343 simply responding to the viral infection or are hijacked by the virus to act in an uncontrolled  
344 rogue manner. Emerging evidence indicates that targeting the cytokines and chemokines  
345 associated with their activation or restoring their innate immunity control may provide the means  
346 to successfully combat COVID-19.

347

348

349 **Dedication**

350 This manuscript is dedicated to G. Warren and Jessie Booz, two gentle, loving, caring, and gifted  
351 individuals, and all of those wonderful and remarkable individuals who were taken from us way  
352 too soon by the COVID-19 pandemic.

353

354

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723 **Figure Legend**

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725 **Figure 1** – Macrophages at the center of the cytokine storm. With inflammation, macrophages, T  
726 cells, endothelial cells and a number of other immune and mesenchymal cells, produce the  
727 monomeric glycoprotein granulocyte-macrophage colony-stimulating factor (GM-CSF) (red  
728 arrows). Besides stimulating the production of granulocytes and monocytes, GM-CSF can serve  
729 as a chemoattractant for the migration of monocytes and neutrophils into the tissue (blue arrows),  
730 and can alter neutrophil receptors. GM-CSF signaling promotes a pro-inflammatory M1  
731 macrophage phenotype and the production of a number of inflammatory cytokines and  
732 chemokines by monocyte-derived or tissue macrophages (black arrows). Macrophages  
733 themselves are direct targets of the SARS-CoV-2 via expression of the receptor for viral binding  
734 ACE2, as well as TMPRSS2 or a furin-like proprotein convertase. The effect of SARS-CoV-2 on  
735 macrophage phenotype is not defined, although inhibition of protective interferon signaling is  
736 reported. Lung macrophages also express the G protein-coupled alpha 7 nicotinic receptors  
737 (nAChRs  $\alpha 7$ ) that signal through JAK-STAT3 and oppose inflammatory signaling by blocking  
738 the translocation of p65/p50 NF- $\kappa$ B into the nucleus upon I $\kappa$ B $\alpha$  (inhibitor of NF- $\kappa$ B)  
739 degradation. See text for additional details. Some of the content is adapted from Servier Medical  
740 Art (<https://smart.servier.com/>).

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