COVID-19: unravelling the clinical progression of nature’s virtually perfect biological weapon

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Abstract: Coronavirus disease 2019 (COVID-19) pandemic has shocked the world and caused morbidity and mortality on an unprecedented level in the era of modern medicine. Evidence generated to-date on the virulence and pathogenicity of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) suggests that COVID-19 may be considered a perfect storm, caused by a nature’s virtually perfect biological weapon. This conclusion is supported by an updated analysis of pathogenesis and clinical progression of this infectious disease. It is now readily apparent that COVID-19 is not a clear-cut disorder, but is instead a gradually evolving pathology, characterized by a series of stages sustained by different molecular and biological mechanisms. The disease can hence be divided in at least five different phases (incubation, respiratory, pro-inflammatory, pro-thrombotic, and death or remission). Whilst the virus triggers direct cytopathic injury during the initial stage of illness, in the following evolving phases, it is the host itself that undergoes an almost suicidal reaction, sustained, amplified and maintained by the immune, complement and hemostatic systems. Another peculiar property making SARS-CoV-2 a devious and vicious pathogen is the biophysical structure of its receptor binding domain, which needs to be primed by human proteases, thus being less efficiently targetable by the host immune system. The unique pathophysiology of COVID-19 requires the customization of therapy by individual patient characteristics and according to the phase-specific, evolving derangement of the multiple biological pathways.

Keywords: Coronavirus disease 2019 (COVID-19); coronavirus; pathogenesis; inflammation; coagulation

Introduction

Coronavirus disease 2019 (COVID-19) pandemic has shocked the world and caused morbidity and mortality on an unprecedented level in the era of modern medicine. Caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), this pathology has already affected millions of people worldwide, causing nearly four hundred thousand deaths to-date (1). Similar to the other two previous coronavirus outbreaks, severe acute respiratory syndrome (SARS) in 2002 to 2003 and Middle-East respiratory syndrome in 2012, COVID-19 may be characterized by a rather unfavorable, often critical, clinical progression. The global death rate of COVID-19 is now approaching 7% according to the regularly updated statistics from the World Health Organization (WHO) (1), with some notable
outliers (e.g., as high as 13.8% in Italy) due to the influence of some demographic, clinical and environmental factors impacting disease progression (2). Based on recent evidence on the virulence and pathogenicity of SARS-CoV-2, the clinical progression of COVID-19 can be divided into as many as five different phases (Figure 1). In the following sections of this article, we will describe the different characteristics of each phase of COVID-19.

Phase 1: incubation

This is the very first phase, which starts from infection up to the first onset of severe symptoms and usually lasts between 2 and 11 days (mean incubation time: 6 days), with patients likely to be infectious 1–3 days before symptom onset (3). Although the true rate of individuals who will remain asymptomatic, or only mildly symptomatic, until terminal viral shedding is still unknown, some evidence suggests that the number could be as high as 50% (4). Importantly, this rate may be even underestimated due to under-testing or under-reporting (4). It seems now reasonable to hypothesize that this pre-symptomatic phase is perhaps the most critical for containing the outbreak. Interestingly, the viral load of asymptomatic, pre-symptomatic, or mildly symptomatic subjects is comparable to that of patients with overt disease (5). This highlights the significant risk of viral transmission throughout this first phase. Evidence suggests that 50–80% of all cases may be attributed to transmission from an asymptomatic or pre-symptomatic individual (6). As such, the relatively long incubation time and the considerably high rate of asymptomatic-paucisymptomatic individuals combine to explain the pandemic and the rise in cases despite public health intervention and public awareness.

Phase 2: progressive respiratory involvement

The second phase, which involves ~30% of all SARS-CoV-2 infected subjects (7-9), is characterized by the onset of the typical COVID-19 symptoms. These symptoms basically include, but are not limited to, fever in ~85% of all patients, dyspnea (~70–80%), cough (~40–80%), olfactory and gustatory dysfunction (~50%), myalgia (~30%), gastrointestinal symptoms (~10–30%) (7-9). The typical symptoms of upper respiratory infections such as sore throat, nasal congestion and rhinorrhea are only present in a minority of patients (<20%) (7-9). The high prevalence of dyspnea in these patients reflects the progressively severe respiratory involvement, which is variably associated with chest computed tomography imaging with ground-glass

![Figure 1 Phases, clinical progression, management and available therapies of COVID-19. ARDS, acute respiratory distress syndrome; SIRS, severe inflammatory response syndrome.](image-url)
Although the precise mechanisms underlying the onset of this disproportionate host response against the virus remain partially elusive, it has now been acknowledged that SARS-CoV-2 infection of dendritic cells and cells of the monocyte/macrophage lineage triggers their activation and active secretion of a vast array of pro-inflammatory cytokines such as interleukin (IL)-6, IL-2, IL-7, monocyte chemoattractant protein 1 (MCP-1), macrophage inflammatory protein 1-α (MIP 1-α), granulocyte colony stimulating factor (GSF), C-X-C motif chemokine 10 (CXCL10) and tumor necrosis factor-α (TNF-α) (13,14). The renin-angiotensin-aldosterone system (RAAS) plays also a very relevant role in this phase. The binding of SARS-CoV-2 to its receptor angiotensin-converting enzyme 2 (ACE2) at the surface of host cells may be associated with profound derangement of RAAS, culminating in the increased activity of angiotensin II (Ang II) and decreased activity of angiotensin 1,7 (Ang 1,7), thus fostering vasoconstrictive, inflammatory, oxidative and fibrotic injuries (15).

The interplay between the virus and the host immune system may be even more complicated as evidence suggests that SARS-CoV-2 is capable of infecting CD4+ and CD8+ T lymphocytes, thus producing direct cytopathic effects and further contributing to generate a completely dysregulated cytokine pattern (16). Importantly, evolving lymphopenia may also be associated with a defect in antiviral and immune regulatory immunity (17). This highly abnormal immunoinflammatory response has now been universally defined as “cytokine storm” and is accompanied by a variety of focal injuries in multiple organs and tissues. More specifically, lung histopathology reveals the presence of activated pneumocytes, capillary congestion, inflammatory infiltrates, endothelial injury, protein-enriched interstitial edema and diffuse alveolar damage, a clear picture reflecting the development of acute respiratory distress syndrome (ARDS) (18). Notably, at this stage of disease, some typical cytopathic signs attributable to direct viral injury and inflammatory infiltrates can also be found in many other organs and tissues, such as the heart, kidney, liver, and intestine.

**Phase 3: focal and systemic inflammation**

The third phase, which develops in around 15% of all SARS-CoV-2 infected subjects (7-9), is perhaps the most challenging and intriguing from a physiopathological perspective. In fact, whilst the respiratory phase is mostly attributable to direct cytopathic lung injury caused by viral replication in pulmonary parenchyma, the pro-inflammatory phase is instead characterized by an abnormal, almost exaggerated, host reaction against the pathogen, either locally (i.e., in the lung) or systemically, thus mimicking the pathogenesis of severe sepsis and severe inflammatory response syndrome (SIRS) (12). Although the precise mechanisms underlying the onset of this disproportionate host response against the virus remain partially elusive, it has now been acknowledged that SARS-CoV-2 infection of dendritic cells and cells of the monocyte/macrophage lineage triggers their activation and active secretion of a vast array of pro-inflammatory cytokines such as interleukin (IL)-6, IL-2, IL-7, monocyte chemoattractant protein 1 (MCP-1), macrophage inflammatory protein 1-α (MIP 1-α), granulocyte colony stimulating factor (GSF), C-X-C motif chemokine 10 (CXCL10) and tumor necrosis factor-α (TNF-α) (13,14). The renin-angiotensin-aldosterone system (RAAS) plays also a very relevant role in this phase. The binding of SARS-CoV-2 to its receptor angiotensin-converting enzyme 2 (ACE2) at the surface of host cells may be associated with profound derangement of RAAS, culminating in the increased activity of angiotensin II (Ang II) and decreased activity of angiotensin 1,7 (Ang 1,7), thus fostering vasoconstrictive, inflammatory, oxidative and fibrotic injuries (15).

The fourth progressive stage of COVID-19, that appears cumulatively the most severe, tends to develop in around 5% of all SARS-CoV-2 infected subjects (7-9), and is characterized by the onset of microvascular and macrovascular thrombosis fostered by an exaggerated focal and/or systemic inflammation (19). The pro-thrombotic effects of hyperinflammation in veins and arteries are now widely recognized, and encompass a certain number of pathogenetic mechanisms (endothelial dysfunction, plaque instability, platelet hyper-activation, activation of complement system and blood coagulation, hypo-fibrinolysis), which have been comprehensively reviewed elsewhere (20,21). The final result is the development of microvascular and macrovascular thrombosis, the former being virtually commonplace in all COVID-19 patients progressing to this fourth stage, and reflected evidence at lung histopathology of large inflammatory infiltration, microvascular thromboembolism and complete destruction of alveolar septae (18). In fact, it may be suspected that microvascular thrombosis begins early in the disease course and contributes significantly to lung injury leading to progression to severe illness. As such, appropriate anticoagulant or fibrinolytic therapy earlier in disease course may prevent progression to phase 4.

Macrovascular complications can also be frequently observed in severely ill COVID-19 patients, manifesting as deep vein thrombosis (DVT), pulmonary embolism (PE), in situ pulmonary thrombosis and/or acute coronary syndrome (ACS) (22-25). Even in younger patients (<50 years old), significant thrombotic events have been reported, including large vessel stroke (originating from carotid and cerebral arteries) (26). Notably, up to one-third of COVID-19 patients who die have evidence of pulmonary thrombosis as a
Besides the complex and mechanistic interplay responsible for direct and indirect host injuries, SARS-CoV-2 has another peculiar property that makes it a devious and vicious pathogen; the biophysical structure of its spike (S)-protein receptor-binding domain (RBD) is highly ingenious. Unlike its homologous precursor, which caused the SARS outbreak and that has now been renamed SARS-CoV-1, the RBD of SARS-CoV-2 displays ineffective receptor binding at rest (32) because it needs to be primed by human proteases such as the transmembrane serine protease 2 (TMPRSS2) or furin (32,33). The S1 subunit on the S-protein RBD is dissociated from the S2, which facilitates the fusion with the host cell membrane (34). Overall, the RDB of SARS-CoV-2 seems to be less efficiently targetable by the immune system, which may struggle to arrange an efficient immune response against the virus. This theory is confirmed by evidence that the process of antibody generation in the course of SARS-CoV-2 infection is atypical and unique. IgM, the first class of immunoglobulins that typically appears in response to infectious pathogens, may be absent in 20–40% of SARS-CoV-2 infected individuals (35), has a typically low titer (36,37), and, importantly, does not specifically target the S-protein RBD, which makes it theoretically ineffective at neutralizing the virus. The capacity to eradicate the infection may hence be delayed until a more specific, robust, and permanent immune response will emerge, specifically encompassing the generation of IgG immunoglobulins (38). Finally, it remains unknown whether long term protective immunity is afforded post-recovery by antibody production and immunological memory.

Conclusions

Irrespective of the origin and intermediate hosts of SARS-CoV-2 (39), the evidence garnered to-date on its virulence and pathogenicity would lead us to conclude that COVID-19 is the portrait of a perfect storm (40) caused by nature’s virtually perfect biological (lethal) weapon. Whilst in the initial stage of illness the virus causes direct cytopathic injury, in the following evolving phases, it is the host itself that undergoes an almost suicidal reaction, sustained, amplified and maintained by the immune, complement and hemostatic systems. It is now readily apparent that COVID-19 is not a clear-cut disorder, but is instead a gradually evolving pathology characterized by a series of stages sustained by different molecular and biological mechanisms, whose understanding is essential for establishing the most appropriate, phase-specific treatment. COVID-19 would appear as a paradigm example...
of personalized medicine, whereby the therapy needs to be tailored not only to individual characteristics (e.g., male sex, older age, presence of co-morbidities), but also according to the evolving derangement of multiple biological pathways.

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Footnote

Conflicts of Interest: All authors have completed the ICMJE uniform disclosure form (available at http://dx.doi.org/10.21037/atm-20-3989). GL serves as an unpaid executive editor-in-chief of Annals of Translational Medicine from Jan 2016 to Jan 2022. FSG serves as an unpaid editorial board member of Annals of Translational Medicine from Mar 2019 to Feb 2021. BMH has no conflicts of interest to declare.

Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved.

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