Review of immune responses correlated with COVID-19 outcomes: the fight, debacle and aftermath in the Indonesian context.

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ABSTRACT

In the current pandemic, the highly contagious nature of the severe acute respiratory syndrome coronavirus type 2 (SARS-CoV-2) leads to an enormous burden for the global health care system and creates challenging socioeconomic problems. Respiratory mucosa, the main entrance of SARS-CoV-2 infection, are equipped with an innate immune defense system as the initial response against infection. Activation of the adaptive immune system facilitates viral clearance as well as providing immunological memory for prevention from subsequent exposure. However, despite repeated efforts at implementing appropriate interventions, severe and fatal cases are continuing to occur and reports of recurrent cases need clarification. Host factors may contribute to the severity of the diseases while viral immune evasion is a common phenomenon leading to severe outcomes and recurrent infection. Discussions of immunological-based tests for screening, herd immunity, along with the possible advantages or potentially futile efforts of development of vaccine and alternative immunotherapy have become a part of daily household conversations. In this review, evidence of innate and adaptive immune responses or lack of them, and immunological problems relevant for SARS-CoV-2 will be summarized. Finally, perspectives for future studies especially in the Indonesian population will be sketched.

Keywords: immune response; SARS-CoV-2; COVID-19; pandemic; Indonesia;

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INTRODUCTION

The 21st century has already seen two outbreaks of coronaviruses. The first on, severe acute respiratory syndrome-coronavirus (SARS-CoV) in 2002 and the second, Middle East respiratory syndrome coronavirus (MERS-CoV) in 2012. Now in 2020 we are in the middle of another viral outbreak called severe acute respiratory syndrome-coronavirus (SARS-CoV-2). The recent emergence of SARS-CoV-2 infection started in the city of Wuhan in early December 2019, hence the disease is called coronavirus disease 2019 (COVID-19). The disease has now become a global pandemic. As of April 28th, 2020 the World Health Organization (WHO) database recorded over 2.9 million confirmed cases globally, including over 200,000 deaths in more than 200 countries.

Coronaviruses, the largest known positive-sense single-stranded ribonucleic acid (ssRNA) viruses, belong to ordo Nidovirales. The family is classified into three groups, based on antigenic relationships of the spike (S), membrane (M) and nucleocapsid (N) proteins and further viral genetic phylogeny. SARS-CoV, MERS-CoV and SARS-CoV-2 are classified in one of the groups. As the name implies, these viruses cause more severe manifestations compared to members of the other group.\(^1\)\(^-\)\(^3\)

The virus particle has a spherical shape that is 60-140 nm in diameter with distinctive spikes about 8 to 12 nm in length.\(^4\) SARS-CoV-2 genomic sequences are closer to those in bats, with similarities around 96%. The sequences analysis showed that the genome of the viruses from different patients are very conserved,\(^5\)\(^,\)\(^6\) suggesting a recent human virus. Phylogenetic network analysis reported by Forster et al.\(^7\) showed 3 clusters of SARS-CoV-2 based on amino acid changes: the A, B, and C clusters. A is the ancestral type, B derived from A, and further derives into C. Clusters A and C are found outside East Asia, while B is common in East Asian countries.

SARS-CoV-2 is transmitted from human to human through droplets containing the virus particles via the mucous membranes of the eyes, mouth or nose. However, there are no reports on the presence of the viral RNA from COVID-19 patients with conjunctivitis. Some studies reported virus shed from the stool (1-2%) in the late phase of infection,\(^8\) suggesting multiple transmission routes of the virus.\(^9\) The receptor binding site of SARS-CoV-2 and SARS-CoV is similar, with variations in some key amino acids.\(^6\) Infection of SARS-CoV-2 to mucosal epithelium depends on the binding of S1 protein to the host's angiotensin-converting enzyme (ACE)-2 protein (FIGURE 1), which is expressed in type II pneumocytes among other cells. Viral entry requires S protein priming by cellular protease, the transmembrane protease, serine 2 (TMPRSS2), which causes the cleavage of S1/S2 and S2' sites. The virus attachment further allows the fusion of virus with the cellular membrane. Similar capacity was already shown by SARS-CoV,\(^10\) but several amino acid differences in the S2 protein cause more efficient infection capacity of SARS-CoV-2 compared to the SARS-CoV.\(^11\)

Compared to the other 2 SARS-related coronavirus, SARS-CoV-2 infection results in milder symptoms. Three stages of SARS-CoV-2 infection are incubation and asymptomatic period with or without detectable virus, detectable virus with common cold and influenza-like illness, and variety of SARS such as pneumonia with high viral load.\(^12\) The first and second stages of infection require decent immune capacity. Once innate and adaptive immune responses breakdown, the virus will propagate and massive destruction of the affected tissues will occur, especially in organs that have high ACE2 expressions such as lungs, intestines and kidneys.\(^13\) As
mentioned earlier, the virus may enter via several routes other than respiratory tract, e.g. digestive tract and olfactory nerves lead to various symptoms.

The immune system plays an important role in eliminating the virus, yet it may also be involved in the disease pathogenesis that leads to severe morbidity and mortality. We are aware that the pathogenesis of SARS-CoV-2 infection may be more complex and related to other systems such as compromising the function of hemoglobin and blood coagulation. However, those topics are beyond the scope of this review. We begin with the main question related to what do we know about the innate and adaptive immune responses to SARS-CoV-2, the host factors that affect immune response, the possibility of viral immune evasion, and the responses of the immune system in fatal cases. Then, we conclude with possible immunological-based therapeutic and prophylactic approaches. Further, we would like to briefly discuss the immune consequences of the pandemic through the perspectives of Indonesian patients' characteristics.

As a novel and very contagious human coronavirus that prevents proper immediate investigation, reliable data on the immune responses against SARS-CoV-2 are still limited. Most studies were speedy reports with limited number of patients or case reports. This minireview is based on the latest publications concerning SARS-CoV-2, in addition to the recent knowledge related to SARS-CoV reports. Some studies using animal models infected with coronavirus or other virus were included in order to explain the possible mechanisms of the immunopathogenesis. Despite being published in reputable journals, many of our recent references (year 2020, in particular) may not be peer-reviewed, due to the sense of urgency to combat SARS-CoV-2 globally. We searched our references by several key words, including: COVID-19 OR SARS-CoV-2 OR SARS-CoV, and immunity* [AND term/specific topic] in PubMed, Google Scholar and World Health Organization (WHO) databases.

**DISCUSSION**

Appropriate innate and adaptive immune responses against coronavirus in mild cases

Upon coronavirus infection, the infected cells detect viral pathogen-associated molecular patterns (PAMPs) via their pattern recognition receptors (PRRs) such as Toll-like receptors (TLRs) (FIGURE 1). As a single-stranded (ss) RNA virus, the principal TLRs that recognize coronavirus nucleic acid are TLR7 and TLR8. In addition, retinoic-acid-inducible gene I (RIG-I) is also able to recognize them. During their replication, coronaviruses generate double-stranded (ds)RNA intermediates which are recognized by melanoma differentiation-associated gene 5 (MDA5). The recognition of viral nucleic acid by PRRs occurs in endosomes, endocytolysosomes and cytoplasm. MyD88 acts as an adapter protein by most TLRs. Coronavirus infection mainly activates the transcription factors such as nuclear factor-kappa beta (NF-κB) and mitogen-activated protein kinases (MAPKs) pathways to induce host responses such as type I interferon (IFN) and inflammatory factors.

Type I IFN via interferon-α/β receptor, in turn, activates the JAK-STAT pathway that initiates the transcription of IFN-stimulated genes under the control of IFN-stimulated response elements containing promoters. As the host's major antiviral molecules, IFNs limit viral spread, and play an immunomodulatory role to promote macrophage phagocytosis, as well as natural killer (NK) cells restriction of infected target cells.
FIGURE 1. SARS-CoV-2 particle, viral entry and induction of innate immune response. Viral spike1 (S1) protein binds to host’s ACE-2 protein. The virus attachment further allows endocytosis of the viral particle. Innate immune system detects viral ssRNA via TLR7 and TLR8, further activates transcription factors such as NF-κB and MAPKs (not drawn) pathways to induce the expression of host’s pro-inflammatory cytokines. Virus ssRNA and double-stranded (ds) RNA generated as intermediate in virus replication, are also recognized by RIG-I and MDA5 which further induce the expression of type I IFN that leads to antiviral state, i.e. restricts virus replication and signals the viral invasion to adaptive immune system.\textsuperscript{3,10,16}

A successful accumulation of this type I IFN response should be able to suppress viral replication and dissemination at an early stage.\textsuperscript{3} However, transmission of this virus is reported from asymptomatic infected individuals. This situation is very likely to occur due to delayed early response of the innate immune response.\textsuperscript{17} Later on, the delayed type I IFN activation leads to influx of neutrophils and monocyte-macrophage and induces tissue inflammation and development of symptoms. The typical symptoms of COVID-19 are fever, cough, generalized malaise or myalgia. Some patients reported loss of smell and taste called anosmia and ageusia due to the inflammation in the olfactory mucosa and tongue, respectively. Laboratory findings indicated upregulated neutrophilia, with increased C-reactive protein (CRP) and interleukin-6 (IL-6).\textsuperscript{18,19} Some infected individuals remain asymptomatic and become silent cases.\textsuperscript{20}

Initiation of adaptive immune responses against SARS-related coronavirus begins with the activation of respiratory dendritic cells (rDCs) after engulfing the viral antigen. Furthermore, rDCs migrate to the draining (mediastinal and cervical) lymph nodes (DLN) and present processed antigens to naïve T cells. T cells then become activated, proliferate and migrate to site of infection (lung).\textsuperscript{21} Lymphopenia may occur as a result of lymphocyte drainage into the site of infection. In those cases, increases on the neutrophil/lymphocyte ratio is reported.\textsuperscript{18,19}

The crucial protective role of T cells immune response in coronavirus infections has been clearly documented in several animal models infected
with various coronaviruses. Cytokine microenvironments generated by antigen presenting cells direct T cells responses. In general, T H1 orchestrates the direction of T cells response followed by cytotoxic T lymphocytes (CTL) that are essential in killing viral infected cells. Intravenous immunization with peptide-loaded dendritic cells (DCs) followed by intranasal boosting with recombinant vaccinia virus (rVV) encoding single immunodominant epitope of SARS resulted in accumulation of virus-specific memory CD8+ T cells in bronchoalveolar lavage fluid, lungs, and spleen. Cytotoxic CD8+ T cells account for about 80% of total infiltrative inflammatory cells in the pulmonary interstitium in SARS-CoV-infected patients and play a vital role in elimination of CoV’s infected cells.

Strong evidence showed that T H1 response is a key for successful control of SARS-CoV and MERS-CoV, and probably for SARS-CoV-2 as well. Depletion of CD8+ T cells do not affect and delay viral replication at the time of infection with SARS-CoV, while depletion of CD4+ T cells is associated with reduced pulmonary recruitment of lymphocytes and neutralizing antibody and cytokine production, resulting in a strong immune-mediated interstitial pneumonitis and delayed clearance of SARS-CoV from lungs.

Strong T cell responses are significantly correlated with higher production of neutralizing antibodies that block the virus from entering healthy host cells and protect the host against viral re-infection. In patients infected with coronavirus, specific immunoglobulin M (IgM) peak occur on the 9th day after infection. Class switching to IgG, assisted by follicular helper T cells (T FH cells) can be detected as early as 4 days but most of the cases were seroconverting at the 2nd week after the onset of the disease. Delayed and weak antibody responses are associated with severe outcomes.

One report of a patient with mild symptoms revealed that increased plasma cells, T FH cells, activated CD4+ T cells, CD8+ T cells, IgM and IgG antibodies that bound to SARS-CoV-2 were detected in blood samples before symptomatic recovery. These immunological changes persisted for at least 7 days following full resolution of symptoms. From a study in Finland, IgM was reported to be detected on day 9 and IgG at day 20 after infection. Specific IgM started to decline at week 7. Immunoglobulin detection was employed as the basis of SARS-CoV rapid test. One study in Wuhan reported 10% SARS-CoV-2 specific IgG positive rate from asymptomatic individuals.

A cohort study of convalescent SARS-CoV patients from Beijing revealed that the concentration of specific serum IgG and neutralizing antibodies were highly correlated, reaching their peak at 4th month after the onset of disease and decreasing gradually thereafter. The titers decreased noticeably after the 16th month, but long-lasting specific IgG and neutralizing antibodies were reported in most of the patients as long as 2 years after. Only 11.8% of patients turned to negative at month 24. In long-term follow-up survivors, IgG was detectable in recovered patients only up to 6 years after SARS-CoV infection, suggesting decreasing level of memory B cells. Shorter antibody persistence may occur due to several factors such as high dose corticosteroid treatment that is known to suppress antibody response.

Coronaviruses immune evasion

Immune evasion is an important strategy utilized by pathogens to overcome the immune response for efficient replication in the host. Immune evasion by influenza virus A via antigenic shift and antigenic drift is a famous phenomenon. As an RNA virus, coronaviruses mutation rates are
higher than DNA viruses due to higher error rate of their RNA polymerases. The high mutation rate leads to higher probability for immune evasion in secondary exposure.

Coronaviruses have evolved several strategies of immune evasion. One of the main strategies used by SARS-CoV to avoid immune detection is by shielding its RNA from PRRs. This can be done by several mechanisms. One approach is by disguising its RNA and replication machinery inside replication organelles. Since concealing its RNA and replication machinery is not unassailable, SARS-CoV further modifies its RNA by adding cap-structure to the 5'-end or by methylating its mRNA to be mediated by SARS-CoV nsp16 protein. Another means is by activating its endonuclease, thus destructing its own RNA at certain stages of infection to avoid the initiation of host's virus-destroying system.

The other major strategies of immune evasion are to restrain the host's innate immune signaling. Coronavirus inhibits recognition by PRRs such as TLRs. Terminating cellular protein expression is one of the effective mechanisms to achieve that purpose. SARS-CoV nsp1 protein arrests the translation process by binding to the ribosomal 40S subunit and inducing host's mRNA degradation. Another mechanism is by manipulating stress granules formation. Stress granules are structures formed by the accumulation of untranslated mRNA as a consequence of the arrest of cellular translation due to virus infection. This accumulation of viral RNA can be detected by the host's immune sensors such as Protein Kinase R (PKR), RIG-I, and MDA5. Other mechanisms of suppressing innate immune signaling include histone modification, preventing phosphorylation and nuclear translocation of IFN-related transcription factors such as STAT1.

To avoid adaptive immune responses, coronaviruses inhibit viral antigen presentation to T cells. MERS-CoV can infect and replicate in macrophages and dendritic cells resulting in the cells malfunction and inability to present antigens to T cells. MERS-CoV also downregulates MHC I and II in macrophages and dendritic cells, thus leading to failure in activation of naïve T cells. MERS-CoV further stimulates T cells apoptosis such that lymphopenia will ensue. Deficiency of activated T and B lymphocytes are associated with persistent infections and lack of virus clearance.

**Immune system in fatal cases of COVID-19**

The mortality rate of SARS-CoV-2 infection is lower compared to the other SARS related coronavirus. However, due to the highly contagious nature of the virus, the number of fatalities is higher than the two previous outbreaks combined. An impaired immune response that fails to eliminate the virus will likely result in severe cases of morbidity and mortality (TABLE 1 and 2). Accordingly, medical researchers are asking the important question from the immunological point of view: What actually drives the unpredictable fatality rate of COVID-19?

In SARS-CoV infections, persistence of the viral infection and spreading induce activation of macrophages and thus, cause increased production of pro-inflammatory cytokines that leads to consistent influx of neutrophils and monocyte-macrophages. The increased cytokine production and associated CRP may cause the condition called as “cytokine storm” which presents in a vicious cycle fashion and may cause more damage to the cells of the lungs. Initial increment of CRP was suggested to be used to predict severe cases. Despite their purpose to fight off the invading pathogen, the inflammatory
cells can mediate further tissue damage by inciting lung epithelial and endothelial apoptosis via Fas-Fas ligand (FasL) or Tumor necrosis factor (TNF)-related apoptosis-inducing-ligand-Death Receptor 5 (TRAIL-DR5)-dependent mechanisms.\textsuperscript{48} Apoptosis compromises the lung microvascular and alveolar epithelial cell barriers resulting in vascular leakage and alveolar edema. Indeed, pathologic lung examination of two COVID-19 patients with adenocarcinoma revealed diffuse alveolar damage (DAD). Pulmonary edema, proteinaceous and fibrin exudates, mononuclear inflammatory cell infiltration with multinucleated giant cells, alveolar wall thickening with pneumocytes hyperplasia and fibroblast proliferation, without prominent hyaline membrane were observed in both patients. The absence of clinical symptoms of pneumonia indicates these pathological findings may depict early lung abnormality of COVID-19.\textsuperscript{49} Later on the DAD could lead to acute respiratory distress syndrome (ARDS), in which respiratory failure is characterized by the rapid onset of widespread inflammation in the lungs and subsequent fatality.\textsuperscript{50} At this late stage, extensive fibrin deposition (fibrosis) and alveolar collapse occur.\textsuperscript{51} Similar mechanisms manifested in lung immunopathology were also shown in mouse models of SARS-CoV infection.\textsuperscript{23} The ‘cytokine storm’ also occurs in SARS-CoV-2 infection particularly in severe cases.\textsuperscript{19,52} Furthermore, the cytokines can spill over to the circulation, resulting in systemic sepsis and multi organ dysfunction and failure\textsuperscript{53} including myocardial complication reported in many fatal cases.

Besides causing massive apoptosis and tissue injury, this ‘cytokine storm’ can diminish T cells response via TNF-mediated apoptosis. In patients with COVID-19, a more pronounced lymphopenia is observed in fatal cases,\textsuperscript{52,54} suggesting that COVID-19 may damage T lymphocytes. Furthermore, the numbers of CD4\textsuperscript{+} and CD8\textsuperscript{+} T cells were decreased,\textsuperscript{50,52,54} with naïve CD4\textsuperscript{+} T cells count elevated but that of memory CD4\textsuperscript{+} T cells decreased.\textsuperscript{52} Considering that the differentiation of naïve T cells into effector and memory cells is fundamental for the immune response, the higher naïve to memory ratio of CD4\textsuperscript{+} T cells in severe cases indicates an impaired and unbalanced immune response caused by SARS-CoV-2 infection. In addition, the production levels of IFN-γ by CD4\textsuperscript{+} T cells, CD8\textsuperscript{+} T cells, and NK cells tend to be lower.\textsuperscript{54} The production of IFN-γ is essential for resistance against infection of various pathogens and IFN-γ was reportedly able to inhibit viral replication of SARS-CoV.\textsuperscript{55} Lower IFN-γ levels in severe cases of COVID-19 might reflect the host’s inability to fight off the virus which increases the likelihood of fatality. Strong T cells’ responses are significantly correlated with higher neutralizing antibody, while more serum T\textsubscript{H}2 cytokines (IL-4, IL-5, IL-10) were detected in the fatal group of SARS-CoV.\textsuperscript{26} Another interesting effect that SARS-CoV-2 infection has on T lymphocytes is that it may induce T cells exhaustion. Lymphopenia was common in patients with SARS-CoV-2 (72\%) as well as in acute phase of patients with SARS-CoV (~80\%).\textsuperscript{12,21} At the infection onset, it was indicated by the drainage of lymphocytes into the infected tissue but in later stages it indicates the impairment of immune system. Beside the decline in T cells number, COVID-19 patients also showed increased expression of natural killer group 2 member A (NKG2A) in CD8\textsuperscript{+} T cells and NK cells.\textsuperscript{54} NKG2A is an inhibitory receptor, which induces immune cells exhaustion in chronic viral infection.\textsuperscript{56} Its increased expression in CD8\textsuperscript{+} T cells suggests functional exhaustion of cytotoxic lymphocytes in COVID-19 patients. The elevated level of interleukin-10 (IL-10) in severe cases of COVID-19 supports this plausible notion.
Although there is no evidence concerning the ability of IL-10 to increase NKG2A expression, however, IL-10 regulates inhibitory receptor (e.g. Programmed cell-death protein 1; PD-1) expression in CD8⁺ T cells⁵⁷ and induces sustained PD-1 expression and T cells exhaustion,⁵⁸ leading to a suppressive environment that prevents viral clearance.

<table>
<thead>
<tr>
<th>Immune responses to SARS-CoV-2 infection</th>
<th>Action</th>
<th>Dysregulation in severe cases of COVID-19</th>
</tr>
</thead>
<tbody>
<tr>
<td>Detection by PRRs (including TLR7, TLR8, RIG-I, MDA5)</td>
<td>Induces NF-κB and MAPK pathways to activate type I interferon system and other pro-inflammatory cytokines.</td>
<td>Coronavirus avoids PRRs detection by concealing its RNA inside replication organelles; adding cap structure or methylating its RNA; activating endonuclease to degrade its own RNA; halting host's translation process; destructing host’s mRNA; or inhibiting stress granules formation.²⁴⁻³⁹</td>
</tr>
<tr>
<td>Type I interferon</td>
<td>Has antiviral activity. IFN-stimulated genes inhibit virus entry, replication and dissemination.</td>
<td>Coronavirus prevents phosphorylation and nuclear translocation of IFN-related transcription factor, inhibit activation of IFN system.⁴¹</td>
</tr>
<tr>
<td></td>
<td>Enhances NK cells activity to kill infected cells and to secrete IFN-γ; promotes macrophage phagocytosis activity and pro-inflammatory cytokines production; stimulates DC maturation and antigen presentation to T cells; increases MHC I surface expression</td>
<td>Delayed or impaired antiviral response of type I IFN results in uncontrolled viral replication and dissemination.</td>
</tr>
<tr>
<td>Natural Killer (NK) cells</td>
<td>NK cells target and kill virus-infected cells and produce type I interferon cytokines.</td>
<td>Decrease IFN-γ and increase NKG2A expression in NK cells, indicating functional cells exhaustion and impaired virus-killing activity.⁵²</td>
</tr>
<tr>
<td>Macrophages/monocytes</td>
<td>Macrophages infiltrate infected sites; phagocytose apoptotic virus-infected cells; mediate antibody-dependent cell cytotoxicity; and secrete pro-inflammatory cytokines.</td>
<td>Hyper-influx of macrophages-monocytes into the lung that leads to apoptosis of lung epithelial and endothelial cells and tissue damage. Coronavirus also can infect macrophages, rendering it malfunction.⁴²⁻⁴⁸</td>
</tr>
<tr>
<td>Pro-inflammatory cytokines</td>
<td>Increase secretion of IL-1R, IL-6, TNF-α, IL-8, IP-10, MCP1, MIP-1α upon virus infection to recruit neutrophils, monocytes/macrophages, and T cells to infected site and augment their antiviral action.</td>
<td>Exaggerated release of pro-inflammatory cytokines or cytokine storm occurs; leads to further tissue damage, ARDS, systemic sepsis, and multi organ dysfunction.⁴⁹⁻⁵⁰</td>
</tr>
<tr>
<td>Neutrophil in tissue inflammation</td>
<td>Neutrophils migrate to infected sites; phagocytose apoptotic virus-infected cells; and secrete pro-inflammatory cytokines to recruit more immune cells.</td>
<td>Neutrophilia; with hyper-influx of neutrophils into the lung that leads to tissue damage.⁴⁹⁻⁵⁰</td>
</tr>
</tbody>
</table>

*Observed in SARS-CoV or MERS-CoV, however, no information available for SARS-CoV-2 yet. ARDS, acute respiratory distress syndrome. DC, dendritic cells. IFN-γ, interferon-gamma. IL-1R, interleukin-1 receptor. IL-6, interleukin-6. IL-8, interleukin-8. IP-10, interferon gamma-induced protein 10. MAPK, mitogen-activated protein kinases. MCP1, monocyte chemoattractant protein 1. MDA5, melanoma differentiation-associated gene 5. MIP-1α, macrophage inflammatory protein 1-alpha. MHC I, major histocompatibility complex class I. NF-κB, nuclear factor-kappa beta. NK, natural killer. NKG2A, natural killer group 2 member A. PRRs, pattern recognition receptors. RIG-I, retinoic-acid-inducible gene I. TLR7, Toll-like receptor-7. TLR8, Toll-like receptor-8. TNF-α, tumor necrosis factor-alpha.
TABLE 2. Adaptive immune response against SARS-CoV-2

| Immune responses to SARS-CoV-2 | Action                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           | Dysregulation in severe cases of COVID-19                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     |
|--------------------------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------- Adam 1,42,43 |
Humoral immune response against SARS-CoV-2 may also contribute to the severity of the disease. COVID-19 patients with higher IgG levels tend to have more severe disease symptoms compared to those with lower IgG levels. It has been proposed that IgG humoral response may contribute to tissue damage by a mechanism known as antibody-dependent enhancement (ADE). ADE has been confirmed in multiple viral infections. ADE occurs when antiviral neutralizing antibodies (Nab) cannot fully neutralize the virus. Instead, the virus-Nab complexes bind to the Fc Receptor (FcR) in macrophages, leading to viral endocytosis and infection of macrophages. This leads to persistent virus replication and exuberant pro-inflammatory cytokines signaling which eventually causes acute fatal lung injury, as has been observed in patients with SARS-CoV infection.

The impaired immune system in severe cases may increase the risk for bacterial co-infection. A study in China reported that 10% of the COVID-19 severe cases suffered from secondary infection. Tissue damage, exuberant inflammatory cell influx observed in cytokine storm phenomenon may render the pro-inflammatory response inadequate, e.g., lymphocytes recruitment to new infected site and macrophages phagocytosis capacity, when challenged with a bacterial pathogen, while enhancing bacterial entry or adherence to host's cell.

Host factors that influence immune responses against SARS-CoV-2

China, Southern Europe and American East Coast took turns as areas with the highest number of cases during the course of the pandemic. In several regions and countries, certain races and foreign-born individuals were disproportionately affected. Whether genetic and epigenetic factors are involved in increasing the morbidity and mortality risks in the most severely affected areas, needs further study. The 1918 H1N1 pandemic reported high mortality in indigenous communities in Australia, Alaska and South America that may be due to genetic factors or lack of cross-immunity. Currently, temperature and humidity as well as frequency of traveling and migration, population density and social background are considered as crucial factors in the SARS-CoV-2 spreading. Increasing reports have discussed older age, male and comorbidity as risks factors for COVID-19. However, detailed demographic data for COVID-19 cases are not yet available. Most of the government agencies have only reported the number of active cases, recovered and death cases without specific case characteristics.

From China, the first epicenter of the pandemic, reports confirmed that older people are more susceptible to SARS-CoV-2 than the younger group. Children and adolescents are less susceptible to SARS-CoV-2, probably due to the low expression of ACE2. The median of the patients' ages ranged from 41 to 57 year. Old age is known as a risk factor for infection as reported for SARS-CoV infection as well as a study in non-human primate infected with SARS-CoV. Prior to infection, CD8+ T cells and B cells were lower in aged monkeys and after infection CD8+ T cells and B cells responses were significantly reduced in the aged host. A study of mice models of West Nile Virus infection found the virus also predominantly infects the old population. Malnutrition that occurs more frequently in elderly may enhance vulnerability to infection during aging. The IgM and IgG response are commonly impaired and this was linked to delayed germinal center formation in the draining lymph node. Older patients showed severe symptoms at hospital admission.
of the patients in China that died were older than 50 years with median age of 70 years old. European countries also reported that the death rate was higher among senior citizens more than 70 years old. Worldometer in April 14, 2020 reported the data from New York that 47% of the patients that died were more than 70 years old. Despite the higher rate of comorbidities found in older individuals, the dysregulated immune response may contribute to the more severe effects in older individuals. This may be due to lower number of Treg, aberrant microbiota composition, and higher proportion of inflammatory cells during aging.

Many reports from SARS-CoV-2 infection in China revealed that 50–75% admission to the hospitals were male. Moreover, more men died with COVID-19 compared to women. The Guardian, April 16, 2020 edition reported that the number of men who died of COVID-19 in England and Wales was about twice the rate of women. In New York, Worldometer at April 14, 2020 revealed that male patients comprised 61% of the mortality from COVID-19 and even higher up to 72% in patients without underlying conditions. The data are in accordance with the predominantly male patients reported in previous case mortality rates of SARS-CoV and MERS-CoV. Despite social factors such as working outside home, healthcare seeking behavior, smoking habit and medical and biological factors such as higher proportion of comorbidities e.g. hypertension and diabetes as well as higher expression of ACE receptors that are correlated with males, some important aspects of immunological background still contribute to the higher proportion and severity of male patients with COVID-19.

The difference in immune response between sex was reported by many reports. Females have lower susceptibility to infection partly due to several genes encoded in X chromosomes. One of the genes is TLR-7 that detects viral RNA, resulting in higher production of antiviral substance type I IFN. Females have higher expression of TLR-7 that is partly mediated by estrogen signaling. Male mice were more susceptible to SARS-CoV infection compared with age-matched females and the degree of sex bias to SARS-CoV infection increased with advancing age. The X chromosome influences the immune system's response against viral infection by acting on many other genes, including TLR-8, CD40L and CXCR3 which can be over-expressed in women. Infection of mice with H1N1 induced antibody responses, CD4 T cells and CD8 T cells memory responses that were greater in females than males.

Males tend to have higher proinflammatory profiles than females, hence increasing the risk of excessive inflammation in severely affected cases. Many regulatory proteins in the immune system such as FOXP3, the transcription factor for Treg are encoded in X chromosomes. In women the production of inflammatory IL-6 after viral infection is lower than in males and is often correlated with a better longevity. A mice study on SARS-CoV infection showed a persistent increase of inflammatory monocytes and neutrophils in SARS-CoV infected male mice but not in female mice. The estrogen signaling in female mice is proposed to reduce the accumulation and function of macrophages in the mice lungs and have a protective effect against viral infection. These data are in accordance with several reports that showed sexual dimorphism in immune response to COVID-19 infection.

Several comorbidities that increase COVID-19 severity has been reported such as hypertension and diabetes that are partly based on the patients' medication or condition that can upregulate ACE2 expression. Higher expression of ACE2 not only permits more viral entrance but
is also augmenting immune response and proinflammatory cytokine production, which correlated with fatal cases.\textsuperscript{79} The impact of the unhealthy Western diet that leads to inadequate immune response and chronic inflammatory state in COVID-19 pathology has also been discussed recently.\textsuperscript{80} The complex pathogenesis of COVID-19 still needs to be elucidated, in which the effect of SARS-CoV-2 in erythrocytes and components of blood coagulation in addition to immune responses, plays an important role in the process. Several interesting observations such as coronaviruses have not been shown to cause a more severe disease in immunosuppressed patients.\textsuperscript{81} need more confirmation in the future.

**Immunological memory and herd immunity**

Reappearance of highly contagious viral infections such as SARS-CoV-2 is a crucial public health problem. Immunological memory is important for the effective response against pathogens that have been encountered previously. It generates immediate protection and greater magnitude of immune response. Whether memory T cells response is sufficient to protect from re-infection of SARS-CoV-2 needs further study. In SARS-CoV-2 infection, the detection of specific IgG against SARS-CoV-2 in the patients' sera\textsuperscript{28} provides early evidence for immunological memory. However, since the COVID-19 pandemic is still ongoing, data on the time-period of antibody peak level and the duration of the antibody detectability are not yet available on a large scale.

Neutralizing antibodies against S protein protect the susceptible host from lethal infection. However, several studies reported that anti-SARS-CoV antibody response is short-lived in patients who have recovered from SARS-CoV.\textsuperscript{23} IgG titer peaks at 3-4 months. It remains detectable 2 years after infection and starts to decline after 2 years.\textsuperscript{31} Immunological memory to one type of coronavirus may not provide protection against another type of coronavirus as shown in the absence of cross-reactivity of CD8\textsuperscript{+} T cells responses against SARS-CoV with MERS-CoV.\textsuperscript{82}

Few reports are available on memory T cells in SARS-CoV-2 infection. After the SARS-CoV outbreak, SARS-CoV-specific memory T cells had been screened in SARS convalescent patients.\textsuperscript{32} All memory T cells detected were directed at SARS-CoV spike (S), membrane (M) or nucleocapsid (N) protein. These memory T cells were found to last up to 6 years after infection\textsuperscript{32} with most of them identified as polyfunctional central memory T cells.\textsuperscript{26} Re-infection with SARS-CoV has demonstrated the protection provided by memory T cells. In immunized mice challenged with lethal doses of SARS-CoV, its virus-specific memory CD8\textsuperscript{+} T cells efficiently produced cytokines including IFN-\(\gamma\), TNF-\(\alpha\), IL-2 and cytolytic molecules, e.g. granzyme B. These responses prevented extensive viral replication and alveolar damages.\textsuperscript{23} Given the short-lived duration of antibody-specific-SARS-CoV, the long-lived memory T cells and its proven protection upon re-infection is of considerable significance. However, it is worth noting that a decrease in memory T cells count in SARS-CoV-2 infection with high naïve to memory T cells ratio\textsuperscript{52} might indicate impaired memory T cells induction.

Even though many uncertainties remain concerning immunological memory towards SARS-CoV-2, several countries endorse the idea of herd immunity. The common notion in herd immunity is that exposure of the virus to the population may induce immunological memory for most people. As more and more people become immune to the virus, an infected individual has less and less chance of coming into contact with an individual
susceptible to infection. Finally, herd immunity becomes pervasive which at that point, the number of cases will start to decline. If herd immunity is achieved, even in the absence of measures designed to slow or contain transmission, the virus will be contained.\textsuperscript{83}

However, there are many concerns regarding herd immunity in SARS-CoV-2. One concern has to do with the duration of the immunological memory and re-infection. Recently, SARS-CoV-2 patients who have been declared recovered by two negative tests for the virus later tested positive again.\textsuperscript{84} If these cases were indeed re-infection, it would cast doubt on the immunological memory developed by the infected individual. Another concern is the possibility of immune enhancement caused by the infection.\textsuperscript{59} If immunity against SARS-CoV-2 leads to exacerbation of the infection rather than prevention of it, more fatalities will emerge. Whether the virus is mutating over time is also an issue that needs to be addressed. If the virus is mutating even at a slow rate, what those changes will do is a bigger question. The mutations could make the virus more easily transmitted or better at evading our immune system. The main question regarding herd immunity in this current pandemic is what percentage of the population is needed to get infected to achieve herd immunity. As of today, it is very likely that the number of COVID-19 cases is underestimated. Also, considering the severity of the disease and its case fatality rate that varies among many countries, the idea of herd immunity may not be appropriate for the current situation.

**Preventive vaccination and immunotherapy against coronavirus**

Considering the significant protection conferred by memory cells and the fact that COVID-19 is still spreading globally with no specific treatment yet available, it is worth noting that probably vaccines offer the best options to control the current pandemic. By vaccination, those who have not been infected by SARS-CoV-2 will have antibodies which will readily generate protection with greater response against SARS-CoV-2. In addition, vaccination also provides protection against re-infection for those who have recovered from SARS-CoV-2 infection. The immediate availability of SARS-CoV-2 genomic sequence\textsuperscript{5} and the recent mapping of SARS-CoV-2 spike (S) protein\textsuperscript{85} may lay the foundation for more rapid development of a specific vaccine. From studies on SARS-CoV vaccines, the S protein on the surface of the virus is an ideal target for a vaccine.\textsuperscript{86}

According to the most recent report, currently 73 SARS-CoV-2 vaccine candidates are undergoing development at preclinical stages and 5 others are entering clinical trials in humans. These 5 vaccine candidates include mRNA-1273 (NCT04283461/Moderna), Ad5-nCoV (NCT04313127/CanSino Biologicals), INO-4800 (NCT04336410/Inovio), LV-SMENP-DC (NCT04276896/Shenzhen Geno-Immune Medical Institute), and pathogen-specific aAPC (NCT04299724/Shenzhen Geno-Immune Medical Institute). As of mid-March 2020, mRNA-1273 (Moderna) and Ad5-nCoV (CanSino Biologicals) have become two of the first SARS-CoV-2 vaccines to be tested on humans.\textsuperscript{87} Significant breakthroughs in vaccine technology have provided more vaccine development platforms. The platforms used in the development of SARS-CoV-2 vaccine range from RNA vaccine, DNA vaccine, viral vector-based vaccine, recombinant protein vaccine, live attenuated vaccine, and inactivated vaccine. Among these platforms, RNA vaccine design gives flexibility on antigen manipulation and improves protein translation efficiency which can induce a more robust immune response.\textsuperscript{88} Additionally, RNA vaccine
is more rapid and less expensive than traditional vaccines\textsuperscript{88} which would give some advantages and benefits especially during the pandemic situation as it is today.

Although there is urgency to provide a vaccine rapidly due to the current pandemic, governments and pharmaceutical companies need to ensure the safety of the vaccine. Lessons need to be learned from the experience of developing a vaccine for SARS-CoV. Several vaccines for SARS-CoV have been tested in animal models. Most of these vaccines protected animals from the challenges of SARS-CoV, but, many did not induce sterilizing immunity. In some cases, vaccination resulted in complications\textsuperscript{89} or led to enhancement of the disease.\textsuperscript{90} Up to this moment, there is no vaccine for SARS-CoV available.

Another aspect that needs to be considered for a more effective SARS-CoV-2 vaccine development is the decline of antibody response over time. Coronavirus infection does not always induce long-lived antibody response. An individual can be re-infected with the same virus after an extended period of time. In individuals that recovered form SARS-CoV infection, antibody titer starts to decline 2 years after illness onset.\textsuperscript{31} Recently, reports emerged that SARS-CoV-2 re-infection occurs days after recovery.\textsuperscript{84} Thus, a SARS-CoV-2 vaccine will need to tackle this issue to provide protection in case the virus becomes endemic and causes recurrent seasonal epidemics.

SARS-CoV-2 infection causes more severe morbidity and more mortality among older individuals. Therefore, it will be important to develop vaccines that offer protection to this population. Unfortunately older individuals typically respond less well to vaccinations because of immune senescence.\textsuperscript{91} This is another concern that needs to be addressed for SARS-CoV-2 vaccine development.

One of the simple therapeutic modalities for severe cases of COVID-19 is convalescent plasma. It has been used during the SARS outbreak\textsuperscript{92} and is currently used to treat COVID-19 patients.\textsuperscript{93} Other treatments that have been used during this ongoing pandemic are intravenous immunoglobulin (IVIG)\textsuperscript{94} and intravenous injection of mesenchymal stem cells.\textsuperscript{95} Since it is very likely that the ‘cytokine storm’ is the main cause of fatal cases of COVID-19, it is worth considering the use of a neutralizing antibody against pro-inflammatory cytokines involved in the pathogenesis of COVID-19. One of those is tocilizumab, a monoclonal Ab against IL-6R. In China, tocilizumab has been approved to treat COVID-19 patients with pneumonia and elevated IL-6 levels.\textsuperscript{96} Corticosteroid is another medication used to suppress inflammatory responses in COVID-19 patients.\textsuperscript{12,19} However, the lack of proven benefit in COVID-19 cases should render health care workers to be cautious on its usage. The strategy to suppress immune responses in severe cases is challenging as the main immune component that should be inhibited is currently not yet clearly identified. The timing to start the therapy without compromising the beneficial host defense is another question that needs an immediate answer. Furthermore, while the aforementioned therapies are currently used to treat a limited number of COVID-19 patients, evaluation and study on their efficacy are still required.

Heterologous immunity is thought to be a contributing factor that reduces severity of an infection in any age. Exposure to one pathogen may alter the subsequent innate and adaptive immune responses to a different pathogen. It will be interesting to study whether history of infectious diseases alters the vulnerability to suffer from severe COVID-19 via the heterologous immunity mechanism. The number of cases in some countries that has nationwide Bacille-Calmett Guerin (BCG) vaccination
program, are less numerous than other countries with no BCG vaccination program such as European countries, leading to the idea that BCG vaccination may give some protection from SARS-CoV-2 infection. This idea is being tested with two clinical trials, although last year a review showed some evidence against it.¹⁹⁷ Some alternative treatments related to immune function have been proposed for COVID-19 patients. Many doctors recognize the role of positive thinking, praying, meditation in the outcome of diseases, including COVID-19. Preliminary studies show a beneficial impact of vagal stimulation, a method that activate autonomic nervous system to suppress immune function on the recovery of COVID-19 patients.⁹⁸ Herbal medicine has also been used for COVID-19 treatment,⁹⁹ yet the efficacy as well as its mechanism of action in the immune system, if any still need more study. Any herbal medicine that may balance immune responses, activate appropriate responses against SARS-CoV-2 while preventing excessive inflammation as discussed for the famous Chinese herbal remedy called Astragalus¹⁰⁰ may be beneficial in the fight against this disease. The prospect of using Indonesian herbal medicine for balancing immune responses also needs to be investigated due to the region's biological diversity.

The aftermath

While currently not and hopefully never considered the epicenter of the outbreak, Indonesia is the 4⁴th largest population with a large variability of ethnic and socioeconomic backgrounds, hence increased vulnerability to any contagious pathogens. As of 2⁸th of April 2020, the WHO database recorded 9096 confirmed cases in Indonesia, including 765 deaths. The mortality rate from COVID-19 in Indonesia is among the highest in the world. High incidence of infectious diseases and the rise of non-communicable diseases as well as the large smoking population in Indonesia may contribute to dysregulation of immune responses. Whether those factors lead to higher incidence of severe cases and mortality rate of COVID-19 in Indonesia remains to be elucidated. Another important issue in the Indonesian fight against SARS-CoV-2 is the late development of a rapid serological test. Many believe that due to the lack of proper test, the true infection and death rate are higher than the reported data. Development of stronger infrastructure and increased pools of personnel for basic research and reduction of dependency on imported medical reagents need to be strategically planned. When will the current global pandemic end? One projection proposed that we still need some form of social distancing until 2⁰² and should be ready for resurgence of the outbreak in 2⁰⁴.¹⁰¹ Keeping all of previous discussion in mind, serological studies are needed to determine the duration of immunity to SARS-CoV-2. The current practice of using rapid tests as COVID-19 infection screening and discussion of using quantitative methods was reviewed recently.² A study in China reported that autoantibodies were found in SARS-CoV-2 patients.¹⁰² Infection is considered as a risk factor for autoimmune diseases via molecular mimicry or bystander activation. On the other hand, infection may inhibit self-reactive T cells via activation of immune regulation to balance the activation of immune system against the pathogen as has been shown in mice infected with coronavirus.¹⁰³ Whether there is any effect of the current pandemic to the rate of autoimmune diseases and other conditions related to self-reactive immune components remains to be elucidated. The hygienic
behavior such as the use of hand sanitizer, washing hands with soap frequently and showering after arriving home from outdoor activities as well as the spraying of disinfectant may disrupt immune regulation via alteration of microbiome profile with several health consequences, including obesity. Healthy diet, vitamin supplementation, sufficient hours for sleeping at night, and regular exercise are considered as protective measures from the development of severe symptoms by strengthening the immune system. Many efforts have been taken to promote a healthy lifestyle campaign in local and social media while people are staying at home during the pandemic. Continuation of the good habits obtained during the pandemic may provide benefits for stronger immune function and general health in the future. On the other hand, uncomfortable feelings while wearing mask and protective equipment, reduced physical activity and stressful situations during social distancing as well as socioeconomic problems related to it, may impose negative consequences for immune system leading to various disorders. These changes of lifestyle and their impact and long-term effects on the immune system warrant subsequent investigation.

TABLE 3. Immunological studies relevant to COVID-19 in Indonesia

<table>
<thead>
<tr>
<th>Studies on Indonesian COVID-19 cases.</th>
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<tr>
<td>Immunological profile of Indonesian COVID-19 patients, the correlation with disease severity and which host factor contributes most in severe cases.</td>
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<td>The effect of comorbidities (infection and non-infection) to the immunological profile related to disease progression.</td>
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<td>Possible alternative route of infection (e.g. gastrointestinal tract, central nervous system) and correlated immune response in designated organs.</td>
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<td>Evaluation of serological-based test used in Indonesia during COVID-19 pandemic (sensitivity, specificity).</td>
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<td>Methods of serological screening: developing Indonesian-based test (rapid test, ELISA based test).</td>
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<td>Evaluation of the current immunotherapy and development of novel immunotherapeutical agent, including Indonesian herbal medicine.</td>
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<td>Do we achieve herd immunity? How many cases of asymptomatic COVID-19 can be found? Which host factor contributes most to appropriate immune response in the asymptomatic cases?</td>
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<td>Study on the prospect of vaccine development suitable for common type of SARS-CoV-2 in Indonesia.</td>
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<th>Study on the effect of changing lifestyle during and after the pandemic on immune-related problem.</th>
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<tr>
<td>Effect of adopted lifestyle and stress during social distancing on immunological profile, especially in individuals with high risk for severe COVID-19 (elderly, male, hypertension, allergy, autoimmune, etc.).</td>
</tr>
<tr>
<td>Correlation of “super-hygienic” lifestyle, microbiota diversity and immunological profile in Indonesian population.</td>
</tr>
</tbody>
</table>
The profound effect COVID-19 pandemic has on all aspects of life has ignited the global scientific community to work and team up to uncover more about this disease and discover the best strategy to overcome it. The same should be happening in Indonesia as currently we are dealing with the consequences conferred by the pandemic. Wide array of research topic in immunology field relevant to the situation in Indonesia is still waiting to be explored (Table 3). The more knowledge we have about this disease, the more chances we have to defeat it.

CONCLUSION

Innate and adaptive immune responses may keep SARS-CoV-2 at bay and prevent severe symptoms. However, several host factors such as genetic and epigenetic factors from old age, male, history of previous infection, and comorbidities as well as the immune evasion mechanism of the virus may influence the immune responses leading to severe and fatal cases.

The highly contagious nature of the virus prompts swift action to contain it from spreading further. Vaccines, though needing time for proper development, might offer the best chance to fight off the virus especially if it becomes a situation involving recurrent seasonal epidemics. However, for now, physical distancing combined with healthy lifestyle provide the best options to keep the virus in check.

As the pandemic situation continues developing, the best available scientific evidence should be used to guide future research aimed to uncover more about the disease, discover the best diagnostic, therapeutic and intervention strategies, curb the worst possible outcomes and prepare for the likely next pandemic. Thus, research collaboration between governments, international organizations, and the private sector is urgently required. Real-time sharing of scientific information about the virus, disease pathogenesis, and human’s immune response will lead to advancement of the knowledge needed to overcome the many challenges from the current pandemic.

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