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EDITORIALS

 Piha-Paul SA, Oh D-Y, Ueno M, et al. Efficacy and safety of pembrolizumab for the treatment of advanced biliary cancer: results from the KEYNOTE-158 and KEYNOTE-028 studies. Int J Cancer 2020;147:2190–2198.

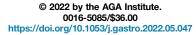
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Conflicts of interest

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Most current article



Rear Window—What Can the Gut Tell Us About Long-COVID?



See "Postacute COVID-19 is characterized by gut viral antigen persistence in inflammatory bowel diseases," by Zollner A, Koch R, Jukic A, et al, on page 495; and "Reduced replication efficacy of severe acute respiratory syndrome coronavirus 2 omicron variant in "mini-gut" organoids" by Miyakawa K, Machida M, Kawasaki T, et al, on page 514.

hile life-threatening acute coronavirus disease 2019 (COVID-19) is mostly linked to respiratory insufficiency or exacerbation of underlying comorbidity, up to 50% of patients with acute COVID-19 may experience mild to moderate gastrointestinal symptoms,¹ because intestinal cells are also prone to infection by severe acute respiratory syndrome coronavirus 2 (SARS-CoV2) and widely express the viral entry receptor angiotensinconverting enzyme 2 (ACE2).² In this issue of Gastroenterology, two articles focus on the intestine and pressing issues in dealing with the COVID-19 pandemic.^{3,4} The article by Miyakawa et al³ addresses the direct infection of intestinal cells by SARS-CoV2. Gastrointestinal symptoms, including diarrhea, abdominal cramping, and pain, are frequently observed in patients with COVID-19,⁵⁻⁸ possibly due to direct infection of the intestinal cells. In fact, stomach cells and epithelial cells of the small and large intestine have been shown to express ACE2 and the coreceptor transmembrane protease, serine 2 (TMPRSS2),^{2,9,10} underlining the possibility that these cells are directly targeted by SARS-CoV2.

The possibility that gastrointestinal cells are infected by SARS–CoV2 was previously demonstrated by studies in intestinal organoids. These studies showed that SARS-CoV2 infection may cause direct infection of epithelial cells, followed by production of infectious viral particles.¹¹ However, the question about potential differences in gastrointestinal infection due to variants of the virus remained to be investigated. In this issue, Miyakawa et al³ now make use of mini-guts as tools to study cell-type specific replication behavior of novel variants of potential concern. Other studies had previously observed altered cell-type specific viral replication efficacies of B.1.529/ Omicron BA.1 compared with B.1.617.2/Delta in lower airway organoids, while upper airway replication was increased.^{12,13}

Extending these findings, Miyakawa et al³ now show reduced viral replication efficacy of Omicron BA.1 and BA.2 compared with the B.1.617.2/Delta variant. This interesting observation correlated to less cell damage as well as to reduced lactose dehydrogenase and high mobility group box 1 protein (HMGB1) release, induced by BA.1 and BA.2. It will be important to follow epidemiology and gastrointestinal symptomatology of the current Omicron wave to assess whether this translates to clinically observed reductions in gastrointestinal symptoms associated to Omicron infection.

For many patients, COVID-19 has been a transient acute illness. However, persistence of symptoms and newly occurring signs of disease not attributable to other causes have emerged in others and lead to an ongoing need to characterize epidemiology, natural history, pathophysiology, diagnosis, and treatment of the phenomenon of post-COVID-19 syndrome, also called post-COVID conditions, long-COVID, postacute COVID-19, or chronic COVID.¹⁴ In addition, it has been observed that the burden of cardiovascular comorbidities is increased in COVID-19 survivors and remains elevated up to 1 year compared with historical control populations,¹⁵ even in patients not hospitalized during acute disease.

There remains an incomplete understanding of the pathophysiology of post-COVID-19 syndrome. Different hypotheses can be formulated to explain ongoing symptoms in the postacute phase:

- 1. Tissue damage that occurred during acute COVID-19 as a result of direct viral toxicity, vessel injury, microthrombosis or macrothrombosis, and ongoing or incomplete healing processes might cause persistent symptoms.^{16,17}
- 2. Incomplete viral clearance and persistent low-level viral replication or continued presence of viral antigens in specialized niches of the body might affect recovery.
- 3. Autoimmune attacks instigated by the immune response to SARS-CoV2 might cause ongoing damage to specific tissues.
- 4. Increased body awareness and the perceived association of diffuse symptoms to COVID might spike reporting of symptoms that usually would remain unnoted by health care providers.

5. Consequences of aggressive therapy—especially in the setting of postcritical illness—might contribute to prolonged or incomplete recovery.

Fecal shedding of viral RNA has been observed to persist longer than oropharyngeal shedding of SARS-CoV2,18,19 putting the gastrointestinal tract into the focus of the study of postacute sequelae of COVID-19. In this issue, Zollner et al⁴ describe substantial novel scientific arguments to this debate by demonstrating persistence of viral antigens in the colon as long as 257 days after initial virus detection, even in the absence of persistent fecal shedding. They made use of various detection methods, including detection of viral RNA by quantitative polymerase chain reaction (RNA-dependent RNA polymerase, nucleocapsid, spike, and envelope proteins) as well as immunofluorescence of colonic biopsy specimens (nucleocapsid protein). They included 46 patients in Austria with inflammatory bowel disease (IBD) who had also experienced a SARS-CoV2 infection from October 2020 to February 2021.

During scheduled routine colonoscopy for control of IBD activity, the study team took tissue samples, and patients who were experiencing ongoing COVID-related symptoms were compared with those with a complete symptom resolution. Interestingly, the authors found persistence of viral antigens in many intestinal tissue samples. Persistence of viral antigen was strongly associated to ongoing symptoms in this study cohort. Owing to the selection of the study cohort, whether these findings are limited to patients with IBD currently remains unclear.⁹ However, Natarajan et al¹⁹ reported the association of persistent viral shedding to persistent gastrointestinal symptoms in a general population.

On the basis of the current evidence, an IBD-related effect thus seems rather unlikely, although one cannot exclude the possibility that effects on viral persistence in patients with IBD are more pronounced due to IBD-related changes of intestinal epithelial cells or accompanying medication (eg, corticosteroid use). Furthermore, whether prolonged persistence of viral components is gut-specific remains unclear. Prolonged detection of viral RNA in feces compared with oropharyngeal swabs does not at all prove an organ-restricted effect. Fecal RNA can be seen as the integral of the entire gastrointestinal tract rather than the direct shedding of the oropharyngeal swab content, and gastrointestinal tissue samples as in the current study are far more easily accessible than lower airway samples.

Taken together, the reported findings by Zollner et al⁴ do not directly prove the functional role of persisting viral antigens as causative for persistent symptoms or the specificity of the gut as a reservoir of viral antigens. However, this report makes an important contribution to the field by showing that persistent viral antigens in the intestine are present in patients with long-COVID and may play a key role the pathogenesis of post-COVID symptoms.

Further studies are clearly needed to further define the molecular mechanisms of post-COVID conditions. However, the ease of tissue sampling in the gut could potentially guide further studies of immunologic mechanisms of prolonged persistence in the context of postacute COVID-19. Colonoscopy thus provides an unexpected "rear window" to the mystery of post-COVID syndrome and provides an easily accessible way to detect viral persistence.

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