

The role of angiotensin-converting enzyme 2 (ACE2) receptor in the intestine in COVID-19: more research needed

Char Leung, Alice PY Wong
Deakin University

(Please cite as: Leung CH, Wong APY. The role of angiotensin-converting enzyme 2 (ACE2) receptor in the intestine in COVID-19: more research needed. *Gastroenterol Hepatol Bed Bench* 2020;13(4):280-281).

Despite being a respiratory disease caused by SARS-CoV-2, a human betacoronavirus, COVID-19 can cause gastrointestinal symptoms such as diarrhea and nausea. While the etiology and pathogenesis of COVID-19 remain not clearly known, it has been suggested that angiotensin-converting enzyme 2 (ACE2), a protein mediating the cell entry of SAR-CoV and SAR-CoV-2, might play a role in the transmission given its surface expression on lung alveolar epithelial cells as well as enterocytes of the small intestine (1). Therefore, a number of studies have purposed to investigate the link between ACE2 and the gastrointestinal symptoms caused by COVID-19 (2,3). However, a number of issues and much debate remain unresolved and, hence, require further research.

Can ACE2 explain the gender differences in susceptibility? A number of clinical studies in China have observed a higher prevalence of COVID-19 in males (4,5). Thus, it has been proposed that ACE2 plays a role as it is highly expressed in Asian men (6). However, ACE2 is also the receptor for SARS-CoV, and a higher prevalence of SARS in females has been observed (7-10). The following questions remain unresolved. Is ACE2 equally expressed in the intestine in males and females? If so, why does it fail to explain the gender differences in susceptibility? Does the gender ratio disparity affect the observed higher

prevalence in males in China? Do clinical findings in Caucasians also demonstrate a higher prevalence in males? Other than ACE2, is any other protein involved in the host cell entry in COVID-19 but not in SARS, although SARS-CoV and SARS-CoV-2 are genetically similar?

Can ACE2 explain the link between the gastrointestinal symptoms observed in a patient and how the patient is infected? The answer to this question is of great importance to infection control, as it helps identify the source of infection by identifying the route of transmission. In an outbreak of SARS-CoV in a residential building in Hong Kong where the faulty sewer system was believed to have spread the virus, the virus RNA was detected in 97% of the patients' stool samples, and 73% of all patients developed diarrhea (10). If the expression of ACE2 in the intestine is proven to contribute to gastrointestinal symptoms, the mode of transmission of SARS-CoV-2 can be further clarified.

To conclude, the gastroenterological aspects of COVID-19 remain an unexplored area. While many studies have delivered important clinical findings for the control and management of COVID-19, little is known about the pathophysiology of diarrhea, vomiting, and nausea in COVID-19 patients. More research should be done, and the role of ACE2 may serve as a hint.

Received: 21 July 2020 Accepted: 18 August 2020
Reprint or Correspondence: Char Leung, MD. Deakin University, Burwood Highway, 3125, Burwood, Australia.
E-mail: char.leung@deakin.edu.au
ORCID ID: 0000-0002-4215-4513

Conflict of interests

The authors declare that they have no conflict of interest.

References

1. Hamming I, Timens W, Bulthuis MLC, Lely AT, Navis GJ, van Goor H. Tissue distribution of ACE2 protein, the functional receptor for SARS coronavirus. A first step in understanding SARS pathogenesis. *J Pathol* 2004;203:631-7.
2. Wong SH, Lui RN, Sung JJ. COVID-19 and the digestive system. *J Gastroenterol Hepatol* 2020;35:744-8.
3. Jin X, Lian JS, Hu JH, Gao J, Zheng L, Zhang YM, et al. Epidemiological, clinical and virological characteristics of 74 cases of coronavirus-infected disease 2019 (COVID-19) with gastrointestinal symptoms. *Gut* 2020;69:1002-9.
4. Novel Coronavirus Pneumonia Emergency Response Epidemiology Team. The epidemiological characteristics of an outbreak of 2019 novel coronavirus diseases (COVID-19) in China. *Chin J Epidemiol* 2020;41.
5. Guan WJ, Ni ZY, Hu Y, Liang WH, Ou CQ, He JX, et al. Clinical Characteristics of Coronavirus Disease 2019 in China. *N Engl J Med* 2020;382:1708-20.
6. Cao Y, Li L, Feng Z, Wan S, Huang P, Sun X, et al. Comparative genetic analysis of the novel coronavirus (2019-nCoV/SARS-CoV-2) receptor ACE2 in different populations. *Cell Discovery* 2020;6:11.
7. Lee ML, Chen CJ, Su IJ, Chen KT, Yeh CC, King CC, et al. Severe acute respiratory syndrome --- Taiwan, 2003. *MMWR Morb Mortal Wkly Rep* 2003;52:461-6.
8. Leung GM, Hedley AJ, Ho LH, Chau P, Wong IOL, Thach TQ, et al. The Epidemiology of Severe Acute Respiratory Syndrome in the 2003 Hong Kong Epidemic: An Analysis of All 1755 Patients. *Ann Intern Med* 2004;141:662-73.
9. Leo YS, Chen M, Heng BH, Lee CC, Paton N, Choo P, et al. Severe acute respiratory syndrome --- Singapore, 2003. *MMWR Morb Mortal Wkly Rep* 2003;52:405-411.
10. Peiris J, Chu C, Cheng V, Chan K, Hung I, Poon L, et al. Clinical progression and viral load in a community outbreak of coronavirus-associated SARS pneumonia: a prospective study. *Lancet* 2003;361:1767-72.6.