

Severe acute respiratory syndrome coronavirus-2 and its structural proteins


To The Editor,

We read with great interest a review that provides a systematic evaluation of the most recent *in vitro* and *in vivo* investigations targeting severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) cell entry, although authors showed that SARS-CoV-2 has five structural proteins including the S (spike), M (membrane), N (nucleocapsid), and E (envelope) proteins, and the HE (hemagglutinin esterase) in fig. 2 (Seyedpour et al., 2021). However, based on the scientific evidence which is described in this letter, SARS-CoV-2 lacks HE so it cannot be considered as a structural protein in SARS-CoV-2 (Chan et al., 2020; Mittal et al., 2020; Zandi et al., 2021).

Recently a novel coronavirus called SARS-CoV-2 causes coronavirus disease 2019 (COVID-19) as a global pandemic (Navand et al., 2020). SARS-CoV-2 is considered the seventh human coronavirus. Coronaviruses belong to *Coronaviridae* family which are classified in four genera: alpha-, beta-, gamma-, and delta-coronaviruses. In coronaviruses, only some beta-coronaviruses contain HE (Chan et al., 2020). There are five subgenera Embecovirus, Sarbecovirus, Merbecovirus, Nobecovirus, and Hibecovirus in beta-coronavirus genus. Both SARS-CoV and SARS-CoV-2 are members of beta-coronaviruses of Sarbecovirus, MERS-CoV is a beta-coronavirus of Merbecovirus, although OC43-CoV and HKU1-CoV are beta-coronaviruses of the Embecovirus (Committee ICoToVE, 2020; Frutos et al., 2021; Zhang et al., 2021).

The genome of coronaviruses is 26–32 kb in length. The genome of SARS-CoV-2 codes four structural proteins: S, E, M, N, and several accessory proteins: ORF3a, 3b, 6, 7a, 7b, 8, 9b, 9c, and 10. SARS-CoV-2 does not contain HE gene (Chan et al., 2020; Zhang et al., 2021). However, other beta-coronaviruses such as HKU1-CoV and OC43-CoV of Embecovirus subgenera bear HE gene and can encode HE (Hulswit et al., 2019).

In conclusion, SARS-CoV-2 as a beta-coronavirus of Sarbecovirus lacks HE and encodes four structural proteins thus HE cannot be used as a drug target against COVID-19.

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REFERENCES

- Chan, J. F., Kok, K.-H., Zhu, Z., Chu, H., To, K. K., Yuan, S., & Yuen, K. Y. (2020). Genomic characterization of the 2019 novel human-pathogenic coronavirus isolated from a patient with atypical pneumonia after visiting Wuhan. *Emerging Microbes & Infections*, 9(1), 221–236.
- Committee ICoToVE. (2020). The new scope of virus taxonomy: Partitioning the virosphere into 15 hierarchical ranks. *Nature Microbiology*, 5(5), 668–674.
- Frutos, R., Serra-Cobo, J., Pinault, L., Lopez Roig, M., & Devaux, C. A. (2021). Emergence of bat-related betacoronaviruses: Hazard and risks. *Frontiers in Microbiology*, 12, 437.
- Hulswit, R., Lang, Y., Bakkers, M., Li, W., Li, Z., Schouten, A., Ophorst, B., van Kuppeveld, F., Boons, G. J., Bosch, B. J., Huizinga, E. G., & de Groot, R. J. (2019). Human coronaviruses OC43 and HKU1 bind to 9-O-acetylated sialic acids via a conserved receptor-binding site in spike protein domain A. *Proceedings of the National Academy of Sciences of the United States of America*, 116(7), 2681–2690.
- Mittal, A., Manjunath, K., Ranjan, R. K., Kaushik, S., Kumar, S., & Verma, V. (2020). COVID-19 pandemic: Insights into structure, function, and hACE2 receptor recognition by SARS-CoV-2. *PLoS Pathogens*, 16(8), e1008762.
- Navand, A. H., Soltani, S., Moghadami, M., Hosseini, P., Nasimzadeh, S., & Zandi, M. (2020). Diabetes and coronavirus infections (SARS-CoV, MERS-CoV, and SARS-CoV-2). *Journal of Acute Disease*, 9(6), 244.
- Seyedpour, S., Khodaei, B., Loghman, A. H., Seyedpour, N., Kisomi, M. F., Balibegloo, M., Nezamabadi, S. S., Gholami, B., Saghadzadeh, A., & Rezaei, N. (2021). Targeted therapy strategies against SARS-CoV-2 cell entry mechanisms: A systematic review of *in vitro* and *in vivo* studies. *Journal of Cellular Physiology*, 236(4), 2364–2392.
- Zandi, M., Behboudi, E., & Soltani, S. (2021). Role of glycoprotein hemagglutinin-esterase in COVID-19 pathophysiology? *Stem Cell Reviews and Reports*, 1–2.
- Zhang, Y., Gargan, S., Lu, Y., & Stevenson, N. J. (2021). An overview of current knowledge of deadly CoVs and their interface with innate immunity. *Viruses*, 13(4), 560.