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SARS-CoV-2 as a betacoronavirus comprises five structural proteins?

Dear editor,

We read with interest a review article by Bakhshandeh et al. (Bakhshandeh et al., 2021). The authors considered hemagglutinin esterase (HE) as one of the structural proteins of SARS-CoV-2 (Bakhshandeh et al., 2021). Although, scientific evidences show that the genome of SARS-CoV-2 lacks the HE gene and it has no hemagglutinin-esterase glycoprotein (Kumar et al., 2020; Anastasopoulou and Mouzaki, 2020; Crawford et al., 2020; Zhang et al., 2020). Thus, HE cannot be considered as an antigenic component in SARS-CoV-2 and it has no role in SARS-CoV-2 infection.

SARS-CoV-2 as a betacoronavirus contains a non-segmented positive-sense, single-stranded RNA (Pal et al., 2020). The genome of SARS-CoV-2 has been sequenced, and based on the genomic sequence, SARS-CoV-2 shared 79.6% sequence identity to SARS-CoV and 96% identity similar to bat coronavirus (Zhou et al., 2020). The genome organization of SARS-CoV-2 is 5'UTR-Rep-S-3a-3b-E-M-6-7a-7b-8-N-10-3'UTR (Abdel-Moneim and Abdelwhab, 2020). Coronaviruses comprise four structural proteins spike (S), envelope (E), membrane protein (M), and nucleoprotein (N) (Fani et al., 2021), however, some betacoronaviruses such as OC43-CoV, Bovine-CoV and HKU1-CoV and, murine hepatitis virus encode hemagglutinin esterase (Lang et al., 2020). Therefore SARS-CoV-2 lacks HE and comprises four structural proteins S, E, M and N.

Declaration of competing interest

The authors declare that there is no conflict of interest for this manuscript entitle "SARS-CoV-2 as a betacoronavirus comprises five structural proteins?"

References

Abdel-Moneim, A.S., Abdelwhab, E.M., 2020. Evidence for SARS-CoV-2 infection of animal hosts. *Pathogens* 9 (7), 529.

- Anastasopoulou, S., Mouzaki, A., 2020. The biology of SARS-CoV-2 and the ensuing COVID-19. *Achaiki Iatriki* 39 (1), 29–35.
- Bakhshandeh, B., Sorbini, S.G., Javanmard, A.-R., Mottaghi, S.S., Mehrabi, M.-R., Sorouri, F., et al., 2021. Variants in ACE2; potential influences on virus infection and COVID-19 severity. *Infect. Genet. Evol.* 104773.
- Crawford, K.H., Eguia, R., Dingens, A.S., Loes, A.N., Malone, K.D., Wolf, C.R., et al., 2020. Protocol and reagents for pseudotyping lentiviral particles with SARS-CoV-2 spike protein for neutralization assays. *Viruses* 12 (5), 513.
- Fani, M., Zandi, M., Ebrahimi, S., Soltani, S., Abbasi, S., 2021. The role of miRNAs in COVID-19 disease. *Futur. Virol.* 16 (4), 301–306.
- Kumar, S., Nyodu, R., Maurya, V.K., Saxena, S.K., 2020. Morphology, genome organization, replication, and pathogenesis of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2). In: *Coronavirus Disease 2019 (COVID-19)*. Springer, pp. 23–31.
- Lang, Y., Li, W., Li, Z., Koerhuis, D., Van Den Burg, A.C., Rozemuller, E., et al., 2020. Coronavirus hemagglutinin-esterase and spike proteins coevolve for functional balance and optimal virion avidity. *Proc. Natl. Acad. Sci.* 117 (41), 25759–25770.
- Pal, M., Berhanu, G., Desalegn, C., Kandi, V., 2020. Severe acute respiratory syndrome coronavirus-2 (SARS-CoV-2); an update. *Cureus* 12 (3).
- Zhang, L.-P., Wang, M., Wang, Y., Zhu, J., Zhang, N., 2020. Focus on the 2019 novel coronavirus (SARS-CoV-2). *Future Microbiol.* 15 (10), 905–918.
- Zhou, P., Yang, X.-L., Wang, X.-G., Hu, B., Zhang, L., Zhang, W., et al., 2020. A pneumonia outbreak associated with a new coronavirus of probable bat origin. *Nature* 579 (7798), 270–273.

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