Infants Younger Than 6 Months Infected With SARS-CoV-2 Show the Highest Respiratory Viral Loads

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## To the Editor:

We read with great interest the article by Ochoa et al. on the role of infants in spreading SARS-CoV-2 infection [1]. Based on their findings, they concluded that children younger than 6 months old displayed higher SARS-CoV-2 loads with less severe presentation compared to other age groups[1]. Although this article provided insights into coronavirus disease 2019 (COVID-19), the findings on the accuracy of ORF1ab Cycle Threshold (Ct) might be compromised due to other predictor variables.

To begin with, there is uncertainty regarding the accuracy of Reverse Transcriptase polymerase chain reaction (RT-PCR) using nasopharyngeal specimens, when viral loads are low; in such situations we recommend supplemental fecal specimen testing to be performed [2].

In addition, the time between COVID-19 infection and PCR testing may affect the PCR test results. For instance, it has been suggested that the later an infected individual is tested after symptom onset, the less likely the test is positive [3]. Early sampling within seven days of symptom onset can decrease the false-negative rate of RT-PCR [4]. In this study, the final tests used to determine viral loads may not be as precise since the samples may have been collected at a different time. The observed ORF1ab Ct may be underestimated. Therefore, we recommend that measures such as testing before washing in the morning may increase diagnostic accuracy in patients with low viral loads during latent infection stages [5].

Finally, although some potential influencing factors have been excluded in this study [1], other predictive variables, such as laboratory data including CCL5 expression, may lead to potential confounding bias. For example, previous studies showed that COVID-19 patients had upregulated innate immunity genes, which were directly associated with SARS-CoV-2 load. [6] In this regard, subgroup analysis of genes would greatly reduce potential confounding bias. Therefore, we suggest genetic testing data of mucosal biomarkers be included in future studies.

In conclusion, several factors that may influence the accuracy of viral load detection should be considered in future studies. For instance, reference fecal sample testing in addition to nasopharyngeal specimens for RT-PCR tests should be included. Likewise, genetic testing, such as CCL5 expression, should be considered as a predictor variable and stratified in further analyses. Lastly, strategies to enhance diagnostic accuracy of PCR tests in latent infection should be considered.

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