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represents in healthy newborns, whether it is related to changing pulmonary vascular resistance, changes in intracardiac or extracardiac shunts, fluid shifts, or all of these processes in concert.

With respect to the statistical analyses performed and our sample size, we created our hypothesis regarding differences in BNP trajectory based on our clinical experience and performed exploratory analyses based on this hypothesis. It is our practice to conduct power analyses in the context of negative findings to ensure that we have not encountered false-negative results. In the receiver operating characteristic curve analyses, we identify robust findings in the accuracy of BNP for prediction of clinical outcome during weeks 3 through 5, with cut-off values that maximize the percent correctly classified, and sensitivity and specificity without bias toward either measure. Despite our sample size, the positive findings for these weeks negates the potential for false-negative results. We are overall reassured that our findings are robust to multiple different analytic approaches.

Elyssa Guslits, MD

Martina A. Steurer, MD, MAS

Department of Pediatrics, Critical Care, University of California San Francisco, San Francisco, California

Hythem Nawaytou, MD

Department of Pediatrics, Cardiology, University of California San Francisco, San Francisco, California

Roberta L. Keller, MD

Department of Pediatrics, Neonatology, University of California San Francisco, San Francisco, California

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Coronavirus disease 2019, multisystem inflammatory syndrome in children, apolipoprotein E4, and race



To the Editor:

Kaushik et al presented a series of 33 children from New York City hospitals diagnosed with multisystem inflammatory syndrome in children (MIS-C); 81% had antibodies to severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2).¹ Similarly, Carter et al, reported on a series of 25 children diagnosed with MIS-C from the United Kingdom; 68% were SARS-CoV-2 seropositive.² The children in both groups exhibited a cytokine profile consistent with a robust innate immune response. At the same time, emerging evidence suggests the hypothesis that the apolipoprotein E4 (apoE4) genotype can predict coronavirus disease 2019 (COVID-19) severity in adults.^{3,4} Accordingly, we propose the hypothesis that the apoE4 genotype may identify children at increased risk of developing MIS-C from SARS-CoV-2 infection.

Classically, the apoE4 genotype has been associated with cardiovascular disease and Alzheimer's disease⁵; however, it has also been associated with an enhanced in vivo innate immune response.⁶ Notably, individuals of African descent may have twice the frequency of the $\epsilon 4$ allele (30%-40%) compared with those of European or Asian descent,⁷ and therefore, they may be more likely to exhibit a stronger innate immune response to the SARS-CoV-2 infection.

In the series presented by Carter et al, the children who were SARS-CoV-2 seropositive exhibited more severe disease; 8 of 9 black children compared with 5 of 10 white children in the series were SARS-CoV-2 seropositive.² Of 21 patients with a depressed systolic ventricular function in the report by Kaushik et al, 11 were black and 1 white (Table 4).¹ This suggests an overrepresentation of black children diagnosed with MIS-C from severe SARS-CoV-2 infection.

Therefore, as seen with adults, the apoE4 genotype may identify children at a greater risk of severe SARS-CoV-2 infection; and in particular, MIS-C.

Mark R. Goldstein, MD, FACP

NCH Physician Group
Center for Healthy Living
Naples, FL

Gregory A. Poland, MD, MACP, FIDSA, FRCP (London)

Mayo Clinic and Foundation
Rochester, MN

Charles W. Graeber, MD

NCH Healthcare System Internal Medicine Residency
Affiliate of the Mayo Clinic School of Medicine and Science
Naples, FL

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RNA is not virus



To the Editor:

Yonker et al demonstrated high levels of viral RNA in nasopharyngeal samples obtained from children.¹ A plausible explanation for the seemingly counterintuitive lack of symptoms and high viral load is the methods employed are detecting high viral RNA, which does not correlate to high viral load as the authors suggest. Establishing a correlation between RNA and virus in the asymptomatic pediatric population with a virus culture is a prerequisite for assertion that RNA positive children carry high viral loads.² An alternative explanation is that the children in the study, and generally most children, mount an effective immune response to subclinical infection. That immune response causes lysis of infected cells, spilling cellular contents including viral RNA and proteins into surrounding tissue or interference with viral assembly inhibiting excretion of infectious virus. High RNA in nasopharyngeal samples obtained from asymptomatic patients is a reflection of an effective immune response rather than high viral load, explaining minimal infectivity of such subjects.³

Gregory I. Sawchyn, MD, MBA

Population Health
Sound Physicians Group
Tacoma, Washington

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Reply



To the Editor:

In our article, we report high viral load in children infected with severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) in the first 2 days of illness. We did not assess the viability of virus from these respiratory secretions and agree that RNA does not necessarily correlate with live virus. However, in hospitalized adults, live virus is readily cultured from respiratory secretions during the first week of symptoms.¹ It is plausible that differences exist between pediatric and adult immune responses to SARS-CoV-2²; distinctions in mucosal immune responses could impact viral detection by reverse-transcription polymerase chain reaction and/or severity of symptoms. Another plausible hypothesis is that SARS-CoV-2 could colonize the upper airways efficiently in both children and adults but children may be less likely to have colonization in the lower respiratory tract. Although we described SARS-CoV-2 serology and show that infected children mount a humoral response following infection, we did not assess mucosal responses or regional differences of viral load within the airways. Research is needed to test these hypotheses.

Lael M. Yonker, MD

Alessio Fasano, MD

Mucosal Immunology and Biology Research Center
Department of Pediatrics
Massachusetts General Hospital
Harvard Medical School
Boston, Massachusetts

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