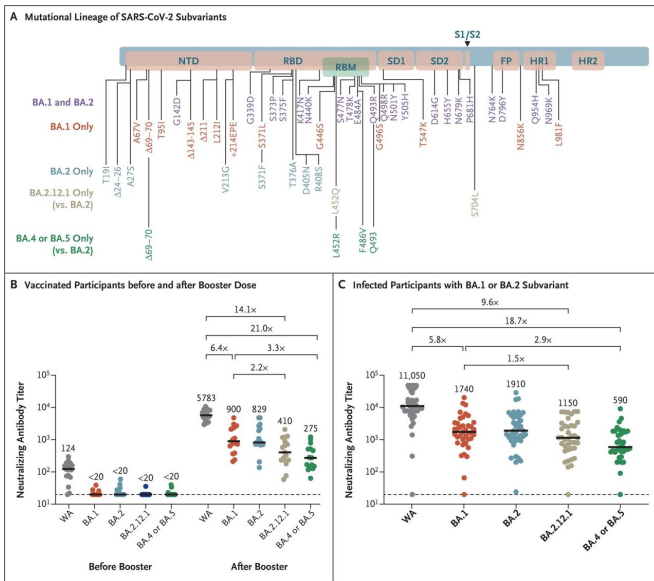


# Newer COVID-19 subvariants are less vulnerable to immunity induced by vaccination and previous infection

23 June 2022, by Jacqueline Mitchell



(black bars) are shown numerically, and factor differences from other subvariants are indicated; the dashed horizontal line indicates the lower limit of detection for the assay. Credit: *New England Journal of Medicine* (2022). DOI: 10.1056/NEJMc2206576

Since the initial highly infectious SARS-CoV-2 omicron variant (officially known as BA.1 or B.1.1.529) of COVID-19 emerged last fall, new subvariants of omicron continue to evolve.

Notably, the omicron subvariants BA.4 and BA.5—not identified in the United States until late April—now account for more than 21% of new cases, according to the Centers for Disease Control and Prevention's (CDC) estimates for the week ending June 11. New variants that emerge may be more transmissible and/or may more effectively bypass the immune protection from prior infection or vaccination.

In a letter published in the *New England Journal of Medicine*, physician-scientists at Beth Israel Deaconess Medical Center (BIDMC) report that the three omicron subvariants currently dominant in the United States—officially known as subvariants BA.2.12.1, BA.4, and BA.5—substantially escape neutralizing [antibodies](#) induced by both vaccination and previous infection.

Barouch and colleagues evaluated [antibody responses](#) to multiple SARS-CoV-2 omicron subvariants in 27 vaccinated and boosted individuals and 27 individuals who had previously contracted COVID-19. Neutralizing antibody responses to BA.4 and BA.5 were approximately 20-fold lower than to the original WA1/2020 strain and were 3-fold lower than to the omicron BA.1 and BA.2 variants.

Omicron Subvariant Mutations and Neutralizing Antibody Responses. Panel A shows the lineage of mutations that have been identified in the omicron BA.1, BA.2, BA.2.12.1, and BA.4 or BA.5 subvariants of SARS-CoV-2, as compared with the reference WA1/2020 isolate. BA.4 and BA.5 have identical sequences of the spike protein and thus have been grouped together. FP denotes fusion peptide, HR1 heptad repeat 1, HR2 heptad repeat 2, NTD N-terminal domain, RBD receptor-binding domain, RBM receptor-binding motif, SD1 subdomain 1, and SD2 subdomain 2. Panel B shows neutralizing antibody titers as determined by luciferase-based pseudovirus neutralization assays in samples obtained from 27 participants 6 months after receipt of the two-dose BNT162b2 messenger RNA vaccine series and 2 weeks after the third (booster) dose. Panel C shows neutralizing antibody titers in participants who had been infected with the BA.1 or BA.2 subvariant. All the infected participants had been vaccinated except for 1 participant who had a negative neutralizing antibody titer. In 9 participants, two or three time points after infection are shown. Neutralizing antibody titers were measured against the SARS-CoV-2 reference isolate WA1/2020 and the omicron BA.1, BA.2, BA.2.12.1, and BA.4 or BA.5 subvariants. In Panels B and C, medians

"Our findings suggest that the [omicron](#) variants have continued to evolve," said senior author Dan H. Barouch, MD, Ph.D., director of the Center for Vaccine and Virology Research at BIDMC. "This has important public health implications and provides the immunologic context for current surges among populations with high rates of vaccinations and previous infection."

**More information:** Nicole P. Hachmann et al, Neutralization Escape by SARS-CoV-2 Omicron Subvariants BA.2.12.1, BA.4, and BA.5, *New England Journal of Medicine* (2022). [DOI: 10.1056/NEJMc2206576](#)

Provided by Beth Israel Deaconess Medical Center

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