OBSERVATION



Neutralization Titers in Vaccinated Patients with SARS-CoV-2 Delta Breakthrough Infections

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ABSTRACT The continuous emergence of SARS-CoV-2 variants with increased transmission and immune evasion has caused breakthrough infections in the vaccinated population. It is important to determine the threshold of neutralizing antibody titers (NT₅₀) that permit breakthrough infections in humans. Here, we tested the neutralization titers of vaccinated patients who contracted Delta variant. All 64 patients with Delta breakthrough infections exhibited NT₅₀ of less than 70. When the breakthrough sera were tested against USA-WA1/2020 (a strain isolated in late January 2020), 82.8%, 15.6%, and 1.6% of them had the NT₅₀ ranges of <20, 20 to 50, and 50 to 69, respectively. When the same breakthrough sera were tested against Delta-spike SARS-CoV-2, 68.7%, 26.6%, and 4.7% of them had the NT₅₀ ranges of <20, 20 to 50, and 50 to 69, respectively. Overall, the results suggest NT₅₀ of 70 as a potential neutralizing threshold required to prevent Delta breakthrough infections. These clinical laboratory results have implications in vaccine strategy and public health policy.

IMPORTANCE Given that neutralizing antibodies play a key role in protection of SARS-CoV-2 infection, it is important to define the neutralization levels in vaccinated individuals when they contracted breakthrough infections. In this study, we analyzed the neutralization levels from 64 vaccinated patients on days 0 to 5 before they tested positive for Delta breakthrough infections. The neutralization titers in these vaccinated individuals were all lower than 70 when they contracted breakthrough infections. The results suggest a neutralization titer of 70 as the potential threshold required to prevent breakthrough infections of Delta variant.

KEYWORDS COVID-19, SARS-CoV-2, breakthrough infection, antibody neutralization, vaccine, variants of concern, vaccine booster

The COVID-19 pandemic is umpired by two dynamic factors: (i) the continuous emergence of SARS-CoV-2 variants with improved transmission and/or immune evasion and (ii) the waning immunity post vaccination and infections (1). This is exemplified by the two recent surges of Delta and Omicron variants, which caused many breakthrough infections in vaccinated individuals. Because antibody neutralization is a key contributor to vaccine protection against symptomatic infection and severe disease (2), it is important to define the neutralization levels in vaccinated individuals who contracted breakthrough infections. Such information is essential to guide vaccine strategy and policy. Here, we characterized the antibody neutralization in vaccinated patients when they acquired Delta variant infections. Editor Sara Cherry, University of Pennsylvania Copyright © 2022 Zou et al. This is an openaccess article distributed under the terms of the Creative Commons Attribution 4.0 International license.

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The authors declare a conflict of interest. X.X. and P.-Y.S. have filed a patent on the reverse genetic system. J.Z., X.X., and P.-Y.S. received compensation from Pfizer for COVID-19 vaccine development. Other authors declare no competing interests.

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To determine the neutralization titers (NT_{50}) in breakthrough patients when they were infected with Delta variant, we collected sera from 64 patients who were vaccinated and subsequently contracted breakthrough infections. The use of human sera for the described research was reviewed and approved by the University of Texas Medical Branch (UTMB) Institutional Review Board (IRB number 20-0070). The deidentified human sera were collected from vaccinated patients who presented COVID-19 symptoms. Table S1 summarizes the patient information and their NT_{50} values. All patients were immunized with two doses of Pfizer or Moderna vaccine or one dose of J&J vaccine. Breakthrough infections were confirmed by positive viral RNA tests. Although the genotypes of individual infecting viruses were not determined by sequencing, they were most likely Delta variant because all infections had occurred from late July to October 2021, when Delta was 100% prevalent in our patient population based on the local SARS-CoV-2 surveillance system (at the University of Texas Medical Branch) and about 98% prevalent in the United States (https://covid.cdc.gov/covid-data-tracker/#variant-proportions). All sera were taken 0 to 5 days before the positive viral nucleic acid tests and within 4 days of symptom onset. We determined the NT₅₀ of each serum using a well-established mNeonGreen reporter USA-WA1/2020 SARS-CoV-2 (3). This neutralization assay has been reliably used to support the BNT162b2 vaccine development (4–6). The $NT_{\rm 50}$ value was defined as the interpolated reciprocal of the dilution yielding a 50% reduction in mNeonGreen-positive cells. Each specimen was tested in duplicates and the geometric mean of the duplicate results is presented. The first serum dilution of the neutralization test was 1:20. The NT₅₀ values of specimens with no detectable neutralizing activities at 1:20 dilution were treated as 10 for plot and calculation purposes (Table S1). The overall results reveal three observations. First, all breakthrough patients had low NT₅₀s of <70 (Fig. 1A). About 82.8%, 15.6%, and 1.6% of the breakthrough patients exhibited the NT₅₀ ranges of < 20, 20 to 50, and 51 to 69, respectively (Fig. 1B). The results suggest NT₅₀ of 70 as a potential neutralizing threshold required to prevent Delta breakthrough infections. In support of our human results, a previous study reported NT_{50} of 50 as the minimal neutralization level required to protect nonhuman primates from SARS-CoV-2 infection (7). Second, senior people appeared to be more vulnerable to breakthrough infections. Approximately 15.6%, 26.6%, and 57.8% of the breakthrough cases were in the age groups of 16 to 40, 41 to 64, and 65 to 97, respectively (Fig. 1C). However, the NT₅₀ differences among the three age groups are not all statistically significant (Fig. 1D). Third, 87.5% of the breakthrough patients had received two doses of Pfizer or Moderna vaccine or one dose of J&J vaccine for more than 120 days (Fig. 1E). However, this observation was not always statistically correlated with the NT₅₀ differences among different time frames post vaccination (Fig. 1F).

The above analysis was performed on the NT₅₀ values against the original mNeonGreen USA-WA1/2020 (3). To directly test the NT₅₀ values against Delta spike, we engineered the complete spike gene from the Delta variant into the backbone of mNeonGreen USA-WA1/2020, resulting in Delta-spike SARS-CoV-2. The construction of the Delta-spike SARS-CoV-2 was previously reported (8, 9). We tested the entire serum panel for NT₅₀ values against the Delta-spike SARS-CoV-2 (Table S1). Compared with the original USA-WA1/2020 (Fig. 1), the Delta-spike SARS-CoV-2 results revealed similar observations (Fig. 2). All breakthrough sera had NT₅₀s of <70 against Delta-spike SARS-CoV-2 (Fig. 2A). Approximately 68.7%, 26.6%, and 4.7% of the breakthrough sera exhibited the NT₅₀ ranges of <20, 20 to 50, and 51 to 69, respectively (Fig. 2B). The NT₅₀ differences among the three age groups (Fig. 2C) or the NT₅₀ differences among the three time frames post vaccination (Fig. 1D) are not always statistically significant.

We previously reported that at 8 months post dose-2 of BNT162b2 vaccine, the neutralization titers against USA-WA1/2020 were 83 and 41 for age groups of 18 to 55 and 65 to 85 years old, respectively (4), suggesting that both age groups, particularly the senior age group, are susceptible to Delta breakthrough infections. The previous result, together with the current data, support the vaccine booster strategy after 6 months of two doses of BNT162b2.

The current study has three weaknesses. The first weakness is even though the serum specimens were collected on 0 to 5 days before viral RNA-positive tests, these sera

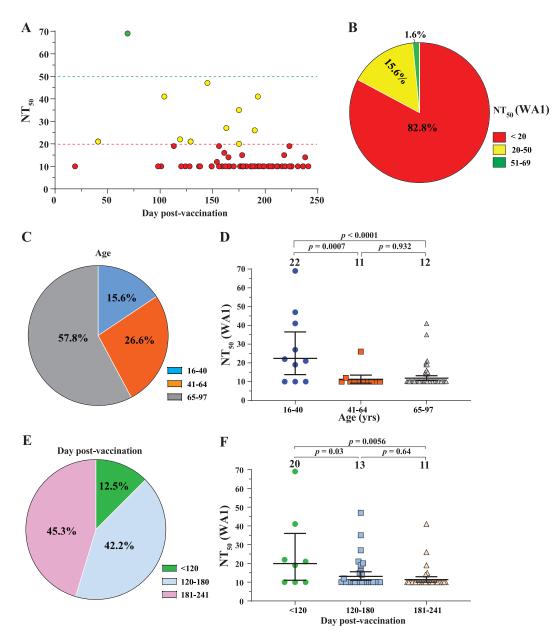


FIG 1 NT₅₀ analysis against USA-WA1/202 SARS-CoV-2. A panel of 64 sera, collected from Delta breakthrough patients, were measured for antibody neutralization titers (NT₅₀) against mNeonGreen USA-WA1/202. All patients were immunized with two doses of Pfizer or Moderna vaccine or one dose of J&J vaccine. All sera were taken 0 to 5 days before viral RNA-positive test results. The sera were heat-inactivated at 56°C for 30 min before the neutralization testing. All patient information and NT₅₀ values are detailed in Table S1. (A) Plot of NT₅₀ values versus days after dose 2 of Pfizer or Moderna vaccine or after 1 dose of J&J vaccine. Each data point represents the geometric mean of NT₅₀ for one serum tested in duplicate assays. Different colors represent different NT₅₀ ranges. Samples with no detectable neutralizing activities were plotted as 10 for calculation purpose. (B) Pie presentation of different NT₅₀ ranges. (C) Age distributions. (D) Plot of NT₅₀ of different age groups. (E) Distribution of breakthrough percentages versus days post vaccination. (F) Plot of NT₅₀ versus days post vaccination. In D and F, the bar heights and the numbers above indicate geometric mean titers. The whiskers indicate 95% confidence intervals. Statistical analysis was performed using the one-way ANOVA with Tukey's correction for multiple-comparison test.

may already contain anamnestic antibodies produced during the presymptomatic period of breakthrough infections, which may lead to an overestimated NT_{50} threshold for breakthrough infections. The second weakness is the small sample size of the study; more breakthrough patient specimens are needed to bolster our conclusion. The third weakness is the NT50 value of vaccinated individuals who resisted Delta breakthrough infections are unknown. Despite the weaknesses, the current results gave a glimpse of

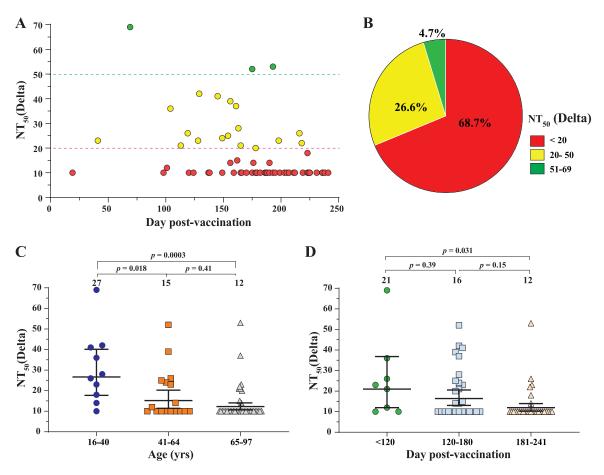


FIG 2 NT_{s0} analysis against Delta-spike SARS-CoV-2. The same sera from Fig. 1 were measured for NT_{s0} values against mNeonGreen Delta-spike SARS-CoV-2. Delta-spike SARS-CoV-2 contains the complete spike gene from the Delta variant in the backbone of mNeonGreen USA-WA1/2020 SARS-CoV-2. The NT_{s0} values are summarized in Table S1. The NT_{s0} values were measured and presented as described in Fig. 1. (A) Plot of NT_{s0} values against Delta-spike SARS-CoV-2. (B) Pie presentation of different NT_{s0} ranges. (C) Plot of NT_{s0} against Delta-spike SARS-CoV-2 of different age groups. (D) Plot of NT_{s0} against Delta-spike SARS-CoV-2 versus days post vaccination. In C and D, the numbers above indicate geometric mean titers. The whiskers indicate 95% confidence intervals. Statistical analysis was performed with the one-way ANOVA with Tukey's correction for multiple-comparison test.

the neutralization status when the vaccinated patients contracted Delta breakthrough infections.

Compared with Delta, the newly emerged Omicron is significantly less susceptible to neutralization by vaccinated or non-Omicron infected human sera (10–13). The reduced neutralization susceptibility, combined with the increased transmissibility of Omicron, may have accounted for the Delta-to-Omicron variant replacement and high breakthrough infections observed in the current Omicron surge. Similar clinical studies are needed to understand the threshold of neutralization required to protect humans from Omicron break-through infections. Laboratory investigations, together with the real-world vaccine effectiveness, have enabled FDA to recommend a bivalent vaccine strategy that includes both the original spike and the current prevalent Omicron BA.4/5 spike.

SUPPLEMENTAL MATERIAL

Supplemental material is available online only. **TABLE S1**, DOCX file, 0.02 MB.

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X.X. and P.-Y.S. have filed a patent on the reverse genetic system. J.Z., X.X., and P.-Y.S. received compensation from Pfizer for COVID-19 vaccine development. Other authors declare no competing interests.

REFERENCES

- 1. Liu Y, Liu J, Shi PY. 2022. SARS-CoV-2 variants and vaccination. Zoonoses 2. https://doi.org/10.15212/ZOONOSES-2022-0001.
- Walsh EE, Frenck RW, Falsey AR, Kitchin N, Absalon J, Gurtman A, Lockhart S, Neuzil K, Mulligan MJ, Bailey R, Swanson KA, Li P, Koury K, Kalina W, Cooper D, Fontes-Garfias C, Shi P-Y, Türeci Ö, Tompkins KR, Lyke KE, Raabe V, Dormitzer PR, Jansen KU, Şahin U, Gruber WC. 2020. Safety and immunogenicity of two RNAbased COVID-19 vaccine candidates. N Engl J Med 383:2439–2450. https:// doi.org/10.1056/NEJMoa2027906.
- Muruato AE, Fontes-Garfias CR, Ren P, Garcia-Blanco MA, Menachery VD, Xie X, Shi PY. 2020. A high-throughput neutralizing antibody assay for COVID-19 diagnosis and vaccine evaluation. Nat Commun 11:4059. https://doi.org/10 .1038/s41467-020-17892-0.
- Falsey AR, Frenck RW, Walsh EE, Kitchin N, Absalon J, Gurtman A, Lockhart S, Bailey R, Swanson KA, Xu X, Koury K, Kalina W, Cooper D, Zou J, Xie X, Xia H, Türeci Ö, Lagkadinou E, Tompkins KR, Shi P-Y, Jansen KU, Şahin U, Dormitzer PR, Gruber WC. 2021. SARS-CoV-2 neutralization with BNT162b2 vaccine dose 3. N Engl J Med 385:1627–1629. https://doi.org/10.1056/NEJMc2113468.
- Sahin U, Muik A, Derhovanessian E, Vogler I, Kranz LM, Vormehr M, Baum A, Pascal K, Quandt J, Maurus D, Brachtendorf S, Lorks V, Sikorski J, Hilker R, Becker D, Eller AK, Grutzner J, Boesler C, Rosenbaum C, Kuhnle MC, Luxemburger U, Kemmer-Bruck A, Langer D, Bexon M, Bolte S, Kariko K, Palanche T, Fischer B, Schultz A, Shi PY, Fontes-Garfias C, Perez JL, Swanson KA, Loschko J, Scully IL, Cutler M, Kalina W, Kyratsous CA, Cooper D, Dormitzer PR, Jansen KU, Tureci O. 2020. COVID-19 vaccine BNT162b1 elicits human antibody and TH1 T cell responses. Nature 586: 594–599. https://doi.org/10.1038/s41586-020-2814-7.
- 6. Vogel AB, Kanevsky I, Che Y, Swanson KA, Muik A, Vormehr M, Kranz LM, Walzer KC, Hein S, Güler A, Loschko J, Maddur MS, Ota-Setlik A, Tompkins K, Cole J, Lui BG, Ziegenhals T, Plaschke A, Eisel D, Dany SC, Fesser S, Erbar S, Bates F, Schneider D, Jesionek B, Sänger B, Wallisch A-K, Feuchter Y, Junginger H, Krumm SA, Heinen AP, Adams-Quack P, Schlereth J, Schille S, Kröner C, de la Caridad Güimil Garcia R, Hiller T, Fischer L, Sellers RS, Choudhary S, Gonzalez O, Vascotto F, Gutman MR, Fontenot JA, Hall-Ursone S, Brasky K, Griffor MC,

Han S, Su AAH, Lees JA, et al. 2021. BNT162b vaccines protect rhesus macaques from SARS-CoV-2. Nature 592:283–289. https://doi.org/10.1038/s41586 -021-03275-y.

- McMahan K, Yu J, Mercado NB, Loos C, Tostanoski LH, Chandrashekar A, Liu J, Peter L, Atyeo C, Zhu A, Bondzie EA, Dagotto G, Gebre MS, Jacob-Dolan C, Li Z, Nampanya F, Patel S, Pessaint L, Van Ry A, Blade K, Yalley-Ogunro J, Cabus M, Brown R, Cook A, Teow E, Andersen H, Lewis MG, Lauffenburger DA, Alter G, Barouch DH. 2021. Correlates of protection against SARS-CoV-2 in rhesus macaques. Nature 590:630–634. https://doi.org/10.1038/s41586-020-03041-6.
- Xie X, Muruato A, Lokugamage KG, Narayanan K, Zhang X, Zou J, Liu J, Schindewolf C, Bopp NE, Aguilar PV, Plante KS, Weaver SC, Makino S, LeDuc JW, Menachery VD, Shi PY. 2020. An infectious cDNA clone of SARS-CoV-2. Cell Host Microbe 27:841–848.e843. https://doi.org/10.1016/j.chom.2020.04.004.
- 9. Liu Y, Liu J, Zou J, Ren P, Weaver SC, Xie X, Shi PY. 2021. Distinct neutralizing kinetics and magnitudes elicited by different SARS-CoV-2 variant spikes. bioRxiv. https://doi.org/10.1101/2021.09.02.458740.
- Xia H, Zou J, Kurhade C, Cai H, Yang Q, Cutler M, Cooper D, Muik A, Jansen KU, Xie X, Swanson KA, Shi PY. 2022. Neutralization and durability of 2 or 3 doses of the BNT162b2 vaccine against Omicron SARS-CoV-2. Cell Host Microbe 30:485–488.e483. https://doi.org/10.1016/j.chom.2022.02.015.
- Zou J, Xia H, Xie X, Kurhade C, Machado RRG, Weaver SC, Ren P, Shi PY. 2022. Neutralization against Omicron SARS-CoV-2 from previous non-Omicron infection. Nat Commun 13:852. https://doi.org/10.1038/s41467-022-28544-w.
- Kurhade C, Zou J, Xia H, Cai H, Yang Q, Cutler M, Cooper D, Muik A, Jansen KU, Xie X, Swanson KA, Shi PY. 2022. Neutralization of Omicron BA.1, BA.2, and BA.3 SARS-CoV-2 by 3 doses of BNT162b2 vaccine. Nat Commun 13: 3602. https://doi.org/10.1038/s41467-022-30681-1.
- Kurhade C, Zou J, Xia H, Liu M, Yang Q, Cutler M, Cooper D, Muik A, Sahin U, Jansen KU, Ren P, Xie X, Swanson KA, Shi P. 2022. Neutralization of Omicron sublineages and Deltacron SARS-CoV-2 by 3 doses of BNT162b2 vaccine or BA.1 infection. Emerg Microbes Infect. 11(1):1828–1832. https://doi.org/10.1080/22221751 .2022.2099305.