

Since January 2020 Elsevier has created a COVID-19 resource centre with free information in English and Mandarin on the novel coronavirus COVID-19. The COVID-19 resource centre is hosted on Elsevier Connect, the company's public news and information website.

Elsevier hereby grants permission to make all its COVID-19-related research that is available on the COVID-19 resource centre - including this research content - immediately available in PubMed Central and other publicly funded repositories, such as the WHO COVID database with rights for unrestricted research re-use and analyses in any form or by any means with acknowledgement of the original source. These permissions are granted for free by Elsevier for as long as the COVID-19 resource centre remains active.



Contents lists available at ScienceDirect

Clinical Immunology



journal homepage: www.elsevier.com/locate/yclim

Seroconversion among rituximab-treated patients following SARS-CoV-2 vaccine supplemental dose

Emily Rose^a, Daniel Magliulo^b, Vasileios C. Kyttaris^{b,*}

^a Department of Medicine, Beth Israel Deaconess Medical Center, Harvard Medical School, Boston, MA, USA
 ^b Division of Rheumatology, Beth Israel Deaconess Medical Center, Harvard Medical School, Boston, MA 02115, USA

ARTICLE INFO

Keywords: Autoimmune diseases COVID-19 Rituximab Hypogammaglobinemia B cells Vaccination

ABSTRACT

Rituximab (RTX) is a very effective treatment for autoimmune rheumatic diseases (AIRD), but it increases infection risk and impairs vaccine responses. Herein we evaluated the antibody response of RTX-treated patients to the supplemental COVID-19 vaccine. After the supplemental dose, 53.1% of patients had detectable antibody titers. Only 36% of patients who did not mount an antibody response after the original vaccine series *did* have detectable antibodies after the supplemental dose (seroconversion). Patients with undetectable CD20⁺ cell levels did not seroconvert while hypogammaglobulinemia was associated with a 15-times decrease in the likelihood of seroconversion. Although we noted 11 COVID-19 infections after the supplemental dose, no patients who received monoclonal antibodies pre-exposure prophylaxis had COVID-19 afterwards. We propose that patients receiving RTX should continue to be prioritized for prophylaxis measures and that vaccination should be timed after B cell recovery wherever possible.

1. Introduction

Rituximab (RTX) is widely used for the treatment of several autoimmune rheumatic diseases (AIRD), including Rheumatoid Arthritis (RA), ANCA-associated Vasculitis (AAV) and a variety of Connective Tissue Diseases (CTD). It a chimeric monoclonal antibody that targets CD20 on B-lymphocytes and induces B cell apoptosis. [1] Although RTX does not directly affect plasma cells, it has been associated with secondary hypogammaglobulinemia [2]. Given its mode action, RTX significantly increases risk of infections including reactivation of hepatitis B, and impairs responses to vaccines [3].

Patients receiving B-cell depleting therapies, such as RTX, have been shown to be vulnerable to COVID-19 and to have poor responses to COVID-19 vaccination [4,5]. Prior studies from our group and others showed that many patients receiving RTX have poor humoral immune responses after vaccination with 2 doses of the BNT162b2 or mRNA-1273 vaccines or 1 dose of Ad26.COV2.SCOVID-19 vaccine. Specifically, we have shown that only a third of rituximab treated patients with AIRD developed measurable titers of IgG anti-SARS-CoV-2 spike antibody after vaccination with the initially recommended doses. One of the main predictors of poor antibody response to anti-SARS-CoV-2 vaccination in that study, was pre-existing hypogammaglobulinemia. [6] Jyssum et al. also showed that most RTX-treated patient did not have an antibody response after 2 vaccine doses. A third dose increased percentages of patients with a serological response, but still less than half responded after 3 doses. T-cell responses though were similar among rituximab and non-rituximab treated patients [7].

On August 13, 2021, the Centers for Disease Control and Prevention (CDC) recommended that immunocompromised individuals receive a supplemental dose (additional primary dose) of COVID-19 vaccine [8]. Herein, we sought to evaluate the effect of the supplemental dose on AIRD patients treated with RTX in relation to vaccine timing, immunological status, infection history and concomitant treatments.

2. Materials and methods

2.1. Study design

We conducted an observational cohort study on adult patients with AIRD treated with RTX at Beth Israel Deaconess Medical Center (BIDMC) in Boston, MA. We measured timing of vaccine administration through chart review and telephone calls to patients. Additional details were also collected regarding disease treatment, COVID-19 infection, demographics and immunologic parameters. The project was approved by

https://doi.org/10.1016/j.clim.2022.109144 Received 18 September 2022; Accepted 4 October 2022 Available online 8 October 2022 1521-6616/© 2022 Elsevier Inc. All rights reserved.

^{*} Corresponding author at: 110 Francis St. Suite 4B, Boston, MA 02215, USA.

E-mail addresses: erose1@bidmc.harvard.edu (E. Rose), daniel.p.magliulo@lahey.org (D. Magliulo), vkyttari@bidmc.harvard.edu (V.C. Kyttaris).

the BIDMC Institutional Review Board.

2.2. Study population

Participants were adult patients (age \geq 18 years). All participants were treated with RTX for an established AIRD, including but not limited to Rheumatoid Arthritis (RA), Antineutrophil Cytoplasmic Antibody Associated Vasculitis (AAV), IgG4-related disease and various connective tissue diseases (CTD, including Systemic Lupus Erythematosus, Mixed Connective Tissue Disease, Anti-synthetase Syndrome). Included patients received at least one dose of Rituximab from January 2020 through to February 2021. Most received subsequent doses during this study.

2.3. Data collection

Medications, indication for RTX by disease, date of last RTX infusion, type of COVID-19 vaccine received, and dates of vaccine administration were collected from a combination of medical records review and patient telephone calls. Post-vaccination serum IgG antibody levels against SARS-CoV-2 spike protein S1 receptor binding domain (RBD), absolute CD19⁺ and CD20⁺ cell counts within 2 months of supplemental dose, and quantitative immunoglobulin levels within one year of supplemental dose were collected from chart review. Documentation of prior SARS-CoV-2 infection and whether requiring hospitalization or intensive care unit level of care was recorded. Hypogammaglobulinemia was defined as laboratory evidence of serum levels of IgG, IgA, or IgM less than the lower limit of normal within 1 year of vaccination date. A prior SARS-CoV-2 infection was determined by either medical record review or by patient reported positive SARS-CoV-2 PCR or rapid test.

2.4. Outcome measures

The primary outcome measure was the proportion of patients receiving RTX treatment who had detectable levels of the anti-spike IgG (seropositive) after vaccination with the supplemental dose of any of BNT162b2 mRNA (manufactured by BioNtech/Pfizer), mRNA-1273 (manufactured by Moderna), and Ad26.COV2.SCOVID-19 (manufactured by Janssen/Johnson & Johnson). Major secondary outcomes include proportion of seropositive patients after vaccination in relation to demographics, immunological parameters at time of vaccination including B-cell counts and immunoglobulin levels, and concomitant medication use. Additional outcomes included incidence of SARS-CoV-2 infection and uptake of pre-exposure prophylaxis.

2.5. Immunogenicity of the vaccine

Serum IgG antibody levels against SARS-CoV-2 spike protein S1 receptor binding domain were measured using anti-spike IgG enzyme immunoassay (Attelica IM COV2G or ADVIA Centaur COV2G, Siemens, Healthineers), which were the tests used at our institution during the study period. The tests were performed via Quest Diagnostics Laboratory. The sensitivity and specificity of these assays are >99%. An Index Value greater than or equal to 1.00 was considered as positive, according to the manufacturer's instruction.

2.6. Statistical analysis

Categorical variables are presented as proportions and continuous variables as median (interquartile range [IQR]). Between group comparisons were done using the chi-square test or Fisher's exact test for categorical variables and the Mann Whitney test to compare continuous variables. Analysis was performed using STATA.

3. Results

3.1. Cohort of rituximab-treated patients with autoimmune rheumatic diseases

In Table 1, we show the demographics, disease status and immunologic profile of the patients that were included in the study. 72 patients

Table 1

Patient demographics, clinical, and immunologic data of autoimmune rheumatic disease patients receiving rituximab.

Patient characteristics Demographics Age (years), median [IQR], $N = 72$ 63 [54.8 71] Gender 48 (66.7) • Female, n (%) 24 (33.3) Underlying disease: 30 (41.7) • Rheumatoid arthritis, n (%) 21 (29.2) • Connective tissue disease; n (%) 32 (4.2) • Connective tissue disease; n (%) 52 (72.2) Supplemental Dose COVD-19 Vaccine® n, (%) 52 (72.2) Supplemental Dose Type 23 (44.2) • BNT162b2 mRNA, n (%) 19 (36.5) • mRNA-1273, n (%) 11 (1.9) • d22.6C0V2.SC0VID-19, n (%) 9 (17.3) Documented history of COVID-19 infection, n (%) 19 (26.4) ⁵ COVID-19 infections after supplemental dose, n (%) 11 (21.2) (Total N = 21) COVID-19 infections fact supplemental dose, n (%) 11 (21.2) (Total N = 21) COVID-19 infections fact supplemental dose, n (%) 20 (28.6) Immunological Parameters B cell counts (measured in $N = 16$): Absolute CD20 + count (#/uL), median [IQR] 0 [0, 9.9] Hypogammaglobulinemia n, (%) 16 (45.7) 12 (43.3) Immunological Parameters	F	
Age (years), median [IQR], N = 72 63 [54.8 71] Gender 48 (66.7) Pemale, n (%) 24 (33.3) • Male, n (%) 24 (33.3) Underlying disease: 30 (41.7) • Rheumatoid arthritis, n (%) 21 (29.2) • Connective tissue disease: n (%) 3 (4.2) • Ig64-related disease, n (%) 3 (4.2) • Ig64-related disease, n (%) 52 (72.2) Supplemental Dose COVID-19 Vaccine [®] n, (%) 52 (72.2) Supplemental Dose COVID-19 vaccine [®] n, (%) 52 (72.2) Supplemental Dose COVID-19 vaccine [®] n, (%) 52 (72.2) Supplemental Dose COVID-19 infection, n (%) 10 (1.9) • Ad26.COV2.SCOVID-19, n (%) 9 (17.3) Unknown, n (%) 10 (26.4) [§] Documented history of COVID-19 infection, n (%) 12 (24.3) (COVID-19 infections leading to hospitalization, n (%) 3 (14.3) (Total N = 21) COVID-19 infections fler supplemental dose, n (%) 11 (21.2) (COVID-19 infections fler supplemental dose, n (%) 10 (0, 10.7] Absolute CD19+ count (#/uL), median [IQR] 0 [0, 10.7] Absolute CD20+ count (#/uL), median [IQR] 0 [0, 9.9] Phypogammaglobul	Patient characteristics	
Age (years), median [IQR], N = 72 63 [54.8 71] Gender 48 (66.7) Pemale, n (%) 24 (33.3) • Male, n (%) 24 (33.3) Underlying disease: 30 (41.7) • Rheumatoid arthritis, n (%) 21 (29.2) • Connective tissue disease: n (%) 3 (4.2) • Ig64-related disease, n (%) 3 (4.2) • Ig64-related disease, n (%) 52 (72.2) Supplemental Dose COVID-19 Vaccine [®] n, (%) 52 (72.2) Supplemental Dose COVID-19 vaccine [®] n, (%) 52 (72.2) Supplemental Dose COVID-19 vaccine [®] n, (%) 52 (72.2) Supplemental Dose COVID-19 infection, n (%) 10 (1.9) • Ad26.COV2.SCOVID-19, n (%) 9 (17.3) Unknown, n (%) 10 (26.4) [§] Documented history of COVID-19 infection, n (%) 12 (24.3) (COVID-19 infections leading to hospitalization, n (%) 3 (14.3) (Total N = 21) COVID-19 infections fler supplemental dose, n (%) 11 (21.2) (COVID-19 infections fler supplemental dose, n (%) 10 (0, 10.7] Absolute CD19+ count (#/uL), median [IQR] 0 [0, 10.7] Absolute CD20+ count (#/uL), median [IQR] 0 [0, 9.9] Phypogammaglobul	Demographics	
Gender 48 (66.7) • Female, n (%) 24 (33.3) • Male, n (%) 24 (33.3) Underlying disease: 30 (41.7) • Rheumatoid arthritis, n (%) 18 (25) • ANCA-associated vasculitis, n (%) 21 (29.2) • Connective tissue disease*, n (%) 3 (4.2) • IgG4-related disease, n (%) 3 (4.2) Supplemental Dose Type 23 (44.2) • BNT162b2 mRNA, n (%) 9 (17.3) • Unknown, n (%) 9 (17.3) Documented history of COVID-19 infection, n (%) 19 (26.4) ⁵ COVID-19 infections leading to hospitalization, n (%) 5 (23.8) (Total N = 21) COVID-19 infections after supplemental dose, n (%) 11 (21.2) (Total N = 21) COVID-19 infections after supplemental dose, n (%) 10 (0, 10.7] Absolute CD19+ count (#/uL), median [IQR] 0 [0, 10.7] Absolute CD20+ count (#/uL), median [IQR] 0 [0, 9.9] Hypogammaglobulinemia n, (%) 12 (34.3) • IgG hypogammaglobulinemia n, (%) 148 (22.5,70] • Immunoglobulin M (mg/dL), median [IQR] 48 (22.5,70] • Immunoglobulin G, median [IQR] <		63 [54 8 71]
$ \begin{array}{ll} & 48 \ (66.7) \\ 24 \ (33.3) \\ & Male, n \ (\%) \\ & Underlying disease: \\ & 30 \ (41.7) \\ & Rheumatoid arthritis, n \ (\%) \\ & 18 \ (25) \\ & ANCA-associated vasculitis, n \ (\%) \\ & 21 \ (29.2) \\ & Connective tissue disease*, n \ (\%) \\ & 21 \ (29.2) \\ & Connective tissue disease*, n \ (\%) \\ & Received Supplemental Dose COVID-19 Vaccine* n, \ (\%) \\ & S2 \ (72.2) \\ & Supplemental Dose Type \\ & 23 \ (44.2) \\ & BNT162b2 mRNA, n \ (\%) \\ & 19 \ (36.5) \\ & mRNA-1273, n \ (\%) \\ & 10 \ (1.9) \\ & A426 \ COV2.SCOVID-19, n \ (\%) \\ & 9 \ (17.3) \\ & Unknown, n \ (\%) \\ & Documented history of COVID-19 infection, n \ (\%) \\ & 19 \ (26.4)^5 \\ & COVID-19 \ infections leading to hospitalization, n \ (\%) \\ & (Total N = 21) \\ & COVID-19 \ infections leading to ICU admission, n \ (\%) \\ & (Total N = 21) \\ & COVID-19 \ infections after supplemental dose, n \ (\%) \\ & (Total N = 21) \\ & COVID-19 \ infections after supplemental dose, n \ (\%) \\ & (Total N = 21) \\ & COVID-19 \ infections after supplemental dose, n \ (\%) \\ & (Total N = 21) \\ & COVID-19 \ infections after supplemental dose, n \ (\%) \\ & (Total N = 21) \\ & COVID-19 \ infections after supplemental dose, n \ (\%) \\ & (Total N = 21) \\ & COVID-19 \ infections after supplemental dose, n \ (\%) \\ & Inmunological Parameters \\ B \ cell \ counts \ (measured in N = 15) \\ & Absolute \ CD20+ \ count \ (\#/uL), median \ [IQR] \ 0 \ (0, 9.9] \\ & Hypogammaglobulinemia \ n, \ (\%) \\ & IgM \ hypogammaglobulinemia \ n, \ (\%) \\ & Immunoglobulin \ (magdL), median \ [IQR] \ 48 \ (22.5, 70] \\ & Hypogammaglobulinemia \ n, \ (\%) \\ & Immunoglobulin \ (measured in N = 35) \\ & IgM \ hypogammaglobulinemia \ n, \ (\%) \\ & Immunoglobulin \ (measured in N = 32 \ patients) \\ & SARS-CoV-2 \ Spike protein \ IgG \ (positive) \ after supplemental dose, n \ (\%) \\ & (ARS, CoV-2 \ Spike protein \ IgG \ (positive) \ after supplemental dose, n \ (\%) \\ & SARS-CoV-2 \ Spike protein \ IgG \ (positive) \ after supplemental dose, n \ (\%) \\ & SARS-CoV-2 \ Spike protein \ IgG \ (positive) \ after supplemental dose, n \ (\%) $		00 [0 110 / 1]
• Female, n (%)24 (33.3)• Male, n (%)30 (41.7)• Rheumatoid arthritis, n (%)18 (25)• ANCA-associated vasculitis, n (%)21 (29.2)• Connective tissue disease*, n (%)3 (4.2)• [gG4-related disease, n (%)3 (4.2)Received Supplemental Dose Type23 (44.2)• BNT162b2 mRNA, n (%)9 (17.3)• MIAnown, n (%)9 (17.3)• Unknown, n (%)9 (17.3)• Unknown, n (%)9 (26.4) ⁵ COVID-19 infections leading to hospitalization, n (%)5 (23.8)(Total N = 21)(COVID-19 infections leading to ICU admission, n (%)3 (14.3)(Total N = 21)(COVID-19 infections after supplemental dose, n (%)11 (21.2)(Total N = 21)(7total N = 21)20 (28.6)Monuological ParametersB cell counts (measured in N = 16): Absolute CD2+ count (#/uL), median [IQR] 0 (0, 10.7] Absolute CD2+ count (#/uL), median [IQR] 	Gender	48 (66 7)
• Male, n (%) Underlying disease: 30 (41.7) • Rheumatoid arthritis, n (%) 18 (25) • ANCA-associated vasculitis, n (%) 21 (29.2) 3 (4.2) • IgG4-related disease, n (%) Received Supplemental Dose COVID-19 Vaccine [®] n, (%) 23 (44.2) • BNT162b2 mRNA, n (%) 19 (36.5) • mRNA-1273, n (%) 10 (1.9) • Ad26.COV2.SCOVID-19, n (%) 10 (0.10.7) • Maknown, n (%) Documented history of COVID-19 infection, n (%) 10 (26.4) [§] COVID-19 infections leading to hospitalization, n (%) 11 (21.2) (COVID-19 infections leading to hospitalization, n (%) 11 (21.2) (Total N = 21) COVID-19 infections after supplemental dose, n (%) 11 (21.2) (Total N = 21) COVID-19 infections after supplemental dose, n (%) 11 (21.2) (Total N = 21) Received monoclonal Ab Pre-exposure prophylaxis, n (%) 20 (28.6) Immunological Parameters B cell counts (measured in N = 16): Absolute CD19 - count (#/uL), median [IQR] 0 [0, 10.7] Absolute CD20 - count (#/uL), median [IQR] 0 [0, 9.9] Hypogammaglobulinemia in the past year n, (%), 12 (34.3) • IgG hypogammaglobulinemia n, (%) • IgG hypogammaglobulinemia n, (%) • IgG hypogammaglobulinemia n, (%) SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (after supplemental dose and SARS-CoV-2 Spike protein IgG measured in N = 32 patients) SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) Medications at the Time of Supplemental Dose (N = 52) Any Steroids at time of Supplemental Dose (N = 52) Any Steroids at time of Supplemental Dose (N = 52) Any Steroids at time of Supplemental Dose (N = 52) Any Steroids at time of Supplemental Dose (N = 52) Any Steroids at time of Supplemental Dose (N = 52) Any Steroids at time of Supplemental Dose (N = 52) Any Steroids at time of Supplemental Dose (N = 52) Any Steroids at time of Supplemental Dose (N = 52) Any Steroids at time of Supplemental Dose (N = 52)	Equals $\pi(0/)$	
Underlying disease: 30 (41.7) 18 (25) ANCA-associated vasculitis, n (%) 21 (29.2) Connective tissue disease*, n (%) 21 (29.2) Connective tissue disease*, n (%) 3 (4.2) IgG4-related disease, n (%) Received Supplemental Dose COVID-19 Vaccine [®] n, (%) 52 (72.2) Supplemental Dose Type 23 (44.2) BNT162b2 mRNA, n (%) 19 (36.5) mRNA-1273, n (%) 1 (1.9) Ad26.COV2.SCOVID-19, n (%) 9 (17.3) Unknown, n (%) Documented history of COVID-19 infection, n (%) 19 (26.4) ^{\$} COVID-19 infections leading to hospitalization, n (%) 5 (23.8) (Total N = 21) COVID-19 infections leading to ICU admission, n (%) 3 (14.3) (Total N = 21) COVID-19 infections after supplemental dose, n (%) 11 (21.2) (Total N = 21) COVID-19 infections after supplemental dose, n (%) 11 (21.2) (Total N = 21) Received monoclonal Ab Pre-exposure prophylaxis, n (%) 20 (28.6) Immunological Parameters B cell counts (measured in N = 16): Absolute CD19+ count (#/uL), median [IQR] 0 [0, 10.7] Absolute CD19+ count (#/uL), median [IQR] 0 [0, 9.9] Hypogammaglobulinemia n, (%) 16 (45.7) 12 (34.3) IgG hypogammaglobulinemia n, (%) 16 (45.7) 12 (34.3) Igg measured in N = 35) IgG hypogammaglobulinemia n, (%) 16 (45.7) 1146.5] Time between supplemental dose and SARS-CoV-2 Spike protein Ig measured in N = 32 patients) SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%), n(%) 146.5] Jime between supplemental dose and SARS-CoV-2 Spike protein Ig measured in N = 32 patients) SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (%) 23.8] Concomitant in Keps patients) SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) 23.8] Concomitant DMARD, n (%) 6 (11.5) Hydroxychloroquine, n (%)		24 (33.3)
30 (41.7)18 (25)• ANCA-associated vasculitis, n (%)21 (29.2)• Connective tissue disease*, n (%)3 (4.2)• [gG4-related disease, n (%)3 (4.2)• BNT162b2 mRNA, n (%)19 (36.5)• mRNA-1273, n (%)10 (36.5)• mRNA-1273, n (%)9 (17.3)• Unknown, n (%)9 (26.4)Documented history of COVID-19 infection, n (%)5 (23.8)(COVID-19 infections leading to hospitalization, n (%)5 (23.8)(Total N = 21)COVID-19 infections leading to hospitalization, n (%)11 (21.2)(Total N = 21)COVID-19 infections after supplemental dose, n (%)11 (21.2)(Total N = 21)COVID-19 infections after supplemental dose, n (%)11 (21.2)(Total N = 21)Received monoclonal Ab Pre-exposure prophylaxis, n (%)20 (28.6)Immunological Parameters8 cell counts (measured in N = 16):Absolute CD19+ count (#/uL), median [IQR]0 [0, 9.9]Hypogammaglobulinemia in the past year n, (%), measured in N = 35)16 (45.7)12 (34.3)• IgG hypogammaglobulinemia n, (%)16 (45.7)1146.5]• Immunoglobulin G, median [IQR]48 [22.5,70]1146.5]• Immunoglobulin G, median [IQR]834 [621.5, 1146.5]1146.5]• SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)13 (25)SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)13 (25)SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)13 (25)SARS-Co		
• Rheumatoid arthritis, n (%)18 (25)• ANCA-associated vasculitis, n (%)21 (29.2)• Connective tissue disease*, n (%)3 (4.2)• IgG4-related disease*, n (%)3 (4.2)Received Supplemental Dose COVID-19 Vaccine* n, (%)52 (72.2)Supplemental Dose Type23 (44.2)• BNT162b2 mRNA, n (%)19 (36.5)• mRNA-1273, n (%)9 (17.3)• Unknown, n (%)9 (17.3)• Unknown, n (%)9 (26.4)Documented history of COVID-19 infection, n (%)19 (26.4)(COVID-19 infections leading to hospitalization, n (%)3 (14.3)(Total N = 21)(COVID-19 infections leading to ICU admission, n (%)3 (14.3)(COVID-19 infections leading to ICU admission, n (%)10 (21.2)(Total N = 21)(COVID-19 infections after supplemental dose, n (%)11 (21.2)(Total N = 21)(COVID-19 infections after supplemental dose, n (%)10 (0, 9.9]Received monoclonal Ab Pre-exposure prophylaxis, n (%)20 (28.6)Immunological ParametersBB cell counts (measured in N = 16):Absolute CD19+ count (#/uL), median [IQR]0 [0, 10.7]Absolute CD29+ count (#/uL), median [IQR]0 [0, 9.9]Hypogammaglobulinemia n, (%)12 (34.3)• IgG hypogammaglobulinemia n, (%)12 (34.3)• IgG hypogammaglobulinemia n, (%)14 (45.5)• Immunoglobulin M (mg/dL), median [IQR]48 [22.5,70]• Immunoglobulin M (mg/dL), median [IQR]834 (621.5, 1146.5]• Immunoglobulin G, median [IQR]834 (621.5, 1146.5]• Immunoglobulin G, m	Underlying disease:	
 ANCA-associated vasculitis, n (%) 21 (29.2) Connective tissue disease", n (%) IgG4-related disease, n (%) Received Supplemental Dose COVID-19 Vaccine[®] n, (%) 52 (72.2) Supplemental Dose Type 23 (44.2) BNT162b2 mRNA, n (%) 19 (36.5) mRNA-1273, n (%) 1 (1.9) Ad26.COV2.SCOVID-19, n (%) 9 (17.3) Unknown, n (%) Documented history of COVID-19 infection, n (%) 19 (26.4)^{\$} COVID-19 infections leading to hospitalization, n (%) 5 (23.8) (Total N = 21) COVID-19 infections leading to ICU admission, n (%) 3 (14.3) (Total N = 21) COVID-19 infections after supplemental dose, n (%) 11 (21.2) (Total N = 21) COVID-19 infections after supplemental dose, n (%) 10 (21.2) (Total N = 21) COVID-19 infections after supplemental dose, n (%) 20 (28.6) Immunological Parameters B cell counts (measured in N = 16): Absolute CD19+ count (#/uL), median [IQR] 0 [0, 10.7] Absolute CD20+ count (#/uL), median [IQR] 0 [0, 9.9] Hypogammaglobulinemia in the past year n, (%), 20 (57.1) (measured in N = 35) IgM hypogammaglobulinemia n, (%) 16 (45.7) 12 (34.3) IgG hypogammaglobulinemia n, (%) Immunoglobulin M (mg/dL), median [IQR] 48 [22.5,70] Immunoglobulin M (mg/dL), median [IQR] 48 [22.5,70] Immunoglobulin M (mg/dL), median [IQR] 64 (45.7) 146.5] SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) 24 (3.8) Are supplemental dose, n (%) SARS-CoV-2 Spike		30 (41.7)
 Connective tissue disease*, n (%) 18 [gG-related disease, n (%) Received Supplemental Dose COVID-19 Vaccine* n, (%) 52 (72.2) Supplemental Dose Type 23 (44.2) BNT162b2 mRNA, n (%) 19 (36.5) mRNA-1273, n (%) 10 (1.9) Ad26.COV2.SCOVID-19, n (%) 9 (17.3) Unknown, n (%) Documented history of COVID-19 infection, n (%) 19 (26.4)^{\$} COVID-19 infections leading to hospitalization, n (%) 5 (23.8) (Total N = 21) COVID-19 infections leading to ICU admission, n (%) 3 (14.3) (Total N = 21) COVID-19 infections after supplemental dose, n (%) 11 (21.2) (Total N = 21) COVID-19 infections after supplemental dose, n (%) 11 (21.2) (Total N = 21) Received monoclonal Ab Pre-exposure prophylaxis, n (%) 20 (28.6) Immunological Parameters B cell counts (measured in N = 16): Absolute CD20+ count (#/uL), median [IQR] 0 [0, 9.9] Hypogammaglobulinemia in the past year n, (%), 20 (57.1) (measured in N = 35) IgM hypogammaglobulinemia n, (%) 16 (45.7) 12 (34.3) IgG hypogammaglobulinemia n, (%) 48 [22.5,70] Immunoglobulin M (mg/dL), median [IQR] 48 [22.5,70] Immunoglobulin M (mg/dL), median [IQR] 48 [22.5,70] Immunoglobulin G, median [IQR] 48 [22.5,70] Masurement in weeks, median [IQR] 48 [22.5,70] ARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) 4.8, 12.3] IgG measurement in weeks, median [IQR] ARS-CoV-2 Spike protein IgG consitive) after supplemental dose, n (%) 6.4,8, 12.3] Medications at the Time of Supplemental Dose (N = 52) Any Steroids at time of supplemental Dose (N = 52) Any Steroids at time of supplemental Do	 Rheumatoid arthritis, n (%) 	18 (25)
 IgG4-related disease, n (%) Received Supplemental Dose COVID-19 Vaccine⁶ n, (%) S2 (72.2) Supplemental Dose Type 23 (44.2) BNT162b2 mRNA, n (%) 19 (36.5) mRNA-1273, n (%) Ad26.COV2_SCOVID-19, n (%) 9 (17.3) Unknown, n (%) Documented history of COVID-19 infection, n (%) (Total N = 21) COVID-19 infections leading to hospitalization, n (%) 3 (14.3) (Total N = 21) COVID-19 infections after supplemental dose, n (%) 11 (21.2) (Total N = 21) COVID-19 infections after supplemental dose, n (%) 11 (21.2) (Total N = 21) Received monoclonal Ab Pre-exposure prophylaxis, n (%) 20 (28.6) Immunological Parameters B cell counts (measured in N = 16): Absolute CD19+ count (#/uL), median [IQR] 0 [0, 10.7] Absolute CD20+ count (#/uL), median [IQR] 0 [0, 9.9] Hypogammaglobulinemia n, (%) 16 (45.7) 12 (34.3) IgG hypogammaglobulinemia n, (%) IgG hypogammaglobulinemia n, (%) IgG hypogammaglobulinemia n, (%) Immunoglobulin M (mg/dL), median [IQR] 48 [22.5,70] Immunoglobulin M (mg/dL), median [IQR] 834 [621.5, 1146.5] Time between supplemental dose and SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, 17 (53.1) n (%) (measured in N = 32 patients) SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%) 23 (23.4) After supplemental dose, n (%) 24 (34.3) Apsteroids at time of supplemental Dose (N = 52) Any Steroids at time of supplemental Dose (N = 52) Any Steroids at time of supplemental Dose (N = 52) Any Steroids at time of supplemental Dose (N = 52) Any Steroids at time of supplemental Dose (N = 52) Any Steroids at time of supplemen	 ANCA-associated vasculitis, n (%) 	21 (29.2)
 IgG4-related disease, n (%) Received Supplemental Dose COVID-19 Vaccine⁶ n, (%) S2 (72.2) Supplemental Dose Type 23 (44.2) BNT162b2 mRNA, n (%) 19 (36.5) mRNA-1273, n (%) Ad26.COV2_SCOVID-19, n (%) 9 (17.3) Unknown, n (%) Documented history of COVID-19 infection, n (%) (Total N = 21) COVID-19 infections leading to hospitalization, n (%) 3 (14.3) (Total N = 21) COVID-19 infections after supplemental dose, n (%) 11 (21.2) (Total N = 21) COVID-19 infections after supplemental dose, n (%) 11 (21.2) (Total N = 21) Received monoclonal Ab Pre-exposure prophylaxis, n (%) 20 (28.6) Immunological Parameters B cell counts (measured in N = 16): Absolute CD19+ count (#/uL), median [IQR] 0 [0, 10.7] Absolute CD20+ count (#/uL), median [IQR] 0 [0, 9.9] Hypogammaglobulinemia n, (%) 16 (45.7) 12 (34.3) IgG hypogammaglobulinemia n, (%) IgG hypogammaglobulinemia n, (%) IgG hypogammaglobulinemia n, (%) Immunoglobulin M (mg/dL), median [IQR] 48 [22.5,70] Immunoglobulin M (mg/dL), median [IQR] 834 [621.5, 1146.5] Time between supplemental dose and SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, 17 (53.1) n (%) (measured in N = 32 patients) SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%) 23 (23.4) After supplemental dose, n (%) 24 (34.3) Apsteroids at time of supplemental Dose (N = 52) Any Steroids at time of supplemental Dose (N = 52) Any Steroids at time of supplemental Dose (N = 52) Any Steroids at time of supplemental Dose (N = 52) Any Steroids at time of supplemental Dose (N = 52) Any Steroids at time of supplemen	 Connective tissue disease*, n (%) 	3 (4.2)
Received Supplemental Dose COVID-19 Vaccine " n, (%)52 (72.2)Supplemental Dose Type23 (44.2)• BNT162b2 mRNA, n (%)19 (36.5)• mRNA-1273, n (%)1 (1.9)• Ad26, COV2. SCOVID-19, n (%)9 (17.3)• Unknown, n (%)Documented history of COVID-19 infection, n (%)19 (26.4)COVID-19 infections leading to hospitalization, n (%)5 (23.8)(Total $N = 21$)COVID-19 infections leading to ICU admission, n (%)3 (14.3)(Total $N = 21$)COVID-19 infections after supplemental dose, n (%)11 (21.2)(Total $N = 21$)Received monoclonal Ab Pre-exposure prophylaxis, n (%)20 (28.6)Immunological ParametersB cell counts (measured in $N = 16$):Absolute CD19+ count (#/uL), median [IQR]0 [0, 10.7]Absolute CD19+ count (#/uL), median [IQR]0 [0, 9.9]12 (34.3)• IgG hypogammaglobulinemia n, (%)16 (45.7)• IgG hypogammaglobulinemia n, (%)12 (34.3)• IgG hypogammaglobulinemia n, (%)9 [4.8, 12.3]• IgG neasurement in weeks, median [IQR]48 [22.5,70]• Immunoglobulin G, median [IQR]834 [621.5,Time between supplemental dose and SARS-CoV-2 Spike protein a (%) (measured in $N = 32$ patients)9 (4.8, 12.3]SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (measured in $N = 32$ patients)13 (25)SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)13 (25)Steroids ≥ 10 mg/day, n (%)2 (3.8)Concomitant DMARD, n (%)9 (17.3)· Mycophenolate, n (%) </td <td></td> <td></td>		
Supplemental Dose Type23 (44.2)• BNT162b2 mRNA, n (%)• mRNA-1273, n (%)1(1.9)• Ad26.COV2.SCOVID-19, n (%)9 (17.3)• Unknown, n (%)Documented history of COVID-19 infection, n (%)19 (26.4)COVID-19 infections leading to hospitalization, n (%)(CoVID-19 infections leading to ICU admission, n (%)(Total $N = 21$)COVID-19 infections after supplemental dose, n (%)(Total $N = 21$)COVID-19 infections after supplemental dose, n (%)(Total $N = 21$)COVID-19 infections after supplemental dose, n (%)(Total $N = 21$)COVID-19 infections after supplemental dose, n (%)(Total $N = 21$)COVID-19 infections after supplemental dose, n (%)20 (28.6)Immunological ParametersB cell counts (measured in $N = 16$):Absolute CD19+ count (#/uL), median [IQR]0 [0, 10.7]Absolute CD20+ count (#/uL), median [IQR]0 [0, 9.9]Hypogaanmaglobulinemia n (%)• IgG hypogaanmaglobulinemia n, (%)• IgG hypogaanmaglobulinemia n, (%)• IgG hypogaanmaglobulinemia n, (%)• Immunoglobulin G, median [IQR]48 [22.5,70]SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%)9 [4.8, 12.3]IgG measured in $N = 32$ patients)SARS-CoV-2 Spike protein IgG changed from negative to positive8/22 (35.4)after supplemental dose, n (%)Medications at the Time of Supplemental Dose ($N = 52$)Any Steroids \geq		52 (72.2)
23 (44.2)• BNT162b2 mRNA, n (%)19 (36.5)• mRNA-1273, n (%)11 (1.9)• Ad26.COV2.SCOUD-19, n (%)9 (17.3)• Unknown, n (%)9 (17.3)Documented history of COVID-19 infection, n (%)19 (26.4)(COVID-19 infections leading to hospitalization, n (%)5 (23.8)(Total $N = 21$)COVID-19 infections leading to ICU admission, n (%)3 (14.3)(Total $N = 21$)COVID-19 infections after supplemental dose, n (%)11 (21.2)(Total $N = 21$)COVID-19 infections after supplemental dose, n (%)20 (28.6)Immunological ParametersEEB cell counts (measured in $N = 16$):Absolute CD19+ count (#/uL), median [IQR]0 [0, 9.9]Absolute CD20+ count (#/uL), median [IQR]0 [0, 9.9]Hypogammaglobulinemia in the past year n, (%), (measured in $N = 35$)20 (57.1)• IgG hypogammaglobulinemia n, (%)16 (45.7)12 (34.3)12 (34.3)• IgG hypogammaglobulinemia n, (%)13 (521.5, 1146.5]Time between supplemental dose and SARS-CoV-2 Spike protein g [4.8, 12.3]9 [4.8, 12.3]IgG measured in $N = 32$ patients)SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, after supplemental dose, n (%)13 (25)Stards-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)13 (25)Steroids ≥ 10 mg/day, n (%)2 (3.8)Concomitar DMARD, n (%)2 (3.8)• Mycophenolate, n (%)9 (17.3)• Hydroxychloroquine, n (%)6 (11.5)• Methotrexate, n (%)6 (11.5		02 (/ 212)
• BNT162b2 mRNA, n (%)19 (36.5)• mRNA-1273, n (%)1 (1.9)• Ad26.COV2.SCOVID-19, n (%)9 (17.3)• Unknown, n (%)9 (17.3)Documented history of COVID-19 infection, n (%)19 (26.4)COVID-19 infections leading to hospitalization, n (%)5 (23.8)(Total N = 21)COVID-19 infections after supplemental dose, n (%)11 (21.2)(Total N = 21)COVID-19 infections after supplemental dose, n (%)11 (21.2)(Total N = 21)COVID-19 infections after supplemental dose, n (%)11 (21.2)(Total N = 21)COVID-19 infections after supplemental dose, n (%)20 (28.6)Immunological ParametersB20 (28.6)B cell counts (measured in N = 16):Absolute CD19+ count (#/uL), median [IQR]0 [0, 10.7]Absolute CD20+ count (#/uL), median [IQR]0 [0, 9.9]Hypogammaglobulinemia in the past year n, (%), (measured in N = 35)20 (57.1)• IgG hypogammaglobulinemia n, (%)16 (45.7)• IgG hypogammaglobulinemia n, (%)11 (46.5]• Immunoglobulin M (mg/dL), median [IQR]48 [22.5,70]• Immunoglobulin G, median [IQR]48 [22.5,70]• Immunoglobulin G, median [IQR]9 [4.8, 12.3]IgG measurement in weeks, median [IQR] (measured in N = 32 patients)9 [4.8, 12.3]SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (measured in N = 32 patients)17 (53.1) n (%)SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%)13 (25)Steroids ≥10 mg/day, n (%)23.8)Concomitant DMARD, n (%)	Supplemental Dose Type	22 (44 2)
• mRNA-1273, n (%) 1 (1.9) • Ad26,COV2.SCOVID-19, n (%) 9 (17.3) • Unknown, n (%) Documented history of COVID-19 infection, n (%) 19 (26.4) [§] COVID-19 infections leading to hospitalization, n (%) 5 (23.8) (Total $N = 21$) COVID-19 infections leading to ICU admission, n (%) 3 (14.3) (Total $N = 21$) COVID-19 infections after supplemental dose, n (%) 11 (21.2) (Total $N = 21$) Received monoclonal Ab Pre-exposure prophylaxis, n (%) 20 (28.6) Immunological Parameters B cell counts (measured in $N = 16$): Absolute CD19+ count (#/uL), median [IQR] 0 [0, 10.7] Absolute CD19+ count (#/uL), median [IQR] 0 [0, 9.9] Hypogammaglobulinemia in the past year n, (%), 20 (57.1) (measured in $N = 35$) • IgM hypogammaglobulinemia n, (%) 16 (45.7) 12 (34.3) • IgG hypogammaglobulinemia n, (%) • Immunoglobulin M (mg/dL), median [IQR] 48 [22.5,70] • Immunoglobulin G, median [IQR] 48 [22.5,70] • Immunoglobulin G, median [IQR] 48 [22.5,70] • Immunoglobulin G, median [IQR] (measured in $N = 32$ patients) SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, 17 (53.1) n (%) (measured in $N = 32$ patients) SARS-CoV-2 Spike protein IgG changed from negative to positive 8/22 (35.4) after supplemental dose, n (%) 13 (25) Steroids ≥10 mg/day, n (%) 2 (3.8) Concomitant DMARD, n (%) 9 (17.3) • Mycophenolate, n (%) 6 (11.5) • Methotrexate, n (%)	$\mathbf{DNT1}(0\mathbf{h}0 \rightarrow \mathbf{DNA} \rightarrow (0/0)$	
 Ad26.COV2.SCOVID-19, n (%) 9 (17.3) Unknown, n (%) Documented history of COVID-19 infection, n (%) 19 (26.4)[§] COVID-19 infections leading to hospitalization, n (%) 5 (23.8) (Total N = 21) COVID-19 infections leading to ICU admission, n (%) 3 (14.3) (Total N = 21) COVID-19 infections after supplemental dose, n (%) 11 (21.2) (Total N = 21) COVID-19 infections after supplemental dose, n (%) 20 (28.6) Immunological Parameters B cell counts (measured in N = 16): Absolute CD19+ count (#/uL), median [IQR] 0 [0, 10.7] Absolute CD20+ count (#/uL), median [IQR] 0 [0, 9.9] Hypogammaglobulinemia in the past year n, (%), 20 (57.1) (measured in N = 35) IgG hypogammaglobulinemia n, (%) Immunoglobulin (mg/dL), median [IQR] 48 [22.5,70] Masser and the form form negative to positive 9 [4.8, 12.3] 12 (34.3) 13 (25) SARS-CoV-2 Spike protein IgG changed from negative to positive 8/22 (35.4) after supplemental dose, n (%) Medications at the Time of Supplemental Dose (N = 52) Any Steroids ≥10 mg/day, n (%) Concomitant DMARD, n (%) Also (23.8) Mycophenolate, n (%) Yethords ≥10 mg/day, n (%) Concomitant DMARD, n (%) Concomitant DMARD, n		
• Unknown, n (%) Documented history of COVID-19 infection, n (%) COVID-19 infections leading to hospitalization, n (%) (Total N = 21) COVID-19 infections leading to ICU admission, n (%) (Total N = 21) COVID-19 infections after supplemental dose, n (%) (Total N = 21) COVID-19 infections after supplemental dose, n (%) (Total N = 21) Received monoclonal Ab Pre-exposure prophylaxis, n (%) 20 (28.6) Immunological Parameters B cell counts (measured in N = 16): Absolute CD19+ count (#/uL), median [IQR] Absolute CD20+ count (#/uL), median [IQR] (measured in N = 35) • IgM hypogammaglobulinemia n, (%) • IngM hypogammaglobulinemia n, (%) • IngM hypogammaglobulinemia n, (%) • Inmunoglobulin G, median [IQR] • IgG hypogammaglobulinemia n, (%) • Immunoglobulin G, median [IQR] • IgG measurement in weeks, median [IQR] • IgG measurement in weeks, median [IQR] • IgG measured in N = 32 patients) SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, 17 (53.1) n (%) (measured in N = 32 patients) SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%) Medications at the Time of Supplemental Dose (N = 52) Any Steroids z 10 mg/day, n (%) • Mycophenolate, n (%) • Mycophenolate, n (%) • Methotrexate, n (%)		
Documented history of COVID-19 infection, n (%)19 (26.4) \$ COVID-19 infections leading to hospitalization, n (%)19 (26.4) \$COVID-19 infections leading to ICU admission, n (%)3 (14.3) (Total N = 21)3 (14.3)COVID-19 infections after supplemental dose, n (%)11 (21.2) (Total N = 21)11 (21.2)Received monoclonal Ab Pre-exposure prophylaxis, n (%)20 (28.6)Immunological ParametersB cell counts (measured in N = 16): Absolute CD19+ count (#/uL), median [IQR]0 [0, 10.7] Absolute CD20+ count (#/uL), median [IQR]0 [0, 9.9]Hypogammaglobulinemia in the past year n, (%), (measured in N = 35)20 (57.1) (measured in N = 35)16 (45.7) 12 (34.3)• IgG hypogammaglobulinemia n, (%)16 (45.7) 12 (34.3)12 (34.3)• IgG hypogammaglobulinemia n, (%)16 (45.5]Time between supplemental dose and SARS-CoV-2 Spike protein IgG measurement in weeks, median [IQR] (measured in N = 32 patients)9 [4.8, 12.3]SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, after supplemental dose, n (%)13 (25)SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)13 (25)Medications at the Time of Supplemental Dose (N = 52) Any Steroids ≥10 mg/day, n (%)2 (3.8) Concomitant DMARD, n (%)2 (3.8) Concomitant DMARD, n (%)• Mycophenolate, n (%)9 (17.3) 7 (13.5)9 (17.3) 7 (13.5)• Methotrexate, n (%)6 (11.5)		9 (17.3)
COVID-19 infections leading to hospitalization, n (%)5 (23.8)(Total $N = 21$)COVID-19 infections leading to ICU admission, n (%)3 (14.3)(Total $N = 21$)COVID-19 infections after supplemental dose, n (%)11 (21.2)(Total $N = 21$)COVID-19 infections after supplemental dose, n (%)11 (21.2)(Total $N = 21$)Received monoclonal Ab Pre-exposure prophylaxis, n (%)20 (28.6)Immunological ParametersB cell counts (measured in $N = 16$):Absolute CD19+ count (#/uL), median [IQR]0 [0, 10.7]Absolute CD20+ count (#/uL), median [IQR]0 [0, 9.9]19Hypogammaglobulinemia in the past year n, (%), (measured in $N = 35$)20 (57.1)• IgG hypogammaglobulinemia n, (%)16 (45.7) 12 (34.3)• IgG hypogammaglobulinemia n, (%)14 (21.5, 1146.5]• Immunoglobulin G, median [IQR]834 [621.5, 1146.5]Time between supplemental dose and SARS-CoV-2 Spike protein IgG measurement in weeks, median [IQR] (measured in $N = 32$ patients)9 [4.8, 12.3]SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (measured in $N = 32$ patients)8/22 (35.4)SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)13 (25)Steroids ≥10 mg/day, n (%) • Mycophenolate, n (%)2 (3.8) Concomitant DMARD, n (%) • (13.5)13 (25)• Mycophenolate, n (%)9 (17.3) 7 (13.5)• Hydroxychloroquine, n (%)6 (11.5)		
(Total $N = 21$) COVID-19 infections leading to ICU admission, n (%) (Total $N = 21$)3 (14.3) (Total $N = 21$)COVID-19 infections after supplemental dose, n (%)11 (21.2) (Total $N = 21$)Received monoclonal Ab Pre-exposure prophylaxis, n (%)20 (28.6)Immunological Parameters8B cell counts (measured in $N = 16$): Absolute CD20+ count (#/uL), median [IQR]0 [0, 10.7] Absolute CD20+ count (#/uL), median [IQR]Newsite CD20+ count (#/uL), median [IQR]0 [0, 9.9]Hypogammaglobulinemia in the past year n, (%), (measured in $N = 35$)20 (57.1) (measured in $N = 35$)• IgG hypogammaglobulinemia n, (%)16 (45.7) 12 (34.3)• IgG hypogammaglobulinemia n, (%)148 [22.5,70]• Immunoglobulin G, median [IQR]834 [621.5, 1146.5]Time between supplemental dose and SARS-CoV-2 Spike protein IgG measurement in weeks, median [IQR] (measured in $N = 32$ patients)9 [4.8, 12.3] 8ARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (measured in $N = 32$ patients)SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)13 (25) Steroids ≥10 mg/day, n (%) 2 (3.8) Concomitant DMARD, n (%) 7 (13.5)• Mycophenolate, n (%)9 (17.3) 7 (13.5)• Mytonychloroquine, n (%)6 (11.5)• Methotrexate, n (%)	Documented history of COVID-19 infection, n (%)	19 (26.4) ^{\$}
COVID-19 infections leading to ICU admission, n (%)3 (14.3)(Total N = 21)(Total N = 21)COVID-19 infections after supplemental dose, n (%)11 (21.2)(Total N = 21)(Total N = 21)Received monoclonal Ab Pre-exposure prophylaxis, n (%)20 (28.6)Immunological ParametersB cell counts (measured in N = 16):Absolute CD19+ count (#/uL), median [IQR]0 [0, 10.7]Absolute CD20+ count (#/uL), median [IQR]0 [0, 9.9]Hypogammaglobulinemia in the past year n, (%), (measured in N = 35)20 (57.1)• IgM hypogammaglobulinemia n, (%)16 (45.7)12 (34.3)12 (34.3)• IgG hypogammaglobulinemia n, (%)16 (45.7)1146.5]1146.5]9 [4.8, 12.3]1146.5]11mmunoglobulin G, median [IQR]48 [22.5,70]11mm between supplemental dose and SARS-CoV-2 Spike protein 1gG measurement in weeks, median [IQR] (measured in N = 32 patients)9 [4.8, 12.3]SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (measured in N = 32 patients)17 (53.1) n (%)SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)13 (25)Steroids ≥10 mg/day, n (%) Concomitant DMARD, n (%)2 (3.8) Concomitant DMARD, n (%)2 (3.8) Concomitant DMARD, n (%)• Mycophenolate, n (%)6 (11.5)•• Methotrexate, n (%)6 (11.5)	COVID-19 infections leading to hospitalization, n (%)	5 (23.8)
(Total N = 21) COVID-19 infections after supplemental dose, n (%) (Total N = 21)11 (21.2) (Total N = 21)Received monoclonal Ab Pre-exposure prophylaxis, n (%) B cell counts (measured in N = 16): Absolute CD19+ count (#/uL), median [IQR] Absolute CD20+ count (#/uL), median [IQR] (measured in N = 35)0 [0, 10.7] 0 [0, 9.9]Hypogammaglobulinemia in the past year n, (%), (measured in N = 35)20 (57.1) (measured in N = 35)• IgG hypogammaglobulinemia n, (%) • Immunoglobulin M (mg/dL), median [IQR] • Immunoglobulin G, median [IQR]48 [22.5,70]• Immunoglobulin G, median [IQR] • Immunoglobulin G, median [IQR]9 [4.8, 12.3]IgG measurement in weeks, median [IQR] (measured in N = 32 patients)9 [4.8, 12.3]SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, after supplemental dose, n (%)17 (53.1) n (%) (measured in N = 32 patients)SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)13 (25)Steroids at time of supplemental Dose (N = 52) Any Steroids at time of supplemental Dose (N = 52) Any Steroids at time of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) <td>(Total $N = 21$)</td> <td></td>	(Total $N = 21$)	
(Total N = 21) COVID-19 infections after supplemental dose, n (%) (Total N = 21)11 (21.2) (Total N = 21)Received monoclonal Ab Pre-exposure prophylaxis, n (%) B cell counts (measured in N = 16): Absolute CD19+ count (#/uL), median [IQR] Absolute CD20+ count (#/uL), median [IQR] (measured in N = 35)0 [0, 10.7] 0 [0, 9.9]Hypogammaglobulinemia in the past year n, (%), (measured in N = 35)20 (57.1) (measured in N = 35)• IgG hypogammaglobulinemia n, (%) • Immunoglobulin M (mg/dL), median [IQR] • Immunoglobulin G, median [IQR]48 [22.5,70]• Immunoglobulin G, median [IQR] • Immunoglobulin G, median [IQR]9 [4.8, 12.3]IgG measurement in weeks, median [IQR] (measured in N = 32 patients)9 [4.8, 12.3]SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, after supplemental dose, n (%)17 (53.1) n (%) (measured in N = 32 patients)SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)13 (25)Steroids at time of supplemental Dose (N = 52) Any Steroids at time of supplemental Dose (N = 52) Any Steroids at time of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) Any Steroids at Ime of supplemental Dose (N = 52) <td>COVID-19 infections leading to ICU admission, n (%)</td> <td>3 (14.3)</td>	COVID-19 infections leading to ICU admission, n (%)	3 (14.3)
COVID-19 infections after supplemental dose, n (%)11 (21.2)(Total N = 21)Received monoclonal Ab Pre-exposure prophylaxis, n (%)20 (28.6)Immunological ParametersB cell counts (measured in $N = 16$): Absolute CD19+ count (#/uL), median [IQR]0 [0, 10.7] 0 [0, 9.9]Absolute CD20+ count (#/uL), median [IQR]0 [0, 9.9]Hypogammaglobulinemia in the past year n, (%), (measured in $N = 35$)20 (57.1) (measured in $N = 35$)• IgG hypogammaglobulinemia n, (%)16 (45.7) 12 (34.3)• IgG hypogammaglobulinemia n, (%)18 (22.5,70]• Immunoglobulin M (mg/dL), median [IQR]834 [621.5, 1146.5]• Imme between supplemental dose and SARS-CoV-2 Spike protein Ig G measurement in weeks, median [IQR] (measured in $N = 32$ patients)9 [4.8, 12.3]SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, after supplemental dose, n (%)13 (25)Steroids at time of supplemental Dose ($N = 52$) Any Steroids at time of supplemental Dose ($N = 52$) Any Steroids at time of supplemental Dose ($N = 52$) Any Steroids at time of supplemental Dose ($N = 52$) Any Steroids at time of supplemental Dose ($N = 52$) Any Steroids at time of supplemental Dose ($N = 52$) Any Steroids at time of supplemental Dose ($N = 52$) Any Steroids at time of supplemental Dose ($N = 52$) Any Steroids at Im (%)2 (3.8) Concomitant DMARD, n (%) 9 (17.3) 7 (13.5)• Hydroxychloroquine, n (%)9 (17.3) 7 (13.5)• Methotrexate, n (%)6 (11.5)	-	
(Total N = 21)Received monoclonal Ab Pre-exposure prophylaxis, n (%)20 (28.6)Immunological ParametersB cell counts (measured in $N = 16$): $Absolute CD19+ count (#/uL), median [IQR]0 [0, 10.7]Absolute CD20+ count (#/uL), median [IQR]0 [0, 9.9]VHypogammaglobulinemia in the past year n, (%),(measured in N = 35)20 (57.1)•IgM hypogammaglobulinemia n, (%)16 (45.7)12 (34.3)•IgG hypogammaglobulinemia n, (%)16 (45.7)12 (34.3)•IgG hypogammaglobulinemia n, (%)16 (45.7)12 (34.3)•IgG magnetic field of the median [IQR]834 [621.5, 1146.5]•Immunoglobulin G, median [IQR]834 [621.5, 1146.5]•Imme between supplemental dose and SARS-CoV-2 Spike protein1g G measurement in weeks, median [IQR] (measured in N = 32patients)9 [4.8, 12.3]SARS-CoV-2 Spike protein IgG (positive) after supplemental dose,n (%) (measured in N = 32 patients)8/22 (35.4)SARS-CoV-2 Spike protein IgG changed from negative to positiveafter supplemental dose, n (%)13 (25)Steroids \geq 10 mg/day, n (%)2 (3.8)Concomitant DMARD, n (%)2 (3.8)• Mycophenolate, n (%)9 (17.3)7 (13.5)• Hydroxychloroquine, n (%)6 (11.5)$		11 (21.2)
Received monoclonal Ab Pre-exposure prophylaxis, n (%)20 (28.6)Immunological ParametersB cell counts (measured in $N = 16$): Absolute CD19+ count (#/uL), median [IQR]0 [0, 10.7] Absolute CD20+ count (#/uL), median [IQR]0 [0, 9.9]Hypogammaglobulinemia in the past year n, (%), (measured in $N = 35$)20 (57.1) (measured in $N = 35$)20 (57.1) (2 (34.3)• IgG hypogammaglobulinemia n, (%)16 (45.7) 12 (34.3)12 (34.3)• IgG hypogammaglobulinemia n, (%)16 (45.7) 12 (34.3)• Immunoglobulin M (mg/dL), median [IQR]48 [22.5,70]• Immunoglobulin G, median [IQR]834 [621.5, 1146.5]• Time between supplemental dose and SARS-CoV-2 Spike protein IgG measurement in weeks, median [IQR] (measured in $N = 32$ 		11 (2112)
Immunological ParametersB cell counts (measured in $N = 16$):Absolute CD19+ count (#/uL), median [IQR]0 [0, 10.7]Absolute CD20+ count (#/uL), median [IQR]0 [0, 9.9]Hypogammaglobulinemia in the past year n, (%), (measured in $N = 35$)• IgM hypogammaglobulinemia n, (%)16 (45.7) 12 (34.3)• IgG hypogammaglobulinemia n, (%)• IgG hypogammaglobulinemia n, (%)• Immunoglobulin M (mg/dL), median [IQR]48 [22.5,70]• Immunoglobulin G, median [IQR]48 [22.5,70]• Immunoglobulin G, median [IQR]9 [4.8, 12.3]IgG measurement in weeks, median [IQR] (measured in $N = 32$ patients)SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (measured in $N = 32$ patients)SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)Medications at the Time of Supplemental Dose ($N = 52$) Any Steroids at time of supplemental dose n, (%)13 (25)Steroids ≥10 mg/day, n (%) 2 (3.8)Concomitant DMARD, n (%) 9 (17.3) 7 (13.5)• Hydroxychloroquine, n (%)• Methotrexate, n (%)		20 (28 6)
B cell counts (measured in N = 16): Absolute CD19+ count (#/uL), median [IQR]0 [0, 10.7] 0 [0, 9.9]Absolute CD20+ count (#/uL), median [IQR]0 [0, 9.9]Hypogammaglobulinemia in the past year n, (%), (measured in N = 35)20 (57.1) (measured in N = 35)• IgM hypogammaglobulinemia n, (%)16 (45.7) 12 (34.3)• IgG hypogammaglobulinemia n, (%)16 (45.7) 12 (34.3)• IgG hypogammaglobulinemia n, (%)16 (45.7) 12 (34.3)• Immunoglobulin M (mg/dL), median [IQR]48 [22.5,70]• Immunoglobulin G, median [IQR]834 (621.5, 1146.5]• Time between supplemental dose and SARS-CoV-2 Spike protein 1gG measurement in weeks, median [IQR] (measured in N = 32 patients)9 [4.8, 12.3]SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (measured in N = 32 patients)17 (53.1) n (%) Medications at the Time of Supplemental Dose (N = 52) Any Steroids at time of supplemental dose n, (%)13 (25) Steroids ≥10 mg/day, n (%) 2 (3.8) Concomitant DMARD, n (%)28 (53.8) 9 (17.3) 7 (13.5)• Hydroxychloroquine, n (%)6 (11.5)• Methotrexate, n (%)6 (11.5)		20 (20.0)
Absolute CD19+ count (#/uL), median [IQR]0 [0, 10.7]Absolute CD20+ count (#/uL), median [IQR]0 [0, 9.9]Hypogammaglobulinemia in the past year n, (%), (measured in $N = 35$)20 (57.1)• IgM hypogammaglobulinemia n, (%)16 (45.7) 12 (34.3)• IgG hypogammaglobulinemia n, (%)16 (45.7) 12 (34.3)• IgG hypogammaglobulinemia n, (%)12 (34.3)• IgG hypogammaglobulinemia n, (%)18 (22.5,70]• Immunoglobulin M (mg/dL), median [IQR]48 [22.5,70]• Immunoglobulin G, median [IQR]834 (621.5, 1146.5]• Imme between supplemental dose and SARS-CoV-2 Spike protein IgG measurement in weeks, median [IQR] (measured in $N = 32$ patients)9 [4.8, 12.3]SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (measured in $N = 32$ patients)17 (53.1) 8/22 (35.4)SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)8/22 (35.4) 13 (25)Medications at the Time of Supplemental Dose ($N = 52$)Any Steroids at time of supplemental dose n, (%)13 (25) Steroids ≥10 mg/day, n (%) 9 (17.3) 7 (13.5)• Hydroxychloroquine, n (%)6 (11.5)• Methotrexate, n (%)6 (11.5)		
Absolute CD20+ count (#/uL), median [IQR]0 [0, 9.9]Hypogammaglobulinemia in the past year n, (%), (measured in $N = 35$)20 (57.1)• IgM hypogammaglobulinemia n, (%)16 (45.7) 12 (34.3)• IgG hypogammaglobulinemia n, (%)12 (34.3)• IgG hypogammaglobulinemia n, (%)834 [621.5, 1146.5]• Immunoglobulin G, median [IQR]834 [621.5, 1146.5]• Immunoglobulin G, median [IQR]834 [621.5, 1146.5]• Immunoglobulin G, median [IQR] (measured in $N = 32$ patients)9 [4.8, 12.3]SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, after supplemental dose, n (%)17 (53.1) 8/22 (35.4)Medications at the Time of Supplemental Dose ($N = 52$) Any Steroids at time of supplemental dose n, (%)13 (25)Steroids ≥10 mg/day, n (%) (Mocomitant DMARD, n (%)28 (53.8) 9 (17.3) 7 (13.5)• Hydroxychloroquine, n (%)6 (11.5)		
Hypogammaglobulinemia in the past year n, (%), (measured in $N = 35$)20 (57.1)• IgM hypogammaglobulinemia n, (%)16 (45.7) 12 (34.3)• IgG hypogammaglobulinemia n, (%)12 (34.3)• IgG hypogammaglobulinemia n, (%)12 (34.3)• Immunoglobulin M (mg/dL), median [IQR]48 [22.5,70]• Immunoglobulin G, median [IQR]834 [621.5, 1146.5]• Time between supplemental dose and SARS-CoV-2 Spike protein IgG measurement in weeks, median [IQR] (measured in $N = 32$ patients)9 [4.8, 12.3]SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (measured in $N = 32$ patients)17 (53.1) 8/22 (35.4)SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)8/22 (35.4)Medications at the Time of Supplemental Dose ($N = 52$) Any Steroids at time of supplemental dose n, (%)13 (25) Steroids ≥10 mg/day, n (%) 9 (17.3) 7 (13.5)• Mycophenolate, n (%)9 (17.3) 7 (13.5)• Hydroxychloroquine, n (%)6 (11.5)		
(measured in $N = 35$)• IgM hypogammaglobulinemia n, (%)16 (45.7) 12 (34.3)• IgG hypogammaglobulinemia n, (%)12 (34.3)• Immunoglobulin M (mg/dL), median [IQR]48 [22.5,70]• Immunoglobulin G, median [IQR]834 [621.5, 1146.5]• Imme between supplemental dose and SARS-CoV-2 Spike protein IgG measurement in weeks, median [IQR] (measured in $N = 32$ patients)9 [4.8, 12.3] SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (measured in $N = 32$ patients)SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)13 (25) Steroids at time of supplemental dose n, (%)Medications at the Time of Supplemental dose n, (%)13 (25) Steroids ≥ 10 mg/day, n (%)2 (3.8) Concomitant DMARD, n (%)• Mycophenolate, n (%)9 (17.3) 7 (13.5)7 (13.5)• Hydroxychloroquine, n (%)6 (11.5)	Absolute CD20+ count (#/uL), median [IQR]	0 [0, 9.9]
• IgM hypogammaglobulinemia n, (%)16 (45.7) 12 (34.3)• IgG hypogammaglobulinemia n, (%)12 (34.3)• IgG hypogammaglobulinemia n, (%)Immunoglobulin M (mg/dL), median [IQR]48 [22.5,70]• Immunoglobulin G, median [IQR]834 [621.5, 1146.5]1146.5]Time between supplemental dose and SARS-CoV-2 Spike protein IgG measurement in weeks, median [IQR] (measured in $N = 32$ patients)9 [4.8, 12.3]SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (measured in N = 32 patients)17 (53.1) after supplemental dose, n (%)SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)8/22 (35.4) after supplemental dose, n (%)Medications at the Time of Supplemental dose n, (%)13 (25) Steroids at time of supplemental dose n, (%)13 (25) Steroids ≥10 mg/day, n (%) 9 (17.3) 7 (13.5)• Hydroxychloroquine, n (%)6 (11.5)• Methotrexate, n (%)5	Hypogammaglobulinemia in the past year n, (%),	20 (57.1)
12 (34.3) • IgG hypogammaglobulinemia n, (%) • Immunoglobulin M (mg/dL), median [IQR] • Immunoglobulin G, median [IQR] • Immunoglobulin G, median [IQR] • Immunoglobulin G, median [IQR] • Immunoglobulin G, median [IQR] • It46.5] Time between supplemental dose and SARS-CoV-2 Spike protein IgG measurement in weeks, median [IQR] (measured in $N = 32$ patients) SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (measured in N = 32 patients) SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%) Medications at the Time of Supplemental Dose (N = 52) Any Steroids at time of supplemental dose n, (%) Steroids $\geq 10 \text{ mg/day}$, n (%) Concomitant DMARD, n (%) • Mycophenolate, n (%) • Methotrexate, n (%)	(measured in $N = 35$)	
 IgG hypogammaglobulinemia n, (%) Immunoglobulin M (mg/dL), median [IQR] Immunoglobulin G, median [IQR] R34 [621.5, 1146.5] Time between supplemental dose and SARS-CoV-2 Spike protein 1gG measurement in weeks, median [IQR] (measured in N = 32 patients) SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (measured in N = 32 patients) SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%) Medications at the Time of Supplemental Dose (N = 52) Any Steroids at time of supplemental dose n, (%) Steroids ≥10 mg/day, n (%) Q (3.8) Concomitant DMARD, n (%) Mycophenolate, n (%) Hydroxychloroquine, n (%) Methotrexate, n (%) 	 IgM hypogammaglobulinemia n, (%) 	16 (45.7)
 IgG hypogammaglobulinemia n, (%) Immunoglobulin M (mg/dL), median [IQR] Immunoglobulin G, median [IQR] R34 [621.5, 1146.5] Time between supplemental dose and SARS-CoV-2 Spike protein 1gG measurement in weeks, median [IQR] (measured in N = 32 patients) SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (measured in N = 32 patients) SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%) Medications at the Time of Supplemental Dose (N = 52) Any Steroids at time of supplemental dose n, (%) Steroids ≥10 mg/day, n (%) Q (3.8) Concomitant DMARD, n (%) Mycophenolate, n (%) Hydroxychloroquine, n (%) Methotrexate, n (%) 		12 (34.3)
• Immunoglobulin M (mg/dL), median [IQR]48 [22.5,70]• Immunoglobulin G, median [IQR]834 [621.5, 1146.5]• Immunoglobulin G, median [IQR]834 [621.5, 1146.5]Time between supplemental dose and SARS-CoV-2 Spike protein IgG measurement in weeks, median [IQR] (measured in $N = 32$ patients)9 [4.8, 12.3]SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (measured in $N = 32$ patients)17 (53.1)SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)8/22 (35.4)Medications at the Time of Supplemental Dose ($N = 52$) Any Steroids at time of supplemental dose n, (%)13 (25)Steroids ≥10 mg/day, n (%) 02 (3.8)Concomitant DMARD, n (%) 9 (17.3) 7 (13.5)9 (17.3) 7 (13.5)• Hydroxychloroquine, n (%)6 (11.5)	• IgG hypogammaglobulinemia n. (%)	
• Immunoglobulin G, median [IQR] 834 [621.5, 1146.5] Time between supplemental dose and SARS-CoV-2 Spike protein 9 [4.8, 12.3] IgG measurement in weeks, median [IQR] (measured in $N = 32$ patients) SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, 17 (53.1) n (%) (measured in $N = 32$ patients) SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%) Medications at the Time of Supplemental Dose ($N = 52$) Any Steroids at time of supplemental dose n, (%) 13 (25) Steroids ≥10 mg/day, n (%) 2 (3.8) Concomitant DMARD, n (%) 9 (17.3) • Mycophenolate, n (%) 6 (11.5) • Methotrexate, n (%)		48 [22.5.70]
1146.5]1146.5]9 [4.8, 12.3]IgG measurement in weeks, median [IQR] (measured in $N = 32$ patients)SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (measured in $N = 32$ patients)SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (measured in $N = 32$ patients)SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)Medications at the Time of Supplemental Dose ($N = 52$)Any Steroids at time of supplemental dose n, (%)13 (25)Steroids ≥ 10 mg/day, n (%)2 (3.8)Concomitant DMARD, n (%)9 (17.3)7 (13.5)Hydroxychloroquine, n (%)6 (11.5)		
Time between supplemental dose and SARS-CoV-2 Spike protein IgG measurement in weeks, median [IQR] (measured in $N = 32$ patients)9 [4.8, 12.3]SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (measured in $N = 32$ patients)17 (53.1) 8ARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)8/22 (35.4) 3 (25)Medications at the Time of Supplemental dose n, (%)13 (25)Steroids ≥ 10 mg/day, n (%)2 (3.8)Concomitant DMARD, n (%)9 (17.3)• Mycophenolate, n (%)6 (11.5)• Methotrexate, n (%)6 (11.5)		
IgG measurement in weeks, median [IQR] (measured in $N = 32$ patients)SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (measured in N = 32 patients)17 (53.1) n (%) (measured in N = 32 patients)SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)8/22 (35.4) after supplemental dose, n (%)Medications at the Time of Supplemental Dose (N = 52) Any Steroids at time of supplemental dose n, (%)13 (25) Steroids ≥10 mg/day, n (%)Concomitant DMARD, n (%)28 (53.8) 9 (17.3) 7 (13.5)• Hydroxychloroquine, n (%)6 (11.5)	Time between sumlemental dass and CADC CaV 2 Caillo motoin	
patients) SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, 17 (53.1) n (%) (measured in N = 32 patients) SARS-CoV-2 Spike protein IgG changed from negative to positive 8/22 (35.4) after supplemental dose, n (%) Medications at the Time of Supplemental Dose (N = 52) Any Steroids at time of supplemental dose n, (%) 13 (25) Steroids $\geq 10 \text{ mg/day}$, n (%) 2 (3.8) Concomitant DMARD, n (%) 28 (53.8) • Mycophenolate, n (%) 9 (17.3) 7 (13.5) • Hydroxychloroquine, n (%) 6 (11.5) • Methotrexate, n (%)		9 [4.8, 12.3]
SARS-CoV-2 Spike protein IgG (positive) after supplemental dose, n (%) (measured in N = 32 patients)17 (53.1)SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)8/22 (35.4)Medications at the Time of Supplemental Dose (N = 52)13 (25)Any Steroids at time of supplemental dose n, (%)2 (3.8)Concomitant DMARD, n (%)28 (53.8)• Mycophenolate, n (%)9 (17.3)7 (13.5)6 (11.5)• Mythorequine, n (%)6 (11.5)	-	
n (%) (measured in N = 32 patients)8/22 (35.4)SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%)8/22 (35.4)Medications at the Time of Supplemental Dose (N = 52)Any Steroids at time of supplemental dose n, (%)13 (25)Steroids ≥ 10 mg/day, n (%)2 (3.8)Concomitant DMARD, n (%)28 (53.8)• Mycophenolate, n (%)9 (17.3)• Hydroxychloroquine, n (%)6 (11.5)• Methotrexate, n (%) $\approx 10^{-1}$		
SARS-CoV-2 Spike protein IgG changed from negative to positive after supplemental dose, n (%) $8/22 (35.4)$ Medications at the Time of Supplemental Dose ($N = 52$)13 (25)Any Steroids at time of supplemental dose n, (%)13 (25)Steroids $\geq 10 \text{ mg/day}$, n (%)2 (3.8)Concomitant DMARD, n (%)28 (53.8)• Mycophenolate, n (%)9 (17.3)• Hydroxychloroquine, n (%)6 (11.5)• Methotrexate, n (%) N		17 (53.1)
after supplemental dose, n (%)Medications at the Time of Supplemental Dose (N = 52)Any Steroids at time of supplemental dose n, (%)13 (25)Steroids $\geq 10 \text{ mg/day, n (%)}$ 2 (3.8)Concomitant DMARD, n (%)28 (53.8)• Mycophenolate, n (%)9 (17.3)• Hydroxychloroquine, n (%)6 (11.5)• Methotrexate, n (%)		
Medications at the Time of Supplemental Dose $(N = 52)$ Any Steroids at time of supplemental dose n, (%)13 (25)Steroids $\geq 10 \text{ mg/day, n (%)}$ 2 (3.8)Concomitant DMARD, n (%)28 (53.8)• Mycophenolate, n (%)9 (17.3)• Hydroxychloroquine, n (%)6 (11.5)• Methotrexate, n (%) \sim	SARS-CoV-2 Spike protein IgG changed from negative to positive	8/22 (35.4)
Any Steroids at time of supplemental dose n, (%)13 (25)Steroids $\geq 10 \text{ mg/day, n}$ (%)2 (3.8)Concomitant DMARD, n (%)28 (53.8)• Mycophenolate, n (%)9 (17.3)• Hydroxychloroquine, n (%)6 (11.5)• Methotrexate, n (%) \sim	after supplemental dose, n (%)	
Steroids ≥10 mg/day, n (%) 2 (3.8) Concomitant DMARD, n (%) 28 (53.8) • Mycophenolate, n (%) 9 (17.3) • Hydroxychloroquine, n (%) 7 (13.5) • Methotrexate, n (%) 6 (11.5)	Medications at the Time of Supplemental Dose ($N = 52$)	
Steroids ≥10 mg/day, n (%) 2 (3.8) Concomitant DMARD, n (%) 28 (53.8) • Mycophenolate, n (%) 9 (17.3) • Hydroxychloroquine, n (%) 7 (13.5) • Methotrexate, n (%) 6 (11.5)	Any Steroids at time of supplemental dose n, (%)	13 (25)
Concomitant DMARD, n (%) 28 (53.8) • Mycophenolate, n (%) 9 (17.3) • Hydroxychloroquine, n (%) 7 (13.5) • Methotrexate, n (%) 6 (11.5)		
 Mycophenolate, n (%) 9 (17.3) 7 (13.5) Hydroxychloroquine, n (%) 6 (11.5) Methotrexate, n (%) 		
7 (13.5) • Hydroxychloroquine, n (%) 6 (11.5) • Methotrexate, n (%)		
 Hydroxychloroquine, n (%) 6 (11.5) Methotrexate, n (%) 	• Mycophenolate, II (70)	
• Methotrexate, n (%)		
	• Hydroxychioroquine, n (%)	ь (11.5)
Weeks from last Rituximab, median [IQR] 24 [18.5, 31]		
	Weeks from last Rituximab, median [IQR]	24 [18.5, 31]

^{*} The connective tissue disease group is composed of systemic lupus erythematosus, inflammatory myopathies, anti-synthetase syndrome, overlap syndromes, and mixed connective tissue disease patients.

 $^{\#}\,$ 3rd doses for patients who received mRNA vaccines or 2nd doses for patients who received JNJ

^{\$} 2 patients were infected twice so there were a total of 21 infections in the cohort documented.

(median [IQR] age, 63 [54.8, 71] years) were included. The most common indication for RTX treatment was RA followed by AAV and CTD. 72.2% patients received a supplemental COVID-19 vaccine from August 2021 to May 2022. Most of the supplemental doses were either the BNT162b2 mRNA vaccine 23 (44.2%) or the mRNA-1273 vaccine 19 (36.5%). One person received Ad26.COV2.SCOVID-19 vaccine as the supplemental dose and 9 have documentation of receiving a supplemental dose but it is unknown which one.

Of the patients who received the supplemental dose, 25% were on steroids and 53.8% were on a concurrent Disease Modifying Anti-Rheumatic Drug (DMARD) at the time of the dose. The median [IQR] time from pre-vaccination RTX infusion to vaccination was 24 [18.5, 31] weeks.

Absolute $CD19^+$ and $CD20^+$ counts were measured in 16 patients within 2 months of the supplemental dose. Most of the patients (56.3%) had undetectable levels of $CD19^+$ and $CD20^+$ measured.

Thirty-five patients had gamma globulins measured within a year of the supplemental dose. 57.1% of patients had hypogammaglobulinemia (low IgG, low IgM or both); 45.7% had IgM hypogammaglobulinemia and 34.3% had IgG hypogammaglobulinemia.

3.2. Antibody response to the vaccine

From August 2021 to May 2022, 32 patients out of 52 (61.5%) had anti-spike antibody serologies measured after receiving the supplemental dose. The median [IQR] time from the supplemental dose to the SARS-CoV-2 spike protein IgG lab draw was 9 [4.8, 12.3] weeks. (Table 1).

Of the 32 patients who had anti-spike antibody serologies measured after the supplemental dose, 53.1% had detectable antibody titers (tested positive). Compared to patients who had positive titers, those who had negative titers had lower absolute CD20⁺ levels (median [IQR] 10.17 [0, 10.71] versus 0 [0,0], p = 0.034). Out of 8 patients with undetectable CD19⁺ and CD20⁺ levels, only 2 had positive titers; whereas out of 5 patients with detectable CD19⁺ and CD20⁺ levels, 4 had positive antibody titers. The 2 patients with undetectable CD19⁺ and CD20⁺ levels but positive spike antibodies also had positive spike antibodies after the first 2 vaccines. Patients with negative titers were more likely to have hypogammaglobulinemia than patients with detectable anti-spike antibodies (85% versus 20%, p = 0.002). There was no difference in gender, age, indication for RTX or type of vaccine received between the patients who had positive titers and those who had negative titers. There was also no difference in corticosteroid use, concomitant DMARDs, time since last RTX dose or cumulative RTX dose. (Table 2).

3.3. Positive antibody response in patients who did not respond to the initial vaccination series (seroconversion)

29 patients had Anti-SARS-CoV-2 Spike Protein IgG measured after both the initial primary series and the supplemental dose. Only 24.1% of those patients had positive titers after the initial primary series. (Table 1) Of the patients who did not mount an antibody response after the original vaccine series, 36% had positive antibodies (seroconverted) after the supplemental dose (Table S1). Patients who seroconverted had higher absolute $CD20^+$ vs the ones who did not mount a detectable antibody response (median [IQR] 10.64 [9.7, 12.42] vs 0 [0,0], p = 0.007). None of the patients with undetectable $CD19^+$ and $CD20^+$ levels seroconverted after the supplemental dose; whereas, 50% of patients with detectable levels did seroconvert. There was a trend toward patients who seroconverted having higher overall IgM levels (76 [52.5, 102.5] versus 34.5 [9.5, 59], *p* = 0.089) and IgG levels (1179.5 [822.5, 1671.5] versus 694 [538.5, 806.5], p = 0.090). Importantly, patients who did not seroconvert were more likely to have IgM hypogammaglobulinemia vs. patients who seroconverted (58% versus 0%, p =0.042]. There was no difference in vaccine type, history of COVID-19 infections, indication for RTX or concomitant rheumatologic

Table 2

Comparisons of patients with positive versus negative titer of Anti-SARS-CoV-2 Spike Protein IgG after the supplemental vaccine dose.

Factor	Spike Titer Negative ¹	Spike Titer Positive ¹	p- value ²
	N = 15	N = 17	
Demographics			
Male Gender, n (%)	7 (47%)	6 (35%)	0.51
Age in years, median, [IQR]	68 [55, 76)]	68 [56, 71]	0.98
Indication for Rituximab, n (%)			
 Rheumatoid Arthritis 	6 (40%)	7 (41%)	0.46
 Vasculitis 	5 (33%)	3 (18%)	
 Connective Tissue Disease 	4 (27%)	5 (29%)	
• IGG4	0 (0%)	2 (12%)	
Initial COVID vaccine received, n (%)			
• BNT162b2	9(64%)	10 (59%)	0.92
• mRNA-1273	4 (29%)	6 (35%)	
 Ad26.COV2.SCOVID-19 	1 (7%)	1 (6%)	
Supplemental COVID vaccine received, n (%)			
• BNT162b2	8 (53%)	10 (59%)	0.11
• mRNA-1273	3 (20%)	6 (35%)	
 Ad26.COV2.SCOVID-19 	0 (0%)	1 (6%)	
 Unknown 	4 (27%)	0 (0%)	
Immunological Parameters			
		10.17 [0,	
CD20 ⁺³ , median [IQR]	0 [0,0]	10.71]	0.034
Hypogammaglobulinemia, ⁴ n (%)	11 (85%)	2 (20%)	0.002
• Low IgM, n (%)	8 (62%)	1 (10%)	0.012
• Low IgG, n (%)	6 (46%)	2 (20%)	0.19
 IgM level, median [IQR] 	32 [10, 54]	69 [53, 106]	0.022
	170.5 [93,	303 [153,	
 IgA level, median [IQR] 	242]	423]	0.099
	702 [546,	1179.5 [724,	
IgG level, median [IQR]	834]	1250]	0.013
Spike Ab titer positive after initial			
series, ^{5,6} n (%)	0 (0%)	7 (47%)	0.003
Spike titer level after initial series, ^{5,6}	0.50.03	0.000	0.004
median [IQR]	0 [0, 0]	0 [0,20]	0.004
Medications at the Time of Supplement		0 (100)	0.14
Steroids, n (%)	6 (40%)	3 (18%)	0.16
Any Concomitant DMARD, n (%)	8 (53%)	8 (47%)	0.72
• Mycophenolate, n (%)	4 (27%)	2 (12%)	0.28
• Hydroxychloroquine, n (%)	1 (7%)	1 (6%)	0.93
• Methotrexate, n (%)	1 (7%)	1 (6%)	0.93
Weeks from last rituximab, median	00 [17 00]	07 [17 46]	0.40
[IQR]	22 [17, 28]	27 [17, 46]	0.42
Rituximab Cumulative Dose, mg,	8570 [5000,	6000 [4000,	0.01
median [IQR]	10,000]	8000]	0.31

¹ Median (IQR) or n(%).

² Mann Whitney or chi-square tests.

³ Within 2 months of supplemental dose if no rituximab in the interim.

⁴ Within 12 months of supplemental dose.

 5 2 doses of the BNT162b2 mRNA or mRNA-1273 vaccines or 1 dose of the Ad26.COV2.SCOVID-19 vaccine.

⁶ Index <1.00 negative, >/= 1.00 positive).

medications between the group that seroconverted and the group that did not (Table S1).

These data show that patients who did not respond to the initial vaccine series were 15.2 times less likely to seroconvert if they were hypogammaglobulinemic. IgM hypogammaglobulinemia in particular was found to be a significant predictor of booster vaccine effectiveness when this is measured by spike antibody levels.

3.4. SARS-CoV-2 infections

In our cohort, there have been 21 infections in 19 patients since the beginning of the pandemic, including 11 infections after the supplemental dose. Five patients (26%) required hospitalization for their COVID-19 infection, including 2 after receiving the supplemental dose. Three of the hospitalized patients (60%) were treated in an ICU. There were no deaths. (Table 1).

The 11 patients with COVID-19 infections after the supplemental

dose were compared to the 41 patients without a COVID-19 infection. There were no differences in RTX indication, type of vaccine received, hypogammaglobinemia or concomitant steroids or DMARDs. Importantly, there were no differences in spike Ab positivity or titer between the group infected after the third dose versus the non-infected group, with the caveat that the time the patients followed after the supplemental dose may not have been sufficient to detect such differences.

3.5. Pre-exposure prophylaxis

In our center, we started using Evusheld (tixagevimab co-packaged with cilgavimab) monoclonal antibodies for pre-exposure prophylaxis in late January 2022. From January through May 2022, 28.6% of RTX-treated patients had received Evusheld and none have had COVID-19 after the prophylaxis. In that same time period, there were 4 COVID-19 infections among the patients that did not receive Evusheld. Patients who received Evusheld and those who did not, were equally likely to have detectable antibody titers after the supplemental dose (46% versus 58%, p = 0.51).

4. Discussion

We report the results of a single center observational cohort study on SARS-CoV-2 supplemental dose vaccination rates and immunogenicity in adult patients with AIRD treated with RTX. Most patients have received the supplemental dose, which for the majority of patients was their third dose. Approximately half (53.1%) of the patients in our study were positive for anti-spike IgG antibodies after the supplemental dose. Other studies have shown that the initial series of one or two doses resulted in seropositivity of about a third. [5,6,9] The supplemental dose has generally been associated with an increase in seropositivity. A recent meta-analysis showed that patients on anti-CD20⁺ therapy had a much lower rate of seroconversion following the booster dose compared to patients receiving non-anti-CD20⁺ therapy (25% vs 81%). [10] In our study, approximately one third of the patients who did not initially mount an antibody response to COVID-19 vaccination, were positive after the supplemental dose. This again supports an increase in seroconversion with additional doses, albeit lower than in patients not on RTX

We found that hypogammaglobulinemia was a strong negative predictor of seroconversion after the supplemental dose. RTX therapy is associated with hypogammaglobulinemia, and this complication has previously been associated with an absence of seropositivity following vaccination and increased rates of serious infection [6,11,12].

Additionally, we found patients with negative spike protein antibody titers had lower absolute CD20⁺ counts. Many patients in this study had undetectable B-cell levels, and those patients were very likely to not respond to the supplemental dose. The B-cell depletion therefore caused by RTX negatively affects antibody responses to vaccinations, as has also been suggested by prior studies. [5,9,13,14] Mrak et al. found that patients with no measurable peripheral B-cells did not develop antibodies after SARS-CoV-2 vaccination during the initial series, but some patients with repopulated B-cells did mount antibody responses. [5] The optimal timing of COVID-19 vaccination in patients treated with rituximab is unclear, but patients may have a better response when they have recovery of B-cell levels. Our study suggests that since none of the patients with undetectable CD20⁺ levels developed new spike protein antibodies following the supplemental dose, those patients may benefit from waiting for B-cell recovery prior to receiving vaccine doses.

Breakthrough infections occurred despite vaccination. Patients with rheumatic diseases have a higher risk of developing SARS-CoV-2 infection and a higher risk of having a poor outcome compared to the general population. [15] Further, patients on B-cell depleting therapy have been shown to be at increased risk of COVID-19-related hospitalization compared with the general population, although subsequent ICU admission or death were infrequent in one study. [16] Another study found that RTX was associated with an increased risk for in-hospital death with COVID-19. [17] While there were no deaths in our cohort, hospitalization including ICU level care for COVID-19, were not uncommon.

Even after infection and vaccination not all patients seroconverted. Prior studies have indicated that patients can recover from COVID-19 infection in the absence of a humoral immune response [18]. Discordant B- and T- cell responses to infection with COVID-19 have been documented. [7] Multiple aspects of the immune system including memory CD4⁺, CD8⁺ T cells, memory B cells and antibodies contribute to long term immunity against SARS-CoV2. [19] Painter et al found that vaccination induced rapid antigen-specific CD4⁺ T cell responses in COVID-19 naive subjects [20]. In patients undergoing B-cell depleting therapy, it is possible that these additional aspects of the immune system may be driving recovery from the infection despite the apparent absence of a humoral response.

-Because of the poor antibody generation, additional strategies are necessary to protect these vulnerable RTX treated patients from COVID-19. One such strategy is monoclonal antibody pre-exposure prophylaxis. Approximately a quarter of RTX-treated patients in this study have received monoclonal antibodies pre-exposure prophylaxis which to date has been highly efficacious in preventing infection. For patients receiving RTX, particularly those with hypogammaglobulinemia or undetectable levels of CD20⁺ cells, pre-exposure prophylaxis can be a good bridge to protect from COVID-19.

This study has several limitations. First, although vaccination status and infection history were identified using multiple data sources, some vaccinations and infections among patient receiving rituximab may have been missed. This could lead to misclassification. Second, prior SARS-CoV-2 infections may influence the captured spike protein percentage. Nucleocapsid antibodies were not measured, so it is possible some of the positive spike antibodies are a response to infection instead of vaccine. Third, the data is obtained from a single health care system which may limit generalizability. Finally, many patients did not receive antibody testing and the timing was inconsistent in the cohort. We do not know the infection prevention behaviors of the patients included.

Despite these limitations, these findings can inform strategies to prevent COVID-19 in patients receiving RTX for autoimmune rheumatic diseases. Following the supplemental dose, additional patients developed antibodies to the spike protein but patients who had hypogammaglobulinemia and/or undetectable B-cell counts oftentimes did not seroconvert. We propose wherever it is feasible to time the vaccination to COVID-19 not only to RTX infusion but most importantly to B cell recovery. In addition, patients who are hypogammaglobulinemic should be strongly considered for pre-exposure prophylaxis. Finally, although B cell depletion impairs vaccine effectiveness, T cell responses are as important: measuring T cell responses to COVID-19 vaccination in patients who are B cell depleted, in a standardized fashion will further assist in risk stratification.

Declaration of Competing Interest

None.

Data availability

Data will be made available on request.

Acknowledgements

This research did not receive any specific grant from funding agencies in the public, commercial, or not-for-profit sectors.

Appendix A. Supplementary data

Supplementary data to this article can be found online at https://doi.

org/10.1016/j.clim.2022.109144.

References

- M.D. Pescovitz, Rituximab, an anti-cd20 monoclonal antibody: history and mechanism of action, Am. J. Transplant. Off. J. Am. Soc. Transplant. Am. Soc. Transplant Surg. 6 (2006) 859–866.
- [2] P. McLaughlin, et al., Rituximab chimeric anti-CD20 monoclonal antibody therapy for relapsed indolent lymphoma: half of patients respond to a four-dose treatment program, J. Clin. Oncol. Off. J. Am. Soc. Clin. Oncol. 16 (1998) 2825–2833.
- [3] S. Aksoy, O. Dizdar, M. Hayran, H. Harputluoğlu, Infectious complications of rituximab in patients with lymphoma during maintenance therapy: a systematic review and meta-analysis, Leuk. Lymphoma 50 (2009) 357–365.
- [4] M.B. Moor, et al., Humoral and cellular responses to mRNA vaccines against SARS-CoV-2 in patients with a history of CD20 B-cell-depleting therapy (RituxiVac): an investigator-initiated, single-Centre, open-label study, Lancet Rheumatol. 3 (2021) e789–e797.
- [5] D. Mrak, et al., SARS-CoV-2 vaccination in rituximab-treated patients: B cells promote humoral immune responses in the presence of T-cell-mediated immunity, Ann. Rheum. Dis. (2021), https://doi.org/10.1136/annrheumdis-2021-220781 annrheumdis-2021-220781.
- [6] D. Magliulo, S.D. Wade, V.C. Kyttaris, Immunogenicity of SARS-CoV-2 vaccination in rituximab-treated patients: effect of timing and immunologic parameters, Clin. Immunol. Orlando Fla 234 (2022), 108897.
- [7] I. Jyssum, et al., Humoral and cellular immune responses to two and three doses of SARS-CoV-2 vaccines in rituximab-treated patients with rheumatoid arthritis: a prospective, cohort study, Lancet Rheumatol. 4 (2022) e177–e187.
- [8] H.E. Fast, Booster and additional primary dose COVID-19 vaccinations among adults aged ≥65 years — United States, august 13, 2021–November 19, 2021, MMWR Morb. Mortal. Wkly Rep. 70 (2021).
- [9] V. Furer, et al., Immunogenicity and safety of the BNT162b2 mRNA COVID-19 vaccine in adult patients with autoimmune inflammatory rheumatic diseases and in

the general population: a multicentre study, Ann. Rheum. Dis. (2021), https://doi. org/10.1136/annrheumdis-2021-220647 annrheumdis-2021-220647.

- [10] A.R.Y.B. Lee, S.Y. Wong, S.H. Tay, Booster COVID-19 vaccines for immunemediated inflammatory disease patients: a systematic review and Meta-analysis of efficacy and safety, Vaccines 10 (2022) 668.
- [11] S.D. Wade, V.C. Kyttaris, Rituximab-associated hypogammaglobulinemia in autoimmune rheumatic diseases: a single-center retrospective cohort study, Rheumatol. Int. 41 (2021) 1115–1124.
- [12] M.Y. Md Yusof, et al., Predicting severe infection and effects of Hypogammaglobulinemia during therapy with rituximab in rheumatic and musculoskeletal diseases, Arthritis Rheumatol. Hoboken NJ 71 (2019) 1812–1823.
- [13] P. Deepak, et al., Effect of immunosuppression on the immunogenicity of mRNA vaccines to SARS-CoV-2: A prospective cohort study, Ann. Intern. Med. 174 (2021) 1572–1585.
- [14] R. Spiera, S. Jinich, D. Jannat-Khah, Rituximab, but not other antirheumatic therapies, is associated with impaired serological response to SARS- CoV-2 vaccination in patients with rheumatic diseases, Ann. Rheum. Dis. 80 (2021) 1357–1359.
- [15] R. Conway, et al., SARS-CoV-2 infection and COVID-19 outcomes in rheumatic diseases: a systematic literature review and Meta-analysis, Arthritis Rheumatol. Hoboken NJ 74 (2022) 766–775.
- [16] L. Boekel, et al., Antibody development and disease severity of COVID-19 in nonimmunised patients with rheumatic immune-mediated inflammatory diseases: data from a prospective cohort study, RMD Open 8 (2022), e002035.
- [17] K.M. Andersen, et al., Long-term use of immunosuppressive medicines and inhospital COVID-19 outcomes: a retrospective cohort study using data from the national COVID cohort collaborative, Lancet Rheumatol. 4 (2022) e33–e41.
- [18] H. Levavi, G. Lancman, J. Gabrilove, Impact of rituximab on COVID-19 outcomes, Ann. Hematol. 1–8 (2021), https://doi.org/10.1007/s00277-021-04662-1.
- [19] J.M. Dan, et al., Immunological memory to SARS-CoV-2 assessed for up to 8 months after infection, Science 371 (2021) eabf4063.
- [20] M.M. Painter, et al., Rapid induction of antigen-specific CD4+ T cells is associated with coordinated humoral and cellular immunity to SARS-CoV-2 mRNA vaccination, Immunity 54 (2021) 2133–2142.e3.