

RESEARCH ARTICLE

Anti-spike antibody response to the COVID vaccine in lymphoma patients

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Data Availability Statement: The study involves a review of the electronic medical records of lymphoma patients who received the COVID vaccine. The research data set has removed most but not all protected health information (PHI). Therefore, the data set contains potentially identifying or sensitive patient information (for example: actual dates of antibody testing were included as needed for analysis but would be considered PHI). The dataset is also considered property of Hackensack Meridian Health and not owned by the investigators. Therefore, for both reasons, the dataset cannot be made openly

Abstract

Patients with hematologic malignancies have poor outcomes from COVID infection and are less likely to mount an antibody response after COVID infection. This is a retrospective study of adult lymphoma patients who received the COVID vaccine between 12/1/2020 and 11/30/2021. The primary endpoint was a positive anti-COVID spike protein antibody level following the primary COVID vaccination series. The primary vaccination series was defined as 2 doses of the COVID mRNA vaccines or 1 dose of the COVID adenovirus vaccine. Subgroups were compared using Fisher's exact test, and unadjusted and adjusted logistic regression models were used for univariate and multivariate analyses. A total of 243 patients were included in this study; 72 patients (30%) with indolent lymphomas; 56 patients (23%) with Burkitt's, diffuse large B-cell lymphoma (DLBCL), and primary mediastinal B-cell lymphoma (PMBL) combined; 55 patients (22%) with chronic lymphocytic leukemia or small lymphocytic lymphoma (CLL/SLL); 44 patients (18%) with Hodgkin and T-cell lymphomas (HL/TCL) combined; 12 patients (5%) with mantle cell lymphoma; and 4 patients (2%) with other lymphoma types. One-hundred fifty-eight patients (65%) developed anti-COVID spike protein antibodies after completing the primary COVID vaccination series. Thirty-eight of 46 (83%) patients who received an additional primary shot and had resultant levels produced anti-COVID spike protein antibodies. When compared to other lymphoma types, patients with CLL/SLL had a numerically lower seroconversion rate of 51% following the primary vaccination series whereas patients with HL/TCL appeared to have a robust antibody response with a seropositivity rate of 77% ($p = 0.04$). Lymphoma patients are capable of mounting a humoral response to the COVID vaccines. Further studies are required to confirm our

available per restrictions imposed by Hackensack Meridian Health IRB. However, upon request, we are willing to share portions of the data for appropriate review. Requests can be made through the corresponding author (Dr Andrew Ip; Email: Andrew.Ip@hmn.org) or directly to representatives of Hackensack Meridian Health IRB (Email: hmhrib@hmn.org; Phone: 201-880-3669).

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Competing interests: The authors have read the journal's policy and have the following competing interests: Lori A. Leslie is a consultant, honoraria, and/or member of the speakers bureau for Seattle Genetics, KitePharma, Celgene, Pharmacyclics, Janssen, AstraZeneca, Beigene, Epizyme, Karyopharm, AbbVie, and TG Therapeutics; and is a member of the advisory board for KitePharma, Seattle Genetics, ADC Therapeutics, Janssen, Pharmacyclics, Beigene, AbbVie, AstraZeneca, TG Therapeutics, and Epizyme. Andre H. Goy is a consultant, honoraria, and/or member of the speakers bureau for Pharmacyclics, Janssen, Celgene, Novartis, Hoffman la Roche, Kite, Pharmacyclics/AbbVie, Elsevier's Practice Update Oncology, Clinical Advances in Hematology & Oncology, Medscape, Physicians Education Resource LLC, Roswell Park, OncLive Peer Exchange, Michael J Hennessey Associates Inc., BMS, Xcenda, MorphoSys, and AstraZeneca; is a member of the advisory board for Janssen, Vincerx Pharma, Kite Pharma, Elsevier's Practice Update Oncology, Gilead, AbbVie, Pharmacyclics, BMS, AstraZeneca, and Alloplex; is on the board of directors of COTA, Genomic Testing Cooperative; has resilience stock in COTA, Genomic Testing Cooperative, Alloplex, and Resilience; and has received research funding from Janssen, Acerta, Celgene, Constellation, Genentech, Hoffman la Roche, Infinity Pharmaceuticals, MorphoSys, Karyopharm, Kite, Pharmacyclics, AstraZeneca, Verastem, Seattle Genetics, and BMS. Tatyana A. Feldman is a consultant, honoraria, and/or member of the speakers bureau for AbbVie, ADC Therapeutics, BMS, Celgene, Janssen, Pharmacyclics, Seattle Genetics, and Takeda; and is a member of the advisory board for AbbVie, ADC Therapeutics, Daiichi Sankyo, Genmab, Karyopharm, KITE, Kyowa Kirin, and Morphosys. Andrew Ip is a consultant, honoraria, and/or member of the speakers bureau for TG Therapeutics and AstraZeneca, and is a member of the advisory board for Secura Bio. There are no patents, products in development or marketed products associated with this research to declare.

findings, including whether T-cell immunity would be of clinical relevance in this patient population.

Introduction

Patients with hematologic malignancies have poor outcomes from COVID infection with associated mortality of up to 30–40% [1,2]. Studies have shown that these patients are less likely to mount an antibody response after COVID infection [3]. The Pfizer-BioNTech and Moderna COVID mRNA vaccines have been shown to be 94% effective in preventing severe disease in the general population [4]. However, it is well known that certain clinically vulnerable populations do not develop protective immunity after completing the COVID vaccine series [5].

Production of antibodies against the SARS-CoV-2 spike protein is the major mechanism of protective immunity induced by COVID vaccination. Patients with lymphoma may be unable to seroconvert following vaccination due to impaired humoral and T-cell immunity secondary to the malignancy itself as well as immunosuppressive treatment [6]. This inability to produce antibodies against the SARS-CoV-2 spike protein is most notable in patients with B-cell lymphomas and chronic lymphocytic leukemia (CLL), with seroconversion rates ranging between 64% to 78% [7]. Treatment with B-cell directed therapies, such as anti-CD20 monoclonal antibodies and Bruton's tyrosine kinase (BTK) inhibitors, further compromises the ability of patients with lymphoma to mount an antibody response following vaccination and persists over time [8]. Vaccine responses in patients with different lymphoma subtypes or receiving B-cell depleting therapies have not been fully elucidated.

With the emergence of the omicron variant and availability of an additional primary shot of the COVID vaccine, it has become increasingly important to describe the efficacy of the COVID vaccines in lymphoma patients and to suggest the optimal timing of vaccination to elicit immunity in patients receiving immunochemotherapy. There have been several reports of patients with lymphoma frequently not achieving serologic response to COVID vaccines, especially after recent treatments with B-cell directed therapies [7–15]. However, these studies have varying limitations, which include the following: lack of serology data following an additional primary shot or third vaccine dose; exclusion of certain patient groups (like those with a history of COVID infection or specific lymphoma subtypes) from the analysis; and less rigorous methods (such as utilizing patient reports or multiple serology assays). In this study, we evaluated antibody levels against the SARS-CoV-2 spike protein in lymphoma patients following receipt of the primary COVID vaccination series and additional primary shot. We hypothesized that patients with lymphoma are not developing a robust immune response following vaccination.

Methods

Study design and cohort selection

This was a retrospective, single center study at the John Theurer Cancer Center (JTCC) within Hackensack Meridian Health (HMH). Electronic health record (EHR)-derived data was utilized to identify patients who received at least 2 doses of the COVID mRNA vaccines or at least 1 dose of the COVID adenovirus vaccine. The primary objective was to evaluate the efficacy of the COVID vaccines following 2 doses of the COVID mRNA vaccines or 1 dose of the COVID adenovirus vaccine.

This does not alter our adherence to PLOS ONE policies on sharing data and materials."

Patients were included in this analysis if they were: 1) Adult patients ≥ 18 years; 2) Had a diagnosis of lymphoma or CLL; 3) Received the COVID vaccine series between December 15, 2020 and November 30, 2021; 4) had a resulted anti-COVID spike protein antibody level following 2 doses of the COVID mRNA vaccines (Pfizer-BioNTech vaccine and/or Moderna COVID mRNA vaccine) or after 1 dose of the COVID adenovirus vaccine (Johnson&Johnson). The Elecsys Anti-SARS-CoV-2 electrochemiluminescence immunoassay (Roche Diagnostics) was used to measure the antibody response.

Institutional Review Board (IRB) approval was obtained under HMH IRB. This trial was conducted under the International Conference on Harmonization Good Clinical Practice guidelines and according to the Declaration of Helsinki. The requirement for patient informed consent (verbal or written) was waived by the IRB as this project represented a non-interventional study utilizing routinely collected data for secondary research purposes.

Data sources

Data was collected from HMH's EHR (Epic) which is utilized throughout the network. Patients treated at JTCC were identified by convenience sampling if they shared their vaccination status at the time of their outpatient visit. Demographics, clinical characteristics, treatments, and outcomes were manually extracted by physicians, pharmacists, nurses, and medical students from JTCC and HMH. Data abstracted by the team was entered into an Excel spreadsheet, and quality control was performed by physicians overseeing data collection.

Demographic data included age, sex, race/ethnicity, co-morbidities, and last anticancer treatment received. Co-morbidities were defined as any of the following present prior to receipt of the COVID vaccine series: chronic obstructive pulmonary disorder (COPD), autoimmune diseases, liver disease, diabetes mellitus, chronic kidney disease (CKD), human immunodeficiency virus (HIV), cardiovascular disease, and smoking history. Lymphoma types were categorized as follows: chronic lymphocytic leukemia (CLL)/small lymphocytic leukemia (SLL); Burkitt's lymphoma/diffuse large B-cell lymphoma (DLBCL)/primary mediastinal B-cell lymphoma (PMBL); indolent B-cell lymphomas; mantle cell lymphoma (MCL); and Hodgkin's lymphoma/T-cell lymphomas (HL/TCL). Indolent lymphomas included follicular lymphoma, marginal zone lymphoma, lymphoplasmacytic lymphoma, and Waldenstrom macroglobulinemia. T-cell lymphomas included peripheral T-cell lymphomas not otherwise specified (NOS), anaplastic large cell lymphoma (ALCL), cutaneous T-cell lymphomas, angioimmunoblastic T-cell lymphoma (AITL), and acute T-cell leukemia/lymphoma (ATLL). Most recent anticancer treatment types (such as cytotoxic chemotherapy, anti-CD20 monoclonal antibody therapy, or small molecule inhibitors), date of last treatment prior to vaccination, and duration of treatment were also collected.

Outcome measures

The primary outcome was a positive anti-COVID spike protein antibody level following the primary COVID vaccination series. A positive antibody level was defined based on the 2 reference laboratories used at HMH—Quest Diagnostics used a cutoff of ≥ 0.8 U/mL, and BioReference Laboratories, Inc. used a cutoff of ≥ 15 AU/mL. The primary vaccination series was defined as either 2 doses of the COVID mRNA vaccines or 1 dose of the COVID adenovirus vaccine. Among patients who had anti-COVID spike protein antibody results, COVID-19 history was collected and included the following (if available): type of COVID vaccine received and number of doses, history of positive COVID infection, and therapies received for COVID treatment (if applicable).

Secondary outcomes included positive anti-COVID spike protein antibody levels in the following patient subgroups: patients who received an additional primary shot; different

lymphoma subtypes; patients who received anti-CD20 monoclonal antibody therapy within the previous 12 months; and patients who received active anticancer treatment. The additional primary shot was defined as either an additional dose (“third dose”) of the COVID mRNA vaccine or an additional dose (“second dose”) of the COVID adenovirus vaccine. Active anticancer treatment was defined as the receipt of any therapy intended to treat cancer that was administered within the six months prior to initiating the primary COVID vaccination series.

Statistical analysis

Demographic and clinical characteristics were summarized using descriptive statistics such as percentiles, median, and interquartile range (IQR) by the primary vaccination series and additional primary shot (third vaccine dose) groups, respectively. Seropositivity rates between demographic and clinical categories were compared using Fisher’s exact test or Pearson’s chi-squared tests. Unadjusted and multivariable adjusted logistic regression models (UVA and MVA, respectively) using variables with $p < 0.05$ in the unadjusted model were then fitted and their odds ratios (OR) and confidence intervals (CI) were computed. A two-sided p -value < 0.05 is considered for the statistical significance. All analyses were performed using R (*ver* 4.1, R Core Team (2021), Vienna, Austria) software.

Results

Patient demographics and comorbidities

Two-hundred forty-three patients were identified and included in this analysis with baseline characteristics as shown in [Table 1](#). Overall, the study population was elderly at a median age of 67 (IQR 59–75) years old, 52% of patients were male, and 78% of patients were white. The most frequent comorbidities were cardiovascular disease (44%) and former smoking history (30%). Lymphoma subtypes in our cohort were: indolent lymphomas (30%), Burkitt’s, DLBCL, PMBL combined (23%), CLL/SLL (22%), Hodgkin’s and T-cell lymphomas (HL/TCL) combined (18%), mantle cell lymphoma (MCL) (5%), and other types of lymphomas (2%). As of November 30, 2021, one-hundred twelve (46%) patients received an additional primary shot.

Testing for anti-COVID spike protein antibodies occurred at a median of 61 (IQR 29–98) days after completion of the primary vaccination series and at a median of 37 (IQR 21–56) days after an additional primary shot. Majority of patients received COVID mRNA vaccines, and we were able to confirm the specific type in 197 (81%) patients. Only 3 patients received the COVID adenovirus vaccine ([Table 1](#)).

Primary outcome

One-hundred fifty-eight patients (65%) developed anti-COVID spike protein antibodies after completing their primary COVID vaccination series ([Fig 1](#) and [Table 2](#)). There were 112 patients who received an additional primary shot, but only 46 of these patients had anti-COVID spike antibody levels resulted at the time of data cut-off. Of these 46 patients, 38 patients (83%) had anti-COVID spike protein antibodies after receiving this dose ([Table 2](#)). Among the 38 patients who were seropositive after the additional primary dose, 18 patients were seronegative after their primary COVID vaccination series and seroconverted only after the additional primary shot of COVID vaccine.

There were 8 patients that did not produce antibodies following both the primary vaccination series and additional primary shot. Of these 8 patients, 6 (75%) were diagnosed with indolent lymphomas, 1 (12.5%) with CLL/SLL, and 1 (12.5%) with MCL. At the time of

Table 1. Baseline characteristics.

Patient characteristics	Primary vaccination series N = 243 n(%)	Additional primary shot N = 112 n(%)
Age, years, median (IQR)	67 (59–75)	67 (59–75)
Male	127 (52)	59 (53)
Race		
Asian	4 (2)	0
Black or African American	11 (4)	3 (3)
Hispanic or Latino	23 (10)	6 (5)
White	189 (78)	98 (88)
Other	11 (4)	3 (3)
Missing	5 (2)	2 (2)
Disease type		
Burkitt's, DLBCL, PMBL	56 (23)	24 (21)
CLL/SLL	55 (22)	24 (21)
Hodgkin's lymphomas (HL)	20 (8)	9 (8)
Indolent lymphomas [†]	72 (30)	36 (32)
Mantle cell lymphoma (MCL)	12 (5)	8 (7)
T-cell lymphomas (TCL) [‡]	24 (10)	8 (7)
Other [§]	4 (2)	3 (3)
Comorbidities		
Cardiovascular disease [¶]	106 (44)	49 (44)
COPD	6 (3)	3 (3)
Autoimmune disease ^{††}	27 (11)	13 (12)
Mild liver disease ^{‡‡}	6 (3)	2 (2)
Diabetes	38 (16)	19 (17)
Chronic kidney disease	10 (4)	7 (6)
HIV/AIDS	6 (3)	2 (2)
Smoking history	8 (3)	4 (4)
Current smoker	74 (30)	33 (30)
Former smoker		
Vaccine received		
Pfizer-BioNTech	115 (47)	66 (59)
Moderna	79 (33)	41 (37)
Johnson&Johnson	3 (1)	0
Unknown	46 (19)	5 (4)

[†]Includes follicular lymphomas, marginal zone lymphoma, lymphoplasmacytic lymphoma, and Waldenstrom macroglobulinemia.

[‡]Includes peripheral T-cell lymphomas not otherwise specified (NOS), anaplastic large cell lymphoma (ALCL), cutaneous T-cell lymphomas, angioimmunoblastic T-cell lymphoma (AITL), and acute T-cell leukemia/lymphoma (ATLL).

[§]Include monoclonal gammopathy of undetermined significance (MGUS), T-cell prolymphocytic leukemia (T-PLL), Essential thrombocytosis, and primary CNS lymphoma.

[¶]Cardiovascular disease defined as coronary artery disease, hypertension, or cerebrovascular accident.

^{††}Includes autoimmune diseases, rheumatological disorders, and connective tissue disease.

^{‡‡}Mild liver disease defined as either chronic hepatitis or cirrhosis without portal hypertension.

DLBCL = diffuse large B-cell lymphoma; PMBL = primary mediastinal B-cell lymphoma; COPD = chronic obstructive pulmonary disease; HIV/AIDS = human immunodeficiency virus/acquired immunodeficiency syndrome.

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vaccination, 4 of these 8 (50%) patients were receiving active anticancer treatment. Active anticancer treatment in these patients consisted of 50% (2/4) BTK inhibitor monotherapy, 25% (1/4) rituximab with chemotherapy, and 25% (1/4) other therapy. One patient experienced progression of chronic graft-versus-host disease (cGVHD) at the time of vaccination and was receiving immunosuppressive treatments, including steroids, etanercept, sirolimus, tacrolimus, and belumosudil, that may have diminished the ability to mount an immune response.

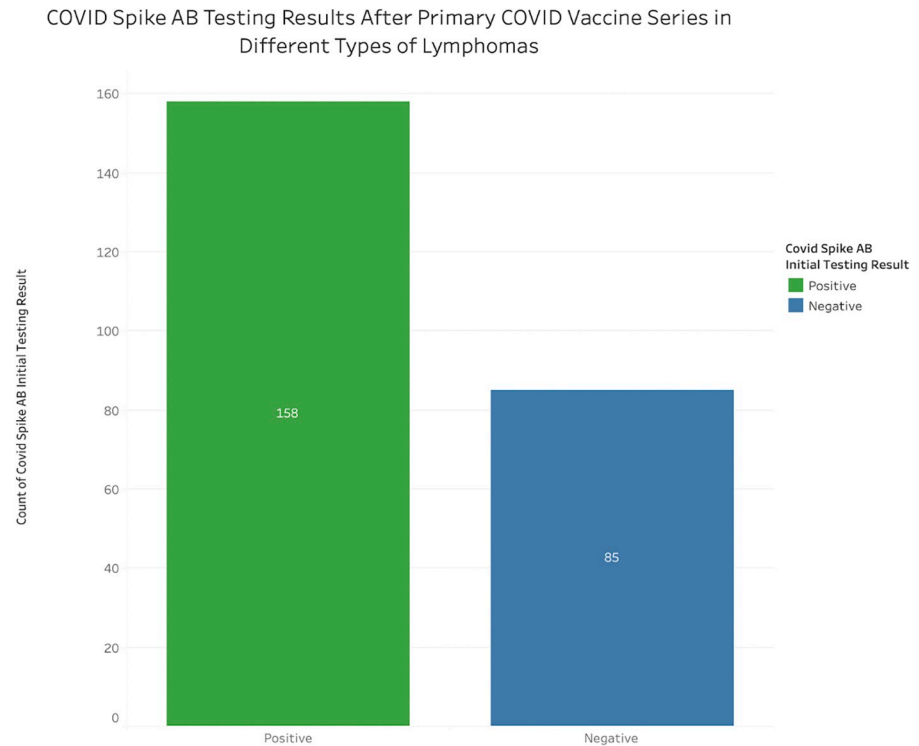


Fig 1. Serologic response against COVID spike protein after primary vaccination series in the overall population.

<https://doi.org/10.1371/journal.pone.0266584.g001>

Secondary outcomes

There were 45 patients who received an anti-CD20 monoclonal antibody-containing regimen in the last 12 months prior to vaccination. Of these patients, 47% (21/45) produced antibodies after the primary COVID vaccination series and 57% (4/7) produced antibodies after the additional primary shot (Table 2). When compared to those who did not receive any anti-CD20 monoclonal antibody therapy, patients who received an anti-CD20 monoclonal antibody-containing regimen in the past 12 months prior to vaccination had a significantly lower rate of seroconversion (UVA OR 0.51 (0.26, 1.00), $p = 0.05$; MVA 3.35 (1.47, 7.63), $p = 0.004$). In contrast, patients who received an anti-CD20 monoclonal antibody more than 12 months prior to vaccination did not have a statistically significant lower rate of seroconversion than patients who were never treated with anti-CD20 antibody therapy (UVA OR 0.62 (0.27, 1.42), $p = 0.26$; MVA OR 2.38 (0.85, 6.70), $p = 0.10$).

There were differences observed in the ability to produce serology with the COVID vaccine amongst lymphoma subtypes. Of 55 patients with CLL/SLL, 28 (51%) produced antibodies after the primary COVID vaccination series (Table 2 and Fig 2). There were 12 patients who received an additional primary shot, of which 92% were seropositive (Fig 3). There were 20 CLL/SLL patients receiving anticancer treatment at the time of vaccination, of which 7 (7/20, 35%) patients were seropositive after the primary vaccination series and 6 (6/7, 86%) after the additional primary shot (Table 3). The most common treatment type in these patients was monotherapy with a Bruton's tyrosine kinase (BTK) inhibitor, where 36% (4/11) of patients produced antibodies. In comparison, there were 35 CLL/SLL patients not receiving active anticancer treatment and had a numerically higher rate of seroconversion at 60% (21/35) following

Table 2. Results.

Patients	Seropositivity against COVID spike protein		p-value; UVA OR (95% CI), p-value; MVA OR (95% CI), p-value
	Primary vaccination series no. of events/ total no. (%)	Additional primary shot [†] no. of events/ total no. (%)	
Overall population	158/243 (65)	38/46 (83)	
<i>Timing of last anticancer treatment</i>			<0.001;
None (ref)	30/46 (65)	9/9 (100)	0.51 (0.24, 1.06), p = 0.072,
Within the last 6 months	45/92 (49)	20/24 (83)	2.01 (0.93, 4.33), p = 0.074;
More than 6 months	83/105 (79)	9/13 (69)	0.52 (0.22, 1.23), p = 0.14, 1.16 (0.43, 3.08), p = 0.77
<i>Timing of anti-CD20 monoclonal antibody</i>			<0.001;
None (ref)	91/144 (63)	29/32 (91)	0.51 (0.26, 1.00), p = 0.05, 3.35 (1.47, 7.63), p = 0.004;
Received rituximab in past 12 months	21/45 (47)	4/7 (57)	0.62 (0.27, 1.42), p = 0.26,
Received rituximab ≥12 months	46/54 (85)	5/7 (71)	2.38 (0.85, 6.70), p = 0.10
<i>Timing of IVIG</i>			0.014;
Received no IVIG. (ref)	145/213 (68)	31/38 (82)	0.36 (0.16, 0.78), p = 0.01;
Received IVIG previous 6 months	13/30 (43)	7/8 (88)	0.39 (0.17, 0.93), p = 0.03
<i>Disease type</i>			0.016;
CLL/SLL (ref)	42/56 (75)	11/12 (92)	2.89 (1.30, 6.46), p = 0.01; 2.71 (1.04, 7.06), p = 0.04.
Burkitt's, DLBCL, PMBL	28/55 (51)	10/10 (100)	3.27 (1.39, 7.91), p = 0.01;
HL/TCL [‡]	34/44 (77)	3/3 (100)	3.29 (1.25, 8.64), p = 0.02.
Indolent lymphomas [§]	44/72 (61)	9/15 (60)	1.52 (0.75, 3.81), p = 0.25;
MCL	6/12 (50)	4/5 (80)	1.41 (0.62, 3.20), p = 0.42.
Other [¶]	4/4 (100)	1/1 (100)	0.96 (0.27, 3.36), p = 0.95; 1.29 (0.33, 5.03), p = 0.71. Inf, p = 0.98; Inf, p = 0.98
<i>History of COVID infection</i>			0.09;
Negative (ref)	139/216 (64)	37/45 (82)	2.6 (0.8, 7.8), p = 0.1; 3.1 (1, 9.9), p = 0.06
Positive	19/24 (79)	1/1 (100)	

[†]In patients with resulted anti-COVID spike protein antibody levels.

[‡]Includes peripheral T-cell lymphomas not otherwise specified (NOS), anaplastic large cell lymphoma (ALCL), cutaneous T-cell lymphomas, angioimmunoblastic T-cell lymphoma (AITL), and acute T-cell leukemia/lymphoma (ATLL).

[§]Includes follicular lymphomas, marginal zone lymphoma, lymphoplasmacytic lymphoma, and Waldenstrom macroglobulinemia.

[¶]Include monoclonal gammopathy of undetermined significance (MGUS), T-cell prolymphocytic leukemia (T-PLL), Essential thrombocytosis, and primary CNS lymphoma.

DLBCL = diffuse large B-cell lymphoma; PMBL = primary mediastinal B-cell lymphoma; CLL = chronic lymphocytic leukemia; SLL = small lymphocytic lymphoma; HL = Hodgkin lymphomas; TCL = T-cell lymphomas; MCL = mantle cell lymphoma; IVIG = intravenous immune globulin.

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receipt of the primary COVID vaccine series. CLL/SLL patients were less likely to mount an antibody response to the COVID vaccine when compared to those with other types of lymphoma (p = 0.016).

For patients with HL/TCL, 34 of 44 (77%) patients produced antibodies after the primary vaccination series and all patients (3/3, 100%) after the additional primary shot (Table 2, Figs 2 and 3). Of 20 HL/TCL patients who received anticancer treatment in the last 6 months, 70% (14/20) produced antibodies at a median level of 120 AU/mL (reference ≥ 15 AU/mL), with 7 patients having a robust response of antibody levels >400 AU/mL. On statistical analysis, HL/TCL patients were more likely to elicit an antibody response to the COVID vaccine when compared to those with CLL/SLL, and this response was significant on UVA (OR 3.27, 95% CI (1.39, 7.91), p = 0.01) and MVA (OR 3.29, 95% CI (1.25, 8.64), p = 0.02) when active anticancer therapy, lymphoma diagnosis, and history of COVID infection were considered in the multivariate analysis.

Covid Spike Antibody Result After Primary Series in Different Types of Lymphomas

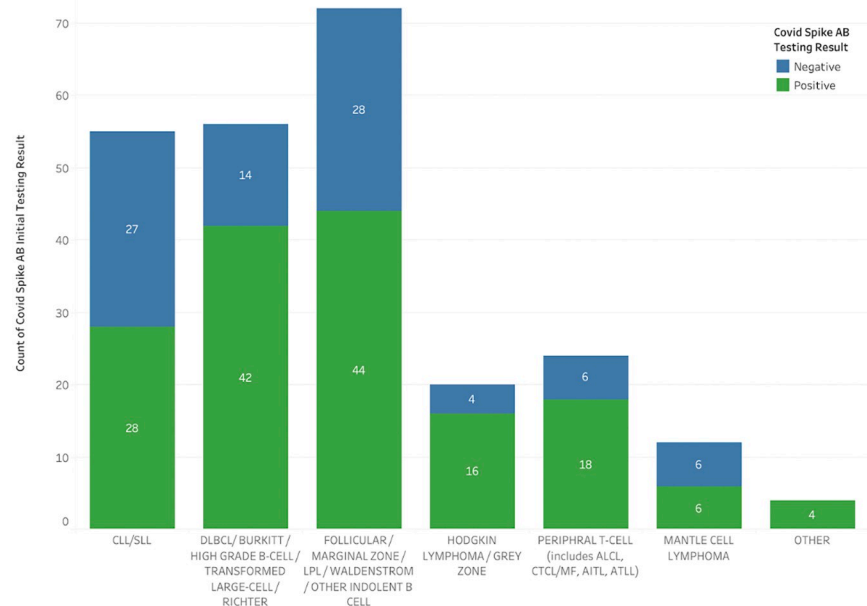


Fig 2. Serologic response against COVID spike protein after primary vaccination series by lymphoma diagnosis.

<https://doi.org/10.1371/journal.pone.0266584.g002>

There were 30 patients who received IVIG in the past 6 months prior to vaccination, and 43% (13/30) of these patients produced antibodies after the primary COVID vaccination series (Table 2). This seroconversion rate was significantly lower than in 68% (145/213) of patients who developed antibodies and had not received IVIG (UVA OR 0.36, 95% CI (0.16, 0.78), $p = 0.01$); MVA (OR 0.39, 95% CI (0.17, 0.93), $p = 0.03$). In patients who received an additional primary shot, 88% (7/8) were able to produce antibodies despite having received IVIG in the previous 6 months (Table 2).

There were 24 patients who had a documented COVID infection. Of these 24 patients, there were 13 patients who had a positive COVID PCR test result prior to receiving the primary COVID vaccine series; 3 patients had positive COVID PCR test results following completion of the primary vaccination series and serology testing; 1 patient had completed the primary vaccine series and had COVID infection prior to serology testing; and 6 patients that were indeterminable due an unknown date of COVID vaccination or COVID PCR test result. In these patients with documented COVID infection, 19 (19/24, 79%) patients were seropositive after the primary vaccination series. This seroconversion rate was not significantly different when compared to patients who did not have a history of COVID infection ($p = 0.09$; UVA OR 2.55 (0.84, 7.81), $p = 0.1$; MVA OR 3.08 (0.96, 9.87), $p = 0.06$) (Table 2). There were 2 patients who received convalescent plasma, 4 patients who received monoclonal antibody therapy, and 1 patient who received both convalescent plasma and monoclonal antibody treatment for COVID treatment.

Discussion

In this retrospective, single center study, lymphoma patients (65%) were capable of mounting a humoral response to the primary COVID vaccination series, and our findings were

COVID Spike AB Testing Results After Additional Primary Shot in Different Types of Lymphomas

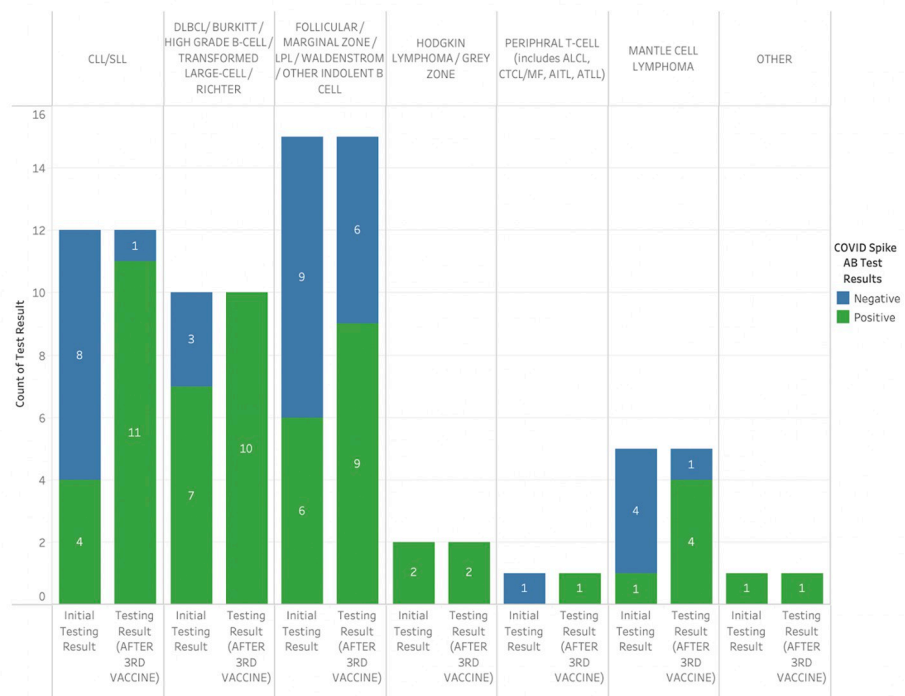


Fig 3. Serologic response against COVID spike protein after additional primary shot by lymphoma diagnosis.

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consistent with similar reports [7–15]. The majority of patients completed their primary vaccination series and additional primary shot with the COVID mRNA vaccines. The ability to produce antibodies to the COVID vaccine appeared to vary according to their lymphoma diagnosis and type of active anticancer treatment received in the past 6 months.

Failure to generate a competent humoral immune response against COVID is associated with an elevated risk of infection and poorer prognosis. Among all patients with hematological malignancies, those with B-cell malignancies generally have suboptimal antibody responses following vaccination against COVID, ranging from 44%-79%, depending on the subtype [7]. This is compared with seropositivity of 91%, 88%, and 95% in vaccinated patients with AML, ALL, and MM, respectively [7]. Among the leukemias, CLL patients have the lowest rate of seroconversion following vaccination, with 64% of patients failing to produce sufficient quantities of anti-spike protein antibodies [7]. The low rate of seroconversion in patients with CLL was also observed in this study. Approximately half (51%) of patients with CLL/SLL produced antibodies following their COVID vaccine, which was considerably lower when compared to most subtypes of lymphoma. In contrast, patients with HL/TCL have been shown to produce antibodies at higher rates 85%-98%, and this may be attributed to differences in disease biology [7]. The results of this study showed patients with HL/TCL had significantly higher rates of seroconversion following the primary vaccination series at 77% compared to those with CLL/SLL (UVA OR 3.27, 95% CI (1.39, 7.91), $p = 0.01$; MVA (OR 3.29, 95% CI (1.25, 8.64), $p = 0.02$).

Seroconversion rates following vaccination are further reduced among patients receiving supportive care agents, like IVIG, and certain anticancer agents, such as BTK inhibitors, anti-CD20 monoclonal antibodies, or the BCL2 inhibitor venetoclax [7]. IVIG is an Ig preparation

Table 3. Serologic response against COVID spike protein in patients who received anticancer treatment in the previous 6 months.

Seropositive against COVID spike protein	N = 92 [†]				
	Burkitt's, DLBCL, PMBL n(%)	CLL/SLL n(%)	HL/TCL [‡] n(%)	Indolent lymphomas [§] n(%)	MCL n(%)
<i>COVID vaccination history</i>					
Primary vaccination series	12/18 (67)	7/20 (35)	14/20 (70)	8/26 (32)	2/6 (33)
Additional primary shot	7/7 (100)	6/7 (86)	2/2 (100)	3/6 (50)	2/2 (100)
<i>Type of anticancer treatment in past 6 mo.</i>					
Anti-CD20 mAb monotherapy	4/4 (100)	-	-	4/11 (36)	2/2 (100)
BTK inhibitor	1/1 (100)	4/11 (55)	-	0/2	0/1
Rituximab + chemotherapy	4/8 (50)	-	-	2/5 (40)	-
Rituximab + BTK inhibitor	-	0/1	-	-	-
Anti-CD20 mAb + venetoclax	-	0/2	-	-	-
CAR T-cell therapy	-	-	1/4 (25)	-	0/1
Auto-HSCT	-	-	6/7 (86)	-	-
Brentuximab vedotin + other	-	0/1	-	-	-
PI3K inhibitor	-	2/3 (67)	-	-	0/2
Clinical trial	3/5 (60)	1/2 (50)	7/9 (78)	2/8 (25)	-
Other	-	-	-	-	-

[†]Note: 2 patients in "other" diagnostic category with T-cell prolymphocytic leukemia (T-PLL) and primary CNS lymphoma.

[‡]Includes peripheral T-cell lymphomas not otherwise specified (NOS), anaplastic large cell lymphoma (ALCL), cutaneous T-cell lymphomas, angioimmunoblastic T-cell lymphoma (AITL), and acute T-cell leukemia/lymphoma (ATLL).

[§]Includes follicular lymphomas, marginal zone lymphoma, lymphoplasmacytic lymphoma, and Waldenstrom macroglobulinemia.

DLBCL = diffuse large B-cell lymphoma; PMBL = primary mediastinal B-cell lymphoma; CLL = chronic lymphocytic leukemia; SLL = small lymphocytic lymphoma; HL = Hodgkin lymphomas; TCL = T-cell lymphomas; MCL = mantle cell lymphoma; mAb = monoclonal antibody; BTK = Bruton's tyrosine kinase; CAR = chimeric antigen receptor; Auto-HSCT = autologous hematopoietic stem cell transplant; PI3K = phosphoinositide 3-kinase.

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that consists of varying amounts of antibody and may lead to interference with the immune response to vaccines. Furthermore, recent IVIG supplementation may imply hypogammaglobulinemia, which has been independently linked with lower response rates to the COVID vaccine in patients with CLL [10,11]. Bruton's tyrosine kinase is a cytoplasmic tyrosine kinase responsible for B-cell maturation that is constitutively activated in CLL. The CD20 antigen is a pan-B cell marker elevated in certain B-cell lymphomas. BCL2 is an anti-apoptotic protein expressed in high quantities in cancer cells that inhibits the intrinsic pathway of apoptosis. Targeted therapy against these three markers have proven to be effective therapeutic strategies against various B-cell malignancies by suppressing their functionality, including the capacity to generate neutralizing antibodies. In our study, patients with CLL/SLL who received IVIG or BTK inhibitors with or without an anti-CD20 monoclonal antibody in the past 6 months had suboptimal antibody responses to the COVID vaccine, and this is consistent with previous findings [7,14,15].

For patients receiving B-cell directed therapies, peripheral blood B-cell counts may begin to recover after 3 months following treatment and may be completely reversed by 9 to 12 months [8]. A meta-analysis of thirty-eight studies in patients treated with anti-CD20 monoclonal antibody therapy demonstrated that patients actively on these therapies may have suboptimal antibody responses to all types of vaccines. This phenomenon was determined to be at least partially ameliorated by 3–6 months following treatment, with repopulation of the B-cell pool in 12 months after treatment [16,17]. Similarly, patients in this study who received rituximab in the previous 12 months had significantly lower seropositivity rates to the COVID vaccine compared to patients who had never received rituximab (UVA OR 0.51 (0.26, 1.00), $p = 0.05$; MVA 3.35 (1.47, 7.63), $p = 0.004$). However, this difference was no longer significant when

comparing patients who received rituximab great than the previous 12 months to those who had not received any rituximab (UVA OR 0.62 (0.27, 1.42), $p = 0.26$; MVA OR 2.38 (0.85, 6.70), $p = 0.10$). Of note, the majority of patients with Burkitt's, DLBCL or PMBL in this study were not actively receiving anticancer therapy in the months prior to COVID vaccination and 75% were able to mount a humoral response to vaccination. This response was significant on UVA (OR 2.89, 95% CI (1.30, 6.46), $p = 0.01$) and MVA (OR 2.71, 95% CI (1.04, 7.06), $p = 0.04$), which suggests the ability of these patients to recover B-cell counts following B-cell directed therapies and the need for optimal timing of vaccination following these therapies. Furthermore, this adds to the complexity of predicting response to vaccines in patients with lymphoma. Not only is it affected by the biology of the disease itself, but also the type of therapy they are receiving as well as its timing.

There are some limitations to this study. Firstly, this was a single-center, retrospective chart review. Some patients received their COVID vaccine at an outside facility, therefore, the exact date and the type of vaccine received was not consistently available. Secondly, the timing of the additional primary shot of COVID vaccine was not uniform among all patients, and not all patients received this additional dose. As the COVID-19 pandemic progressed and the Center for Disease Control updated recommendations based on available evidence, some patients received the additional primary shot of the COVID vaccine closer to the recommended 28 days after completion of the primary vaccination series than others, and some have yet to receive their additional primary shot. Third, lymphoma patients included in this study were on a wide spectrum in terms of their cancer-related care, with many patients not receiving active anticancer treatment in the previous 6 months. Consequently, not all patients were frequently visiting the outpatient cancer center for follow-up visits due to lack of active treatment or timing of remission scans, and there were fewer opportunities to test patients for anti-COVID spike protein antibody levels. Lastly, although there were 243 patients evaluated in the overall analysis, some subgroups of particular interest had smaller numbers of patients. Only 3 patients included in this study received the COVID adenovirus vaccine; despite this small number, these patients remained in the overall analysis to reflect the outcomes of a real-world population of lymphoma patients. These patients who received the COVID adenovirus vaccine had CLL or indolent lymphomas, which are lymphoma types that have shown suboptimal seroconversion rates following vaccination. Of note, these patients were not actively receiving anticancer treatment at the time of vaccination. We observed that these 3 patients were able to mount an immune response to the COVID adenovirus vaccine. This finding shows that some patients with CLL or indolent lymphomas may still derive benefit from vaccination, and it remains vitally important to vaccinate these patients.

In conclusion, lymphoma patients are capable of mounting a humoral response to the COVID vaccines. Although seroconversion is lower than in the normal population, it is still significant and warrants administration of COVID vaccine without delay, particularly during continuing high rate of infection. CLL/SLL appears predictive of a negative antibody response to the COVID vaccine, while HL/TCL histologies appeared to correlate to a positive antibody response, even with treatment within 6 months of vaccination. Our study suggests anti-CD20 monoclonal antibody therapy in the last 12 months may affect the ability to produce antibodies towards a COVID vaccine. Further studies are required to confirm our findings, including whether T-cell immunity would be of clinical relevance in this patient population.

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