SARS-CoV-2 Antibody Response is Associated with Age and Body Mass Index in Convalescent Outpatients


Department of Pediatrics, UMPC Children’s Hospital of Pittsburgh, Pittsburgh, PA
Department of Family Medicine, University of Pittsburgh, Pittsburgh, PA
Department of Microbiology and Immunology, Tulane University School of Medicine, New Orleans, LA
Center for Vaccine Research, University of Pittsburgh, Pittsburgh, PA

Abstract

COVID-19 has had an unprecedented global impact on human health. Understanding the antibody memory responses to infection is one tool needed to effectively control the pandemic. Among 173 outpatients who had virologically confirmed SARS-CoV-2 infection, we evaluated serum antibody concentrations, microneutralization activity, and enumerated SARS-CoV-2 specific B cells in convalescent human blood specimens. Serum antibody concentrations were variable, allowing for stratification of the cohort into high and low responders. Neither participant sex, the timing of blood sampling following the onset of illness, nor the number of SARS-CoV-2 spike protein specific B cells correlated with serum antibody concentration. Serum antibody concentration was positively associated with microneutralization activity and participant age, with participants under the age of 30 showing the lowest antibody level. These data suggest that young adult outpatients did not generate as robust antibody memory, compared with older adults. Body mass index was also positively correlated with serum antibody levels. Multivariate analyses showed that participant age and body mass index were independently associated with antibody levels. These findings have direct implications for public health policy and current vaccine efforts. Knowledge gained regarding antibody memory following infection will inform the need for vaccination in those previously infected and allow for a better approximation of population-wide protective immunity.
Introduction

The ongoing COVID-19 pandemic caused by SARS-CoV-2 has resulted in over 250 million cases and 5 million deaths worldwide as of December 2021. With a case fatality rate near 2%, most of those infected survive and go on to generate immune memory against the virus. Numerous studies have shown that greater than 95% of those infected generate antibodies that recognize SARS-CoV-2 proteins in the months immediately following infection, as reviewed (1). SARS-CoV-2 infection results in neutralizing antibody production in most of those infected, although the half-life of neutralizing antibody may be a relatively short 2 months (1). Neutralizing antibody production is thought to be protective against re-infection in about 90% of individuals (1). Many studies to date have focused on those hospitalized with severe COVID-19 although several groups have published investigations into the antibody response to SARS-CoV-2 infection in those with less severe disease (2–8). Comparison of the antibody response in patients with severe disease with those with mild/moderate disease or the response induced by vaccination have shown that severe illness results in robust and greater antibody levels than the other settings (9, 10).

Although SARS-CoV-2 infection has been shown to produce a significant antibody response that initially protects against re-infection, it has been shown to wane over the course of months. This waning immune memory suggests a need for immune boosting by vaccination (9). Further, the impact of age, sex, and body mass index (BMI) on antibody response to SARS-CoV-2 is inconsistent in the literature and may play a key role in defining which patients are most in need of boosting for sustaining adequate immune memory as reviewed in (11). The goal of the present study was to better understand the antibody response to SARS-CoV-2 infection in adults with mild to moderate illness who sought outpatient care using well characterized and controlled assays for SARS-CoV-2 spike (S1, receptor binding domain (RBD), S2 domains) and nucleoprotein (N) specific IgG antibody. SARS-CoV-2 antibody concentrations were correlated with in vitro virus neutralization activity to demonstrate the functional relevance of high and low antibody concentration responders. In addition, the presence of IgG producing memory B cells was examined by flow cytometry in participants with high serum antibody responses to SARS-CoV-2.

Methods

Cohort

All participant samples were collected as part of the US Influenza Vaccine Effectiveness (FLUVE) Network – Pittsburgh site. This network is an annual prospective study of outpatients who seek care for acute respiratory illness. This study was approved by the University of Pittsburgh Institutional Review Board. All 173 participants were adults (≥18 years of age) and were SARS-CoV-2 positive by molecular testing. Relevant participant information was subsequently extracted from the medical record. One participant did
not disclose sex, one participant did not have the timing of blood sampling relative to illness recorded, and two participants did not have BMI recorded. These participants were excluded for the relevant analyses. Participant age groupings were selected based upon the observation of raw antibody data which revealed lower levels in participants between ages 18–29; additional groups were then established using 15 year intervals.

**Blood Processing**

Blood was collected in Becton Dickinson SST (serum) or CPT (peripheral blood mononuclear cells (PBMC)) and processed following manufacturer’s instructions. Serum was aliquoted and stored at −80° C prior to analyses. PBMC were aliquoted, frozen, and stored in liquid nitrogen prior to analyses.

**Antibody Assay**

SARS-CoV-2 specific antibody concentrations were determined using a commercially available Biorad Bioplex Kit; Bio-Plex Pro Human IgG SARS-CoV-2 N/RBD/S1/S2 4-Plex Panel (catalog #12014634). Analyses of serum samples were performed by manufacturer’s instructions. Banked pre-COVID-19 serum samples were utilized as negative controls and were all below the limit of detection (n=5). SARS-CoV-2 antibody standards were provided by Biorad, with concentrations of U/ml. To control for plate-to-plate variability, five samples were repeated on all three assay plates run. Concordance was very high between assay runs ($r^2$ of 0.85–0.96), suggesting that combining data was appropriate (Supplemental Figure 1A–C). In addition, World Health Organization SARS-CoV-2 antibody reference samples were assayed to compare with values obtained using the Bioplex kit. The COVID-19 convalescent plasma panel (NIBSC 20/118) was obtained from the National Institute for Biological Standards and Control, United Kingdom. Concentrations derived using the Bioplex kit correlated with the reported WHO reference sample concentrations (Supplemental Figure 1D–E).

**Neutralization Assay**

Antibody neutralization of SARS-CoV-2 was performed as published (12). Spots in Vero E6 cell cultures were imaged, counted, and processed using an ImmunoSpot Counter (CTL). Foci were counted in experimental wells and compared to control wells. The dilution of serum at which 80% of foci are neutralized is reported as the FRNT$_{80}$.

**B Cell Flow Cytometry**

Frozen PBMCs were rapidly thawed in a water bath at 37° C, rinsed with 1ml flow sorter buffer (1x phosphate buffered saline, 2% newborn calf serum, and 0.1% sodium azide) and centrifuged at 500 x g for 5 minutes, 4° C. Cells were resuspended in 100 μl of sorter buffer with 2.5 μg of human Fc Block (clone Fc.3216) for 10 minutes at room temperature. Next, cells were incubated with 1 mM PE-Cy5 decoy for 5 minutes at room temperature followed by 1 mM SARS-CoV-2 Spike-PE B cell tetramer for 25 minutes on ice in the dark using decoy and tetramer created in the lab as previously described (13, 14). Cells were then washed and incubated with anti-PE magnetic beads (Miltenyi) and passed over a LS column on a quadroMACS magnet. Eluted cells were stained for 30 minutes at 4° C in the dark.
with the following anti-human antibodies (clone): CD19 (SJ25C1), CD3 (UCHT1), CD14 (HCD14), CD16 (B73.1), CD21 (Bu32), CD27 (M-T271), IgD (IA6–2), IgM (MHM-88) and IgG (G18–145). Next, cells were washed with 1 ml flow sorter buffer, centrifuged at 500 × g for 5 minutes, and fixed with 200 μl 2% paraformaldehyde at 4° C for 10 minutes. Cells were washed and resuspended in flow sorter buffer and collected on an LSRFortessa (Beckton-Dickson). Data was analyzed using FlowJo software (TreeStar, Ashland, OR). As a negative control, 4 banked pre-pandemic PBMC samples were analyzed.

Statistics

Statistical analyses were performed using GraphPad Prism Software (San Diego, CA). Correlation data was analyzed by simple linear regression with an alpha of 0.05 for significance. Differences between two means were analyzed by unpaired t-test with an alpha of 0.05 for significance. Differences between more than two means were analyzed by one-way ANOVA followed by Tukey post-hoc test with an alpha of 0.05 for significance. Cohort data sets are shown as violin plots with the median and quartiles indicated. Alternatively, data sets are displayed as dot plots with mean and standard error of the mean displayed. Multivariable linear regression models were run separately for each SARS-CoV-2 spike and nucleoprotein specific IgG antibodies and assessed the association of age on SARS-CoV-2 antibody response adjusting for BMI, and vice versa. Statistical significance of two-sided tests was set at type I error (alpha) equal to 0.05. These analytical procedures were performed using SAS® 9.4 (Cary, NC).

Results

To examine the antibody response to mild SARS-CoV-2 infection, we recruited 173 convalescent outpatients for blood sampling through the FLUVE network. The patient cohort included 105 females with a mean age of 38.3 (range = 19–79) years and 67 males with a mean age of 42.7 (range = 20–78) years. One participant did not disclose their sex; this patient was excluded from birth sex analyses. The convalescent blood sample was drawn at an average of 44.5 (range = 22–131) days post symptom onset for females and 42.3 (range = 21–89) days for males. BMI was a mean 30.2 for females (range = 17.9–59.8) and a mean of 30.9 for males (range = 19.3–72.0). There were no significant differences in age, timing of sampling, or BMI by participant sex.

SARS-CoV-2 antibodies were detectable in most participant samples. Concentrations ranged from 0 to 2×10⁶ U/ml. Low responders were defined as antibody concentration less than 5 × 10⁴ U/ml. Approximately one-fifth of participants had low antibody levels against N, RBD, and S1 proteins (Table I). Incidence of low response to S2 was lower than the other antigens tested at 6.9% of participants. Approximately 6% of participants had very low antibody responses (less than 10⁴ U/ml) to N, RBD, and S1, while only a few had very low antibody levels against S2. Of the 18 with the lowest antibody concentrations, 13 were females. Of these 18, low responses were noted against 1 of 4 antigens in 6 persons, 2 antigens in 7 persons, 3 antigens in 3 persons and all 4 antigens in 2 persons.

We examined correlations between antibody levels for pairs of SARS-CoV-2 proteins. The highest correlation (R² = 0.74) in antibody levels was for S1 and RBD, which would be
predicted since RBD is part of S1 (Figure 1). Significant correlations between antigen antibody levels were found for all 4 antigens tested. Interestingly, N protein antibody levels correlated at the lowest levels in all comparisons. Despite statistically significant correlations, the relationship between individual antigen antibody levels were not overtly linear, as was the case for S1 and RBD.

To determine if measurable differences in SARS-CoV-2 antibody concentration was relevant to functional antibody, microneutralization assays were performed, on a subset of participants with the most divergent antibody levels. This subset of 56 participants was selected to assess both high and low responders: based upon high antibody response (greater than $7 \times 10^5$ U/ml for at least 1 antigen) and low antibody response (less than $1 \times 10^5$ U/ml for at least 1 antigen). Evidence of neutralizing antibody activity (FRNT$_{80}$) was uncommon in the low responder group with only 9 of 28 participants achieving neutralizing titer $\geq 1:20$ (Figure 2). No high titer neutralizing activity was seen in the low responder group (greatest observed 1:80). Conversely, all but 1 high responder displayed neutralizing activity, with 18 of 28 showing a neutralizing titer greater than or equal to 1:160. Further, microneutralization titer significantly correlated with the concentration of antibody determined by Bioplex (Supplemental Figure 2). Correlation was strongest between microneutralization titer and spike protein antigens (e.g., $r^2$ of 0.36 for RBD).

Flow cytometry was then utilized to determine if SARS-CoV-2 specific B, memory B, and IgG producing B cells were detectable in PBMC from high responders. We employed a SARS-CoV-2 spike tetramer and a gating strategy to differentiate total spike-specific B cells, memory B cells, and class switched IgG producing B cells (Supplemental Figure 3). We successfully detected spike specific B cells in all three gates from all high responders. The number of spike-specific B cells did not correlate with microneutralization titer, nor did memory B cell or IgG producing B cell numbers (Figure 3). No differences were seen in the number of spike-specific B cells, spike-specific memory or IgG producing B cells by patient sex (Figure 4). Participant age was then compared with the number of spike-specific B cells; again no significant correlations were observed (Figure 4). There was no correlation between B cell numbers and the timing of sampling post-illness (spike-specific B cells $R^2 = 0.0406$, $p = 0.3814$, spike-specific memory B cells $R^2 = 0.0054$, $p = 0.7518$, IgG producing B cells $R^2 = 0.0192$, $p = 0.5497$).

We examined the impact of sample timing and/or biologic variables on SARS-CoV-2 antibody responses in the entire cohort. The mean sampling time was approximately 6 weeks post onset of illness, which was utilized to divide the group into those sampled before and after 42 days post illness onset. There was no significant difference in antibody levels by time of sampling for any of the 4 antigens tested based (Figure 5). Further, there was no significant correlation between sample timing and antibody concentrations against N, RBD, and S1 antigens (Supplemental Figure 4A–D). S2 antibody levels were positively correlated with longer timing of sampling post-illness. There was no difference in antibody concentration against the SARS-CoV-2 antigens tested when analyzed by patient sex (Figure 6).
We broke the cohort into 4 groups of 18–29, 30–45, 46–59 and over 60 years. When examined by age groups (18–29, 30–45, 46–59 and over 60 years), antibodies against SARS-CoV-2 N, RBD, and S1 were significantly lower in participants under the age of 30 years compared with participants over age 45 years (Figure 7). Further, patient age significantly positively correlated with SARS-CoV-2 antibody concentration (Supplemental Figure 4E–H). However, there was no correlation between older age and serum neutralizing activity. While all participants were recruited as outpatients, we attempted to characterize the severity of disease by measuring the number of days until recovery and if the patient returned to normal activity at the time of convalescent sampling. The mean days to recovery was significantly longer in participants over 45 compared with those in the 18–29 and 30–45 groups (Table II). Similarly, a greater proportion of participants had not yet returned to normal activity in the over 45 group compared with the two young participant groups.

Finally, we compared participant BMI with serum antibody levels. BMI significantly positively correlated with antibody concentrations for all four antigens tested (Figure 8). We next examined the relationship between patient age and BMI since both parameters correlated with antibody levels. BMI was not significantly associated with patient age ($R^2 = 0.0115$, $p = 0.1624$). To test for interaction between these variables, multivariable regression was performed to determine the association of age upon SARS-CoV-2 antibody response adjusting for BMI. In adjusted models, age was significantly related to SARS-CoV-2 antibody response (Table III). BMI adjusted for age was also significantly associated with SARS-CoV-2 antibody. There was no evidence of an interaction between age and BMI for any SARS-CoV-2 antibody outcome.

Discussion

This study focused on a cohort of COVID-19 outpatients with mild or moderate illness. Many published serology studies have focused on more severe disease, with mild illness limited to a comparator for the robust antibody response in hospitalized participants. More than ten studies have shown that antibody response in severe COVID-19 is greater than in mild cases, as reviewed (11). These findings have been extended by similar observations regarding the presence of neutralizing antibodies in serum (11). While study of severe disease is important, most cases of COVID-19 do not require hospitalization and thus, most individuals infected with SARS-CoV-2 would be expected to have antibody responses similar to those observed in our study. Our finding that 98.8% (171/173) of our outpatient cohort developed measurable antibody against at least one SARS-CoV-2 antigen is consistent with findings in many other studies that suggest that natural infection of variable severity induces humoral immunity against the virus (1).

Given our focus on COVID-19 outpatients, we were able to stratify these patients into groups of low and high responders to perform functional antibody assays and to examine the presence of SARS-CoV-2 specific memory B cells. Microneutralization assays revealed that relatively high and low antibody concentration detected in the Bioplex assay correlated with the amount of functional antibody present. In the low responder group, the highest FRNT80 titer observed was 1:80, while nearly all high responders tested were equal to or greater than this level. Surprisingly, two participants had high antibody concentration, but no or low
microneutralizing activity, suggesting that antibody level alone may not be protective for some patients. Correlation of microneutralization titer with antibody concentration revealed a positive relationship between high spike protein antibody and functional ability, as one would predict given that spike is the target of neutralizing antibodies. The correlation between neutralizing activity and N protein antibody was less strong. Others have shown that SARS-CoV-2 specific memory IgG producing B cells are formed in both mild and severe patients (13, 15). In severe COVID-19 patients, memory B cells were found in all patients studied, however only 80% had neutralizing antibody. In our study, all antibody high responders had detectable spike-specific memory B cells, although the number of memory B cells did not correlate with antibody concentration. All participants tested for memory B cells had serum neutralizing activity of variable titer. These data suggest that high responding outpatients form memory B cells which may provide longer lasting protection against SARS-CoV-2 than serum antibody concentration would indicate.

By measuring IgG against 4 four SARS-CoV-2 antigens, we were able to characterize the relative immunodominance of the different antigens. Antibody response to N protein only partially correlates with spike protein antibody responses. Our data showed that some participants responded to N better than to spike proteins and vice versa, with the relationship between N and S1 antibodies being the strongest positive correlation. S1 and S2 antibody response was positively correlated, but not entirely linear suggesting differential immune recognition of these spike protein domains among the cohort. Interestingly, the incidence of a low antibody response was lowest for the S2 domain of the spike protein when compared with S1, RBD, or N proteins. In the rare occurrence of very low antibody response to 1 or more antigens tested, females were more likely than males to display this poor response phenotype.

Our study did not show an impact of sample timing or biologic sex on serum antibody concentrations. Numerous studies have shown a degree of antibody waning 2 to 3 months after infection. Our cohort spanned 1 to 4 months after the onset of symptoms with a median near 42 days. In outpatients, we did not observe a decrease in serum antibody in this time period, nor did we see a negative correlation between sample timing post-illness and antibody concentration. Several studies have shown no effect of biologic sex on serum antibody titer, however neutralizing antibody levels have been shown to correlate with male sex, as reviewed (11, 16, 17). This is consistent with our findings of antibody concentration; however, we did not see a significant difference in neutralizing antibody titer and sex (data not shown). The effect of sex on SARS-CoV-2 antibody response does not appear to be a primary driver of response in outpatients in this study.

Importantly, a predictor of antibody response in our study was participant age. Participants aged 18–29 had significantly lower antibody concentration for N, S1, and RBD proteins compared with older participants. We also found a significant positive correlation between patient age and serum antibody levels. This is consistent with some studies that have shown a correlation between IgG levels and age, with lower antibody concentration in younger adults (16, 18–20). In those studies, conflicting results were found regarding neutralizing antibody titers as one study found an increase in older adults, while the other showed higher levels in younger adults. Other research did not show a relationship between age and

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antibody concentration, as reviewed (11, 17). One possible explanation for the age effect observed would be the severity of disease in our cohort. To address this issue, we examined the number of days until recovery and if a return to normal was seen at the time of sampling. We did show that recovery and a lack of return to normal activities was significantly longer in participants over 45 years, compared with younger participants. However, these parameters were not different in participants 18–29 or those from 30–45, while antibody concentration was lower between these two groups. These data suggest that measures of outpatient severity do not likely explain the lower antibody response in young participants in our study. It is possible that younger participants had better controlled viral replication, and thus, there was less viral antigen to stimulate ongoing immune responses.

Finally, our study showed a significant positive correlation between BMI and serum antibody levels. The impact of obesity on COVID-19 remains unclear as some have found a positive correlation between BMI and severity, while others have not (21, 22). Similarly, the relationship between BMI or metabolic syndrome and antibody levels has varied. Poor metabolic health (defined by low adiponectin:leptin ratio) was shown to correlate with increased cross-reactive SARS-CoV-2 strain antibodies (22). Further, BMI and metabolic syndrome were shown to correlate with higher antibody levels (23, 24). However, others have shown either no or the opposite relationship (25, 26). In our study, multivariate analyses of age and BMI revealed that they are independent predictors of antibody response to SARS-CoV-2 in outpatients. Again, it is unclear why BMI may be associated with antibody levels, but differences in viral load, co-morbidities, or subtle variability in disease severity may be involved.

In summation, we characterized serum antibody and B cell responses to SARS-CoV-2 in a cohort of COVID-19 outpatients. We did so using well validated assays to generate reliable antibody concentration, microneutralization, and flow cytometry data. The primary findings were that participant age and BMI independently affected serum antibody concentration, with younger and lower BMI participants producing lower antibody responses. We also showed that antibody concentration was predictive of microneutralization activity, but not the number of spike-specific B cells in blood. In whole, these data inform current vaccination policy supporting that young and low BMI previously infected people likely require vaccine boosting for long term protection. Further, antibody concentration may not directly relate to the presence of SARS-CoV-2 specific memory B cells capable of providing protection against infection and/or severity of disease. As serum antibody levels wane following infection, memory B cells may persist longer to provide sustained protection. These questions will need to be answered at later time points following infection. In addition, our study was conducted prior to SARS-CoV-2 vaccine availability or the SARS-CoV-2 delta and omicron variant waves, which may differentially impact antibody responses.

**Supplementary Material**

Refer to Web version on PubMed Central for supplementary material.

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References


Key Points

1. Participant age positively associated with SARS-CoV-2 antibody levels in outpatients.

2. Participant BMI positively correlated with SARS-CoV-2 antibody levels in outpatients.

3. Antibody levels were associated with microneutralization activity against SARS-CoV-2.
Correlation between SARS-CoV-2 antigen specific IgG concentrations. Samples in the cohort were analyzed for 4 SARS-CoV-2 antigens (N=173). Concentration values (U/ml) were compared using simple linear regression, $R^2$ values are denoted, $P \leq 0.0001$ for all comparisons. Red depicts the highest correlation (1.0), green the lowest (0) with color gradients in between.

![Table showing correlation values](image-url)
Figure 2 --
In vitro SARS-CoV-2 microneutralization titers in high and low responding participants. A subset of participants were stratified by relative antibody response into two groups of lowest (A) and highest (B) antibody concentrations (N=28 each). The dilution of serum at which 80% of viral foci are neutralized is reported as the FRNT$_{80}$ (1:X titer). The limit of detection for the assay is 1:20.
Figure 3 -. Correlation of spike-specific B cells and serum microneutralization activity. Flow cytometry was performed on high responder participant PBMC using a spike-specific tetramer and B cell markers (N=21). Spike B cells (A) were defined as CD19\(^+\) tetramer\(^+\) cells. Spike memory B cells (B) were defined as CD19\(^+\) tetramer\(^+\) CD27\(^+\) cells. Spike-specific IgG B cells (C) were defined as CD19\(^+\) tetramer\(^+\) IgD\(^-\) IgM\(^-\) IgG\(^+\) cells. Data were compared using simple linear regression.
Figure 4 – Effect of biologic sex or participant age on SARS-CoV-2 spike specific B memory and IgG producing B cells. Flow cytometry was performed on high responder participant PBMC using a spike-specific tetramer and B cell markers (N=21). Spike B cells were defined as CD19+ tetramer+ cells (A, D). Spike memory B cells were defined as CD19+ tetramer+ CD27+ cells (B, E). Spike-specific IgG B cells were defined as CD19+ tetramer+ IgD− IgM− IgG+ cells (C, F). Data were compared using unpaired t-test or simple linear regression.
Figure 5 –.
The impact of sample timing on SARS-Cov-2 antibody concentration. Participants were stratified by when blood samples were collected after the onset of COVID-19 symptoms with a split at 42 days (A-D) (N=97 less than or equal to 42, 75 greater than 42). Antibody concentrations were determined by Bioplex assay and compared using unpaired t-test.
Figure 6.
The impact of biologic sex on SARS-Cov-2 antibody concentration. Participants were stratified by sex (A-D) (N=105 female, 67 male). Antibody concentrations were determined by Bioplex assay and compared using unpaired t-test.
Figure 7 –.
The impact of participant age on SARS-Cov-2 antibody concentration. Participants were stratified by age (A-D) (N=58 18–29, 55 age 30–45, 36 age 46–59, 24 age 60 and over). Antibody concentrations were determined by Bioplex assay and compared using one-way ANOVA followed by Tukey post-hoc test. Serum neutralization activity and age were compared using simple linear regression (E).
Figure 8 — The association between BMI and SARS-CoV-2 antibody concentration. Simple linear regression was performed on participant BMI and antibody levels (A-D) (N=171). Antibody concentrations were determined by Bioplex assay.
Table I.
Outpatients with lower antibody concentration against SARS-CoV-2

<table>
<thead>
<tr>
<th>Antibody</th>
<th>Female (N=105)</th>
<th>Male (N=67)</th>
<th>Total (N=173)</th>
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<tr>
<td></td>
<td>N</td>
<td></td>
<td></td>
</tr>
<tr>
<td>&lt;10^3</td>
<td>7 (6.7%)</td>
<td>4 (6.0%)</td>
<td>11 (6.4%)</td>
</tr>
<tr>
<td>&lt;5x10^3</td>
<td>25 (23.8%)</td>
<td>13 (19.4%)</td>
<td>39 (22.5%)</td>
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<tr>
<td>RBD</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>&lt;10^3</td>
<td>9 (8.6%)</td>
<td>2 (3.0%)</td>
<td>11 (6.4%)</td>
</tr>
<tr>
<td>&lt;5x10^3</td>
<td>26 (24.8%)</td>
<td>12 (17.9%)</td>
<td>39 (22.5%)</td>
</tr>
<tr>
<td>S1</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>&lt;10^3</td>
<td>9 (8.6%)</td>
<td>3 (4.5%)</td>
<td>12 (6.9%)</td>
</tr>
<tr>
<td>&lt;5x10^3</td>
<td>26 (24.8%)</td>
<td>12 (17.9%)</td>
<td>39 (22.5%)</td>
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<tr>
<td>S2</td>
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<td></td>
<td></td>
</tr>
<tr>
<td>&lt;10^3</td>
<td>2 (1.9%)</td>
<td>1 (1.5%)</td>
<td>3 (1.7%)</td>
</tr>
<tr>
<td>&lt;5x10^3</td>
<td>5 (4.8%)</td>
<td>6 (9.0%)</td>
<td>12 (6.9%)</td>
</tr>
</tbody>
</table>

1The number of patients with antibody concentrations < 10^3 U/ml or < 5x10^3 U/ml.
Table II.

Recovery from SARS-CoV-2 symptoms by patient age.

<table>
<thead>
<tr>
<th></th>
<th>18–29</th>
<th>30–45</th>
<th>46–59</th>
<th>60+</th>
</tr>
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<tbody>
<tr>
<td>Full Recovery (Days)</td>
<td>11.1 ± 0.6 (n=51)</td>
<td>12.7 ± 0.7 (n=43)</td>
<td>15.4 ±1.9 (n=19) *</td>
<td>16.2 ± 1.8 (n=17) **</td>
</tr>
<tr>
<td>Return to Normal (No)</td>
<td>9/57</td>
<td>15/52</td>
<td>16/32 **</td>
<td>12/24 *</td>
</tr>
</tbody>
</table>

* The number of days to full recovery and if the patient had returned to normal activity by the time of sampling.

** p < 0.01 when compared to 18–29 group.
Table III.

Association of age and BMI on SARS-COV-2 antibody

<table>
<thead>
<tr>
<th>Variable</th>
<th>Beta</th>
<th>Standard Error</th>
<th>p value</th>
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</thead>
<tbody>
<tr>
<td>N</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Age</td>
<td>9038.1094</td>
<td>2691.2749</td>
<td>0.0010*</td>
</tr>
<tr>
<td>BMI</td>
<td>24357.6913</td>
<td>4870.0729</td>
<td>&lt;0.0001*</td>
</tr>
<tr>
<td>RBD</td>
<td></td>
<td></td>
<td></td>
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<tr>
<td>Age</td>
<td>8318.1547</td>
<td>2595.7786</td>
<td>0.0016*</td>
</tr>
<tr>
<td>BMI</td>
<td>11804.7176</td>
<td>4697.2647</td>
<td>0.0129*</td>
</tr>
<tr>
<td>S1</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Age</td>
<td>7755.0474</td>
<td>2480.7334</td>
<td>0.0021*</td>
</tr>
<tr>
<td>BMI</td>
<td>13769.9620</td>
<td>4489.0816</td>
<td>0.0025*</td>
</tr>
<tr>
<td>S2</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Age</td>
<td>6922.26184</td>
<td>3016.9755</td>
<td>0.0230*</td>
</tr>
<tr>
<td>BMI</td>
<td>10904.20579</td>
<td>5459.4536</td>
<td>0.0474*</td>
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</tbody>
</table>


* p < 0.05