

# Microbiology Spectrum

# Hundred-fold increase in SARS-CoV-2 spike antibody levels over three years in a hospital clinical laboratory.

Patrizio Caturegli, Oliver Laeyendecker, Aaron Tobian, and David Sullivan

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# **Transaction Report:**

(Note: With the exception of the correction of typographical or spelling errors that could be a source of ambiguity, letters and reports are not edited. The original formatting of letters and referee reports may not be reflected in this compilation.)

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August 15, 2023

Dr. David J Sullivan Johns Hopkins University Bloomberg School of Public Health Molecular Microbiology & Immunology 615 N. Wolfe St., Rm. W 4606 Baltimore, MD 21205

Re: Spectrum02183-23 (Hundred-fold increase in SARS-CoV-2 spike antibody levels over three years in a hospital clinical laboratory.)

Dear Dr. David J Sullivan:

Minor modifications are required before publication. Please correct the manuscript according to the suggestions of reviewers 1 and 2.

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Bar-On Yotam

Editor, Microbiology Spectrum

Journals Department American Society for Microbiology 1752 N St., NW Washington, DC 20036 E-mail: spectrum@asmusa.org

Reviewer comments:

Reviewer #1 (Comments for the Author):

This is an interesting and significant descriptive study of the changes in the level of antibodies against spikes throughout the evolution of the COVID-19 pandemic. The authors, through a simple analytical approach, attempt to support the use of CCP donors as an anti-SARS-CoV-2 treatment, in particular for immunocompromised patients.

I appreciate the opportunity to review this interesting manuscript by Caturegli et al.

The authors analyzed spike antibodies measured by a tertiary hospital clinical immunology laboratory over a 3-year period, (April 2020 - February 2023), using the dilutional "Anti-SARS-CoV-2 ELISA" assay (Euroimmun US, Mountain Lakes, NJ). The study showed that spike IgG levels rose markedly over time, from a median of 0.13 ELISA arbitrary units (AU) in period 1 to 48.7 in period 5 (p<0.0001). The spike IgG 80th percentile distribution threshold was 0.55, 8.1, 9.6, 64.9, and 151 AU in each of these periods.

This manuscript is an interesting piece of work, but it is not in a final state to be published yet. Therefore, the manuscript requires corrections before considering its acceptance.

Comments:

The abstract is well-written and informative. It clearly states the purpose of the study, the methods used, the main findings, and the implications of the findings.

Methods: the study is well-designed. The authors used a large sample size and a validated assay.

The findings of the study are important and have implications for the use of COVID-19 convalescent plasma therapy. The authors mention that the FDA uses a threshold of 3.5 AU to qualify CCP donors. However, they do not discuss why this threshold was chosen and why the 55 AU was also chosen in the newer assay.

The authors suggested that restricting CCP donors to those with high titre spike antibodies may be more effective in protecting immunocompromised patients from variants. However, they do not provide any data to support this suggestion. It would be helpful to see data on the clinical effectiveness of CCP therapy in patients with different levels of spike antibodies. Other comments are highlighted in the attached Pdf.

Staff Comments:

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Thank you for submitting your paper to Microbiology Spectrum.

Hundred-fold increase in SARS-CoV-2 spike antibody levels over three years in a hospital clinical laboratory.

In this study, the authors evaluated the increase in neutralizing antibodies against SARS-CoV-2 over a three-year period in the setting of a clinical study. This time period was divided into five intervals defined by the U.S. population prevalence of major SARS-CoV-2 mutations. The authors found an increase in antibody levels against the RBD region of the spike. The increase was even more evident when the 80<sup>th</sup> percentile of the five intervals were compared. Based on these results, the authors suggest redefining threshold levels of antibodies that could be used to identify COVID-19 convalescent plasma (CCP) donors.

This is an interesting and significant descriptive study of the changes in the level of antibodies against spikes throughout the evolution of the COVID-19 pandemic. The authors, through a simple analytical approach, attempt to support the use of CCP donors as an anti-SARS-CoV-2 treatment, in particular for immunocompromised patients. A major limitation of the study is the lack of a neutralization test. These tests could have been performed on a subset of data from each of the time intervals. A correlation between antibody levels and neutralization was previously demonstrated [1]. The present study could have determined the correlation between the neutralization tests and the categorization that is proposed based on the level of antibodies. Or the authors could have used the results of the two different tests to infer the CCP donor selection thresholds. Although the authors' proposal to use the upper 20% of the distribution of antibody levels is consistent with what was previously suggested by other authors [2], this categorization is not supported from a quantitative point of view. A second point that could have enriched this work would have been the identification of variables associated with the number of antibodies. Indeed, given the large volume of data, it would have been interesting to identify clinical, epidemiological, or biological variables of the virus that are associated with the number of antibodies. This exploratory analysis could identify criteria to take into account in the definition of thresholds for the selection of CCP donors.

## Suggestions

- The authors could add a table (plot) describing the population analyzed. This could be organized by time intervals; in each of them, it could be indicated, for example, the number of women and men, the number of people vaccinated, the number of people with previous SARS-CoV-2 infections, and the number of seropositive patients. This table could be cited in the text. - In addition to the median, it can be informative to add to the text the minimum and maximum values of the number of antibodies.

- The authors could evaluate if there are significant statistical differences between the median number of antibodies of the different time intervals.

- The authors could evaluate if there are significant statistical differences between the median of seropositives of the different time intervals.

- The authors could evaluate if there are significant statistical differences in the number of antibodies between sexes for the same time interval or between different intervals for the same sex.

The results of these tests could support some of the claims made in the text.

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- <u>https://doi.org/10.1128/mbio.03523-22</u>
   https://doi.org/10.1371/journal.pone.0273223

- Hundred-fold increase in SARS-CoV-2 spike antibody levels over three years in a hospital clinical
  laboratory.
- 4 Running title- Hundred-fold increase in spike antibody levels
- 5
- 6 <sup>1</sup>Patrizio Caturegli, <sup>2</sup>Oliver Laeyendecker, <sup>1</sup>Aaron A.R. Tobian, <sup>4</sup>David J. Sullivan
- 7
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- 16 Abstract 249 (limit 250 words)
- 17 Word count is 1139
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- 19 References 17
- 20 Conflict of interest- Authors declare no competing financial interests
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# 26 Abstract

27	Natural infection with the SARS-CoV-2 virus and spike protein immunization increase the serum spike
28	antibody levels. These levels, which differ among the various commercial assays, are used by the FDA
29	to qualify individuals as potential COVID-19 convalescent plasma (CCP) donors. Over a 3-year
30	period, (April 2020 – February 2023), we analyzed spike antibodies measured by a tertiary hospital
31	clinical immunology laboratory using the dilutional "Anti-SARS-CoV-2 ELISA" assay (Euroimmun
32	US, Mountain Lakes, NJ). The three year interval was arbitrarily classified into five periods based on
33	the SARS-CoV-2 strain variant epidemiology. A total of 15,820 sera, derived from 11,022 individuals
34	(6,362 females, mean age 50±21 years), ranging from severe immunocompromised state to routine
35	health visits, were tested for spike IgG antibodies. Spike IgG levels rose markedly over time, from a
36	median of 0.13 ELISA arbitrary units (AU) in period 1 to 48.7 in period 5 (p<0.0001). The spike IgG
37	80 <sup>th</sup> percentile distribution threshold was 0.55, 8.1, 9.6, 64.9, and 151 AU in each of these periods.
38	Using the 3.5 AU threshold the FDA uses to qualify CCP donors with this assay, the percentage of
39	subjects eligible for CCP donation would have been 11%, 44%, 61%, 81%, and 91% in the five time
40	periods. Antibody levels have risen more than hundred-fold while variants have become resistant to
41	clinically available monoclonal antibodies. As high-titer CCP is most effective against variants,
42	restricting CCP donors to those with spike antibody levels in the upper two deciles may allow
43	protection against variants when transfused to the immunocompromised.
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51 Importance-112 words (150 limit)

52	Despite the evolution of SARS-CoV-2 variants of concern and ongoing transmission, COVID-19
53	hospitalization and mortality rates continue to decline. Both the percent seropositive and antibody
54	levels have risen over the past three years. Here we observe more than 90% seropositivity as well as
55	more than a hundred-fold increase in spike IgG levels in a tertiary hospital clinical immunology
56	laboratory setting. Antibody effector functions (such as neutralization, opsonization, and complement
57	activation) and cell mediated immunity all contribute to protection from COVID-19 progression to
58	hospitalization, and all correlate to the total SARS-CoV-2 antibody levels. We recommend therapeutic
59	COVID-19 convalescent plasma be restricted to the top 20% of potential donors to maintain activity
60	against ongoing SARS-CoV-2 variant evolution.
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62	Keywords
63	COVID-19, SARS-CoV-2, serologic population kinetics, plasma donor transfusions
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68	Considering that populations have undergone multiple vaccinations and reinfections with
69	evolving SARS-CoV-2 variants, we analyzed the validity of the Euroimmun 3.5 arbitrary unit (AU)
70	threshold using a patient cohort of clinical provider ordered serology, seen at a tertiary hospital
71	between April 10, 2020 and Feb 28, 2023. This 3-year time interval was classified into five periods to
72	reflect the epidemiology of the SARS-CoV-2 strains: up to January 2021 for the original strain;
73	January to June 2021 for the alpha variants with partial vaccinations; July to November 2021 for the
74	delta variants; December 2021 to June 2022 for the omicron BA.1 and 2; and July 2022 to January
75	2023 for the omicron BA.4/5 with BQ.1 and XBB.
76	A total of 11,022 subjects (6,362 females and 4,660 males, 50±21 years of age) contributed
77	15,820 samples measured for spike IgG antibodies. The study population ranged from
78	immunocompromised to vaccinated immunocompetent individuals. SARS-CoV-2 antibody levels
79	increased significantly over time, from a median AU of 0.13 in the pre-vaccination period to 1.8, 5.3,
80	11.7, and 48.8 AU in the following periods (p<0.0001, Figure 1). During the first period, most sera
81	(2,763 of 3,109, 89%) tested negative (i.e., value <1.23 AU), while during the last period most tested
82	positive (2,306 of 2,422, 95%). The 80 <sup>th</sup> percentile of the spike IgG distribution was 0.55, 8.1, 9.6,
83	64.9, and 151 AU in the five time periods, indicating a greater than 250-fold increase when comparing
84	last to first periods. The increasing trend in spike antibody levels was confirmed when longitudinal
85	data analysis was performed in the subset of subjects (2,571 of 11,022, 23%) who were tested two or
<mark>86</mark>	more times (Figure 2A). Using the FDA COVID-19 convalescent plasma (CCP) eligibility threshold of
87	3.5 AU would have classified eligible CCP donors, 346 of the 3109 sera (11%) in period 1, 1324 of
88	3039 (44%) in period 2, 2263 of 3724 (61%) in period 3, 2849 of 3526 (81%) in period 4, and 2213 of
89	2422 (91%) in period 5. Thus, the vast majority of individuals currently qualify as potential CCP
90	donors since they have spike antibody levels exceeding the FDA criteria for "high titer" CCP.
91	The newer Euroimmun spike IgG assay features a standard curve composed of 6 calibrators
92	(ranging in concentration from 1 to 120 RU/mL), rather than the single calibrator found in the original

93	assay. Although the newer assay was not used in this study, we compared the two assays in a subset of
94	subjects (552 of 11,022, 5%). Upon transforming the raw antibody results to the log10 scale, the two
95	assays showed a highly significant linear correlation (adjusted r-squared of 0.861, p<0.0001, Figure
96	2B): for every unit increase in the value of the original assay, the value of the newer assay increased 14
97	RU/mL (95% CI from 13.1 to 15.7). A total of 58% of the subjects (243 of 417) evaluated for the assay
98	comparison in June 2021 (gray symbols) would qualify as CPP donors because having spike antibody
99	levels $>3.5$ AU in the original assay and $>55$ RU/mL in the newer assay. On the contrary, all 135
100	random patient subjects tested in February 2023 (red symbols) would meet the qualification criteria.
101	This study demonstrated that most individuals now qualify as a potential CCP donor. During
102	the initial COVID-19 pandemic months, therapeutic CCP was selected based on the donor's
103	symptomatology and any positive SARS-CoV-2 laboratory test(1). By February 2021, the FDA had
104	established therapeutic threshold values for the most commonly used serum spike antibody assays(2),
105	such as the 3.5 AU for the original Euroimmun "Anti-SARS-CoV-2 ELISA" assay (approved for the
106	US market on May 4, 2020), or the near equivalent threshold 55 RU/mL for the later version ("Anti-
107	SARS-CoV-2 Curve ELISA", approved on October 5, 2021)(3). Since ideal donors for therapeutic
108	CCP are those with the highest spike antibody levels, we suggest increasing the threshold as to include
109	only those who have antibodies in the upper two deciles of the spike antibody distribution. Vaccine
110	efficacy metrics often uses the reference of "COVID-19 convalescent plasma" levels(4, 5), which is a
111	broad range as seen here. We advocate that CCP units used for therapy comprise the upper quintile as
112	the goal at present and in the future for therapeutic CCP. The volume of distribution for CCP
113	approaches 3-5 L with 15-to-20-fold dilution from 250 milliliters of plasma(6). An outpatient treatment
114	CCP study qualified the top 60% of donors and demonstrated in the top 30%, early plasma
115	administration reduced hospital risk 92%(95% CI 41%-99%, p=.014)(7). The high titer plasma quintile
116	retains potent virus neutralizations against current and future variants for months(8, 9).

117	As monoclonal antibody therapy has become ineffective, there is increased interest in
118	polyclonal CCP to complement small molecule antiviral drug therapy, especially for the
119	immunocompromised patients and those at highest risk of hospitalization(10, 11). There have been
120	offsetting trends. On one hand with more vaccine boosters and cumulative COVID-19 incidence, the
121	levels of neutralizing antibodies (not the total spike antibody measured here) are trending to ten times
122	the geometric means from the original CCP from unboosted WA-1 or pre-alpha COVID-19(9, 12). On
123	the other hand new Omicron variants like XBB and BQ are more than ten times resistant to virus
124	neutralization compared to WA-1 with preBQ or preXBB vaccine AND recent Omicron plasma(13).
125	The present blood donor qualification system in the past has tolerated the rough correlation 📻 tal
126	spike, S-1 or RBD antibodies to virus neutralization(14, 15).
127	True identical match to circulating variants like XBB with collection and rapid dispensing may
128	not be achievable. However, primary virus neutralization data and two systemic reviews show CCP
129	from clinical cohorts (not qualified donor units) collected after previous variants are still able to
129 130	from clinical cohorts (not qualified donor units) collected after previous variants are still able to neutralize both existing and future variants(12, 13). High titer viral mismatch neutralizes to the same $\overline{\mathcal{V}}$
130	neutralize both existing and future variants(12, 13). High titer viral mismatch neutralizes to the same $\overline{\checkmark}$
130 131	neutralize both existing and future variants(12, 13). High titer viral mismatch neutralizes to the same extent as medium titer match. There is early data from vaccination cohorts, that despite the fold drop in
130 131 132	neutralize both existing and future variants(12, 13). High titer viral mismatch neutralizes to the same extent as medium titer match. There is early data from vaccination cohorts, that despite the fold drop in virus neutralizations by XBB.* and BQ.1, hospital rates are still low (CDC Tracker Feb 12, 2023 at
<ul><li>130</li><li>131</li><li>132</li><li>133</li></ul>	neutralize both existing and future variants(12, 13). High titer viral mismatch neutralizes to the same extent as medium titer match. There is early data from vaccination cohorts, that despite the fold drop in virus neutralizations by XBB.* and BQ.1, hospital rates are still low (CDC Tracker Feb 12, 2023 at 1.1 hospitalizations per 100,000 falling from 2/100,000 on Jan 5, 2023) with current vaccinations.
<ul> <li>130</li> <li>131</li> <li>132</li> <li>133</li> <li>134</li> </ul>	neutralize both existing and future variants(12, 13). High titer viral mismatch neutralizes to the same extent as medium titer match. There is early data from vaccination cohorts, that despite the fold drop in virus neutralizations by XBB.* and BQ.1, hospital rates are still low (CDC Tracker Feb 12, 2023 at 1.1 hospitalizations per 100,000 falling from 2/100,000 on Jan 5, 2023) with current vaccinations. A strength of this analysis is the more than 15 thousand samples over a three-year period on a
<ul> <li>130</li> <li>131</li> <li>132</li> <li>133</li> <li>134</li> <li>135</li> </ul>	neutralize both existing and future variants(12, 13). High titer viral mismatch neutralizes to the same extent as medium titer match. There is early data from vaccination cohorts, that despite the fold drop in virus neutralizations by XBB.* and BQ.1, hospital rates are still low (CDC Tracker Feb 12, 2023 at 1.1 hospitalizations per 100,000 falling from 2/100,000 on Jan 5, 2023) with current vaccinations. A strength of this analysis is the more than 15 thousand samples over a three-year period on a single Euroimmun serologic assay platform. Limitations include variable number of
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<ol> <li>130</li> <li>131</li> <li>132</li> <li>133</li> <li>134</li> <li>135</li> <li>136</li> <li>137</li> <li>138</li> </ol>	neutralize both existing and future variants(12, 13). High titer viral mismatch neutralizes to the same extent as medium titer match. There is early data from vaccination cohorts, that despite the fold drop in virus neutralizations by XBB.* and BQ.1, hospital rates are still low (CDC Tracker Feb 12, 2023 at 1.1 hospitalizations per 100,000 falling from 2/100,000 on Jan 5, 2023) with current vaccinations. A strength of this analysis is the more than 15 thousand samples over a three-year period on a single Euroimmun serologic assay platform. Limitations include variable number of immunocompromised antibody deficient individuals and consolidation of data points near 10 and 100 AU from not fully diluting all the samples both which may underestimate population increases in antibody levels.

- 142 antibodies(16), the upper quintile of available CCP donor units will provide the highest effective viral
- 143 specific antibody dose for the longest duration against both matched and mismatched SARS-CoV-2
- 144 variants.
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217 Authorship Contribution: DS & MC designed data analysis from existing hospital data. MC analyzed

- 218 data, DS drafted manuscript with input from all (OL and AT) authors and all authors approved the final
- 219 version of the manuscript.

220 **Conflict-of-interest disclosure:** All authors report no relevant disclosures.

221

222 Figure legends

223	Figure 1. Increasing trend of spike IgG antibodies in a tertiary hospital patient population. The 3-
224	year period between April 2020 to February 2023 was divided into five time periods to represent the
225	alpha variant (up to January 2021) the alpha variants with partial vaccinations (to June 2021), the delta
226	variants (to December 2021), the omicron BA.1 and 2 (to July 2022). and the omicron BA.4/5 with
227	BQ.1 and XBB variants (to January 2023). A total of 15,820 sera are shown by the individual points,
228	contributed by 11,022 patients. Most patients (8,451, 77%) were tested only once during the 3 years,
229	while the remaining 2,571 underwent sequential measurements. The box plot in each of the five time
230	periods represent the middle 50% of the observations, bordered at the 25 <sup>th</sup> and 75 <sup>th</sup> percentiles, and
231	contain a line indicating the median antibody value. The 3.5 ELISA optical density value suggested by
232	the FDA as cutoff for COVID-19 convalescent plasma donations is shown by the red dashed line.
233	
234	Figure 2A. Sequential spike IgG antibody levels in 49 of the total 2,571 patients with sequential
235	measurements. The 49 patients were tested between 7 and 13 times, contributed 381 sera, and had a
236	mean follow-up time of 381 days. The shaded area represents the 95% confidence interval around the
237	linear fit (which is not shown).
238	

Figure 2B. Comparison of the Euroimmun spike antibody ELISA assays, the original version
released in May 2020 and the newer version with a standard curve released in October 2021. The
dotted lines indicated the FDA recommended threshold for COVID-19 convalescent plasma donation:
3.5 AU for the original assay on y-axis and 55 RU/mL for the newer assay on x-axis. The comparison
was made in June 2021 for the grey and February 2023 for the red symbols.

Dr. Bar-On Yotam Editor, Microbiology Spectrum ybaron@technion.ac.il

#### Dear Dr. Bar-On Yotam,

Thank you very much for the effort you and the reviewers dedicated to our paper [Spectrum 02183-23]. We were pleased to read that only minor modifications are required before publication. The two reviewers made several insightful comments, which have helped us to improve the paper. Our point-by-point rebuttal is presented here below. The corresponding changes in the text are also indicated.

## From Reviewer 1

1. The study could have included virus neutralization tests in a subset of patients from each of the time intervals, as to correlate the total serum antibody levels with their neutralization function.

The point is well taken. However, it is well-established that antibody concentrations positively correlate with neutralization function. For example, Otter et al in Microbiology Spectrum compared 138 convalescent blood donor sera using the same Euroimmun assay we used to measure total antibody levels and a neutralization assay. The authors reported a highly significant positive correlation (Pearson r coefficient of 0.83, p<0.0001) [Figure 5 in (1)]. Numerous other studies have described increases in neutralization titer over the course of the pandemic in association with increased total antibody levels. For example: a 9-fold increase from the 311 geometric mean titer from 27 studies against the WA-1 strain to the 2753 titer after both pre-omicron COVID-19 and vaccination in 19 studies; and a 10-fold rise (from 15 to 192) with Omicron BA.1 tested in parallel in the same convalescent plasma studies (2). Convalescent plasma after vaccination in 12 studies harvested in 2022 during Omicron has GMT neutralizations near 6000 essentially a 20-fold increase(3). Finally, it is important to remember the design of the present study: our was a retrospective study aimed to analyze the total spike IgG levels measured in a routine clinical laboratory, a laboratory not equipped with the research tools that are needed to perform viral neutralization assays.

2. The authors could have enriched the study by associating the spike antibody results with other variables such as clinical, epidemiological, or biological variables of the virus.

Thank you for the suggestion. Routine clinical laboratories do not have access to epidemiological data or biological characteristics of the virus, but can integrate laboratory results with basic clinical characteristics, such as age and gender. We have integrated the reviewer's suggestion in the revised manuscript, as detailed in our responses to the specific suggestions made by the reviewer.

3. The authors could add a table (plot) describing the population analyzed. This could be organized by time intervals; in each of them, it could be indicated, for example, the number of women and men, the number of people vaccinated, the number of people with previous SARS-CoV- 2 infections, and the number of seropositive patients. This table could be cited in the text. We have now added a table (Supplemental Table 3) that captures the contemporaneous cumulative SARS-CoV-2 cases in the State of Maryland, along with the percentage of population vaccinated, seroprevalence from CDC Seronet, and the CDC Tracker data from the blood donor seroprevalence studies. Corresponding changes have been integrated in the manuscript.

4. In addition to the median, it can be informative to add to the text the minimum and maximum values of the number of antibodies.

We have now calculated median, minimum and maximum for the five COVID-19 time periods and presented the results in a new table (Supplemental Table 2).

5. The authors could evaluate if there are significant statistical differences between the median number of antibodies of the different time intervals.

Thank you for the suggestion. Spike antibody levels increase significantly from one period to the next one, both in a crude (unadjusted) analysis and after adjusting for gender and age. We have presented these new results in the revised text and in Supplemental Table 2.

6. The authors could evaluate if there are significant statistical differences in the number of antibodies between sexes for the same time interval or between different intervals for the same sex.

We have now performed this statistical analysis. Gender was indeed strongly associated with spike antibody levels, in keeping with the notion that the number of B lymphocytes and the levels of circulating antibodies are higher in women (4). We have integrated these new findings in the revised text and in Supplemental Table 1.

# From Reviewer 2

We thank the reviewer for considering our study interesting and having important implications for the use of COVID-19 convalescent plasma therapy.

1. The authors mention that the FDA uses a threshold of 3.5 AU to qualify CCP donors. However, they do not discuss why this threshold was chosen and why the 55 AU was also chosen in the newer assay.

The FDA first issued the Emergency Use Authorization (EAU) for administering CCP to hospitalized patients in August 2020, qualifying donors based on a documented history of COVID-19 or a positive spike antibodies Orthos test. The EAU was later revised in Feb 2021 to include several commercial spike antibody assays, such as the Euroimmun assay used in this study. In this assay, the positivity threshold recommended by the manufacturer is 1.1 ELISA optical density (OD) ratio (the ratio derived by dividing the optical density obtained with the patient serum by the density obtained with the single calibrator provided with the kit). The positivity threshold validated in our laboratory was slightly higher, at 1.23 (5); and the qualification threshold for CCP donors chosen by the FDA was 3.5. About one year later, in February 2022, the FDA authorized the use of a newer version of the Euroimmun assay, called QuantiVac. This newer version features 6 calibrators (rather than just one), thus allowing the derivation of a standard curve and the expression of results as arbitrary, relative units per mL (rather than as a ratio of optical densities). The FDA selected a value of 55 RU/mL as the qualification threshold for CCP donations, thus suggestion of conversion factor of 15.7 between the two Euroimmun assays (that is, a value of 1 OD ratio in the original assay corresponds to a value of 15.7 RU/mL in the QuantiVac assay). In this study (Figure 2B), we showed that the conversion factor between the two assays is, in our hands, 14, thus similar to the 15.7 in the FDA data. But the bottom line is that the qualification threshold for CCP donations selected by the FDA when using

Euroimmun assays (either the 3.5 of the original assay or the corresponding 55 in the newer assay) has not essentially changed over the past 2 years. Our data (Figure 1) show that while using these thresholds would have qualified only the top 15% of donors during the initial period (March 2020 – December 2020), they would have qualified almost everybody as donor (95%) during the last period (July 2022 – February 2023). This selection strategy is detrimental to recipients of CCP because studies have shown that the benefits of CCP transfusions are greater when the levels of spike antibodies they contain are higher (see point 2 below for references).

2. The authors suggested that restricting CCP donors to those with high titer spike antibodies may be more effective in protecting immunocompromised patients from variants. However, they do not provide any data to support this suggestion. It would be helpful to see data on the clinical effectiveness of CCP therapy in patients with different levels of spike antibodies.

Among outpatients with COVID-19, performing a meta-analysis of 5 randomized trials 2,620 patients, Levine and colleagues reported that CCP was more effective in reducing all-cause hospitalization when given within 5 days of symptom onset and contained high (top half of research donor units) antibody titers. (6). Similarly, in hospitalized patients, Joyner and colleagues noted reductions in death when using CCP of higher antibody levels (7). A large meta-analysis in hospitalized patients noted better outcomes with higher antibody doses in those already hospitalized (8). Overall, data suggest that is best to administer CCP that contains high antibody titers. As far as the immunosuppressed population is concerned, clinical data are sparse during the period of this study, so we have revised the final statement of the Abstract accordingly.

3. Abstract, line 34, yellow highlight: (6,362 females, mean age 50±21 years) We have now revised this line by specifying the numbers in both genders.

*4. Abstract, lines 41-43, yellow highlight* We have now revised the English to make the conclusion clearer.

5. Importance, line 51, yellow highlight

We have now removed the word limit.

6. Text, lines 85-86, yellow highlight

We have changed the text to "who were tested longitudinally two or more times".

# 7. Text, lines 101-103, yellow highlight

We have now better explained the spike antibody thresholds the FDA had chosen to consider donors eligible for CCP donations.

8. Text, lines 111-112, yellow highlight We have replaced "upper quintile" with "upper decile" (also throughout text).

*9. Text, line 15, yellow highlight ("this sentence is not really clear").* We have deleted this sentence in the revision.

# 10. Text, lines 130-131 and 136, yellow highlight.

This paragraph has been largely re-written.

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August 29, 2023

Dr. David J Sullivan Johns Hopkins University Bloomberg School of Public Health Molecular Microbiology & Immunology 615 N. Wolfe St., Rm. W 4606 Baltimore, MD 21205

Re: Spectrum02183-23R1 (Hundred-fold increase in SARS-CoV-2 spike antibody levels over three years in a hospital clinical laboratory.)

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