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SARS-CoV-2 neutralizing antibody response in vaccinated and non-vaccinated hospital healthcare workers with or without history of infection

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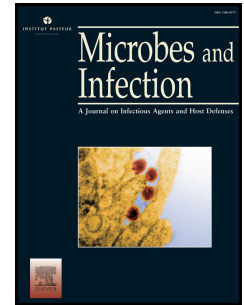
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1 **SARS-CoV-2 neutralizing antibody response in vaccinated and**
2 **non-vaccinated hospital healthcare workers with or without**
3 **history of infection**

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22 **Abstract**

23 Between March 2021 and February 2022, SARS-CoV-2 neutralizing antibodies dynamics was
24 investigated in a prospective observational study in 903 healthcare workers of a hospital in
25 Switzerland. A surrogate neutralization assay measuring the competitive inhibition of the angiotensin
26 converting enzyme 2 (ACE2) binding to the spike protein (S) of the SARS-CoV-2 wild type virus and to
27 five variants of concern (Alpha, Beta, Gamma, Delta, Omicron) was used. We observed a broad
28 distribution of neutralization activity among participants and substantial differences in neutralizing
29 titers against variants. Participants were grouped based on combinations of vaccination status (1, 2 or
30 3 doses) and/or prior or subsequent SARS-CoV-2 infection/reinfection. Triple vaccination resulted in
31 the highest neutralization response, as did double vaccination with prior or subsequent infection.
32 Double vaccination without infection showed an intermediate neutralization response while SARS-
33 CoV-2 infection in non-vaccinated participants resulted in poor neutralization response. After triple
34 vaccination or double vaccination plus infection, additional vaccination and/or reinfection had no
35 impact on neutralizing antibody titers over the observed period. These results strongly support the
36 booster dose strategy, while additional booster doses within short time intervals might not improve
37 immunization. However, dynamics of neutralizing antibodies titers needs to be monitored individually,
38 over time and include newly emerging variants.

39

40 **Keywords:** epidemiology, neutralization test, SARS-CoV-2, vaccination

41

42 1. Introduction

43 End of 2019, a new virus named Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) was
44 reported in Wuhan City (China) which is responsible of the still ongoing pandemic of Coronavirus
45 Disease 2019 (COVID-19). In March 2022, half a billion individuals have been infected worldwide and
46 2.5 billions have been vaccinated (WHO, 2022). In Switzerland, 2.5 millions infections were recorded
47 and 70% of the population was vaccinated with two doses of mRNA vaccine (Pfizer-BioNTech or
48 Moderna) while 42% received a third booster dose [1]. In addition, a vaccine dose was recommended
49 to individuals who had been infected before vaccination. Re-infections post-vaccination were reported
50 with a frequency of 1% to 15% depending on the type of study and the viral variant under investigation
51 [2-4]. Individuals with a positive SARS-CoV-2 serology can be categorized into three major groups:
52 vaccinated without infection, convalescent after infection, vaccinated plus infection. These groups can
53 be subdivided depending on the number of vaccination doses received and the timing of infection
54 before or after vaccination.

55 Serological investigations look for the presence of virus-specific antibodies as a marker of previous
56 infection or vaccination [5]. However, these analyses do not assess whether the detected antibodies
57 display a protective antiviral activity [6]. In SARS-CoV-2, the viral spike protein (S) is the primary target
58 for neutralizing antibodies, which inhibit its binding to the host angiotensin-converting enzyme 2
59 (ACE2) receptor, the trigger of cell membrane fusion between the virus and the human cell [7-10]. As
60 neutralizing antibodies play a key role for viral clearance, the quantification of their activity provides a
61 good estimate of immune protection [11-13]. The gold standard for measuring SARS-CoV-2 neutralizing
62 antibodies activity relies on quantification of the reduction of virus-induced cytopathic effects after
63 infection of ACE2-expressing cells with live virus but simpler cell-free neutralization assays have been
64 developed [14]. In the present study, we used a surrogate neutralization assay measuring the
65 competitive inhibition of ACE2 binding to a trimeric S protein loaded on beads [15]. This method is

66 quantitative, high throughput and allows the simultaneous evaluation of the neutralization activity
67 targeting spike protein encoded by different SARS-CoV-2 variants of concern [16, 17]

68 Here, we investigated at four time points, between March 2021 and February 2022, the dynamics of
69 SARS-CoV-2 neutralizing antibodies against the original Wuhan wild type virus and five major variants
70 of concern (Alpha, Beta, Gamma, Delta and Omicron BA.1). This prospective observational study was
71 conducted in health-care workers at the “Ensemble Hospitalier de la Côte” (EHC), a public hospital in
72 Morges, Western Switzerland with 1’800 employees, 240 acute beds and 85 post-acute beds. The
73 objective of the investigation was to quantify the neutralization activity of anti-SARS-CoV-2 antibodies
74 in seropositive participants according to their vaccination and convalescent status.

75 **2. Materials and methods**

76 **Study design:** A prospective observational study was proposed to all EHC employees, Morges,
77 Switzerland (n=1’800). Participants over 18 years old were included on a voluntary basis after written
78 informed consent at one of the following study visits: March 2021, June 2021, September 2021, and
79 February 2022. Volunteers had the opportunity to be recruited or drop out at any of the four visits.

80 **Questionnaire:** All participants filled in a questionnaire with demographic characteristics, history and
81 date of positive SARS-CoV-2 RT-PCR or antigen (AG) tests as well as date of first, second and/or third
82 vaccination (Supplementary File 1). Questionnaires were manually digitalized.

83 **Serum sampling:** Blood was obtained at the inclusion and follow-up visits (10 ml Monovette[®] without
84 anticoagulant) and processed as previously described [18].

85 **Serological Method:** The samples were analyzed with a standard serological test for IgG anti-Spike
86 (anti-S) and IgG anti-Nucleocapsid (anti-N) SARS-CoV-2 antibodies using the Luminex[®] system [19] as
87 previously described [18]. Samples with a positive serology were further investigated using a surrogate
88 neutralization test [15]. Dilutions of serum samples in PBS were added to plate wells containing S
89 proteins-coupled beads. Variant investigated include the sequence of the wild type Wuhan, Alpha,

90 Beta, Gamma, Delta and Omicron BA.1. The positive control for 100% neutralization consisted of a
91 cocktail of two neutralizing antibodies binding distinct epitopes on the SARS-CoV-2 Spike protein. In
92 absence of neutralizing antibodies, a tagged ACE2-Fc can freely bind to the S protein and induce
93 maximal fluorescence intensity (MFIs). Neutralizing antibodies bind to the S protein and compete with
94 its binding to ACE2: this inhibition effect can be quantified by reduced fluorescence intensities. Results
95 are presented as IC_{50} of the calculated inhibition curve. Neutralization responses were classified in four
96 categories: <50: undetectable neutralizing activity, 50-100: low neutralizing activity, >100-150:
97 moderate neutralizing activity, >150: strong neutralizing activity. All sera were processed at the
98 Laboratory of Immunology and Allergy, Lausanne University Hospital (CHUV), Switzerland.

99 **Group definition:** Participants were grouped according to data extracted from questionnaires and
100 serological results. In absence of a history of documented infection (positive RT-PCR or AG tests)
101 volunteers with a positive anti-S SARS-CoV-2 serology prior to the first vaccination (n=32) or with a
102 positive anti-N serology (n=58) were excluded from the group vaccinated only. As the date of infection
103 was unknown they were not included in the group vaccination/infection. Time course representation,
104 in convalescent subjects and in those vaccinated with two doses or three doses, t=0 was defined as
105 the date of the first positive RT-PCR or AG test, of the second or the third vaccination dose,
106 respectively. The time interval in days elapsed between t=0 and the date of the study visit is
107 represented. For vaccinated individuals (two or three doses), the status of infection before or after
108 vaccination was determined using respectively, the second or third vaccination date as reference. For
109 participants vaccinated with a single dose, the date of the first dose was used. For participants with
110 two reported SARS-CoV-2 infection episodes, the first date was used unless otherwise specified in the
111 text.

112 **Statistical analysis:** All analyses were performed with R version 4.0.2. Local polynomial regression
113 fitting was performed using `stat_smooth` method `loess`. Graphs were drawn with `ggplot2`. Median and
114 interquartile ranges were used to describe continuous variables. Kruskal-Wallis test by rank was used

115 to compare the different groups (pairwise.wilcox.test). The significance level was set with two-sided
116 $p < 0.05$.

117 **Ethics:** The Cantonal Ethical Review Board for Human Research (CER-VD, Commission cantonale
118 d'éthique de la recherche sur l'être humain) approved the study protocol and the participants'
119 informed consent form (Authorization Nr 2020-02300).

120 **3. Results**

121 **3.1 Demographics of study volunteers**

122 A total of 903 volunteers participated to this prospective observational study, representing half of the
123 1'800 hospital employees. 191 participated to all four visits, 147 to three visits, 207 to two visits and
124 358 to one visit for a total of 1977 sera. The majority of study participants were women (84%) and the
125 median age was 43 years (IQR 33-52) (Fig. 1A and Fig. S1A). 74.5% of participants were vaccinated with
126 two doses (or one dose if they had a previously documented infection). Half of them received a third
127 (respectively second) booster dose. 39.6% of participants reported a history of SARS-CoV-2 infection
128 documented by a positive RT-PCR or AG test. The majority of these infections occurred during the
129 epidemic waves preceding the first vaccination campaign in March 2021 (Fig. 1A and 1B). At the first
130 visit, 45.3% of volunteers had a positive SARS-CoV-2 serology, 75% at the second visit, 85% at the third
131 and 94.8% at the last visit (Table S1). A positive anti-S serology was observed in both convalescent and
132 vaccinated individuals, while the anti-N serology was only positive after a natural infection. Re-
133 infections post-vaccination occurred in 15.9% of volunteers, almost exclusively with the Omicron BA.1
134 strain in December 2021-January 2022, prior to the last visit (Fig. S1B).

135 **3.2 Neutralization activity across variants**

136 The neutralization activity was investigated against the S protein of the Wuhan SARS-CoV-2 wild type
137 and of the variants of concern Alpha, Beta, Gamma, Delta and Omicron. The neutralization assay was
138 available at the time of the third study visit when variants Alpha, Beta and Gamma were circulating

139 worldwide while Delta was emerging and Omicron was absent. Therefore, for the first, second and
140 third visits, neutralization tests were performed on the SARS-CoV-2 wild type plus Alpha, Beta, and
141 Gamma. At the fourth visit, Alpha, Beta and Gamma variants had disappeared while the Delta was
142 being progressively replaced by Omicron (BA.1). Hence, neutralization tests were performed on the
143 SARS-CoV-2 wild type plus Delta and Omicron. Overall, we observed a broad distribution of
144 neutralization activity among participants indicating an important variability in inter-individual
145 humoral immune responses and among viral variants (Fig. 1C). The Beta and Gamma variants escaped
146 significantly the neutralizing activities of anti-S antibodies while the SARS-CoV-2 wild type showed the
147 highest response to neutralizing antibodies (Fig. 1C). A progressive increase of neutralization titers was
148 observed across the four visits mirroring the increasing number of volunteers who were vaccinated
149 and/or reported an infection during the study period (Fig. 1D). The relative low increase in
150 neutralization response between the second (June 2021) and third (September 2021) visit is likely
151 linked to the absence of vaccination campaign and low prevalence of infections during the summer
152 2021 (Fig. 1B and 1D). We observed no significant difference in neutralizing activity among age groups
153 (Fig. 1E).

154 **3.3 Neutralization antibody titers in convalescent and vaccinated individuals**

155 Participants with a positive serology (n=773) were classified according to the vaccination status (one
156 dose, two doses, or three doses) and/or the convalescent status (history of SARS-CoV-2 infection and
157 reinfection documented by a positive RT-PCR or AG test) (Table 1). We first investigated the dynamics
158 of the serological response in volunteers vaccinated with two or three doses without history of
159 infection and in convalescent individuals without history of vaccination. These three groups displayed
160 a simple immunization event (second vaccination, third vaccination or infection) that was used as
161 reference time point to follow the dynamics of the neutralizing antibodies response. The date of the
162 second or third vaccination and of the first positive RT-PCR or AG test was set as $t=0$ for vaccinated and
163 convalescent volunteers, respectively. The group of participants who were vaccinated with one single

164 dose is not shown as most volunteers registered only for the second visit and then dropped out of the
165 study. A local polynomial regression-fitting model was used to display an average neutralization
166 response curve across variants and for each group. Convalescent participants showed the lowest
167 neutralization activity after a three to four-month time interval following the reported infection (Fig.
168 2A-C). Vaccination with two doses without a history of infection resulted in a robust neutralization
169 response that slowly decreased over a three to six-month time interval (Fig. 2D-F). The third booster
170 dose of SARS-CoV-2 mRNA vaccine was followed by a rapid and significant increase in neutralization
171 titers (Fig. 2G-I). As the booster dose was made available shortly before the last visit, the dynamic of
172 antibody titers after the vaccination was not recorded. The increases in neutralization titers observed
173 beyond 250 days after a documented infection or a second vaccination likely represent reinfections,
174 that were in part asymptomatic (Fig. 2A-F). The highest neutralization response was observed after
175 triple vaccination and an intermediate response after two vaccinations. An increase of neutralization
176 titers was observed after a second infection in convalescent only individuals in whom the response
177 was comparable to that observed after double vaccination (Fig. 3A). Although we observed substantial
178 differences comparing the neutralization activities against the variants, the overall trends remained
179 similar: convalescent individuals had the lowest neutralization response and a progressive increase of
180 the neutralization activity was observed after double and triple vaccination (Fig. 3B-D and Fig. S2A-F).

181 **3.4 Neutralization antibody titers in vaccinated individuals with documented COVID-19 infections**

182 We investigated the different combinations of vaccinations (one, two, or three doses) with a
183 documented SARS-CoV-2 infection occurring prior or post vaccination. A two-dose vaccination
184 administered prior or after a natural infection (three immunization events) resulted in a neutralization
185 response comparable to that obtained after a triple vaccination (three immunization events) (Fig. 4A-
186 C). The same neutralization dynamics was observed against the SARS-CoV-2 wild type, Delta and
187 Omicron variant, although with Omicron differences were less pronounced likely due to its ability to
188 escape humoral response. Among the volunteers, 39 reported two COVID-19 episodes confirmed by

189 positive RT-PCR or AG test more than 60 days apart. All were vaccinated with either two or three doses
190 and were re-infected recently with the Omicron BA.1 variant. Four (2 vaccinations and 2 infections) or
191 five (3 vaccinations and 2 infections) immunization events did not further boost the neutralization
192 titers compared to those observed after three immunization events (3 vaccinations or 2 vaccinations
193 and 1 infection) (Fig. 4A-C, Table S2). Of note, volunteers with two infections showed the lowest
194 neutralization titers at the study visit preceding the second infection, which was in agreement with a
195 higher probability of getting re-infected (Fig. S3A-B). The same variation trends were observed against
196 all variants.

197 **4. Discussion:**

198 In this one-year prospective observational study in 903 hospital employees, we observed important
199 variations in neutralizing anti-SARS-CoV-2 antibodies activities in seropositive individual [13]. At the
200 group level, overall trends in viral neutralization could be predicted based on the history of vaccination
201 and/or infection. However, large inter-individual differences highlight the difficulty to accurately
202 predict the level of protection and illustrate the value of individual assessments of neutralizing
203 antibodies.

204 As reported in other studies [20, 21], neutralization titers differed significantly against the tested SARS-
205 CoV-2 variants. Antibodies showed the highest neutralizing activity against the Wuhan wild type virus
206 related to its spike protein being used for the development of mRNA vaccines. In convalescent non-
207 vaccinated individuals, mostly exposed to the Alpha, Delta and Omicron variants, the neutralization
208 activity was highest against the Wuhan wild type virus; although after natural infection, a variant-
209 specific increase in neutralization titers would be expected. This discrepant observation could suggest
210 that immunity to variants is not solely due to neutralization antibodies against a mutated S-protein but
211 also to a more complex interplay with the immune system.

212 Significant differences in neutralizing antibodies activity were also observed among the different
213 groups based on history of vaccination (one dose, two doses, or three doses) and SARS-CoV-2 infection.

214 We showed that individuals with a convalescent status had a significantly lower neutralization
215 response compared to vaccinated individuals [22, 23]. SARS-CoV-2 antibody response after infection
216 was previously shown to correlate with the severity of the disease [24-26]. In the present study, only
217 volunteers with low to mild COVID-19 infections (only one volunteer reported a hospitalization) were
218 investigated, which represents a good estimate of the immunization profile in the general population.

219 In individuals vaccinated with two doses, we observed significant neutralization titers followed by a
220 progressive decrease beyond three months after the second dose. A third booster dose resulted in a
221 significant rebound of neutralization activity. A similar boosted neutralization response was observed
222 in individuals with SARS-CoV-2 breakthrough infection after two vaccine doses [27] and in convalescent
223 individuals who received two vaccine doses after infection. These observations suggest that the
224 sequential order of different immunological stimulations (vaccination followed by infection or
225 *viceversa*) does not significantly impact the level of neutralization antibodies [28, 29]. A maximal
226 neutralization response was observed after three immunological stimuli (triple vaccination or double
227 vaccination preceded or followed by infection) while double vaccinated or convalescent individuals
228 showed significantly lower neutralization titers. After three immunization events, additional
229 vaccination or reinfection had limited impact on the neutralization activity. This suggests that
230 additional boosters (four or five immunization events) after reaching a neutralization antibodies titers
231 plateau might be of limited value in the following three to six months period while the persistence of
232 neutralization activity beyond this period remains to be investigated. Indeed, as a significant decrease
233 in neutralizing antibody titers was observed over time in double vaccinated individuals, a similar
234 decline might occur after three immunization events. In addition, significant differences might occur
235 over time among triple vaccinated individuals and those with hybrid immunity (vaccination and natural
236 infection). These heterologous immunization regimens can result in different long-term neutralization
237 responses [28, 30], as shown in a recent study showing that booster durability was longer in
238 participants who had breakthrough infection [31]. Interestingly, we only observed reinfections in

239 vaccinated individuals, but with more than 75% of volunteers being vaccinated the significance of this
240 observation is unclear.

241 Among limitations of the present study, the used neutralization test is a proxy of the immune response
242 measuring solely the antibodies activity on the interaction between the S protein and its ACE2 receptor
243 *in vitro*. The complex interplay of humoral and cellular immune response to infection and/or
244 vaccination was not investigated. While we observed no significant correlation between neutralizing
245 antibody titers and age, other studies showed a consistent decrease of immunity in older individuals
246 [26, 32, 33]. As the present investigation was restricted to working individuals younger than 65 years,
247 we are unable to draw any conclusions on the duration of immunity in the elderly.

248 In conclusion, a triple vaccination or a natural infection prior or after a double vaccination result in a
249 robust neutralization response strongly supporting a booster dose strategy. Although additional
250 booster doses within three to six months after a third immunization event (infection or vaccination)
251 might not boost immunity, the decline of neutralizing antibodies titers beyond this time window needs
252 to be monitored on an individual basis by including new variants in neutralization assays. Yet the
253 protection conferred by neutralizing antibodies against emerging variants is unpredictable: for
254 example, immunity was shown to be strongly reduced against the most recent Omicron BA.4 and BA.5
255 variants [4] and appears to more rapidly decline [34]. A booster vaccine dose integrating newly
256 appearing variants might contribute to meet the complex challenge of maintaining an effective
257 immunity over time.

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277 **Conflict of interest declaration**

278 G. Greub is medical advisor of Resistell, a startup active in the development of a new instrument to
279 faster antibiotic susceptibility results. G. Greub has a research agreement with Becton Dickinson
280 (USA) and Resistell (Switzerland), both unrelated to the present work. The rest of the authors have
281 no conflict of interest to declare.

282 **Figures and legends**

283 **Figure 1**

284 **A.** Schematic representation of the study period: study visits, circulating SARS-CoV-2 variants, key
285 epidemiological and immunization events. BA.1 and BA.2 are from the Omicron lineage. **B.** Positive
286 SARS-CoV-2 RT-PCR and AG tests in volunteers throughout the study. **C.** Aggregate data of neutralizing
287 antibodies titers against the Wuhan SARS-CoV-2 wild type and different variants (Alpha, Beta, Gamma,

288 Delta, Omicron) throughout the study. **D.** Dynamics of neutralization response against the different
289 variants at each of the four visits. Omicron was only investigated at the fourth visit, because it emerged
290 after the third visit. Alpha, Beta and Gamma were only investigated at the first, second and third visit,
291 as they disappeared at the time of the fourth visit. **E.** Age of volunteers versus neutralization response
292 to the different variants.

293 **Figure 2**

294 Dynamics of neutralizing antibodies titers over time against the Wuhan SARS-CoV-2 wild type, Delta
295 and Omicron variants. As Omicron was only investigated at the last visit, fewer points are represented.
296 Dashed lines: < 50 (red): no neutralizing activity; 50 – 100 (blue): low neutralizing activity; 100 – 150
297 (black): moderate neutralizing activity; > 150: high neutralizing activity [24]. **A-C.** Convalescent non-
298 vaccinated volunteers. Time 0 corresponds to the reported date of the first positive SARS-CoV-2 RT-
299 PCR or AG test as documentation of infection. **D-F.** Volunteers vaccinated with two doses and without
300 history of COVID-19. Time 0 corresponds to second vaccination date. **G-I.** Volunteers vaccinated with
301 three doses and without history of COVID-19. Time 0 corresponds to the third vaccination date.

302 **Figure 3**

303 **A.** Neutralizing antibodies titers against the Wuhan SARS-CoV-2 wild type, Delta and Omicron variants
304 in convalescent individuals after the first or second infection and in vaccinated individuals after the
305 second or third vaccination. Only the serological data after either the first or second infection and the
306 second or third vaccination are shown in the graph. **B-D.** Neutralization response against the Wuhan
307 SARS-CoV-2 wild type, Delta and Omicron variants in convalescent individuals and in individuals
308 vaccinated with two or three doses.

309 **Figure 4**

310 Neutralization responses after different numbers of immunization events including infections and/or
311 vaccinations. **A.** Wuhan SARS-CoV-2 wild type. **B.** Delta variant. **C.** Omicron variant.

312 Table 1

313 Groups of volunteers according to their vaccination status (no, one, two or three vaccine doses) and/or
314 convalescent status (no, one or two reported positive SARS-CoV-2 PCR or AG tests as a documentation
315 of infection/reinfection). Number of volunteers / number of serological data throughout the study
316 visits are reported. Of importance, volunteers could be included and drop out at any of the four study
317 visits.

318 Supplementary materials**319 Figure S1**

320 **A.** Distribution of age of study participants. **B.** Second SARS-CoV-2 positive RT-PCR and AG test
321 reported by the participants: re-infections occurred mostly in December 2021 and January 2022 with
322 the Omicron BA.1 variant.

323 Figure S2

324 Neutralizing antibodies titers against Alpha, Beta, and Gamma variants at first, second and third study
325 visits. **A-C.** Convalescent volunteers. Time 0 corresponds to the date of the first positive SARS-CoV-2
326 RT-PCR or AG test. **D-F.** Volunteers vaccinated with two doses. Time 0 corresponds to the second
327 vaccination date. No participant received a third booster dose. **G.** Convalescent individuals either after
328 the first or second infection and vaccinated individuals after the second dose. Only the serological data
329 after either the first or second infection or the second vaccination are shown in the graph as these
330 variants disappeared before the third vaccination campaign.

331 Figure S3

332 Neutralization response after different numbers of vaccination doses followed by one or two
333 infections. Neutralizing titers are represented before and after infection. The other variants are not
334 represented as they were not investigated across all visits and are therefore lacking some time points.
335 **A.** Wuhan SARS-CoV-2 wild type. **B.** Delta variant.

336 **Table S1**

337 Percentages of positive serology throughout the four study visits.

338 **Table S2**

339 Kruskal-Wallis non-parametric analyses to address if there are significant difference in neutralization
340 activities between the groups. The significance level was set with $p < 0.05$.

341 **References**

342 [1] FOPH FOoPH. COVID-19 Switzerland. 2022.

343 [2] Levin-Rector A, Firestein L, McGibbon E, Sell J, Lim S, Lee EH, et al. Reduced Odds of SARS-CoV-2
344 Reinfection after Vaccination among New York City Adults, June–August 2021. medRxiv
345 2021;2021.12.09.21267203.

346 [3] Malhotra S, Mani K, Lodha R, Bakhshi S, Mathur VP, Gupta P, et al. SARS-CoV-2 Reinfection Rate
347 and Estimated Effectiveness of the Inactivated Whole Virion Vaccine BBV152 Against Reinfection
348 Among Health Care Workers in New Delhi, India. *Jama Network Open* 2022;5.

349 [4] Hachmann NP, Miller J, Collier AY, Ventura JD, Yu J, Rowe M, et al. Neutralization Escape by SARS-
350 CoV-2 Omicron Subvariants BA.2.12.1, BA.4, and BA.5. *N Engl J Med* 2022;387:86-8.

351 [5] Caruana G, Croxatto A, Coste AT, Opota O, Lamoth F, Jatun K, et al. Diagnostic strategies for SARS-
352 CoV-2 infection and interpretation of microbiological results. *Clin Microbiol Infect* 2020;26:1178-82.

353 [6] Okba NMA, Muller MA, Li W, Wang C, GeurtsvanKessel CH, Corman VM, et al. Severe Acute
354 Respiratory Syndrome Coronavirus 2-Specific Antibody Responses in Coronavirus Disease Patients.
355 *Emerg Infect Dis* 2020;26:1478-88.

356 [7] Liu LH, Wang PF, Nair MS, Yu J, Rapp M, Wang Q, et al. Potent neutralizing antibodies against
357 multiple epitopes on SARS-CoV-2 spike. *Nature* 2020;584:450-+.

358 [8] Cerutti G, Guo YC, Zhou TQ, Gorman J, Lee M, Rapp M, et al. Potent SARS-CoV-2 neutralizing
359 antibodies directed against spike N-terminal domain target a single supersite. *Cell Host & Microbe*
360 2021;29:819-+.

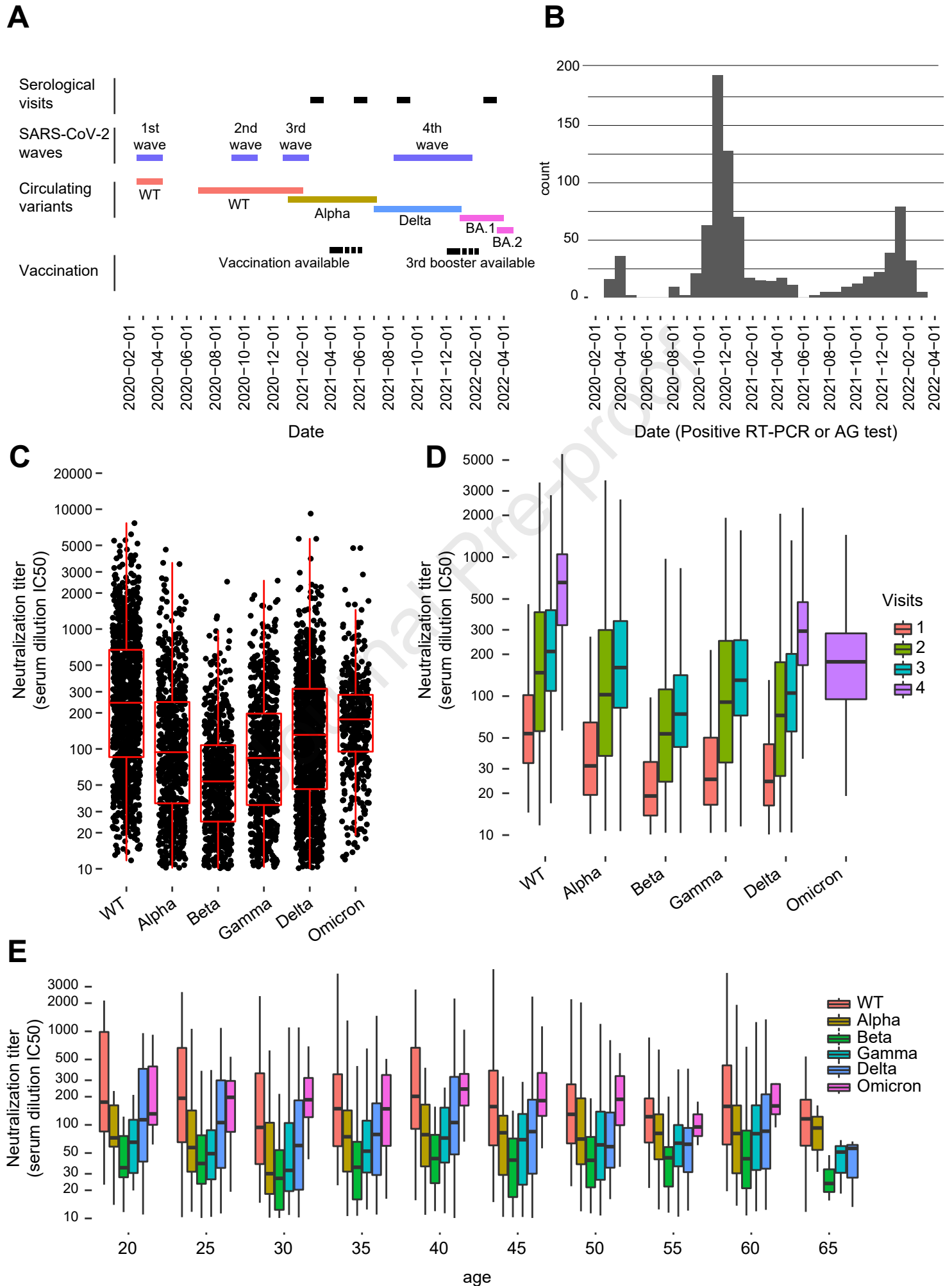
- 361 [9] Barnes CO, Jette CA, Abernathy ME, Dam KMA, Esswein SR, Gristick HB, et al. SARS-CoV-2
362 neutralizing antibody structures inform therapeutic strategies. *Nature* 2020;588:682-+.
- 363 [10] Benton DJ, Wrobel AG, Xu PQ, Roustan C, Martin SR, Rosenthal PB, et al. Receptor binding and
364 priming of the spike protein of SARS-CoV-2 for membrane fusion. *Nature* 2020;588.
- 365 [11] Pang NYL, Pang ASR, Chow VT, Wang DY. Understanding neutralising antibodies against SARS-
366 CoV-2 and their implications in clinical practice. *Military Med Res* 2021;8.
- 367 [12] Khoury DS, Cromer D, Reynaldi A, Schlub TE, Wheatley AK, Juno JA, et al. Neutralizing antibody
368 levels are highly predictive of immune protection from symptomatic SARS-CoV-2 infection. *Nat Med*
369 2021;27:1205-11.
- 370 [13] Chia WN, Zhu F, Ong SWX, Young BE, Fong SW, Le Bert N, et al. Dynamics of SARS-CoV-2
371 neutralising antibody responses and duration of immunity: a longitudinal study. *Lancet Microbe*
372 2021;2:E240-E9.
- 373 [14] Lu YY, Wang J, Li QL, Hu H, Lu JH, Chen ZL. Advances in Neutralization Assays for SARS-CoV-2.
374 *Scand J Immunol* 2021;94.
- 375 [15] Fenwick C, Turelli P, Pellaton C, Farina A, Campos J, Raclot C, et al. A multiplexed high-
376 throughput neutralization assay reveals a lack of activity against multiple variants after SARS-CoV-2
377 infection. 2021;2021.04.08.21255150.
- 378 [16] Malik JA, Ahmed S, Mir A, Shinde M, Bender O, Alshammari F, et al. The SARS-CoV-2 mutations
379 versus vaccine effectiveness: New opportunities to new challenges. *J Infect Public Health*
380 2022;15:228-40.
- 381 [17] Harvey WT, Carabelli AM, Jackson B, Gupta RK, Thomson EC, Harrison EM, et al. SARS-CoV-2
382 variants, spike mutations and immune escape. *Nat Rev Microbiol* 2021;19:409-24.
- 383 [18] Jacot D, von Rotz U, Blondet F, Aebischer O, Matthieu P, De Rham M, et al. SARS-CoV-2
384 seroprevalence in hospital healthcare workers in Western Switzerland at the end of the second
385 pandemic wave. *J Med Microbiol* 2022;71.

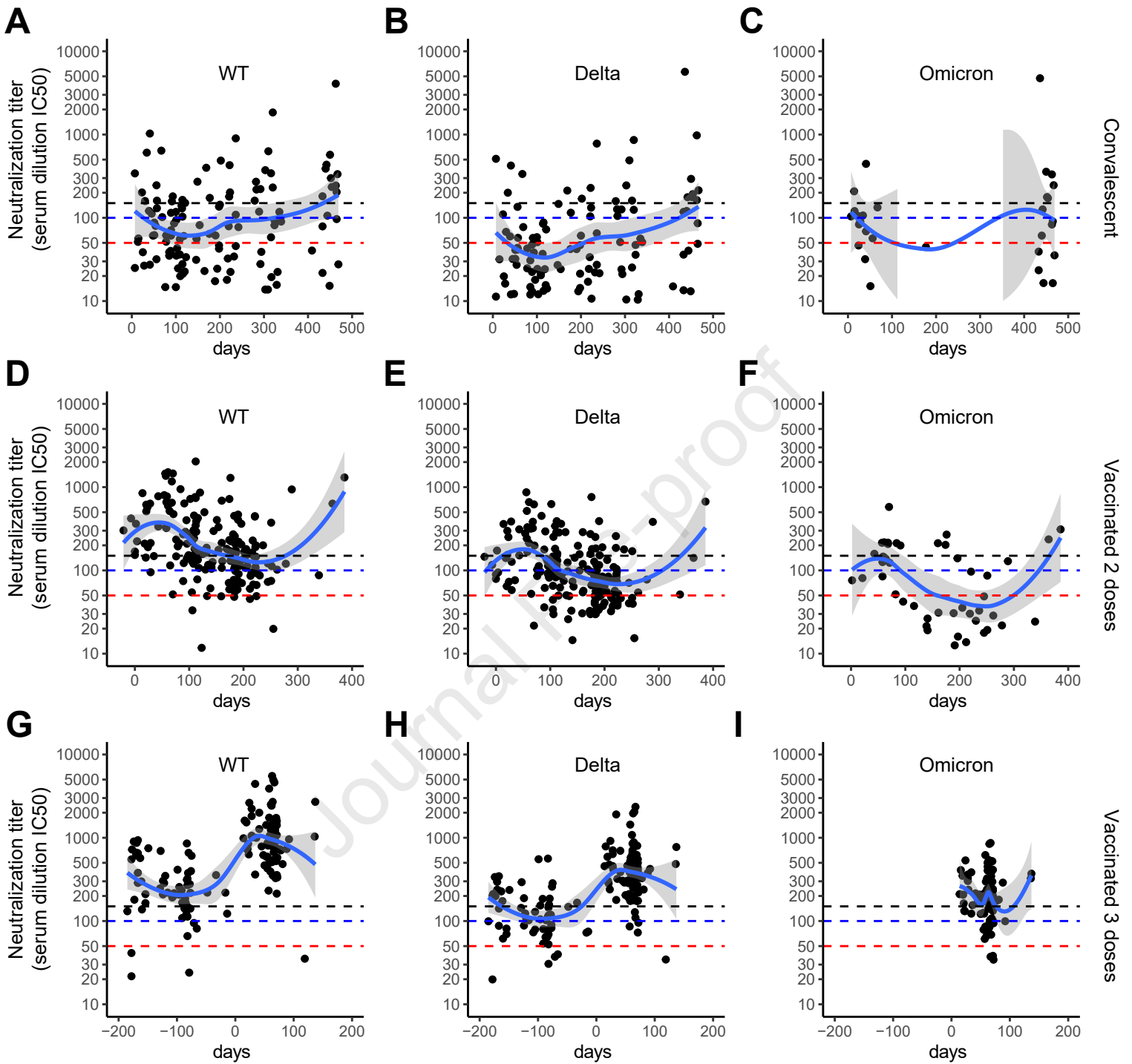
- 386 [19] Fenwick C, Croxatto A, Coste AT, Pojer F, Andre C, Pellaton C, et al. Changes in SARS-CoV-2 Spike
387 versus Nucleoprotein Antibody Responses Impact the Estimates of Infections in Population-Based
388 Seroprevalence Studies. *J Virol* 2020.
- 389 [20] Bates TA, Leier HC, Lyski ZL, McBride SK, Coulter FJ, Weinstein JB, et al. Neutralization of SARS-
390 CoV-2 variants by convalescent and BNT162b2 vaccinated serum. *Nat Commun* 2021;12:5135.
- 391 [21] Stamatatos L, Czartoski J, Wan YH, Homad LJ, Rubin V, Glantz H, et al. mRNA vaccination boosts
392 cross-variant neutralizing antibodies elicited by SARS-CoV-2 infection. *Science* 2021.
- 393 [22] Kuzmina A, Khalaila Y, Voloshin O, Keren-Naus A, Boehm-Cohen L, Raviv Y, et al. SARS-CoV-2
394 spike variants exhibit differential infectivity and neutralization resistance to convalescent or post-
395 vaccination sera. *Cell Host Microbe* 2021;29:522-8 e2.
- 396 [23] Cavanaugh AM, Spicer KB, Thoroughman D, Glick C, Winter K. Reduced Risk of Reinfection with
397 SARS-CoV-2 After COVID-19 Vaccination - Kentucky, May-June 2021. *MMWR Morb Mortal Wkly Rep*
398 2021;70:1081-3.
- 399 [24] Fenwick C, Turelli P, Pellaton C, Farina A, Campos J, Raclot C, et al. A high-throughput cell- and
400 virus-free assay shows reduced neutralization of SARS-CoV-2 variants by COVID-19 convalescent
401 plasma. *Science Translational Medicine* 2021;13.
- 402 [25] Shrock E, Fujimura E, Kula T, Timms RT, Lee IH, Leng Y, et al. Viral epitope profiling of COVID-19
403 patients reveals cross-reactivity and correlates of severity. *Science* 2020;370.
- 404 [26] Garcia-Beltran WF, Lam EC, Astudillo MG, Yang DN, Miller TE, Feldman J, et al. COVID-19-
405 neutralizing antibodies predict disease severity and survival. *Cell* 2021;184:476-+.
- 406 [27] Evans JP, Zeng C, Carlin C, Lozanski G, Saif LJ, Oltz EM, et al. Neutralizing antibody responses
407 elicited by SARS-CoV-2 mRNA vaccination wane over time and are boosted by breakthrough
408 infection. *Sci Transl Med* 2022;14:eabn8057.
- 409 [28] Bates TA, McBride SK, Leier HC, Guzman G, Lyski ZL, Schoen D, et al. Vaccination before or after
410 SARS-CoV-2 infection leads to robust humoral response and antibodies that effectively neutralize
411 variants. *Sci Immunol* 2022;eabn8014.

- 412 [29] Walls AC, Sprouse KR, Bowen JE, Joshi A, Franko N, Navarro MJ, et al. SARS-CoV-2 breakthrough
413 infections elicit potent, broad, and durable neutralizing antibody responses. *Cell* 2022;185:872-80 e3.
- 414 [30] Crotty S. Hybrid immunity. *Science* 2021;372:1392-3.
- 415 [31] Qu P, Faraone JN, Evans JP, Zheng YM, Yu L, Ma Q, et al. Durability of Booster mRNA Vaccine
416 against SARS-CoV-2 BA.2.12.1, BA.4, and BA.5 Subvariants. *N Engl J Med* 2022.
- 417 [32] Bates TA, Leier HC, Lyski ZL, Goodman JR, Curlin ME, Messer WB, et al. Age-Dependent
418 Neutralization of SARS-CoV-2 and P.1 Variant by Vaccine Immune Serum Samples. *Jama-J Am Med*
419 *Assoc* 2021;326:868-+.
- 420 [33] Yang HS, Costa V, Racine-Brzostek SE, Acker KP, Yee J, Chen ZM, et al. Association of Age With
421 SARS-CoV-2 Antibody Response. *Jama Network Open* 2021;4.
- 422 [34] Lyke KE, Atmar RL, Islas CD, Posavad CM, Szydlo D, Paul Chourdury R, et al. Rapid decline in
423 vaccine-boosted neutralizing antibodies against SARS-CoV-2 Omicron variant. *Cell Rep Med*
424 2022;3:100679.
- 425

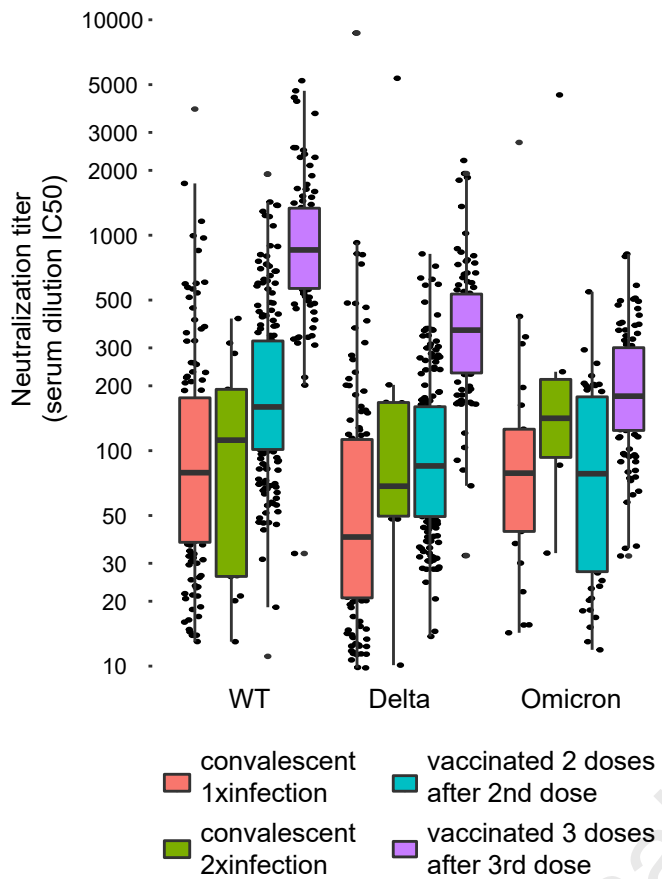
Table 1

Immune status	One reported positive test	Two reported positive tests	No reported positive RT-PCR or AG tests	Total
Convalescent	89/139	6/15	96/118	
Vaccination 1 dose	30/75	10/22	57/75	
Vaccination 2 doses	68/172	17/48	183/296	
Vaccination 3 doses	82/227	6/19	129/255	
Total	269	39	465	773/1461

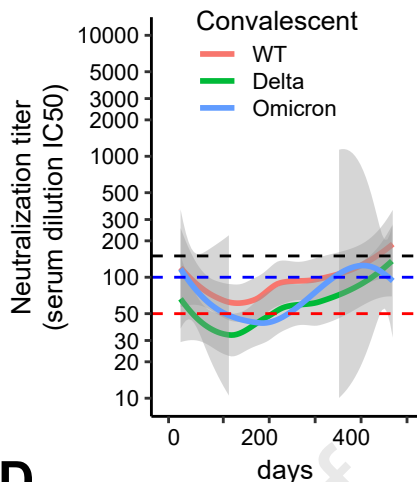




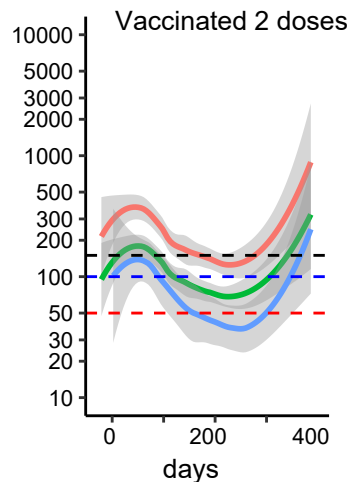
A



B



C



D

