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2 Antibody Correlates of Protection for COVID-19 Convalescent Plasma Associated with Reduced
3 Outpatient Hospitalizations
4

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127 ABSTRACT

128 SARS-CoV-2 antibody levels associated with reduced hospitalization risk remain undefined. Our
129 outpatient COVID-19 convalescent plasma (CCP), placebo-controlled trial observed SARS-
130 CoV-2 antibody levels decreasing 22-fold from matched donor units into post-transfusion
131 seronegative recipients. Unvaccinated recipients were jointly stratified by a) early or late
132 transfusion (≤ 5 or >5 days from symptom onset) and b) high or low post-transfusion SARS-
133 CoV-2 antibody levels ($<$ or \geq geometric mean). Early treatment with high post-transfusion
134 antibody levels reduced hospitalization risk-0/102 (0%) compared to all other CCP recipients-
135 17/370 (4.6%; Fisher exact $p=0.03$) and to all control plasma recipients-35/461 (7.6%; Fisher
136 exact $p=0.001$). A similar donor upper/lower antibody level and early late transfusion stratified
137 analyses indicated significant hospital risk reduction. Pre-transfusion nasal viral loads were
138 similar in CCP and control recipients regardless of hospitalization outcome. Therapeutic CCP
139 should comprise the upper 30% of donor antibody levels to provide effective outpatient use for
140 immunocompromised and immunocompetent outpatients.

141 INTRODUCTION

142 The threshold of antibody correlating with reduced severe disease progression varies and needs
143 to be independently determined for diverse infectious diseases, from protozoans, like
144 *Plasmodium falciparum*^{1,2} to viruses, like SARS-CoV-2^{3,4}. Correlates of protection for antibody
145 levels demarcating vaccine efficacy for COVID-19 hospitalization risk reduction have been
146 represented in relation to COVID-19 convalescent plasma (CCP) equivalents⁵⁻⁷. Likewise, a
147 model⁸ utilizing vaccine, monoclonal antibody, and CCP data from clinical trials was created
148 based on the reference CCP equivalents for 50% hospitalization risk reduction⁹⁻¹¹. Monoclonal
149 antibody outpatient randomized controlled trials (RCTs) demonstrate 50%-80% efficacy in
150 hospital risk reduction¹²⁻¹⁵. However, the same monoclonal antibody doses failed to show disease
151 progression reduction in COVID-19 inpatients^{14,16-18}. Like the monoclonal antibodies,
152 intravenous remdesivir demonstrated greater efficacy in outpatients versus inpatients^{19,20}. In
153 summary, both early timing and dose of antiviral antibodies and drugs matter for outpatient
154 reduction of hospitalization risk.

155
156 Convalescent persons recovering from pre-Alpha period COVID-19 have varying antibody
157 levels spanning a 3-log₁₀ range from undetectable antibody to positive after a 3 to 5 thousand
158 dilution²¹. Individuals with both boosted vaccines and recent COVID-19 (i.e., hybrid immunity)
159 have considerably higher antibody levels (i.e., detectable even over 50,000 inverse dilutional
160 geometric mean titers) than either individuals with antibodies from infection or vaccination alone
161 ^{22,23}. Our CCP outpatient treatment within 9 days of symptom onset, showed greater than 50%
162 hospitalization relative risk reduction, which increased to more than 80% risk reduction (point
163 estimate) when given within 5 days of symptom onset²⁴. Hospitalized participants had the same
164 donor unit range of SARS-CoV-2 antibody levels as those not hospitalized.

165
166 A prespecified analysis from the parent outpatient CCP RCT aimed to compare antibody levels
167 in donor-recipient pairs to define dilutional decrease and to correlate both donor and recipient
168 antibody levels to disease progression culminating in hospitalization. In this substudy, we
169 analyzed the risk of hospitalization among unvaccinated COVID-19 outpatients, comparing
170 control plasma recipients to four different CCP groups jointly stratified by early or late treatment

171 (i.e., ≤ 5 or >5 days from symptom onset), and by high or low post-transfusion recipient or donor
172 antibody levels (i.e., $<$ or \geq geometric mean, GM).

173 RESULTS

174 Trial population

175 At 23 sites throughout the USA, the previously reported outpatient CCP clinical trial (CSSC-004,
176 NCT04373460) transfused 1181 COVID-19 outpatients over 16 months (June 3, 2020 to October
177 1, 2021), to show that receipt of CCP, compared to control plasma, reduced the risk of
178 hospitalization for COVID-19 by 54%²⁴. Written and signed informed consent was obtained
179 from all participants. Hospitalizations occurred in 53 of 964 unvaccinated and 1 of 58 partially
180 vaccinated participants. None of the 159 fully vaccinated individuals were hospitalized²⁴. We
181 focused our correlation of antibody level analysis and hospitalization on those unvaccinated with
182 evaluable antibody data, for which 479 received control plasma and 472 CCP (Table 1). The
183 unvaccinated participant's mean age was 44, with more females than males, approximately 40%
184 obese with BMI 30 or over, and approximately 40% with one or more pre-existing comorbidities
185 for severe COVID-19 risk. Unvaccinated CCP recipients, both screen seropositive or
186 seronegative were further jointly stratified by early (≤ 5 days of symptom onset) or late (>5 days
187 of symptoms onset) time to transfusion as well as post-transfusion antibody levels (measured
188 within 30 minutes after transfusion completion) in the upper or lower half of screened
189 seronegative CCP recipients (Table 1).

190

191 Donor CCP and recipient screen and post-transfusion antibody levels in unvaccinated 192 participants

193 Approximately 40% of all potential CCP study donors were excluded for low antibody levels,
194 thus transfusion units represented the upper 60% of all convalescent plasma donors. Because of
195 the greater correlation with virus neutralization antibody (nAb), we measured the plasma anti-
196 Spike-Receptor Binding Domain (S-RBD) IgG antibodies levels by dilutional titer and the more
197 precise area under the curve (AUC) on both the donors at collection and in over 5,000 recipient
198 samples at pre-transfusion screen, post-transfusion and follow-up visits²¹. The anti-S-RBD IgG
199 titer threshold for seronegative was 180 titer or below. The donor anti-S-RBD IgG GM titer of
200 6,678 and anti-S-RBD IgG AUC of 3,172 separated donor antibody levels' upper and lower half
201 (Fig. 1A). The donor control plasma was either collected in 2019 (211 units) or blood bank tested
202 SARS-CoV-2 seronegative when collected in 2020-2021.

203

204 To experimentally determine the dilution factor associated with the single ~ 200 mL CCP
205 administration in 338 unvaccinated seronegative recipients, measurements of matched donor and
206 recipient anti-S-RBD IgG AUC were performed. The matched transfusion donor antibody levels
207 proportionately decreased 22-fold (approximately 5%) from the GM donor anti-S-RBD IgG
208 3286 AUC to recipient post-transfusion antibody levels at GM 147 AUC measured within 30
209 minutes of transfusion completion (Fig. 1B). The decrease was parallel such that matched donor
210 antibody levels below 1000 AUC were also consistently below 150 AUC post-transfusion in
211 recipients. Similarly, 15 seronegative hospitalized CCP recipients had post-transfusion antibody
212 levels 19 times lower than their matched donors (Fig. 1C).

213

214 The unvaccinated study participants included a screen seropositive participant subset (199/951
215 (21%) (107 in the CCP and 92 in the control plasma groups), with 83% of the 199 seropositive

216 participants already having pre-transfusion antibody levels above the GM 150 AUC for
217 seronegative post-transfusion CCP recipients (Fig. 1D). The pre-transfusion seropositive
218 transfusion antibody levels were near 500 to 1000 GM anti-S-RBD IgG AUC which is 3 to 6
219 times lower than the donor anti-S-RBD IgG antibody levels of 3172 AUC. The low post-
220 transfusion recipient antibodies (early or late) were predictably lower in frequency and GM AUC
221 levels. Segregating the unvaccinated seropositive recipients by days from symptom onset to
222 transfusion revealed a trend of a higher frequency of participants being seropositive after day 3,
223 with higher GM antibody levels approximating 1000 anti-S-RBD IgG AUC, which was
224 associated with the early development of virus-specific immunity also observed by Wolfel²⁵
225 (Fig. 1E).

226

227 Recipient antibody level post-transfusion benchmarks of protection that prevents 228 hospitalization

229 To correlate post-transfusion antibody levels among unvaccinated participants (both screen
230 seronegative and seropositive) to hospital outcome, we stratified the post-transfusion anti-S-RBD
231 IgG AUC by early or late treatment and high or low antibody levels for the CCP group (Fig. 2,
232 Extended Table 1). The recipient antibody geometric mean of 150 AUC delineated high versus
233 low antibody levels. Among the CCP participants treated early (symptom onset within 5 days)
234 with high post-transfusion recipient antibody levels, there were no hospitalizations, 0/102
235 participants. In those unvaccinated CCP recipients treated early with low post-transfusion
236 antibody levels (anti-S-RBD IgG AUC below 150 AUC), we observed 5/104 (5%) participant
237 hospitalizations (Fig. 2). There were 12 total hospitalizations in the two groups receiving
238 transfusions late—6/164 (3.7%) recipients with high antibody levels and 6/102 (6%) recipients
239 with antibodies below anti-S-RBD IgG 150 AUC. We compared the proportion of
240 hospitalizations of the early treatment with high post-transfusion antibody level CCP group with
241 that of the other three CCP groups (17/270, 4.6%; Fishers exact p=0.03), the control plasma
242 group (36/479, 7.5%; Fishers exact p=0.001), and the early control plasma group (24/207,
243 11.6%; Fishers exact p=0.00005; Extended Table 1). Similarly, delineating high and low post-
244 transfusion recipient antibody levels above the geometric mean of 540 anti-S-RBD IgG titer,
245 although less precise than AUC, revealed that participants receiving early treatment with high
246 titer CCP also had no hospitalizations (0/45, 0%; Extended Fig. 1 and Table 1).

247

248 Donor antibody level pre-transfusion benchmarks of protection that prevent 249 hospitalization

250 To correlate matched CCP donor units in the unvaccinated recipients to hospital outcome, we
251 next investigated if the corresponding donor antibody levels would also translate into protection
252 from hospitalization. The unvaccinated CCP recipients were again delineated by symptom onset
253 to transfusion, but instead of post-transfusion antibody level representation, the matched
254 corresponding donor antibody levels were stratified by donor anti-S-RBD IgG antibodies at GM
255 3286 AUC for all 592 recipients (Fig. 3A) and virus neutralization antibody (nAb) levels ≥ 60
256 AUC (Fig. 3B). In all antibody level metrics, we compared the GM upper to lower as high and
257 low antibody levels, respectively. The numbers remained statistically significant by the Fishers
258 exact test compared to both control plasma groups (Fig. 3 and Extended Table 1). Similar to the
259 analyses with AUC values, donor anti-S-RBD IgG and nAb GM titer values, although less
260 precise, revealed their respective recipients with early treatment and high donor antibody titers
261 having the lowest proportion of hospitalizations (Extended Fig. 2 and Table 2). We also stratified

262 donor antibody levels by the commercial assay EUROIMMUN ratio of optical density expressed
263 above or below 6 AU or 600 BAU/mL (Extended Fig 3 and Extended Table 1), a level that also
264 demarcates the upper portion of study donors²⁴.

265

266 **Screen seropositive hospitalizations in the unvaccinated**

267 The screen seropositive recipients with subsequent hospitalizations were investigated, despite
268 low incidence. 7/54 unvaccinated participants, subsequently hospitalized, were seropositive at
269 the initial screen, two in the CCP group and five in the control group (Extended Table 2). Three
270 of the five hospitalized participants from the control group had screen anti-S-RBD IgG antibody
271 titers at 540 and AUC less than 150, fitting the early/low seronegative data described above. In
272 the unvaccinated CCP group, 2/107 (1.9%) who were screen seropositive were hospitalized
273 versus 15/378 (4%) seronegative at screen (Chi square $p=0.29$; CCP seropositive versus
274 seronegative). Among the control group of whom were hospitalized, there were 5/92 (5.4%)
275 screen seropositive versus 29/381 (7.6%) screen seronegative (chi square $p=0.47$ control-
276 seropositive to seronegative). Of those screened seropositive, there was no statistical difference
277 in hospitalization between those who received CCP and control plasma (chi square $p=0.17$; CCP-
278 seropositive to control-seropositive). The single screen seronegative and partially vaccinated
279 hospitalized participant transfused with control plasma received their first vaccine dose the same
280 day as symptom onset. Among the 159 fully vaccinated and not hospitalized, 158 (99%) were
281 screen seropositive, while 30/58 (69%) partially vaccinated participants were seropositive.

282

283 **Pre-transfusion nasal SARS-CoV-2 viral RNA loads**

284 Nasal viral load might independently determine risk of hospitalization. Quantitative RT-PCR
285 was performed on the screen nasal swabs collected before plasma transfusion to correlate the
286 SARS-CoV-2 nasal viral RNA load to hospitalization. All unvaccinated individuals subsequently
287 receiving either control plasma or CCP had indistinguishable screen viral loads (i.e., near 10,000
288 copies) (Fig. 4A). Analyzing nasal viral loads of early/high, early/low, late/high, and late/low
289 recipient groups revealed, as expected, that early transfusions closer to symptom onset were
290 associated with higher viral loads than late transfusions. The CCP early/high and early/low
291 recipient pre-transfusion nasal viral loads were indistinguishable. The viral load was similar
292 among unvaccinated at screen, regardless of subsequent hospitalization outcome (Fig. 4B).
293 While the inclusion criteria required a documented positive molecular SARS-CoV-2 test (87%
294 by RNA detection and 13% by antigen detection), the interval between the study inclusion
295 SARS-CoV-2 test (performed outside the study) and the research study nasal swab collection
296 prior to transfusion may have been up to a week.

297

298 Stratifying nasal viral load by days from symptom onset to transfusion time among unvaccinated
299 participants infected prior to the Delta variant, which has different viral load kinetics^{26,27},
300 indicated a decline in pre-transfusion viral burden among seronegative individuals after 5 days
301 from symptom onset (Fig. 4C). Seropositive individuals had lower viral loads compared to
302 seronegative individuals at all timepoints, with a majority of seropositive viral loads below the
303 limit of detection (330 copies) by day 4 post-symptom onset (Fig. 4D). The screen viral RNA
304 copy data suggest that seropositivity at the time of transfusion correlated with lower viral loads,
305 but differences in viral RNA load did not impact hospitalization outcome.

306

307 Differences in antibody levels weeks after transfusion

308 Passive transfer of SARS-CoV-2 specific antibodies was postulated to possibly down-modulate
309 subsequent SARS-CoV-2 antibody responses. This clinical study was designed to measure
310 antibody responses up to 90 days post-transfusion to compare antibody kinetics between CCP
311 and control plasma recipients. There were no differences in antibody levels between CCP and
312 control plasma recipients at or beyond 14 days post-transfusion (Fig. 5). Longitudinal differences
313 in antibody levels observed were due to hospitalization rather than treatment. The multivariate
314 linear mixed-effects regression, adjusted for variant, age, sex, and BMI, showed no differences in
315 antibody levels between CCP and control plasma recipients beyond 14 days post-transfusion
316 (Fig. 5). There were neither sex, age, BMI, nor comorbidity differences in antibody levels
317 between CCP and control groups.

318
319 Within the control group, there were eight immunocompetent recipients with undetectable
320 SARS-CoV-2 antibody levels during all follow-up visits. Three of the 8 had persistent COVID-
321 19 symptoms after 90 days of study follow-up. All CCP recipients had detectable antibodies at or
322 after the day 14 visit.

323 324 Variant period differences in screen seropositivity, nasal viral load and day 90 antibody levels

325 We investigated how SARS-CoV-2 variants influenced seropositivity, nasal viral load and 3-
326 month antibody levels in the setting of ongoing vaccination after December 2020. We separated
327 participants into three periods: pre-Alpha (June 3, 2020 to January 31, 2021), Alpha (February 1,
328 2021 to July 15 2021), and Delta (July 16, 2021 to October 1, 2021). Participants recruited in the
329 Delta period were younger, with fewer medical conditions, and more than 60% fully vaccinated
330 (Extended Table 3). Screen seropositivity rates among unvaccinated individuals were low (~
331 20%), during pre-Alpha and Alpha periods rising to 46% during the Delta period. The baseline
332 screen anti-S-RBD AUC GM antibody levels in the unvaccinated seropositive participants were
333 529, 482, and 7042 anti-S-RBD AUC during the pre-Alpha, Alpha and Delta periods,
334 respectively compared to the 3286 AUC in the mainly pre-Alpha study donors (Extended Fig.
335 4A).

336
337 Pre-transfusion nasal viral loads were similar between CCP and control during the pre-Alpha,
338 Alpha, and Delta periods (Extended Fig. 4B). When unvaccinated participants were stratified
339 based on seropositivity at the time of transfusion, viral loads were consistently lower in
340 seropositive individuals compared to seronegative individuals regardless of the time period in
341 which CCP was administered (Extended Fig. 4C). During the Delta period, unvaccinated
342 seropositive individuals also had lower viral loads compared to seronegative individuals, but the
343 participant numbers stratified by symptom onset day were low (Extended Fig. 5A, B). Fully
344 vaccinated and seropositive participants at screen also showed a drop in viral load when
345 transfusion occurred more than 5 days after symptom onset (Extended Fig. 5C).

346
347 Comparing CCP and control day 90 antibody levels in those unvaccinated at screen (excluding
348 the 165 vaccinated during the follow-up visits), indicated no difference between treatment
349 groups or groups stratified by early or high CCP antibody treatment (Extended Figure 6A).
350 Those unvaccinated participants, subsequently hospitalized (41/54, 76%) during the pre-Alpha
351 period, had a GM of 13007 AUC with very few having antibody levels below the mean for
352 unvaccinated participants in the pre-Alpha period at day 90 compared to the non-hospitalized

353 unvaccinated pre-Alpha participants GM 2691 AUC antibody level, 5 times less than those
354 hospitalized (Extended Fig. 6B). Unvaccinated recipients during the Alpha period had similar
355 day 90 antibody levels (geometric mean AUC=6683) to those infected during the Delta period
356 (geometric mean AUC=5929), but near double the pre-Alpha period participants. Fully
357 vaccinated recipients at screen who had breakthrough infection during Delta period showed a
358 GM AUC of 53813, which was 20 times greater than the GM AUC of day 90 pre-Alpha period
359 recipients.

360 Discussion

361 This jointly stratified study analysis by both time to transfusion and antibody levels found that
362 post-transfusion anti-S-RBD IgG AUC levels at or over 150 AUC and transfusion within 5 days
363 of symptom onset, resulted in no hospitalization in unvaccinated recipients. In seronegative
364 recipients, the post-transfusion upper 50% antibody levels were proportionately matched to
365 upper 50% donor antibody levels measurements by three different test methodologies-anti-S-
366 RBD IgG ELISA, direct live virus neutralization, and EUROIMMUN anti-SARS-CoV-2 ELISA.
367 Given that we selected the upper 60% of donors for study qualification and by all antibody
368 measurements for stratified analysis, the upper half of qualified donors which equals the upper
369 30%, had similar impact on hospitalization risk reduction. The four-quadrant graphic depiction
370 of hospitalizations was similar in the recipient antibody levels compared to the 3 separate donor
371 antibody level measurements. This implicates the upper 30% of all potential pre-Alpha CCP
372 donors as effective for reducing hospitalizations prior to the Delta variant.

373
374 An individual's circulating plasma volume is, on average, 3 liters, which predicts a 15-fold
375 proportional antibody dilution for a 200 mL CCP transfusion. A prior infection prevention study
376 transfused CCP into healthy nonSARS-CoV-2 infected individuals and similarly measured 20-
377 fold SARS-CoV-2 antibody dilution²⁸. This suggests that the antibody compartment volume is
378 similar between non-infected individuals and those with COVID-19 during the first week of
379 illness. We did not observe a measurably higher volume of distribution, suggesting that
380 extravascular transfer of high levels of antibodies to lower the post-transfusion antibody levels
381 occurred. The donor and recipient antibody levels are directly proportional in Fig. 1B-C where
382 the lines are parallel with the 22-fold dilution. From the recipient post-transfusion levels, one can
383 predict matched donor levels are proportionately greater by a factor of 20. The screen
384 seronegative participants had a clean negative antibody background. Those who were screen
385 seropositive in Fig. 1D already had greater antibody levels than more than half of the post-
386 transfusion seronegative participants. We have not explored whether CCP has better neutralizing
387 abilities than the early host immune response when antibodies are just starting to measurably
388 increase after 4 days from symptom onset for screen seropositive participants (low antibodies in
389 the 500 to 1000 range) (Fig. 1E).

390
391 Antivirals (antibodies or small molecule antivirals) administered early in the COVID-19 course
392 protect from progression to severe disease requiring hospitalization. At least three factors interact
393 to predict risk of outpatient hospitalization: 1) viral load and variant properties; 2) human host
394 risk factors for severe disease (age, obesity, or medical comorbidities), and 3) quantity and
395 quality (degree of match to variants) of SARS-CoV-2 specific antibodies at disease onset, due to
396 active vaccination or immune status (i.e., hybrid immunity, breakthrough, etc.).

397

398 We found no differences in nasal viral loads between control and CCP recipients at screening to
399 account for differences in hospitalization rates. Importantly, those participants who were
400 subsequently hospitalized had similar viral loads at the time of transfusion when compared to
401 those not hospitalized. Unvaccinated recipients who were seropositive at screening had lower
402 viral loads than seronegative participants. We also observed lower viral loads in those screened
403 late. Specifically, those who were screened and transfused late (≥ 5 days after symptom onset)
404 had lower viral loads compared to those who were transfused early, suggesting that the
405 administration of antiviral drugs or CCP early (i.e., during the period of higher viral loads
406 measured by nasal sampling) maximizes the therapeutic effect. Viral load dynamics in the lung
407 or other tissues may differ, and treatment timing should be explored further.

408
409 The jointly stratified subgroup analysis was matched for patients' age, demographics, obesity
410 and one or more medical conditions for severe COVID-19 progression to hospitalization. Our
411 study population was also younger, less obese and with fewer risk factors for disease progression
412 than other published RCTs of CCP outpatient use²⁹. Our study population also had a higher
413 prevalence of seronegative participants upon transfusion than the Emergency Department based
414 study (C3PO) where the seropositive rate was near 50%, which may be indicative of a different
415 patient population³⁰.

416
417 In this study, there are COVID-19 participants with a 100-fold range of antibody levels. CCP
418 units from the top 30% of all convalescent donors are necessary to confer protection against
419 hospitalization with severe COVID-19, such that when antibody levels are diluted approximately
420 20-fold, the recipient antibody levels remain above the CCP GM. These data suggest that
421 convalescent plasma has been underdosed in prior infectious disease outbreaks and that future
422 pandemic patients should receive higher therapeutic donor antibody levels. The current study
423 introduces the concept that therapeutic donor CCP levels must be in the upper half or higher to
424 fully account for the dilution effect in the recipient. Plasma from approximately 330 unique CCP
425 donations were transfused into 592 CCP participants. These donor units were previously
426 characterized for full-length anti-Spike IgG with GM titers of 13,053 and more precise AUC GM
427 of 7938 which equals 243 BAU/mL using the international standards³¹. The median nAb titer
428 was 80 with GM titer of 58 and nAb AUC of 51, equaling GM 27 IU/mL. The commercial
429 EUROIMMUN arbitrary units (AU) mean was 6 for the unique units. The donor anti-S-RBD
430 IgG GM titer was 6,678 and anti-S-RBD IgG GM AUC of 3,172. These thresholds separate the
431 upper and lower portion of donor antibody levels and was also approximately 50% of total full
432 length Spike titer antibody levels, but 100 times the nAb GM titer levels.

433
434 Unvaccinated participants who received CCP after 5 days (late) or received units with low
435 antibody levels still had near significant reduction in hospital risk of about 4% rather than 7.6%
436 in all controls. The parent study included fully vaccinated and partially vaccinated participants in
437 the analyses, which lowered risk of hospitalization to 6.3%. Here, we observed an 11.6% risk of
438 hospitalization among unvaccinated controls transfused early within 5 days of symptom onset.
439 The effect of early CCP transfusion and high post-transfusion antibody is even greater among
440 this subgroup of unvaccinated recipients.

441
442 The antibody level that affords protection is not absolute, as not even monoclonal antibody or
443 small molecule antiviral therapy affords 100% reduction in risk of hospitalization. There were

444 two screen seropositive participants in the CCP group and two in the control group with antibody
445 levels also above the anti-S-RBD IgG antibody threshold of inverse dilution titer over 540 and
446 AUC over 150, presumptively with newly formed antibodies. The finding that early
447 administration of antibodies to SARS-CoV-2 is beneficial in reducing progression of disease is
448 consistent with the observation that those who mount early antibody responses have lower rates
449 of death after hospitalization^{32,33} and that inducement of humoral immunity by vaccination
450 reduces severity of disease and death³⁴.

451
452 When CCP was first deployed in 2020, there were concerns that specific antibody administration
453 to individuals in the early stages of COVID-19 could interfere with the development of
454 endogenous immune responses³⁵. However, our findings show that transfusion of CCP, as
455 compared to control plasma, was not associated with differences in the development of a
456 humoral immune response in recipients, reassuring for the immunological safety of CCP in
457 humans. The C3PO convalescent plasma study also did not see an antibody level difference
458 between CCP and saline infusions^{30,36}.

459
460 While our study had predominately SARS-CoV-2 naïve recipients enrolled prior to the Omicron
461 variant who were largely unvaccinated, the findings are applicable to immunocompromised
462 patients today who lack SARS-CoV-2 antibodies. Another limitation is the low number of
463 seronegative participants transfused within 5 days of symptom onset with post-transfusion above
464 the geometric mean donor antibody levels in our study population (approximately 100
465 participants). The parent study was not powered to look at these stratified quadrants.

466
467 In summary, our results support and reconfirm the adage that for antibody therapy to be
468 effective, sufficient amounts of pathogen specific antibody should be dosed early³⁷. No
469 hospitalizations were observed in those treated at ≤ 5 days of symptom onset with high titer CCP
470 indicating that this is the optimal combination for effective CCP use. Early treatment alone is
471 insufficient as hospitalizations were still observed in the group treated ≤ 5 days with lower titer
472 units, necessitating both early treatment and adequate antibody dosing for optimal efficacy. Our
473 results provide evidence for the best use of CCP. We advocate that CCP units used for therapy
474 comprise the upper 30% of donor antibody levels. These levels should set the threshold for future
475 therapeutic CCP. When humanity faces its next pandemic, there is a high likelihood that
476 convalescent plasma will be used again until better specific therapies become available. Our data
477 provide a roadmap for optimal early, high dose convalescent plasma deployment in such future
478 emergencies.

479

480 Online Methods

481 Study Ethics

482 Johns Hopkins served as the single-IRB (sIRB). For the Center for American Indian Health sites,
483 the protocol was also independently reviewed and approved by the Navajo Nation Health Human
484 Research Review Board and the National Indian Health Service IRB. The protocol was also
485 approved by the Department of Defense (DoD) Human Research Protection Office (HRPO).
486 An independent medical monitor who was unaware of the trial group assignments reviewed all
487 serious adverse events, and an independent panel of three physicians who were unaware of the
488 trial-group assignments adjudicated Covid 19 related hospitalizations and severity. An

489 independent data and safety monitoring board provided interim safety and efficacy reviews. The
490 trial was conducted in accordance with the principles of the Declaration of Helsinki, the Good
491 Clinical Practice guidelines of the International Council for Harmonisation, and all applicable
492 regulatory requirements.

493

494 Study Population

495 In this multicenter, double-blind, randomized, controlled trial, we evaluated the efficacy and
496 safety of COVID-19 convalescent plasma, as compared with control plasma, in symptomatic
497 adults (≥ 18 years of age) who had tested positive for severe acute respiratory syndrome
498 coronavirus 2, regardless of their risk factors for disease progression or vaccination status.
499 Participants were enrolled within 8 days after symptom onset and received a transfusion within 1
500 day after randomization. The primary study outcome (reported previously) was COVID-19–
501 related hospitalization within 28 days after transfusion. There were no obvious imbalances
502 between the trial groups in the parent trial with respect to baseline characteristics, including
503 coexisting conditions, COVID-19 vaccination status, vital signs, and clinical laboratory results.

504

505 Study Center(s):

506 Anne Arundel Medical Center; Ascada Research; Baylor College of Medicine; Johns Hopkins
507 Center for American Indian Health; Johns Hopkins Bloomberg School of Public Health; Johns
508 Hopkins University; Lifespan/Brown University Rhode Island Hospital; Mayo Clinic, Phoenix;
509 MedStar Washington Hospital Center; NorthShore University Health System; The Bliss Group;
510 The Next Practice Group; University of California, Los Angeles Health; University of Alabama
511 at Birmingham; University of California, Irvine Health; University of California, San Diego;
512 University of Cincinnati Medical Center; University of Massachusetts Worcester; University of
513 Miami; University of New Mexico; University of Rochester; University of Texas Health Science
514 Center at Houston; University of Utah Health; Vassar Brothers Medical Center; Wayne State
515 University; Western Connecticut Health Network, Danbury Hospital; Western Connecticut
516 Health Network, Norwalk Hospital.

517

518 Study Plasma

519 The study qualified donor plasma with SARS-CoV-2 positive antibodies after a 1:320 dilution
520 under FDA IND 19725 protocol. After July 2021, the transfused plasma donor units met the
521 existing FDA Emergency Use Authorization (EUA) criteria for high titer at EUROIMMUN
522 arbitrary unit (AU) over 3.5. Many identical apheresis donor plasma units were transfused into 2,
523 3, or 4 separate recipients. Plasma from 333 unique CCP donations was transfused into the 592
524 CCP participants. Seventy-five percent of the donor collections were before September 2020
525 with more than 90% by January 2021 and the last 25 collections by March 2021. These donor
526 units were previously characterized for full-length anti-Spike IgG geometric mean (GM) titers of
527 13,053, which corresponded with a more precise area under the curve (AUC) geometric mean of
528 7938, equaling 243 BAU/mL using the international standards²⁴. The median neutralizing
529 antibody (nAb) titer was 80, with a geometric mean titer of 58, and nAb AUC of 51, equaling
530 GM 27 IU/mL²⁴. The commercial EUROIMMUN arbitrary units (AU) mean was 6 for the
531 unique donor units²⁴.

532

533 Study visits and time periods

534 In these studies, antibody levels were measured at screen before transfusion, within 30 minutes
535 of transfusion, and various timepoints up to 90 days post-transfusion. Participants were
536 transfused during pre-Alpha (June 3, 2020 to January 31, 2021), Alpha (February 1, 2021 to July
537 14, 2021), and Delta (July 15 to October 1, 2021) variant periods. There were just three
538 participants transfused from July 2 to July 9, 2021 which decreased the number of false
539 designations. The first Alpha (B1.1.7) confirmed by sequencing was from a participant
540 transfused February 18, 2021.

541

542 **EUROIMMUN ELISA Assay**

543 The EUROIMMUN anti-SARS-CoV-2 ELISA for IgG (cat. EI2606-9601G) was validated in a
544 Clinical Laboratory Improvement Amendments-certified (CLIA-certified) laboratory for donor
545 qualification as positive after a 1:320 dilution as per IND 19725 protocol. The assay was
546 performed according to the manufacturer's specifications. In a separate research laboratory, the
547 donor optical density (OD) was measured at 1:101 dilution. The ratio calculated by dividing the
548 sample OD by the OD of the calibrator from that run constitutes the AU. To measure anti-SARS-
549 CoV-2 IgG binding, each plate had the following components: 100 μ L plasma (1:101 dilution,
550 per the manufacturer's protocol) and 100 μ L undiluted positive, negative, and calibrator controls.
551 Plates were washed 3 times, followed by the manufacturer's protocol for addition of conjugate
552 and substrate. Ratios of 0.8 or higher were considered positive.

553

554 **Indirect ELISA**

555 The ELISA protocol was adapted from a protocol published by the Florian Krammer
556 laboratory³⁸. The 96-well plates (Immulon 4HBX, Thermo Fisher Scientific-Cat#-3855) were
557 coated with S-RBD of the parent strain at a volume of 50 μ L of 2 μ g/mL diluted antigen in
558 filtered, sterile 1 \times PBS (Thermo Fisher Scientific) at 4°C overnight. The coating buffer was
559 removed, and the plates were washed 3 times with 300 μ L 1 \times PBS plus 0.1% Tween-20 (PBST)
560 wash buffer (Thermo Fisher Scientific) and then blocked with 200 μ L PBST with 3% nonfat
561 milk (milk powder, American Bio) by volume for 1 hour at room temperature. All plasma
562 samples were heat-inactivated at 56°C on a heating block for 1 hour before use and diluted 1:2 in
563 PBS. Negative control samples were prepared at 1:10 dilutions in PBST in 1% nonfat milk and
564 plated at a final dilution of 1:100. A mAb against the SARS-CoV-2 Spike protein was used as a
565 positive control (1:5000 dilution; Sino Biological, 40150-D001). Plasma samples were prepared
566 in 3-fold serial dilutions starting at 1:20 in PBST in 1% nonfat milk. Blocking solution was
567 removed, and 100 μ L diluted plasma was added in duplicate to the plates and incubated at room
568 temperature for 2 hours. Plates were washed 3 times with PBST wash buffer, and 50 μ L of
569 secondary antibody was added to the plates and incubated at room temperature for 1 hour.
570 Antihuman secondary antibody, Fc-specific total IgG HRP (1:5000 dilution; Thermo Fisher
571 Scientific, Invitrogen, A18823), was prepared in PBST plus 1% nonfat milk. Plates were washed,
572 and all residual liquid was removed before the addition of 100 μ L SIGMAFAST OPD (o-
573 phenylenediamine dihydrochloride) solution (MilliporeSigma) to each well, followed by
574 incubation in darkness at room temperature for 10 minutes. To stop the reaction, 50 μ L 3M HCl
575 (Thermo Fisher Scientific) was added to each well. The OD of each plate was read at 490 nm
576 (OD₄₉₀) on a SpectraMax i3 ELISA Plate Reader (BioTek Instruments). The positive cutoff
577 value for each plate was calculated by summing the average of the negative values and 3 times
578 the SD of the negatives. Limits of detection (LOD) were set to half the lowest AUC value at or

579 below 20 titer. The anti-S-RBD IgG titer threshold for seronegative was 180 titer or below. The
580 seropositive anti-S-RBD IgG ELISA titers represent 3-fold dilutions from 540 to 393,660.

581

582 **SARS-CoV-2 Viral Load**

583 Nasopharyngeal specimens obtained at screen were stored in 5 mL of virus transport media at -
584 70°C on site, then shipped to the central storage facility at Johns Hopkins University. RNA was
585 extracted from 200 µL transport media with either the Qiagen viral RNA extraction kit (Qiagen,
586 Hilden, Germany), or the chemagic Viral RNA/DNA 300 H96 kit with chemagic 360 nucleic
587 acid extraction system (Perkin Elmer), according to manufacturer recommended protocols. Real-
588 time reverse transcriptase quantitative PCR (RT-qPCR) assays targeting the SARS-CoV2
589 nucleocapsid (N) gene and the human RNaseP gene were performed based on the methods
590 described by the US CDC³⁹. Primer and FAM-labelled probe sets for CDC nCoV_N1 and
591 RNaseP assays were purchased from IDT (Integrated DNA Technologies) as part of the SARS-
592 CoV2 Research Use Only RUO qPCR primer and probe kit (part number 10006713, 2019_nCoV
593 RUO kit). Single-plex assays with equivalent volumes of RNA (or Positive Control, Plasmid-
594 RNA Standards or Nuclease Free H₂O for No Template Controls (NTCs)) were performed using
595 the TaqPath 1-Step RT-qPCR MasterMix (Applied Biosystems, ThermoFisher Scientific) in a
596 QuantStudio 5 Real-Time PCR system (ThermoFisher Scientific). The SARS-CoV-2 nCoV-N
597 control plasmid comprised the complete nucleocapsid gene of SARS-CoV-2 isolate Wuhan-Hu-
598 1, complete genome (GenBank: NC_045512.2), and the HsRPP30 Positive control contained a
599 portion of the RNaseP (RPP30) gene. Both plasmid controls were purchased from IDT.
600 Standards for quantitative analysis were prepared from serial dilutions of the nCoV-N and
601 HsRPP30 plasmid controls for which target copy number was known. The range covered was
602 200,000 copies to 320 copies. Standard curve analysis of nCoV_N1 Ct values was performed by
603 the QuantStudio Design and Analysis software to determine RNA copies of viral genome. Only
604 samples with quantities within the standard curve range were given a COVID-19 call/score
605 “positive”. A Ct value for the RNaseP gene was used to verify that human RNA was present in
606 each specimen. For samples that did not amplify viral genome or any host cell RNA, a repeat
607 RT-qPCR was performed and subsequently assigned as “undetermined”.

608

609 **SARS-CoV-2 Virus Neutralization Assay**

610 Plasma neutralizing antibodies were determined against WA-1 (SARS-CoV-2/USA-WA1/2020
611 EPI_ISL_404895), which was obtained from BEI Resources, as described previously^{22,40}. Two-
612 fold dilutions of plasma (starting at a 1:20 dilution) were made and infectious virus was added to
613 the plasma dilutions at a final concentration of 1×10^5 TCID₅₀/mL (100 TCID₅₀ per 100 µL).
614 The samples were incubated with the virus for 1 hour at room temperature, and then 100 µL of
615 each dilution was added to 1 well of a 96-well plate of VeroE6-TMPRSS2 cells in hexaplicate.
616 The cells were incubated for 6 hours at 37°C, 5% CO₂. The inocula were removed, fresh
617 infection media (IM) was added, and the cells were incubated at 37°C, 5% CO₂ for 2 days. The
618 cells were fixed by the addition of 100 µL of 4% formaldehyde per well, incubated for at least 4
619 hours at room temperature, and then stained with Naphthol Blue Black (MilliporeSigma). The
620 neutralizing antibodies titer was calculated as the highest serum dilution that eliminated the
621 cytopathic effect in 50% of the wells (NT50), and the AUC was calculated using Graphpad
622 Prism.

623

624 **Statistical analysis**

625 Spearman correlation was used to calculate strength of association between titer and AUC units
626 for antibody measurement. Fold drops in anti-S-RBD IgG AUC between donor and
627 corresponding recipients were calculated by dividing the geometric mean of donors by that of
628 recipients. Statistical differences between donor and post-transfusion recipient antibody levels
629 were determined by non-parametric Mann-Whitney tests. Using viral load or antibody data,
630 multiple comparisons across vaccination status, variant, serostatus, or treatment groups were
631 performed using non-parametric Kruskal-Wallis tests with Dunn's post-hoc corrections. Fishers
632 exact tests were performed to compare the association of timing of treatment (early or late) and
633 antibody levels (high or low) with hospitalization using unvaccinated, CCP recipient post-
634 transfusion (D0) antibody data. Longitudinal recipient antibody data were first log₁₀-transformed
635 and analyzed using a linear mixed-effects regression model, adjusted for variant, age, sex, and
636 BMI. An interaction term was included to examine how antibody levels changed over time by
637 treatment (control or CCP) and hospitalization status for unvaccinated participants. Marginal
638 effects were graphed with 95% confidence intervals. Statistical analyses were performed using
639 GraphPad Prism 8 (GraphPad Software) and Stata 17 (StataCorp).

640 [Data Availability](#)

641 Source data for figures included in supplementary source data file. Data is available from individual
642 authors upon request with reply expected in 14 days. Deidentified data from clinical trial will be
643 deposited in the Vivli server for public access before end of 2023.

644 [Code Availability](#)

645 Unique software or computational code was not created for this study.

646 [References](#)

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769
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779 J.G., J.P., P.B, W.R., ME.C, J.H., B.G., V.C., D.C., K.O., M.A., L.H., C.S., D.N.F., M.Z., E.C.,
780 J.R., S.G.K., C.M., M.R., A.Yarava, K.L., N.M., A.G., N.K., A.S., D.E.F, D.A.J., L.J.A.,
781 D.M.S., B.L., S.E., S.B., T.G., A.Z., D.H., A.C., S.S., E.B., K.G., A.T., O.L., A.P., S.L.K., D.J.S
782 conducted/contributed to the clinical study and/or collected clinical data. HS.P., C.B., A.Yin,
783 J.L., C.C., M.L., S.Y., I.S., A.J., M.R., R.F., O.B., J.S., J.O., T.G., P.C., D.H., A.C., K.G., A.T.,
784 O.L., A.P., S.L.K., D.J.S contributed to data processing and analyses specific to this work.
785 HS.P., C.B., A.Yin, A.C., A.P., S.L.K., D.J.S drafted the manuscript. All authors provided final
786 approval of the version to be published.

787 Competing Interests

788 TG- Paid consultant and employee of Fenwal, a Fresenius Kabi company; AC- Scientific
789 Advisory Board of Sabtherapeutics (cow-derived human immunoglobulins COVID-19 treatment
790 and other infectious diseases) and Ortho Diagnostics Speakers Bureau; EB- member of the FDA
791 Blood Products Advisory Committee; SS reports research grants; F2G, Cidara, Ansun, Zeteo,
792 Emergent Biosolutions: personal fees as consultant, advisory board, data safety monitoring board
793 member; Celltrion, Adagio, Immunome, Adamis, Karyopharm, Intermountain Health: Stock
794 options: Immunome; CS: Centers for Disease Control and Prevention, Merck, Pfizer: Research
795 Grants. All other authors report no relevant disclosures.

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798 **Tables**

799 Table 1 Baseline characteristics for unvaccinated control and CCP recipients and the joint
 800 stratification based upon symptom onset to transfusion as early or late transfusion (≤ 5 or >5 days
 801 from symptom onset) and high or low antibody levels above or below recipient anti-S-RBD IgG
 802 at 150 AUC.

| Unvaccinated | Control | CCP | CCP early high | CCP early low | CCP late high | CCP late low | CCP late (high&low) and early low | Control early |
|------------------------------------|------------|------------|----------------|---------------|---------------|--------------|-----------------------------------|---------------|
| Number | 479 | 472* | 102 | 104 | 164 | 102 | 370 | 207 |
| Median age (IQR) | 44 (33-55) | 42 (32-54) | 42 (30-51) | 46 (33-56) | 41 (31-55) | 44 (33-54) | 44 (32-55) | 43 (32-56) |
| Mean Age | 44 | 43 | 41 | 45 | 43 | 44 | 44 | 45 |
| Age Category, n (%) | | | | | | | | |
| 18-49 yr | 285 (60) | 310 (66) | 74 (73) | 63 (61) | 107 (64) | 66 (65) | 236 (64) | 127 (61) |
| 50-85 yr | 194 (40) | 162 (34) | 28 (27) | 41 (39) | 57 (34) | 36 (35) | 134 (36) | 80 (39) |
| Sex, n (%) | | | | | | | | |
| Female | 283 (59) | 255 (54) | 58 (57) | 52 (50) | 90 (55) | 55 (54) | 197 (53) | 115 (56) |
| Male | 196 (41) | 217 (46) | 44 (43) | 52 (50) | 74 (45) | 47 (46) | 173 (47) | 92 (44) |
| Race, n (%) | | | | | | | | |
| Asian | 17 (4) | 18 (4) | 0 (0) | 5 (5) | 10 (6) | 3 (3) | 18 (5) | 13 (6) |
| Black | 66 (14) | 74 (16) | 12 (12) | 14 (13) | 34 (20) | 14 (14) | 62 (17) | 30 (14) |
| American Indian Pacific Islander | 11 (2) | 9 (2) | 1 (1) | 3 (3) | 3 (2) | 2 (2) | 8 (2) | 8 (4) |
| Other race/not reported | 2 (0) | 2 (0) | 0 (0) | 0 (0) | 1 (1) | 1 (1) | 2 (1) | 1 (0) |
| White | 7 (1) | 5 (1) | 2 (2) | 1 (1) | 1 (1) | 1 (1) | 3 (1) | 4 (2) |
| Ethnicity, n (%) | | | | | | | | |
| Hispanic or Latino | 386 (81) | 367 (78) | 86 (84) | 83 (80) | 117 (70) | 81 (79) | 281 (76) | 159 (77) |
| BMI Category, n (%) | | | | | | | | |
| BMI <30 | 261 (58) | 274 (61) | 64 (65) | 56 (57) | 97 (63) | 57 (59) | 210 (60) | 112 (57) |
| BMI ≥ 30 | 190 (42) | 172 (39) | 34 (35) | 43 (43) | 56 (37) | 39 (41) | 138 (40) | 85 (43) |
| Any Comorbidity, n (%) | | | | | | | | |
| Any Comorbidity | 196 (41) | 193 (41) | 45 (44) | 45 (43) | 56 (34) | 47 (46) | 148 (40) | 88 (43) |
| Hypertension, n (%) | | | | | | | | |
| Hypertension | 117 (24) | 119 (25) | 27 (26) | 27 (26) | 33 (20) | 32 (31) | 92 (25) | 52 (25) |
| Diabetes, n (%) | | | | | | | | |
| Diabetes | 47 (10) | 37 (8) | 5 (5) | 8 (8) | 14 (8) | 10 (10) | 32 (9) | 19 (9) |
| Asthma, n (%) | | | | | | | | |
| Asthma | 59 (12) | 51 (11) | 13 (13) | 11 (11) | 17 (10) | 10 (10) | 38 (10) | 25 (12) |
| HIV, n (%) | | | | | | | | |
| HIV | 12 (3) | 11 (2) | 2 (2) | 4 (4) | 4 (2) | 1 (1) | 9 (2) | 4 (2) |
| Pregnancy, n (%) | | | | | | | | |
| Pregnancy | 0 (0) | 2 (0) | 0 (0) | 1 (1) | 0 (0) | 1 (1) | 2 (1) | 0 (0) |
| Serostatus at Screen, n (%) | | | | | | | | |
| Seronegative* | 368 (77) | 361 (76) | 74 (73) | 97 (93) | 97 (59) | 93 (91) | 287 (78) | 167 (81) |
| Seropositive* | 92 (19) | 107 (23) | 28 (27) | 6 (6) | 66 (40) | 7 (7) | 79 (21) | 30 (14) |
| No screen bloods | 19 (4) | 4 (1) | 0 (0) | 1 (1) | 1 (1) | 2 (2) | 4 (1) | 10 (5) |

803 *13 CCP participants were excluded due to missing post-transfusion anti-S-RBD IgG levels.

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Extended Table 1 Segregation of time (early or late) and antibody level (high or low) to hospitalizations in both unvaccinated groups. Early is symptom onset to transfusion within 5 days and late is after 5 days. The antibody levels are measured by recipient anti-S-RBD IgG inverse dilutional titer and AUC as well as donor anti-S-RBD IgG titer and AUC along with virus neutralizations from donor with titers above the median and AUC at or above the geometric mean. Data are n (%). P values by Fisher's exact tests were calculated by comparing proportion of early and high hospitalizations/non-hospitalizations to the remaining CCP participants, to all unvaccinated controls and to the unvaccinated controls transfused within 5 days of symptom onset.

| | Early/ high CCP | Early/ low CCP | Late/ high CCP | Late/ low CCP | Early/low and Late/high and low | Unvaccinated control plasma | Unvaccinated early control plasma |
|---|-----------------------|----------------------|----------------------|---------------------|--|-----------------------------------|---|
| Recipient anti-S-RBD IgG AUC ≥ 150 | | | | | p=0.03 | p=0.001 | p=0.00005 |
| Hospitalized, n (%) | 0 (0) | 5 (4.8) | 6 (3.7) | 6 (5.9) | 17 (4.6) | 36 (7.5) | 24 (11.6) |
| Total | 102 | 104 | 164 | 102 | 370 | 479 | 207 |
| Donor anti-S-RBD IgG AUC ≥ 3286 | | | | | p=0.21 | p=0.02 | p=0.0007 |
| Hospitalized, n (%) | 1 (1) | 4 (3.6) | 6 (4.8) | 5 (3.6) | 15 (4.0) | 36 (7.5) | 24 (11.6) |
| Total | 96 | 110 | 125 | 140 | 375 | 479 | 207 |
| Donor nAb AUC ≥ 58 | | | | | p=0.016 | p=0.0006 | p=0.00002 |
| Hospitalized, n (%) | 0 (0) | 5 (5.6) | 6 (3.8) | 5 (4.8) | 16 (4.4) | 36 (7.5) | 24 (11.6) |
| Total | 113 | 90 | 159 | 104 | 353 | 479 | 207 |
| Donor EUROIMMUN AU ≥ 6 | | | | | p=0.052 | p=0.003 | p=0.0002 |
| Hospitalized, n (%) | 0 (0) | 5 (4.2) | 5 (4.8) | 6 (3.8) | 16 (4.2) | 36 (7.5) | 24 (11.6) |
| Total | 86 | 119 | 105 | 158 | 382 | 479 | 207 |
| Recipient anti-S-RBD IgG titer > 540 | | | | | p=0.39 | p=0.06 | p=0.01 |
| Hospitalized, n (%) | 0 (0) | 5 (3.1) | 3 (3.1) | 9 (3.7) | 17 (5.2) | 36 (7.5) | 24 (11.6) |
| Total | 45 | 161 | 97 | 169 | 427 | 479 | 207 |
| Donor anti-S-RBD IgG titer > 4860 | | | | | p=0.33 | p=0.02 | p=0.0011 |
| Hospitalized, n (%) | 1 (1.1) | 4 (3.5) | 6 (5.3) | 5 (3.3) | 15 (3.9) | 36 (7.5) | 24 (11.6) |
| Total | 91 | 115 | 114 | 151 | 380 | 479 | 207 |
| Donor nAb titer ≥ 80 | | | | | p=0.03 | p=0.001 | p=0.00005 |
| Hospitalized, n (%) | 0 (0) | 5 (4.9) | 6 (4.1) | 5 (4.3) | 16 (4.4) | 36 (7.5) | 24 (11.6) |
| Total | 101 | 102 | 148 | 115 | 365 | 479 | 207 |
| Donor EUROIMMUN BAU ≥ 600 | | | | | p=0.052 | p=0.003 | p=0.0002 |
| Hospitalized, n (%) | 0 (0) | 5 (4.2) | 5 (4.8) | 6 (3.8) | 16 (4.2) | 36 (7.5) | 24 (11.6) |
| Total | 86 | 119 | 105 | 158 | 382 | 479 | 207 |

Extended Table 2

Unvaccinated screen seropositive participants who were hospitalized

| Participant No. | Treatment group | Screen nasal viral load | Days from symptom onset to transfusion | Variant | Screen RBD Titer | Screen RBD AUC |
|------------------------|------------------------|--------------------------------|---|----------------|-------------------------|-----------------------|
| 1 | Control | 320 | 5 | pre-Alpha | 540 | 134 |
| 2 | Control | 320 | 4 | pre-Alpha | 4860 | 1605 |
| 3 | Control | 23271 | 4 | pre-Alpha | 540 | 87 |
| 4 | Control | 12168 | 3 | Alpha | 540 | 123 |
| 5 | Control | 498 | 9 | Alpha | 14580 | 4302 |
| 6 | CCP | 320 | 7 | pre-Alpha | 1620 | 406 |
| 7 | CCP | 4219 | 7 | Alpha | 1620 | 239 |

Extended Table 3.

Full study participant characteristics during variant periods

| | Pre-Alpha n (%) | Alpha n (%) | Delta n (%) | Total n (%) |
|-------------------------------------|----------------------------|------------------------|------------------------|------------------------|
| Number | 693 | 260 | 228 | 1181 |
| Treatment | | | | |
| CCP | 349 (50) | 128 (49) | 115 (50) | 592 (50) |
| Control | 344 (50) | 132 (51) | 113 (50) | 589 (50) |
| Median Age (IQR) | 47 (35-58) | 40 (31-49) | 36 (29-47) | 43 (32-54) |
| Mean Age | 47 | 40 | 38 | 44 |
| Age Category | | | | |
| 18-49 yr | 388 (56) | 197 (76) | 185 (81) | 770 (65) |
| 50-85 yr | 305 (44) | 63 (24) | 43 (19) | 411 (35) |
| Sex | | | | |
| Female | 384 (55) | 151 (58) | 140 (61) | 675 (57) |
| Male | 309 (45) | 109 (42) | 88 (39) | 506 (43) |
| Race | | | | |
| Asian | 25 (4) | 11 (4) | 8 (4) | 44 (4) |
| Black | 78 (11) | 45 (17) | 40 (18) | 163 (14) |
| American Indian | 16 (2) | 0 (0) | 1 (0) | 17 (1) |
| Pacific Islander | 3 (0) | 1 (0) | 0 (0) | 4 (0) |
| Other/Not Reported | 8 (1) | 2 (1) | 6 (3) | 16 (1) |
| White | 560 (81) | 201 (77) | 173 (76) | 934 (79) |
| Hispanic or Latino | 79 (11) | 40 (15) | 51 (22) | 170 (14) |
| BMI Category | | | | |
| BMI<30 | 395 (57) | 139 (53) | 145 (64) | 679 (57) |
| BMI =>30 | 250 (36) | 116 (45) | 79 (35) | 445 (38) |
| Any Comorbidity | 310 (45) | 98 (38) | 51 (22) | 459 (39) |
| Hypertension | 199 (29) | 52 (20) | 25 (11) | 276 (23) |
| Diabetes | 72 (10) | 18 (7) | 9 (4) | 99 (8) |
| Asthma | 84 (12) | 27 (10) | 21 (9) | 132 (11) |
| HIV | 17 (2) | 7 (3) | 1 (0) | 25 (2) |
| Pregnant | 1 (0) | 0 (0) | 2 (1) | 3 (0) |
| Vaccination Status at Screen | | | | |
| Unvaccinated | 632 (91) | 200 (77) | 79 (35) | 911 (77) |
| Seronegative* | 501 (82) | 154 (77) | 42 (54) | 697 (78) |
| Seropositive* | 110 (18) | 46 (23) | 36 (46) | 192 (22) |

| | | | | |
|--------------------------------|----------|---------------|----------|----------|
| No screening samples | 21 | 0 | 1 | 22 |
| Partly Vaccinated | 20 (3) | 28 (11) | 9 (4) | 57 (5) |
| Seronegative* | 6 (30) | 9 (32) | 2 (22) | 17 (30) |
| Seropositive* | 14 (70) | 19 (68) | 7 (78) | 40 (70) |
| Fully Vaccinated | 0 (0) | 21 (8) | 138 (61) | 159 (13) |
| Seronegative* | 0 (0) | 0 (0) | 1 (1) | 1 (1) |
| Seropositive* | 0 (0) | 21 (100) | 129 (99) | 150 (99) |
| No screening samples | 0 | 0 | 8 | 8 |
| Vaccinated During Study | 100 (14) | 60 (23) | 5 (2) | 165 (14) |
| Hospitalized | 41(6) | 11 (4) | 2 (1) | 54 (5) |
| Seronegative* | 35 (90) | 8 (1-PV) (73) | 1 (100) | 44 (86) |
| Seropositive* | 4 (10) | 3 (27) | 0 | 7 (14) |
| No screening samples | 2 | 0 | 1 | 3 |

Figures

Fig. 1 Screen seropositive participants and post-transfusion seronegative participants. A) Spearman correlation of unique donor anti-S-RBD titer and AUC (n=318) B) Unvaccinated and not hospitalized recipient screen seronegative anti-S-RBD AUC (n=338) with corresponding donor unit antibody levels transfused into matched recipients and C) Seronegative recipients (n=15) subsequently hospitalized after transfusion. D) Antibody levels in the 199 (21%) of unvaccinated, but seropositive recipients at screening before transfusion in the jointly stratified groups. There were no significant differences in screening antibody levels between control and CCP recipients. E) The same 199 unvaccinated seropositive participant antibody levels stratified by days from symptom onset to transfusion. All point estimates are shown with error bars indicating the geometric mean with geometric SD. Numbers above the x-axis represent geometric mean (GM), the number in the group (n). ***p<0.001 by non-parametric Kruskal-Wallis multiple comparisons test with Dunn's post-hoc corrections or Mann-Whitney test. The dashed line in B-E represents the upper post-transfusion 150 AUC recipient's threshold.

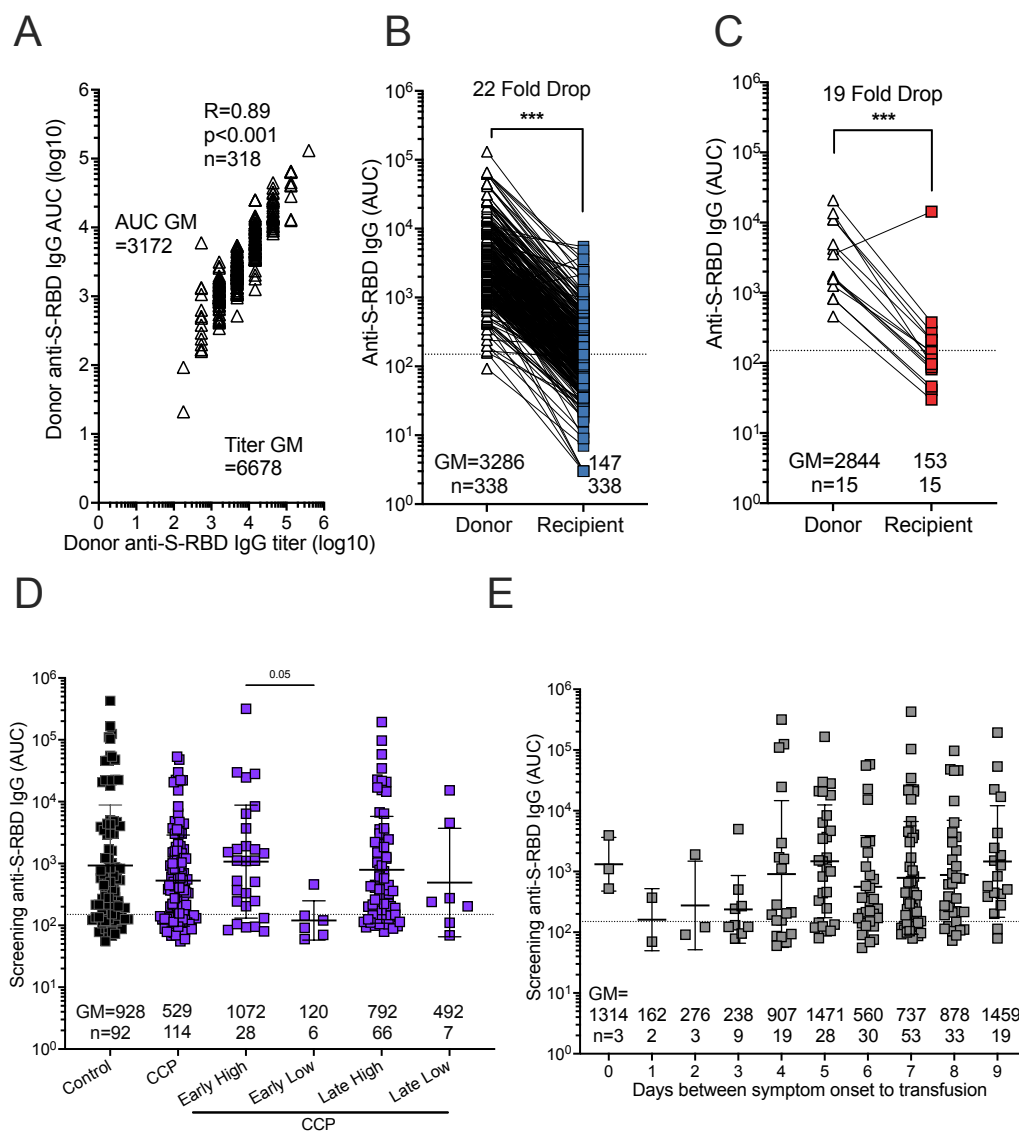


Fig. 2 CCP participant post-transfusion recipient antibody levels stratified by duration from symptom onset to transfusion. Post-transfusion recipient antibody levels segregated by duration from symptom onset to transfusion. Recipient post-transfusion anti-S-RBD IgG AUC levels of subsequently hospitalized (red dots) and non-hospitalized within 28 days from transfusion (blue dots) were plotted by days from symptom onset to transfusion, segregating by early (≤ 5 days of symptom onset) or late (>5 days after symptom onset) transfusion and high (≥ 150 AUC) or low (<150 AUC) antibody levels. Early transfusion with high antibody levels measured 30 minutes post-transfusion (D0) among unvaccinated, CCP recipients had the lowest proportion of hospitalizations (0%) whereas late transfusion with low antibody levels had the greatest proportion of hospitalizations (5.9%). Numbers in each quadrant represent the proportion of hospitalization and sample size for each category.

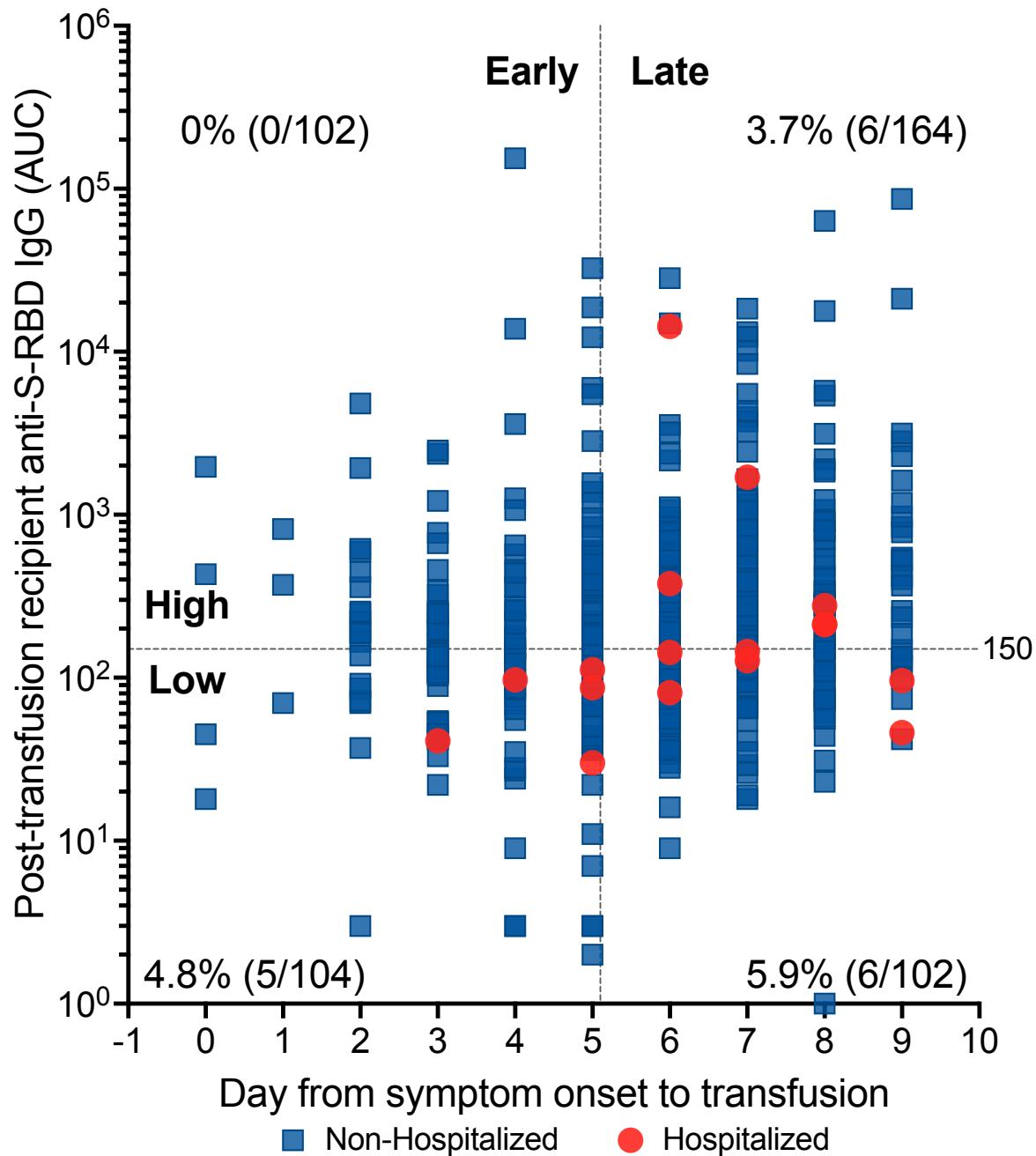


Fig. 3 Mapped recipients with donor unit antibody levels stratified by duration from symptom onset to transfusion. Donor unit antibody levels that correspond with unvaccinated, CCP recipient antibody data segregated by duration from symptom onset to transfusion. A) Early recipients with donor anti-S-RBD IgG AUC levels over 3286 were found to correlate with the lowest proportion of hospitalizations (1.1%). B) Early recipients with donor neutralization antibody (nAb) AUC levels greater than the geometric mean of 58 also had the lowest proportion of hospitalizations (0%). The percentage of recipients hospitalized is indicated in each quadrant of the graph. Numbers in each quadrant represent the proportion of hospitalization and sample size for each category.

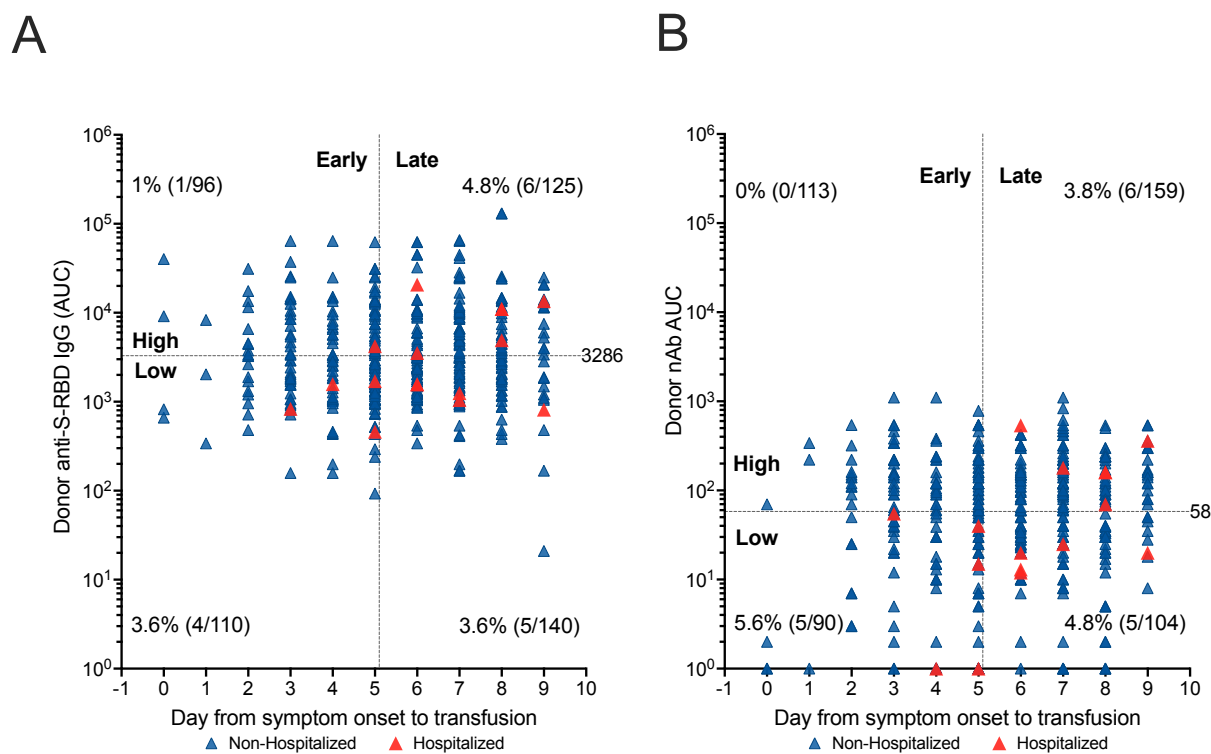


Fig. 4 Screen pre-transfusion nasal swab viral load determinations segregated by A) the unvaccinated jointly stratified early/late or high/low groups. B) CCP or control plasma administration in those hospitalized or not hospitalized. Unvaccinated pre-Delta period participants were segregated into C) seronegative and D) seropositive populations by symptom duration in days to transfusion. Numbers above the x-axis represent geometric mean (GM), the number in the group (n), and percentage of PCR-positive samples (%) for each category. *** p<0.001, ** p=0.002 and * p=0.033 by non-parametric Kruskal-Wallis multiple comparisons test with Dunn's post-hoc corrections. All point estimates are shown with error bars indicating the geometric mean with geometric SD. The dashed lines indicate samples below the limit of detection of 330 viral copies.

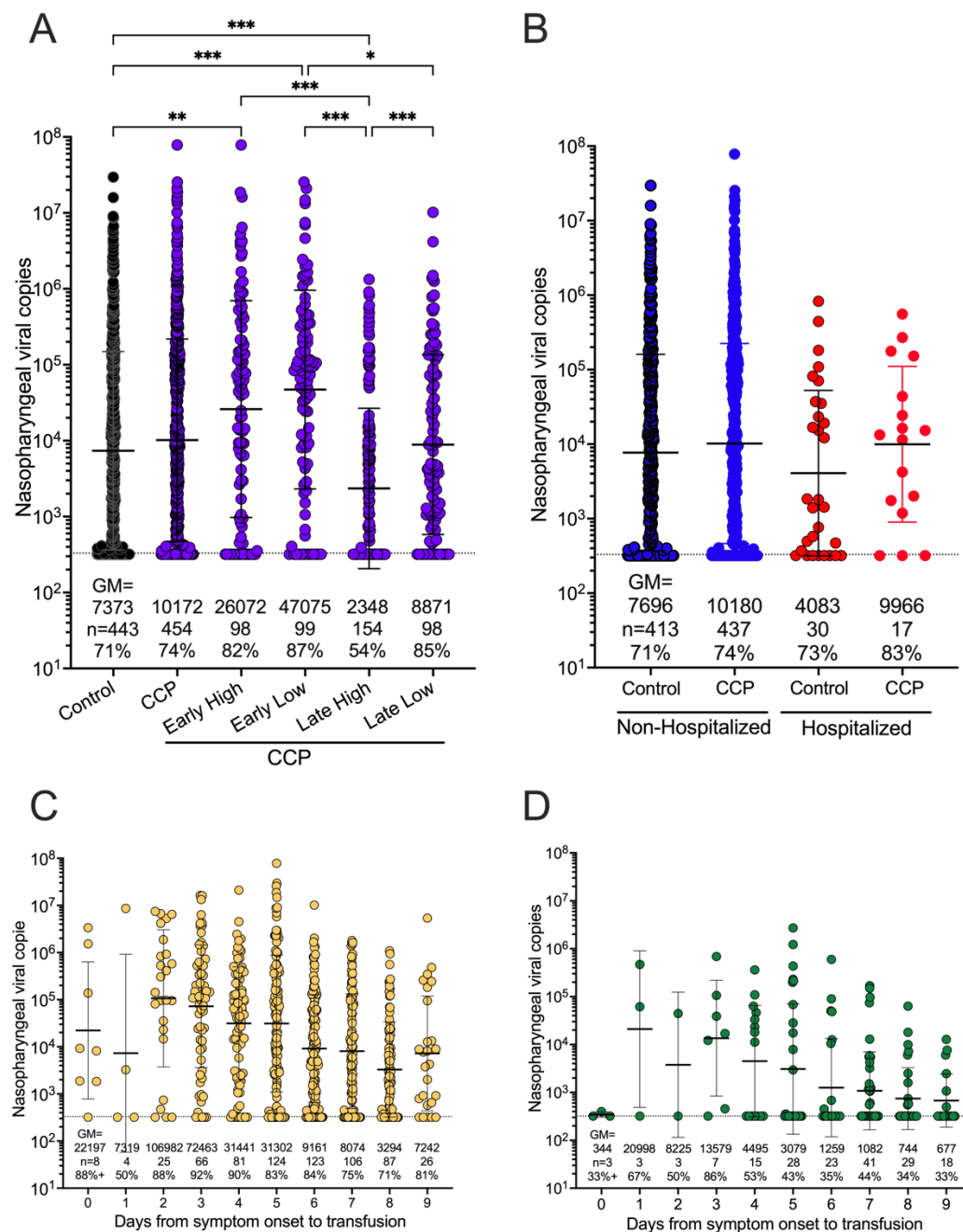
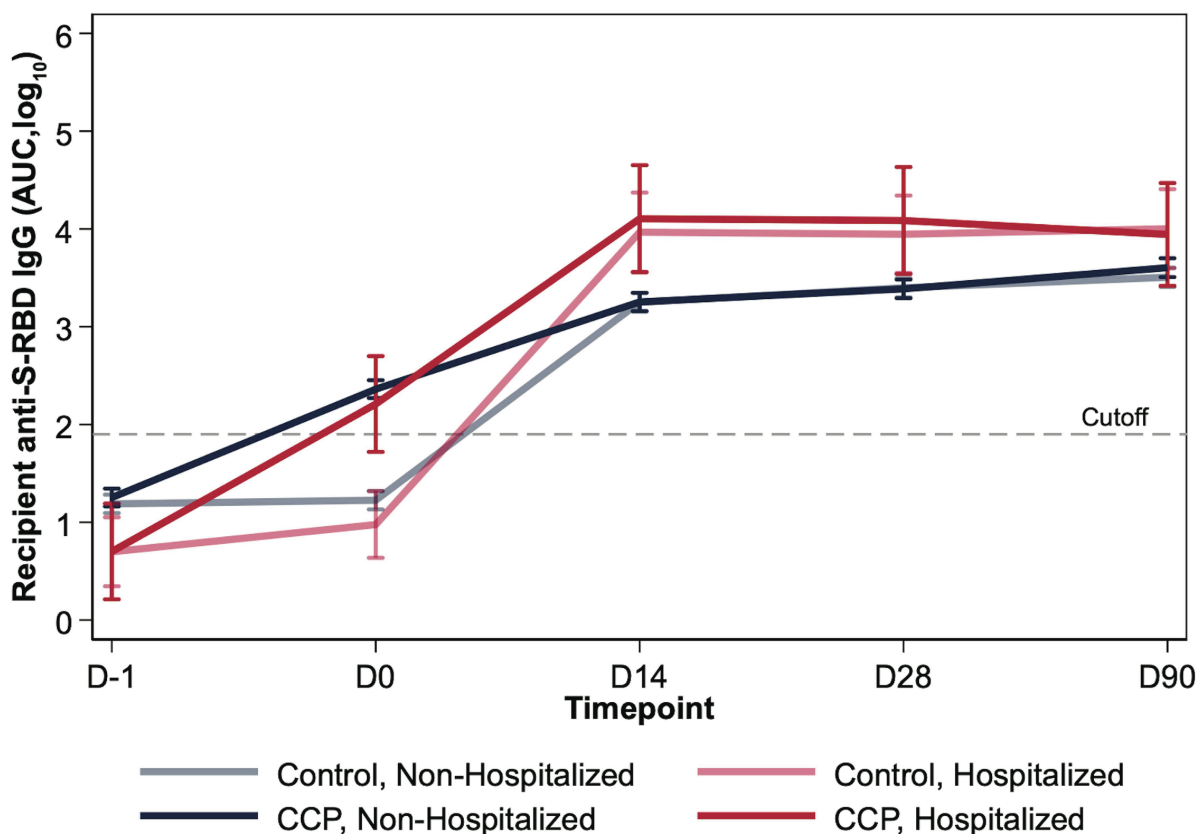


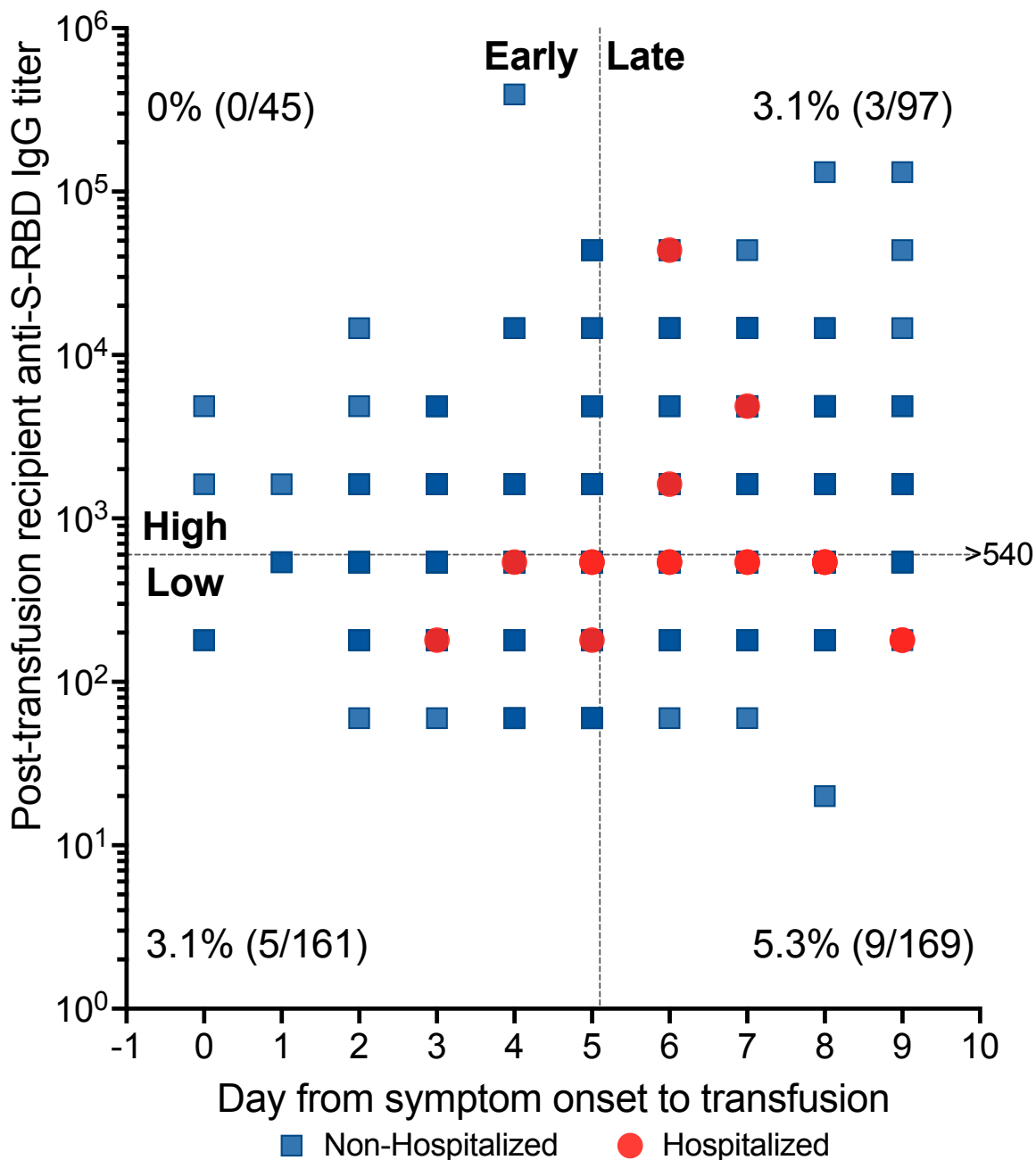
Fig. 5 Antibody levels over three months past transfusion in those hospitalized or not further segregated by CCP or control plasma. Log₁₀-transformed antibody levels up to 90 days post-transfusion were segregated by treatment and hospitalization status of recipients using a linear mixed effects regression model, adjusted for variant, age, sex, and BMI. CCP recipients have greater AUC levels on D0, but by D14, the hospitalized recipients have greater AUC levels than non-hospitalized. The average days from transfusion to hospitalization was 3.05 days, with all post-transfusion hospitalizations occurring between D0 and D14. The dashed line represents the log-transformed cutoff (1.924) for seropositivity. This diagnostic threshold equals the anti-S-RBD IgG log₁₀-transformed value at a 180 titer.



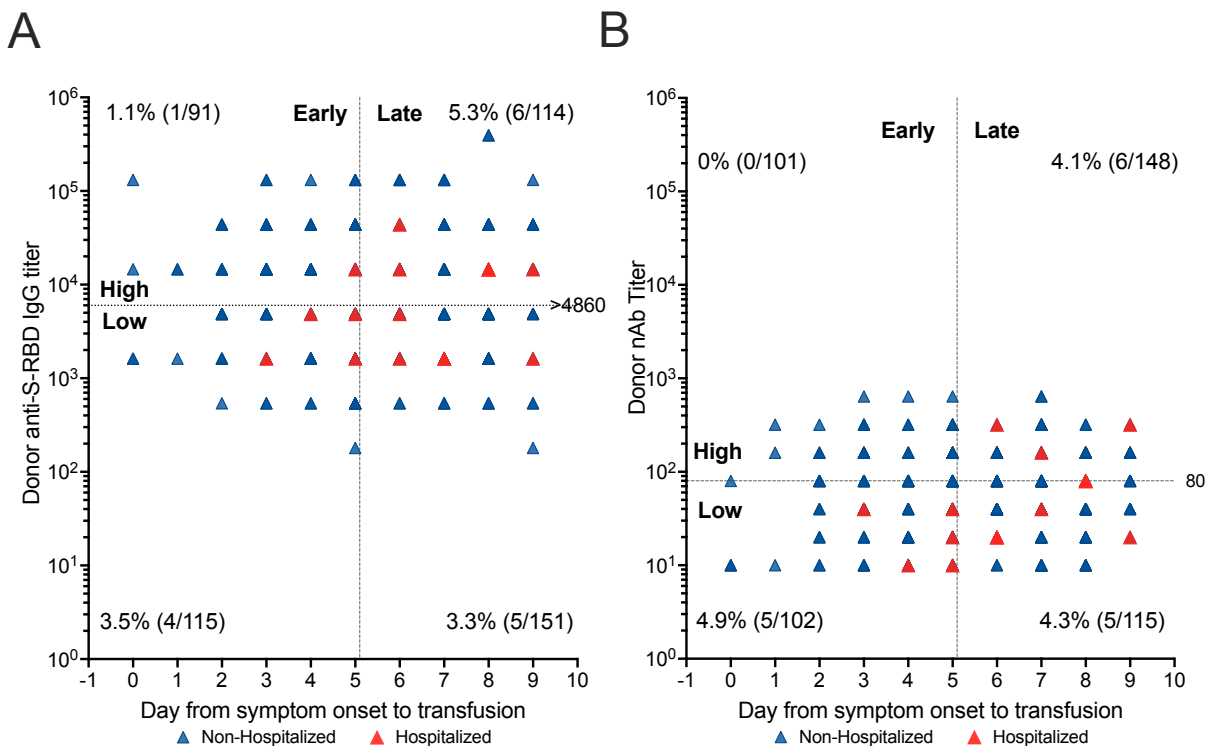
| Group | Comparison | Timepoint | Contrast | P Value |
|------------------|-----------------------------------|-----------|----------|---------|
| CCP | Hospitalized vs. Non-Hospitalized | D0 | -0.144 | 0.573 |
| Control | Hospitalized vs. Non-Hospitalized | D0 | -0.243 | 0.180 |
| Hospitalized | CCP vs. Control | D0 | 1.23 | <0.001 |
| Non-Hospitalized | CCP vs. Control | D0 | 1.13 | <0.001 |
| CCP | Hospitalized vs. Non-Hospitalized | D14 | 0.819 | 0.003 |
| Control | Hospitalized vs. Non-Hospitalized | D14 | 0.686 | 0.001 |
| Hospitalized | CCP vs. Control | D14 | 0.129 | 0.702 |
| Non-Hospitalized | CCP vs. Control | D14 | -0.003 | 0.963 |
| CCP | Hospitalized vs. Non-Hospitalized | D28 | 0.649 | 0.020 |
| Control | Hospitalized vs. Non-Hospitalized | D28 | 0.576 | 0.005 |
| Hospitalized | CCP vs. Control | D28 | 0.061 | 0.856 |
| Non-Hospitalized | CCP vs. Control | D28 | -0.012 | 0.857 |

Extended Figures

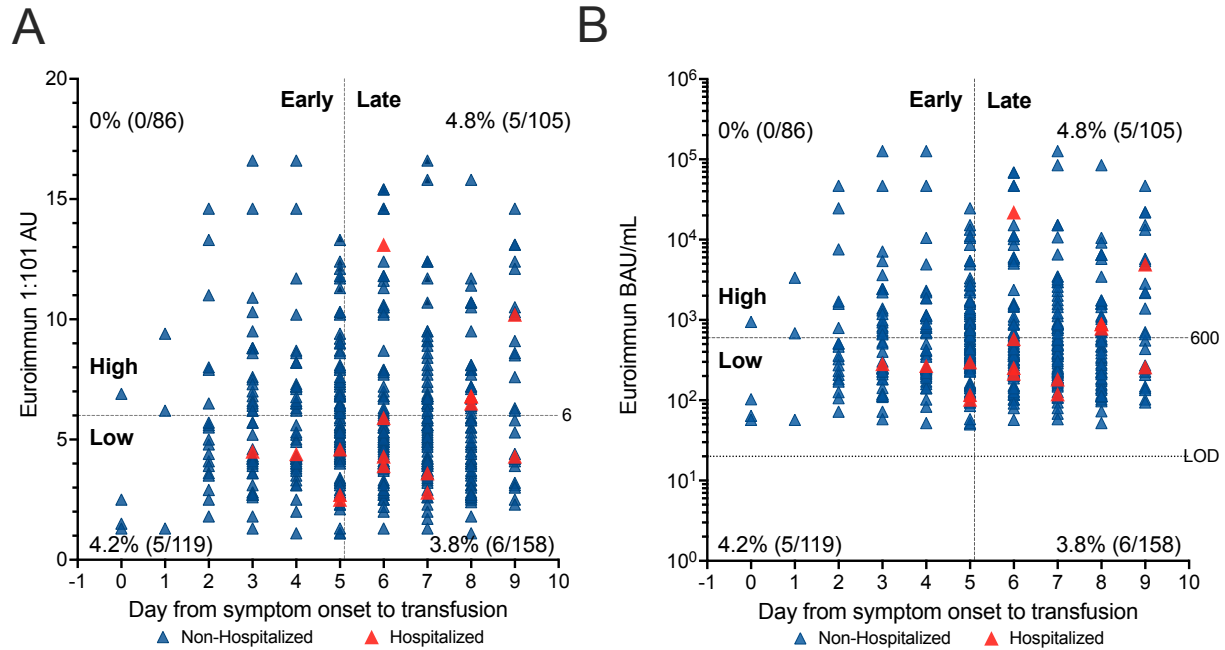
Extended Fig. 1 **Post-transfusion recipient antibody levels stratified by duration from symptom onset to transfusion.** Anti-S-RBD IgG titer levels 30 minutes post-transfusion for unvaccinated CCP recipients are plotted by recipients' days from symptom onset to transfusion, colored by hospitalized (red dots) or non-hospitalized (blue dots). Early (≤ 5 days of symptom onset) or late (>5 days after symptom onset) transfusion is indicated by a dashed line on the x-axis. High (>540 AUC) or low (≤ 540 AUC) levels of antibody are indicated by the dashed line on the y-axis. CCP recipients with early transfusion with high measured anti-S-RBD IgG titer levels shortly after transfusion had the lowest proportion of hospitalization (0%) whereas those with late transfusion and low measured antibody levels had the greatest (5.3%), consistent with the data in AUC units.



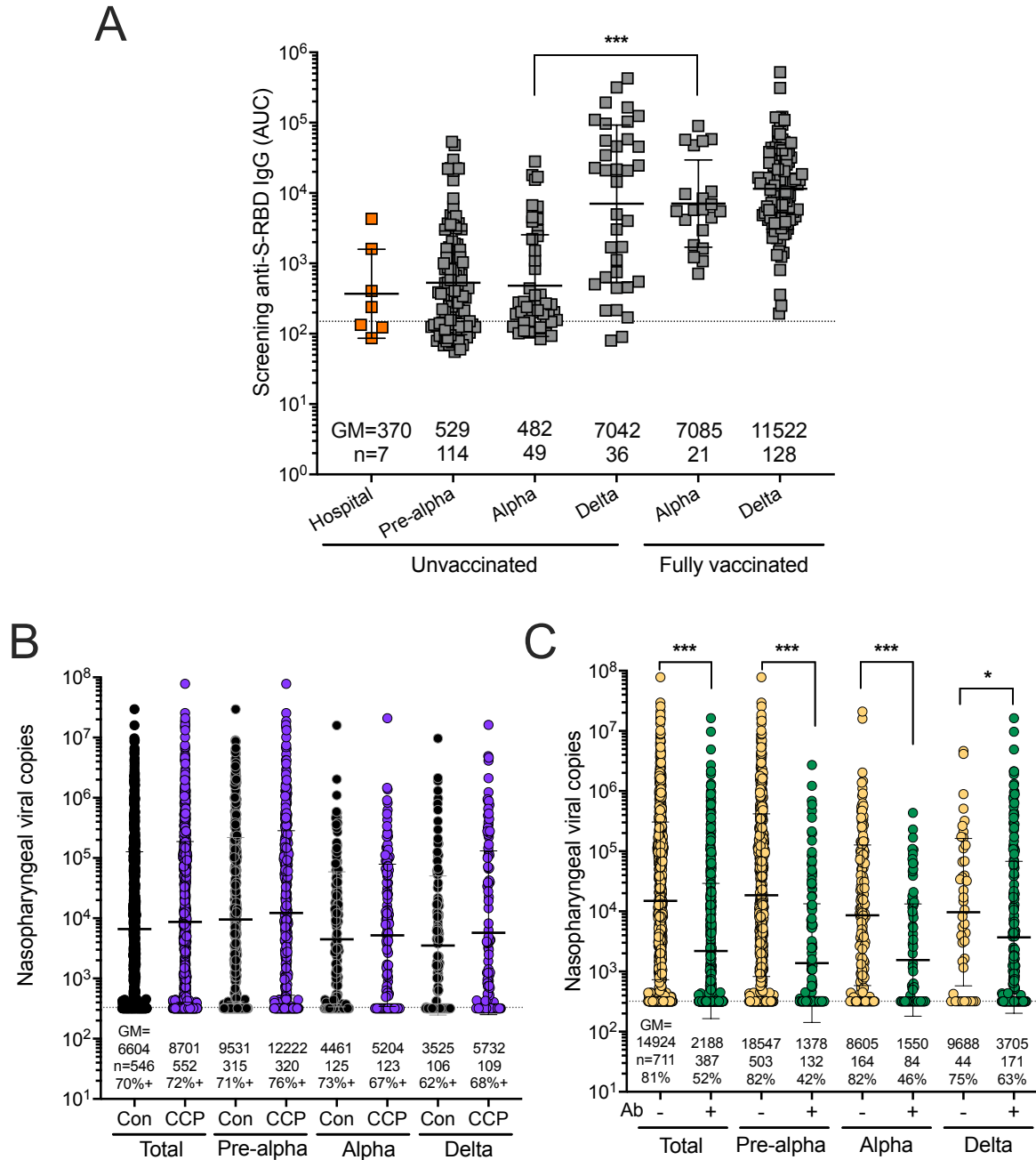
Extended Fig. 2 **Recipients stratified by duration from symptom onset to transfusion matched to donor unit titer antibody levels.** Donor unit antibody levels that correspond with unvaccinated, CCP recipient antibody data segregated by duration from symptom onset to transfusion. A) Early recipients with donor anti-S-RBD IgG titer levels over 4860 were found to correlate with the lowest proportion of hospitalizations (1.1%). B) Early recipients with donor neutralization antibody (nAb) titer levels greater than the median of 80 also had the lowest proportion of hospitalizations (0%). The percentage of recipients hospitalized is indicated in each quadrant of the graph. Numbers in each quadrant represent the proportion of hospitalization and sample size for each category.



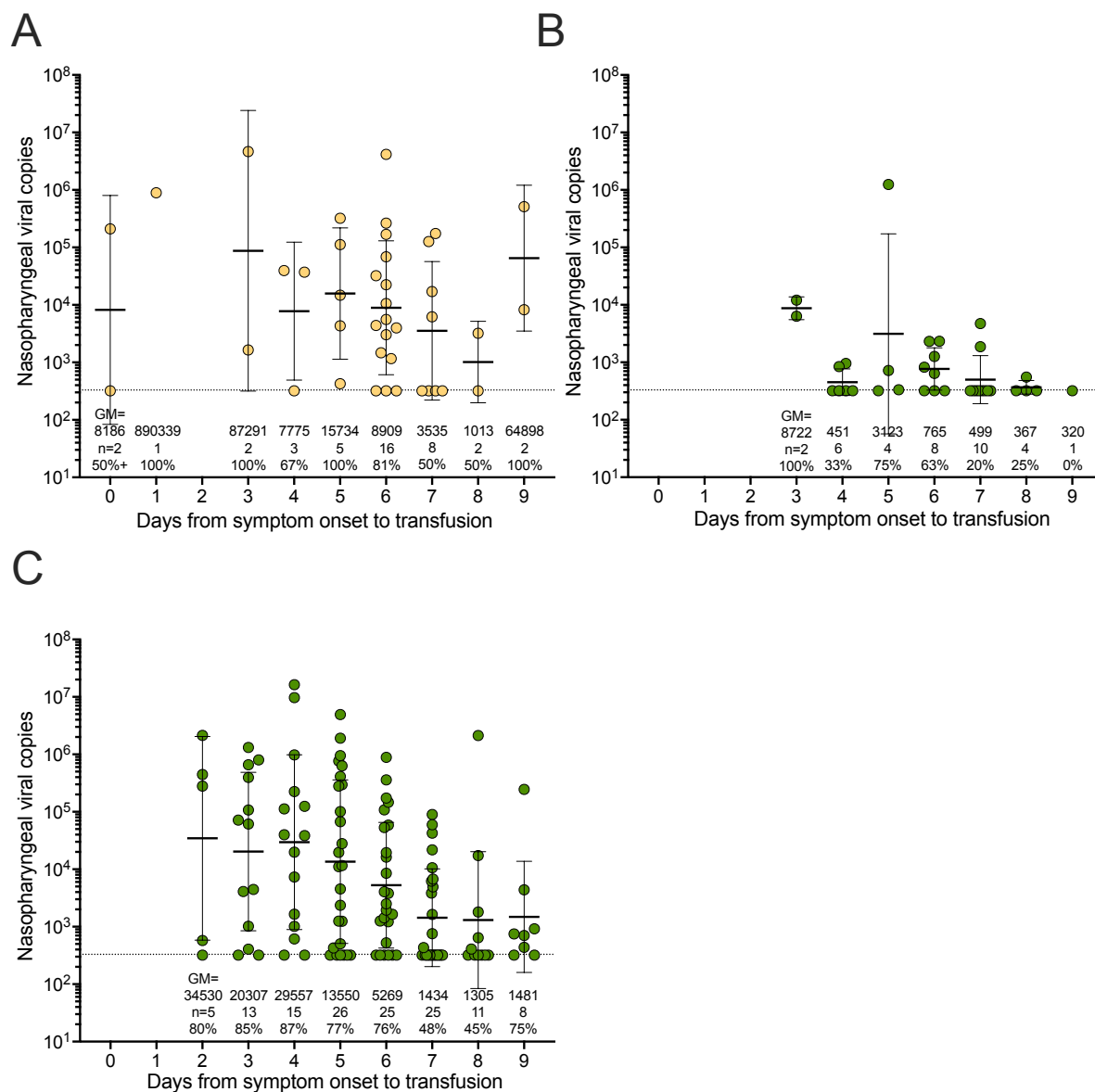
Extended Fig. 3 **Recipients segregated by duration from symptom onset to transfusion matched to donor unit antibody levels measured by EUROIMMUN.** Donor unit antibody levels that correspond with unvaccinated, CCP recipient antibody data segregated by duration from symptom onset to transfusion. A) Early recipients with donor EUROIMMUN over mean of 6 AU were found to correlate with the lowest proportion of hospitalizations (0%). B) Early recipients with EUROIMMUN BAU/mL greater than the geometric mean of 600 also had the lowest proportion of hospitalizations (0%). The percentage of recipients hospitalized is indicated in each quadrant of the graph. Numbers in each quadrant represent the proportion of hospitalization and sample size for each category.



Extended Fig 4 **Antibody levels and viral loads by variant period** A) Unvaccinated and fully vaccinated, but seropositive recipient antibody levels (anti-S-RBD AUC) at screening pre-transfusion as well as fully vaccinated seropositive participants. B) Nasal swab viral load determinations for both vaccinated and unvaccinated segregated by B) CCP or control plasma administration C) serostatus of vaccinated and unvaccinated participants. *** $p < 0.001$ and * $p = 0.033$ by non-parametric Kruskal-Wallis multiple comparisons test with Dunn's post-hoc corrections. All point estimates are shown with error bars indicating the geometric mean with geometric SD. Numbers above the x-axis in B and C represent the geometric means (GM), the number in the group (n) and percentage of samples PCR positive (%). The dashed line in A represents the upper portion post-transfusion 150 AUC recipient's threshold. The dashed lines indicate samples below the limit of detection of 330 viral copies.



Extended Fig. 5 **Screen viral loads during the Delta period.** During the Delta period there were only 77 (34%) participants unvaccinated to segregate into A) seronegative (n=42) and B) seropositive (n=35) groups by duration from symptom onset to transfusion. C) During the Delta period fully vaccinated participants (n=128) were antibody positive with an additional single recipient fully vaccinated, but seronegative with nasal viral load on day 0 of 320 (not graphed). All point estimates are shown with error bars indicating the geometric mean with geometric SD. Numbers above the x-axis represent geometric means (GM), the number in the group (n), and the percentage of samples PCR positive (%). The dashed lines indicate samples below the limit of detection of 330 viral copies.



Extended Fig. 6 **Antibody levels three months post-transfusion.** recipients anti-S-RBD AUC antibody levels at Day 90 post-transfusion (excluding the 165 vaccinated during the follow-up visits) separated by A) Jointly stratified unvaccinated early and late treatment with high or low post-transfusion antibody levels B) both CCP and control recipients by SARS-CoV-2 variant period and vaccination status. Clear squares indicate donor, red squares indicate hospitalized recipients, and gray squares indicate both CCP and control non-hospitalized recipients. *** $p < 0.001$ and * $p = 0.033$ by non-parametric Kruskal-Wallis multiple comparisons test with Dunn's post-hoc corrections. All point estimates are shown with error bars indicating the geometric mean with geometric SD. Numbers above the x-axis represent each category's geometric mean (GM) and number in the group (n). The dashed line in A, B represents the upper portion post-transfusion 150 AUC recipient's threshold, GM donor 3286 AUC and GM donor 6678 titer.

