

1 A Quick Displacement of the SARS-CoV-2 variant Delta with Omicron: Unprecedented Spike in
2 COVID-19 Cases Associated with Fewer Admissions and Comparable Upper Respiratory Viral Loads
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24

25 **Abstract**

26 **Background**

27 The increase in SARS-CoV-2 infections in December 2021 in the United States was driven primarily by
28 the Omicron variant which largely displaced the Delta over a three week span. Outcomes from infection
29 with the Omicron remain uncertain. We evaluate whether clinical outcomes and viral loads differ between
30 Delta and Omicron infections during the period when both variants were co-circulating.

31 **Methods**

32 Remnant clinical specimens from patients that tested positive for SARS-CoV-2 after standard of care
33 testing between the last week of November and the end of December 2021 were used for whole viral
34 genome sequencing. Cycle threshold values (Ct) for viral RNA, the presence of infectious virus, and
35 levels of respiratory IgG were measured, and clinical outcomes were obtained. Differences in each
36 measure were compared between variants stratified by vaccination status.

37 **Results**

38 The Omicron variant displaced the Delta during the study period and constituted 95% of the circulating
39 lineages by the end of December 2021. Patients with Omicron infections (N= 1121) were more likely to
40 be vaccinated compared to patients with Delta (N = 910), but were less likely to be admitted, require ICU
41 level care, or succumb to infection regardless of vaccination status. There was no significant difference in
42 Ct values based on the lineage regardless of the vaccination status. Recovery of infectious virus in cell
43 culture was reduced in boosted patients compared to fully vaccinated without a booster and unvaccinated
44 when infected with the Delta lineage. However, in patients with Omicron infections, recovery of
45 infectious virus was not affected by vaccination.

46 **Conclusions**

47 Omicron infections of vaccinated individuals are expected, yet admissions are less frequent. Admitted
48 patients might develop severe disease comparable to Delta. Efforts for reducing the Omicron transmission
49 are required as even though the admission risk is lower, the numbers of infections continue to be high.

50 **Research in context**

51 **Evidence before this study**

52 The unprecedented increase in COVID-19 cases in the month of December 2021, associated with the
53 displacement of the Delta variant with the Omicron, triggered a lot of concerns. An understanding of the
54 disease severity associated with infections with Omicron is essential as well as the virological
55 determinants that contributed to its widespread predominance. We searched PubMed for articles
56 published up to January 23, 2022, using the search terms (“Omicron”) AND (“Disease severity”) as well
57 as (“Omicron”) AND (“Viral load”) And/ or (“Cell culture”). Our search yielded 3 main studies that
58 directly assessed the omicron’s clinical severity in South Africa, its infectious viral load compared to
59 Delta, and the dynamics of viral RNA shedding. In South Africa, compared to Delta, Omicron infected
60 patients showed a significant reduction in severe disease. In this study, Omicron and non-Omicron
61 variants were characterized based on S gene target failure using the TaqPath COVID-19 PCR (Thermo
62 Fisher Scientific). In the study from Switzerland that assessed the infectious viral load in Omicron versus
63 Delta, the authors analyzed only 18 Omicron samples that were all from vaccinated individuals to show
64 that compared to Delta, Omicron had equivalent infectious viral titers. The third study that assessed the
65 Omicron viral dynamics showed that the peak viral RNA in Omicron infections is lower than Delta. No
66 published studies assessed the clinical discrepancies of Omicron and Delta infected patients from the US,
67 nor comprehensively assessed, by viral load and cell culture studies, the characteristics of both variants
68 stratified by vaccination status.

69 **Added value of this study**

70 To the best of our knowledge, this is the only study to date to compare the clinical characteristics and
71 outcomes after infection with the Omicron variant compared to Delta in the US using variants
72 characterized by whole genome sequencing and a selective time frame when both variant co-circulated. It
73 is also the first study to stratify the analysis based on the vaccination status and to compare fully
74 vaccinated patients who didn't receive a booster vaccination to patients who received a booster
75 vaccination. In addition, we provide a unique viral RNA and infectious virus load analyses to compare
76 Delta and Omicron samples from unvaccinated, fully vaccinated, and patients with booster vaccination.

77 **Implications of all the available evidence**

78 Omicron associated with a significant increase in infections in fully and booster vaccinated individuals
79 but with less admissions and ICU level care. Admitted patients showed similar requirements for
80 supplemental oxygen and ICU level care when compared to Delta admitted patients. Viral loads were
81 similar in samples from Omicron and Delta infected patients regardless of the vaccination status. The
82 recovery of infectious virus on cell culture was reduced in samples from patients infected with Delta who
83 received a booster dose, but this was not the case with Omicron. The recovery of infectious virus was
84 equivalent in Omicron infected unvaccinated, fully vaccinated, and samples from patients who received
85 booster vaccination.

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93 **Introduction**

94 The SARS-CoV-2 Omicron variant was first identified in South Africa and reported to the World Health
95 Organization (WHO) on November 24, 2021 (1). A large number of mutations and amino acid changes
96 were noted across the Omicron genome, 15 of which are within the receptor binding domain (RBD) of the
97 spike (S) protein (2). Some of the Omicron RBD characterized mutations raised concerns of a significant
98 impact on the transmissibility, immunity secondary to vaccination or prior infection, and efficacy of
99 therapeutic monoclonal antibodies. Hence, the WHO designated the Omicron as a variant of concern
100 (VOC) on November 26, 2021 (3, 4). Initial reports from South Africa (5) and the UK (6) suggested that
101 the Omicron variant may be more transmissible but cause less severe infection. However, the US
102 population differs from both the South African and UK populations in significant ways; most importantly,
103 the percentage of the population that is vaccinated is significantly lower in the US than the UK and prior
104 infection is significantly lower in the US compared to South Africa. Thus, significant questions remain as
105 to the impact Omicron will have on the US population. Most importantly, how might outcomes,
106 particularly hospitalization, differ between patients infected with Omicron compared to variants such as
107 Delta. Additionally, there remains uncertainty as to whether enhanced evasion of pre-existing immunity
108 or some other biological mechanisms drive the higher rate of Omicron transmission.

109 The Omicron variant was first identified in Maryland in the last week of November 2021, and became
110 predominant in a matter of 3 weeks. In this study, we evaluated the biological differences between viral
111 variants collected as a part of routine clinical care between November 22nd and December 31st in the
112 Johns Hopkins Health System, as well as the clinical outcomes in patients infected with Omicron and
113 Delta. We focused on this time frame as it witnessed the detection of the first Omicron case in our system
114 in the last week of November, and the switch between Delta predominance in the beginning of December
115 to Omicron predominance at the end of December 2021. We provide a comparison of clinical,
116 demographic and virological load between Delta and Omicron infected individuals and stratify our results
117 based on vaccination status.

118 **Methods**

119 **Ethical considerations and Data availability**

120 Research was conducted under Johns Hopkins IRB protocol IRB00221396 with a waiver of consent.
121 Remnant clinical specimens from patients that tested positive for SARS-CoV-2 after standard of care
122 testing were used for whole genome sequencing. Whole genomes were made publicly available at
123 GISAID.

124 **Specimens and Patient Data**

125 Nasopharyngeal or lateral mid-turbinate nasal swabs after standard of care diagnostic or screening testing
126 were collected and used for genome sequencing. At Johns Hopkins Medical System, SARS-CoV-2
127 clinical testing is performed for inpatients and outpatients (five acute care hospitals and more than 40
128 ambulatory care offices) as well as standard of care screening particularly prior to scheduled surgeries.
129 Molecular assays used include primarily the NeuMoDx SARS-CoV-2 (Qiagen), Cobas SARS-CoV-2
130 (Roche), Xpert Xpress SARS-CoV-2/Flu/RSV (Cepheid), in addition to the RealStar® SARS-CoV-2 RT-
131 PCR (Altona Diagnostics), ePlex Respiratory Pathogen Panel 2 (Roche), Aptima SARS-CoV-2 (Hologic),
132 and Accula SARS-CoV-2 assays (ThermoFisher Scientific) (7-10). Each sample in our cohort represents a
133 unique patient. Table 1 summarizes the numbers of charts and samples used for each part of the study.

134 **Clinical data analysis**

135 Clinical and vaccination data for patients whose samples were characterized by whole genome
136 sequencing was bulk extracted as previously detailed in (11). Repeat tests from the same patient were
137 excluded as were results with uncharacterized genomes due to insufficient quality (n=347). The first
138 Omicron infection was collected from a patient in the Johns Hopkins Medical System during the last
139 week of November, 2022. Notably, samples are collected randomly from the whole system for whole
140 genome sequencing for surveillance. A total of 1121 Omicron and 910 Delta infected patients diagnosed
141 between November 22nd 2021 and December 31st 2021 were included in the study. COVID admission

142 relatedness was determined based on presenting complaints, admission diagnoses, reason for testing, and
143 timing of testing. Patients admitted with no COVID-related complaints and no COVID-related diagnoses
144 that were tested after admission as part of an asymptomatic screening protocol that is administered to all
145 admitted patients were considered non-COVID-related admissions. Additionally, patients who developed
146 symptoms after admission or who tested positive on regular asymptomatic surveillance of inpatients were
147 also considered non-COVID-related admissions. In our vaccinated patients' population, 68.6% received
148 Pfizer/BioNTech, followed by the Moderna mRNA-1273 (26.6%), then the J&J/Janssen COVID-19
149 vaccines (4.8%). Full vaccination was based on the CDC definition of positive test results more than 14
150 days post the second shot for pfizer/BioNTech BNT162b2 and Moderna mRNA-1273 or 14 days after the
151 J&J/Janssen.

152 **Ct value analysis**

153 To ensure comparable Ct values for viral load analyses, samples were retested with the PerkinElmers
154 SARS-CoV-2 kit (<https://www.fda.gov/media/136410/download>) and Ct values of the N gene were used
155 for comparisons.

156 **Amplicon based Sequencing**

157 Specimens preparation, extractions, and sequencing were performed as described previously (12, 13).
158 Library preparation for this cohort was performed using the NEBNext® ARTIC SARS-CoV-2
159 Companion Kit (VarSkip Short SARS-CoV-2 # E7660-L). Sequencing was performed using the
160 Nanopore GridION and reads were basecalled with MinKNOW, and demultiplexed with guppybarcoder
161 that required barcodes at both ends. Alignment and variant calling were performed with the artic-
162 ncov2019 medaka protocol with thresholds set to a minimum of 90% coverage and 100 mean depth.
163 Query mutations were manually confirmed with Integrated Genomics viewer (IGV) (Version 2.8.10),
164 clades were determined using Nextclade beta v 0.12.0 (clades.nextstrain.org), and lineages were
165 determined with Pangolin COVID-19 lineage Assigner ([COG-UK \(cog-uk.io\)](https://cog-uk.org)).

166 **ELISA**

167 Respiratory samples were tested, undiluted, with the EUROIMMUN Anti-SARS-CoV-2 ELISA (IgG)
168 following the package insert (<https://www.fda.gov/media/137609/download>) as we described previously
169 (11). The assay detects antibodies to the SARS-CoV-2 S1 domain of the spike protein. The value 1.1 was
170 used as a cut off for positives.

171

172 **Cell culture**

173 Vero-TMPRSS2 cells were cultured and infected with aliquots of swab specimens as previously described
174 for VeroE6 cells (14). Cultures were incubated for 6 days and SARS-CoV-2 cytopathic effect (CPE) was
175 confirmed by reverse transcriptase PCR.

176

177 **Statistical analysis**

178 Statistical analyses were conducted using GraphPad prism. Chi-square and Fisher Exact tests were used
179 for categorical variable comparisons and t-test and one-way ANOVA were used for comparing
180 continuous independent variables.

181 **Results**

182 **SARS-CoV-2 positivity and variants trends November- December 2021.**

183 The SARS-CoV-2 positivity rate increased markedly in December 2021 (25.5%, Figure 1A). The large
184 increase in SARS-CoV-2 positivity in the month of December was the highest recorded since the
185 beginning of the pandemic (Figure S1 and (15)). The spike in SARS-CoV-2 positivity was particularly
186 evident during the last two weeks of December for both symptomatic and asymptomatic patients (Figure
187 1A). The increase in the positivity correlated with an increase in the detection of the Omicron variant, that
188 went from less than 1% of sequenced strains in the beginning of December to the dominant variant in less

189 than 3 weeks (Figure 1B and C). Table 2 shows the total numbers tested and sequenced and the detailed
190 positivity per respiratory target.

191 **Patient characteristics and infection outcomes in Omicron and Delta infections.**

192 A total of 7,353 samples tested positive for SARS-CoV-2 of a total 45,856 tested in the Johns Hopkins
193 Laboratories between November 22nd 2021 and December 31st 2021. Of these, 2,378 were randomly
194 selected for whole genome sequencing. After excluding repeat tests in patients and results that were
195 unable to be characterized due to low quality, a total of 2,031 patients, 1121 Omicron and 910 Delta, were
196 included in the study. Patients infected with the Omicron variant were younger (median age 32 years vs
197 35, $p < 0.001$, Table 3) and significantly more likely to be vaccinated than patients with Delta infections
198 (Table 3). However, compared to patients with Delta, patients with Omicron were significantly less likely
199 to be admitted (3 % vs 13.8 %, $p < 0.00001$), require ICU level care (0.5% vs 3.5%, $p < 0.00001$), or
200 expire (0.1 vs 1.1, $p = 0.004$, Table 3).

201 In the vaccinated population, patients with either Omicron or Delta were likely to be older, though this
202 was only significant in the Omicron group (Table 4). No significant differences in comorbidities or
203 disease severity was noted when all fully vaccinated were compared to unvaccinated for both Delta and
204 Omicron groups (Table 4). However, in fully vaccinated patients who had received a booster, patients
205 with Delta infections were significantly more likely to have comorbidities including immunosuppression
206 and lung disease and were more likely to be admitted than patients with Omicron (Table S1). While
207 patients admitted with Omicron infections were more commonly fully vaccinated (58.8 % vs 23.8 %, $p =$
208 0.0001), we didn't notice a significant difference in the need for supplemental oxygen, ICU level care, or
209 mortality (Table 5).

210 **Omicron and Delta variants cycle threshold (Ct) values in upper respiratory samples.**

211 To determine if the Ct values in upper respiratory samples were different in Omicron versus Delta
212 infected individuals, we compared the Ct values collected for all the groups regardless of the days to the

213 onset of symptoms or the status of the patients being symptomatic or asymptomatic (Omicron = 419,
214 Delta N = 660). No difference in the mean or median Ct values were notable (Figure 2A). No differences
215 were noted when Ct values were compared between vaccinated and unvaccinated patients from both
216 groups as well (Omicron vaccinated = 235, unvaccinated = 170, Delta vaccinated = 240, unvaccinated =
217 411, Figure 2A). When the Ct analysis was correlated to the days from the onset of symptoms for
218 symptomatic patients, no significant differences were detected between Omicron and Delta fully
219 vaccinated, boosted, or unvaccinated (Figure 2B and C and Figure S2). We conclude that there were no
220 significant differences in viral RNA loads between Omicron and Delta infected individuals.

221 **Recovery of infectious virus in Omicron versus Delta groups**

222 To assess the recovery of infectious virus from upper respiratory tract specimens of individuals infected
223 with Omicron versus Delta variants, samples from 219 Omicron (N, Table 1) and 153 Delta (N, Table 1)
224 infections were used to inoculate Vero-TMPRSS2 cells. Recovery of infectious virus (positive cytopathic
225 effects; CPE) was noted from more specimens from the Delta group as compared to the Omicron group
226 (Delta 78%, Omicron 61%; Figure 3A, $p = 0.0006$). Specimens from the boosted Delta group showed
227 significant reduction in the recovery of infectious virus as compared to the fully vaccinated or the
228 unvaccinated Delta groups (62% vs 87 and 82 %, $p = 0.009$ and 0.04 , Figure 3A). This was not the case in
229 the Omicron groups which had no statistically significant differences in infectious virus recovery between
230 the boosted, fully vaccinated and unvaccinated groups (55%, 69%, and 60%, Figure 3A). A significant
231 increase in the recovery of infectious virus from specimens of patients infected with the Delta variant as
232 compared to the Omicron was noted for both fully vaccinated (87% vs 69%, $p < 0.04$) and unvaccinated
233 (82% vs 60%, $p = 0.008$) groups (Figure 3A). Consistent with our previously published reports (Figure
234 3C and (11, 14)), and since lower Ct values have been associated with positive CPE, we compared
235 samples with Ct values less than 20 for both Delta and Omicron groups (N; Table 1). Our analysis
236 showed that the Delta infection was associated a significant increase in samples with positive CPE
237 compared to Omicron (87% vs 70%, $p = 0.001$) however, no significant differences were seen between

238 boosted, fully vaccinated, or unvaccinated groups (Figure 3B). Taken together, Omicron infections had
239 lower numbers of samples with infectious virus when compared to Delta virus infections, indicating that a
240 higher infectious virus load alone was not driving the higher transmissibility seen in Omicron infections.
241 Notably, no significant differences were noted in the specimen collection time frame in relation to the
242 onset of symptoms in all groups (Figure 3D). Of note, 17.5% of samples with infectious virus were
243 collected after 5 days from symptoms.

244 **Localized SARS-CoV-2 IgG in nasal specimens**

245 We previously showed that SARS-CoV-2 IgG levels in the upper respiratory tract are higher in samples
246 from vaccinated individuals and correlate with less recovery of infectious virus on cell culture (11). To
247 compare the anti-SARS-CoV-2 IgG levels between patients who received a booster and fully vaccinated
248 patients, ELISA was performed on upper respiratory samples from the Omicron and Delta infected
249 groups. As expected, a significant increase in localized IgG levels was observed in patients who received
250 a booster (Figure 4A, $p < 0.0001$) and IgG levels were higher in samples with no detectable infectious
251 virus compared to those with infectious virus (Figure 4A, $p < 0.0001$). The anti-SARS-CoV-2 IgG levels
252 were higher in the boosted and Omicron-infected group compared to the Omicron-infected and fully
253 vaccinated groups but this was not significant in the Delta-infected groups (Figure 4B, $p < 0.0001$). Both
254 Omicron and Delta infected patients that had infectious virus showed a significant decrease in anti-SARS-
255 CoV-2 IgG levels when compared to specimen with no infectious virus (Figure 4C, $p < 0.05$). The data
256 indicate that anti-SARS-CoV-2 antibody levels in the upper respiratory tract can be increased after
257 booster vaccination but the presence of infectious virus with either Omicron or Delta infection is
258 associated with lower local levels of vaccination induced antibodies.

259

260 **Discussion**

261 In this study, we provide a comparison between Omicron and Delta infected patients from the transition
262 from Delta to Omicron dominance. Using this tight time frame also controls for the timing of samples'
263 collection in relation to vaccinations and booster doses between the two groups in addition to the
264 implemented community measures of infection control, including masking and social distancing. Our
265 data showed that Omicron infected patients were associated with higher infection rates in vaccinated
266 individuals and those who received booster vaccinations but admissions, ICU level cares, and mortality
267 were significantly less. Samples collected from Omicron infected patients had equivalent viral loads when
268 compared to samples collected from Delta infected individuals regardless of the vaccination status.
269 Recovery of infectious virus on cell culture was less frequent in the Omicron group with no significant
270 impact of vaccination with the exception of the Delta individuals who received a booster dose, this group
271 showed a significantly reduced recovery of infectious virus when compared to the Delta fully vaccinated
272 and unvaccinated groups. Consistent with our previously reported observations (11), samples with
273 successful recovery of infectious virus on cell culture correlated with less IgG levels in the respiratory
274 samples. In this study, we also report a significant increase of IgG in samples collected from patients who
275 received a booster dose.

276 The Omicron variant was first reported from South Africa early in November 2021. The first case was
277 reported from the US on December 1st 2021, a few days after the WHO classified it as a variant of
278 concern (16). The first case we identified as a part of our SARS-CoV-2 genomic surveillance was from a
279 patient who developed symptoms during the last week of November 2021. A very quick increase in
280 Omicron detection correlated with a marked increase in the overall SARS-CoV-2 positivity to reach an
281 average of close to 50% in symptomatic patients in the last week of December. Notably, the Omicron
282 detection and rapid increase occurred during a spike in Delta circulation. In contrast, the Delta
283 displacement of Alpha variant occurred in a time of markedly low circulation of the latter (Figure S1 and
284 (11)). The rapid increase in Omicron cases could be explained by either an increase in overall
285 transmissibility of this variant or due to the enhanced immune evasion of Omicron through multiple

286 mutations in the Spike protein. In a study that compared the house hold contacts of Delta and Omicron
287 infected patients, the secondary attack rate was higher with Omicron, particularly in fully vaccinated and
288 boosted contacts (17). Notably, 53.1% of the Omicron infected patients from our cohort were fully
289 vaccinated or received booster doses and there was no difference in the presence of Omicron infectious
290 virus in either the unvaccinated, vaccinated or boosted groups, suggesting little effect of vaccination on
291 infectious virus load. Multiple studies have shown that the neutralization of the Omicron is reduced
292 compared to Delta or prior variants (18-21). Interestingly, there were fewer numbers of specimen
293 containing infectious virus in the Omicron group compared to the Delta group, indicating that the
294 presence of infectious virus alone may not explain the higher transmissibility of Omicron.

295 The omicron genome contains 32 amino acid changes in the spike protein, within its NTD, RBD, and
296 close to the furin cleavage region, some of which are shared with prior variants of concern and were
297 previously characterized (22, 23). Interestingly, those changes are expected to impact the binding to the
298 host receptor ACE2 and alter membrane fusion (24). This likely explains the notable phenotypic changes
299 of Omicron in cell culture and animal studies and the change in the viral tropism. Omicron was shown to
300 cause a mild disease in animals and replicate less efficiently in the lungs (25). In addition, the use of the
301 Vero-TMPRSS2 cell line which initially showed an enhanced sensitivity for the recovery of SARS-CoV-
302 2 (26), was also impacted by changes within Omicron. The slower viral growth might indicate an
303 alternative entry pathway that is less dependent on TMPRSS2 (27). This is consistent with our
304 observations that samples from the Omicron infected group were associated with less recovery of
305 infectious virus on this cell line, yet, using this cell line allowed the comparisons between different groups
306 based on their vaccination status. Our data indicates that the recovery of infectious virus with Omicron is
307 not impacted by booster vaccination, which was not the case with Delta infected patients. Even though,
308 when we strictly limited our analysis to samples with Ct values less than 20, the recovery of infectious
309 virus from the Delta infected, boosted group was equivalent to other groups. IgG levels were significantly
310 higher in samples from boosted, vaccinated patients and those with no infectious virus. Taken together,

311 we believe that Omicron evasion of preexisting immunity contributes to lessen the impact of booster
312 vaccination on the recovery of infectious virus, which might contribute to the increased transmission,
313 even in individuals who receive booster vaccination.

314 The most recent CDC guidelines for infection control and isolation indicates that in a contingency status,
315 health care workers can quarantine for 5 days from the onset of symptoms if asymptomatic or with mild
316 to moderate symptoms (28). Our study showed that 17.5% of CPE positive samples were collected after 5
317 days from symptoms' onset. Our data indicates that it is not uncommon to recover infectious virus after 5
318 days from symptoms regardless of the vaccination status when patients have symptoms. Hence care
319 should be taken when making a release from quarantine decisions, especially when patients are showing
320 symptoms.

321 The limitations of our study include the retrospective nature of data collections which doesn't allow the
322 collection of baseline serum and respiratory IgG levels. Antibody neutralization assays and quantification
323 of viruses from clinical samples were not conducted as a part of this study nor were Omicron or Delta
324 specific IgG assays. Clinical data was compiled from patients tested and admitted to the Johns Hopkins
325 Health System. It is possible that patients who tested positive in our system then sought additional clinical
326 care, including admission, at a different hospital not affiliated with Johns Hopkins. While this could
327 artificially lower our admission rates, we have no reason to think that this was different in the first half of
328 December, when Delta was predominant, compared to the end of December.

329 In conclusion, we show a significant reduction in disease severity with Omicron when compared to Delta,
330 yet we show that Omicron associates with significant increases in infections of fully and booster
331 vaccinated individuals. It is important to note that admitted patients didn't show a significant difference in
332 the use of supplementary oxygen, ICU level care, stressing the importance of taking infection control
333 precautions and raising an awareness that Omicron infections should not be underestimated.

334 **Figure Legends**

335 Figure 1. SARS-CoV-2 positivity and variants trends November- December 2021. A) SARS-CoV-2 daily
336 positivity rates between November and December 2021 for both symptomatic and asymptomatic testing.
337 B) SARS-CoV-2 clade distribution between November and December 2021 relative to the 7 day rolling
338 average positives from Johns Hopkins system. C) Percent SARS-CoV-2 clades in November and
339 December 2021.

340 Figure 2. **Omicron and Delta variants cycle threshold (Ct) values in upper respiratory samples.**

341 A) Ct values of Omicron and Delta from all samples with available Ct values (N gene) stratified by
342 vaccination status. B) Ct values of Omicron and Delta from samples collected 5 days or less from the
343 onset of symptoms. For this analysis, samples from asymptomatic patients were not included. C) Ct
344 values of Omicron and Delta variants broken down by vaccination status and associated with days after
345 the onset of symptoms. Vax, fully vaccinated patients who didn't receive a booster dose (panel A only,
346 Vax includes boosted patients); Unvax, unvaccinated; boost, patients with booster dose.

347 Figure 3. **Recovery of infectious virus from respiratory samples of patients infected with Delta or**

348 **Omicron.** A) Percent CPE positives and negatives for Delta and Omicron; total, patients who received a
349 booster, fully vaccinated, and unvaccinated groups. Chi-squared test * $p < 0.05$, ** $p < 0.001$. B) Percent
350 CPE positives and negatives for Delta and Omicron; total, patients who received a booster, fully
351 vaccinated, and unvaccinated groups with Ct values less than 20. Chi-squared test * $p < 0.05$. C) Ct range
352 and medians of Delta and Omicron samples CPE positive and negative. One-way ANOVA * $p < 0.05$ D)
353 Distribution of sample collection time from each group in relation to days from the onset of symptoms.

354 Figure 4. **SARS-CoV-2 IgG levels in upper respiratory samples of infected vaccinated patients.**

355 Boost, patients with booster dose; Vax, fully vaccinated patients who didn't receive a booster dose.
356 Dashed lines demarcate the limit of borderline and negative ELISA results as specified per assay's
357 package insert.

358 **Figure S1. SARS-CoV-2 positivity and variants trends March 2020- December 2021.** SARS-CoV-2
359 clade distribution between March 2020 and December 2021 relative to the 7 day rolling average positives
360 from Johns Hopkins system.

361 **Figure S2. Omicron and Delta variants cycle threshold (Ct) values in upper respiratory samples.**

362 Ct values of Omicron and Delta variants broken down by vaccination status and associated with days after
363 the onset of symptoms in A) Unvax, unvaccinated, B) Vax, fully vaccinated patients who didn't receive a
364 booster, and C) boost, patients with booster vaccinations.

365 **Author contributions**

366 AF, RE, JS. data collection and data interpretation. CPM. data collection and analysis OA, JMN, NG, NS,
367 ML, MF, DCG. data acquisition and collection. AP. cell culture, scientific and manuscript revision. EK.
368 clinical data collection and scientific and manuscript revision. HHM. study design, data collection and
369 analysis, data interpretation, writing, fund acquisition.

370 **Declaration of interests**

371 We declare no relevant competing interests

372 **Data sharing**

373 Whole genome data were made available publicly and raw genomic data requests could be directed to
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492

493

	Omicron				Delta			
Total Patients	1121				910			
	Fully Vaccinated	With booster dose	Unvaccinated	Partially vaccinated	Fully Vaccinated	With booster dose	Unvaccinated	Partially vaccinated
Charts	461	134	488	38	282	83	518	27
Ct analysis	184	51	170	14	195	45	411	9
Patients with known days from the onset of symptoms	159	51	135	12	112	16	225	5
IgG	152	43	0	11	183	43	0	7
Cell culture (total)	55	76	83	5	52	45	56	0
Cell culture (Ct ≤ 20)	50	41	56		49	27	44	

494 Table 1. Charts and samples used for the study.

495

496

	November	December	November	December
SARS-CoV-2 (All testing)	726	7078	2.67	18.65
SARS-CoV-2 (Symptomatic only)	592	6236	5.70	25.50
SARS-CoV-2 whole genome sequencing			%	%
Total sequenced	552	2179	76.03	30.79
21I (Delta)	24	36	4.35	1.65
21J (Delta)	468	791	84.78	36.30
21K (Omicron)	1	1208	0.18	55.44
Other or low QC	59	144	10.69	6.61

497

498 Table 2. SARS-CoV-2 testing and positivity in November and December 2021 and total sequenced.

499 SARS-CoV-2 testing is performed by different molecular assays as detailed in the methods section and

500 the total tested reflects testing by all assays. Sequence counts were up to the time of writing this

501 manuscript.

502

503

	Omicron		Delta		
Total	1121		910		P
Collection range	11/25/2021 to 12/31/21	%	11/22/21 to 12/31/21	%	
Fully Vaccinated (No booster)	461	41.1	282	31.0	< 0.00001
Booster	134	12.0	83	9.1	0.04
Partially vaccinated	38	3.4	27	3.0	
Symptomatic	1037	92.5	837	92.0	
Asymptomatic	84	7.5	72	7.9	0.7
Age, median (stdev)	32 (18.5)		35 (23.4)		0.001
Comorbidities					
Hypertension	242	21.6	283	31.1	< 0.00001
Pregnancy	84	7.5	53	5.8	0.2
Lung Disease	225	20.1	206	22.6	0.2
Kidney Disease	69	6.2	126	13.8	< 0.00001
Immunosuppression	121	10.8	156	17.1	0
Diabetes	106	9.5	138	15.2	0.0001
Heart Failure	29	2.6	62	6.8	< 0.00001
Atrial Fibrillation	17	1.5	46	5.1	< 0.00001
Smoker	115	10.3	142	15.6	0.0004
Cerebrovascular Disease	46	4.1	69	7.6	0.001
Cancer	251	22.4	208	22.9	0.8
Coronary Artery Disease	85	7.6	151	16.6	< 0.00001
Outcome					
COVID Related Admission	34	3.0	126	13.8	< 0.00001
ICU Level Care	6	0.5	32	3.5	< 0.00001
Died	1	0.1	10	1.1	0.004

504

505 Table 3. Clinical and metadata of the Omicron and Delta infected patients. Statistics for ages were

506 calculated by *t* test and all other statistics were calculated by Chi-squared test.

507

	Omicron				p	Delta				p
	Fully vaccinated including boosted		Unvaccinated			Fully vaccinated including boosted		Unvaccinated		
Total	595	%	488	%		365	%	518	%	
Symptomatic	547	91.93	454	93.03	0.60	329	90.14	385	74.32	0.3
Asymptomatic	48	8.07	34	6.97		36	9.86	32	6.18	
Age, median (stdev)	33 (18.2)		31.8 (18.8)		0.02	37 (23.2)		34 (23.6)		0.09
Comorbidities										
Hypertension	127	21.34	109	22.34	0.70	126	34.52	149	28.76	0.07
Pregnancy	50	8.40	30	6.15	0.20	24	6.58	27	5.21	0.50
Lung Disease	107	17.98	111	22.75	0.06	84	23.01	115	22.20	0.80
Kidney Disease	36	6.05	30	6.15	1.00	50	13.70	72	13.90	1.00
Immunosuppression	66	11.09	51	10.45	0.80	65	17.81	86	16.60	0.60
Diabetes	60	10.08	43	8.81	0.50	60	16.44	74	14.29	0.40
Heart Failure	15	2.52	13	2.66	1.00	30	8.22	30	5.79	0.20
Atrial Fibrillation	9	1.51	8	1.64	1.00	18	4.93	27	5.21	0.90
Smoker	64	10.76	45	9.22	0.40	53	14.52	84	16.22	0.50
Cerebrovascular Disease	24	4.03	20	4.10	1.00	26	7.12	41	7.92	0.70
Cancer	142	23.87	103	21.11	0.30	91	24.93	112	21.62	0.30
Coronary Artery Disease	51	8.57	31	6.35	0.20	60	16.44	88	16.99	0.80
Outcome										
COVID Related Admission	23	3.87	9	1.84	0.07	43	11.78	80	15.44	0.13
ICU Level Care	3	0.50	2	0.41	1.00	8	2.19	24	4.63	0.07
Died	0	0.00	1	0.20	0.40	4	1.10	6	1.16	1.00

Table 4. Clinical and metadata of Delta and Omicron vaccinated and unvaccinated patients. Statistics for ages were calculated by t test and all other statistics were calculated by Chi-squared test. stdev; standard deviation.

	Omicron						p (boosted to unvaccinated, Omicron)	Delta						p (boosted to unvaccinated, Delta)	P (boosted Omicron to boosted Delta)
	Fully Vaccinated		Boosted		Unvaccinated			Fully Vaccinated		Boosted		Unvaccinated			
Total	461	%	134	%	488	%		282	%	83	%	518	%		
Symptomatic	420	91.1	127	94.8	454	93.0		255	90.4	74	89.2	385	74.3		
Asymptomatic	41	8.9	7	5.2	34	7.0		27	9.6	9	10.8	32	6.2		
Age (median)	32 (17.9)		34 (19.3)		31.8 (18.8)			36 (22.6)		39 (25.1)		34 (23.6)			
Comorbidities															
Hypertension	95	20.6	32	23.9	109	22.3	0.7	98	34.8	28	33.7	149	28.8	0.3	0.12
Pregnancy	37	8.0	13	9.7	30	6.1	0.2	21	7.4	3	3.6	27	5.2	0.8	0.114
Lung Disease	85	18.4	22	16.4	111	22.7	0.1	62	22.0	22	26.5	115	22.2	0.4	0.08
Kidney Disease	30	6.5	6	4.5	30	6.1	0.5	34	12.1	16	19.3	72	13.9	0.2	0.0008
Immunosuppression	51	11.1	15	11.2	51	10.5	0.9	45	16.0	20	24.1	86	16.6	0.1	0.01
Diabetes	45	9.8	15	11.2	43	8.8	0.4	43	15.2	17	20.5	74	14.3	0.1	0.07
Heart Failure	14	3.0	1	0.7	13	2.7	0.3	24	8.5	6	7.2	30	5.8	0.6	0.01
Atrial Fibrillation	7	1.5	2	1.5	8	1.6	1.0	12	4.3	6	7.2	27	5.2	0.4	0.01
Smoker	54	11.7	10	7.5	45	9.2	0.6	42	14.9	11	13.3	84	16.2	0.6	0.0002
Cerebrovascular Disease	20	4.3	4	3.0	20	4.1	0.8	21	7.4	5	6.0	41	7.9	0.7	0.3
Cancer	104	22.6	38	28.4	103	21.1	0.1	65	23.0	26	31.3	112	21.6	0.1	0.6
Coronary Artery Disease	41	8.9	10	7.5	31	6.4	0.7	45	16.0	15	18.1	88	17.0	0.8	0.03
Outcome															
COVID Related Admission	20	4.3	3	2.2	9	1.8	0.7	30	10.6	13	15.7	80	15.4	1.0	0.0007
ICU Level Care	2	0.4	1	0.7	2	0.4	0.5	6	2.1	2	2.4	24	4.6	0.6	0.6
Died	0	0.0	0	0.0	1	0.2	1.0	2	0.7	2	2.4	6	1.2	0.3	0.1

Table S1. Clinical and metadata of Delta and Omicron vaccinated and unvaccinated patients. Statistics for ages were calculated by t test and all other statistics were calculated by Chi-squared test. stdev; standard deviation.

	Omicron admissions		Delta admissions		
Total	34	%	126	%	p value
Fully vaccinated	20	58.8	30	23.8	0.0001
Booster	3	8.8	13	10.3	1
Median age (stdev)	48 (25.8)		58 (21.1)		0.130
Comorbidities					
Hypertension	17	50.0	82	65.1	0.11
Pregnancy	4	11.8	4	3.2	0.06
Lung Disease	11	32.4	38	30.2	0.8
Kidney Disease	18	52.9	61	48.4	0.7
Immunosuppression	20	58.8	59	46.8	0.24
Diabetes	13	38.2	51	40.5	0.84
Heart Failure	10	29.4	25	19.8	0.24
Atrial Fibrillation	7	20.6	27	21.4	1
Smoker	8	23.5	32	25.4	1
Cerebrovascular Disease	7	20.6	26	20.6	1
Cancer	12	35.3	46	36.5	1
Coronary Artery Disease	15	44.1	64	50.8	0.56
Outcome					
Supplemental O2	23	67.6	92	73.0	0.5
ICU Level Care	6	17.6	32	25.4	0.5
Died	1	2.9	10	7.9	0.5

508

509 Table 5. Clinical and metadata of Delta and Omicron admitted patients. Statistics for ages were calculated by

510 t test and all other statistics were calculated by Chi-squared test.

511

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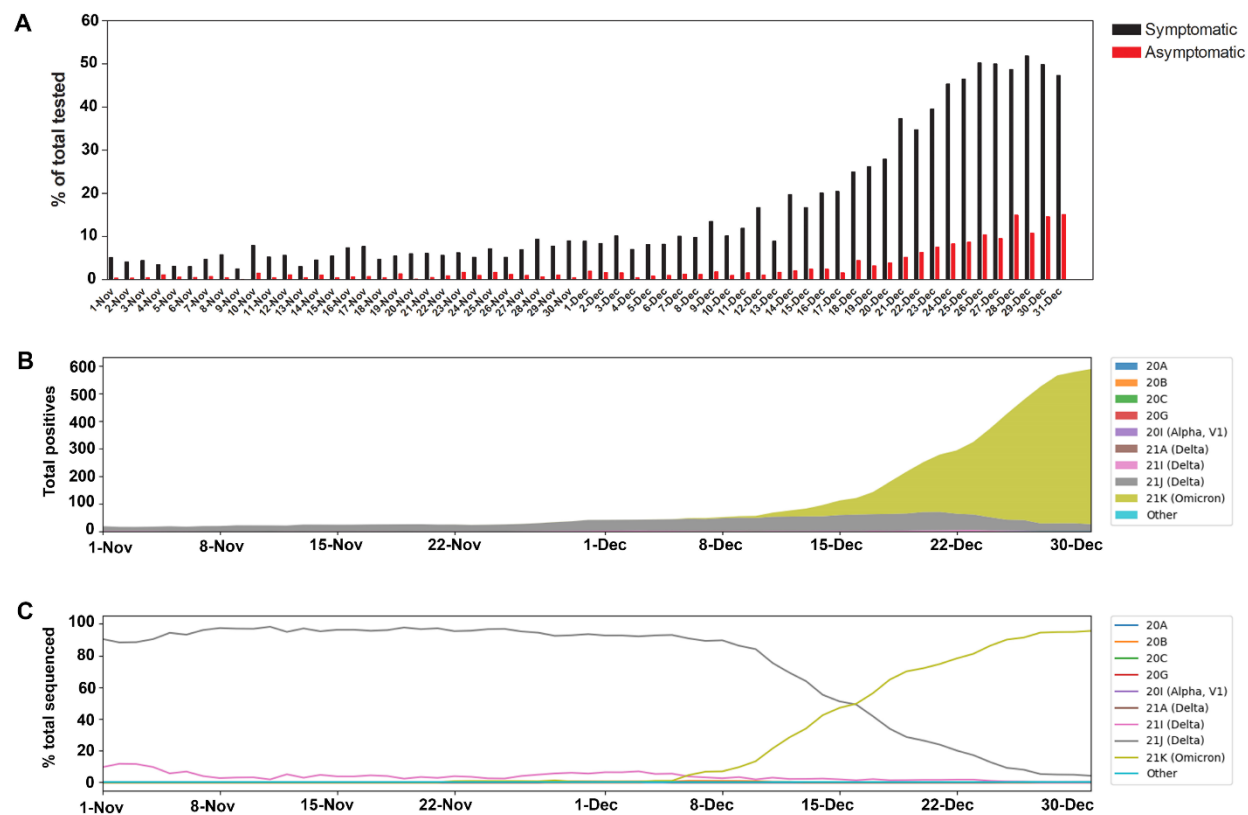
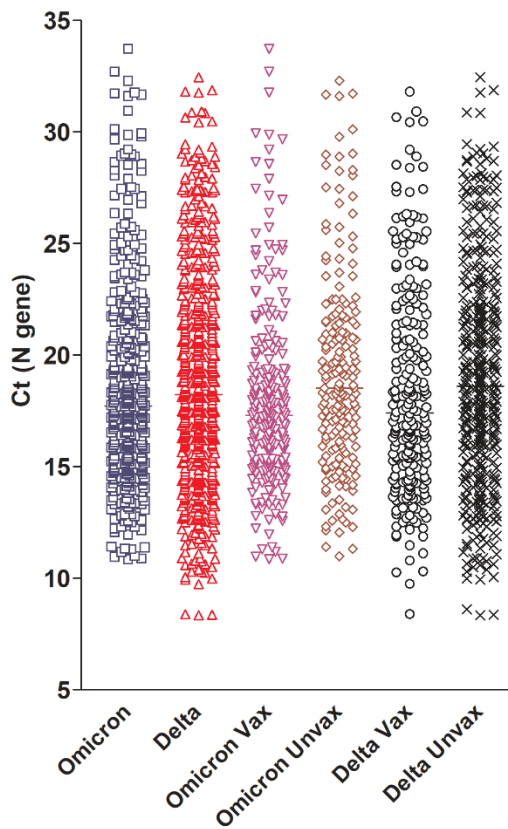
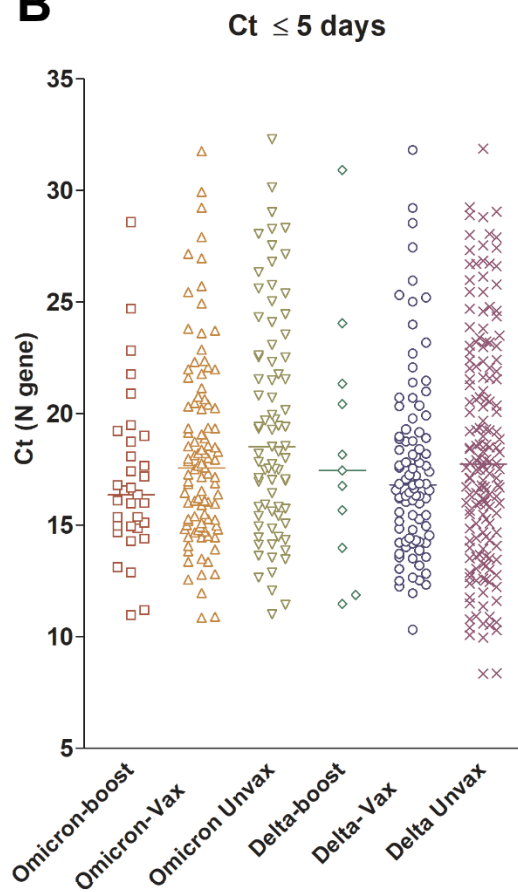


Figure 1. SARS-CoV-2 positivity and variants trends November- December 2021. A) SARS-CoV-2 daily positivity rates between November and December 2021 for both symptomatic and asymptomatic testing. B) SARS-CoV-2 clade distribution between November and December 2021 relative to the 7 day rolling average positives from Johns Hopkins system. C) Percent SARS-CoV-2 clades in November and December 2021.

A



B



C

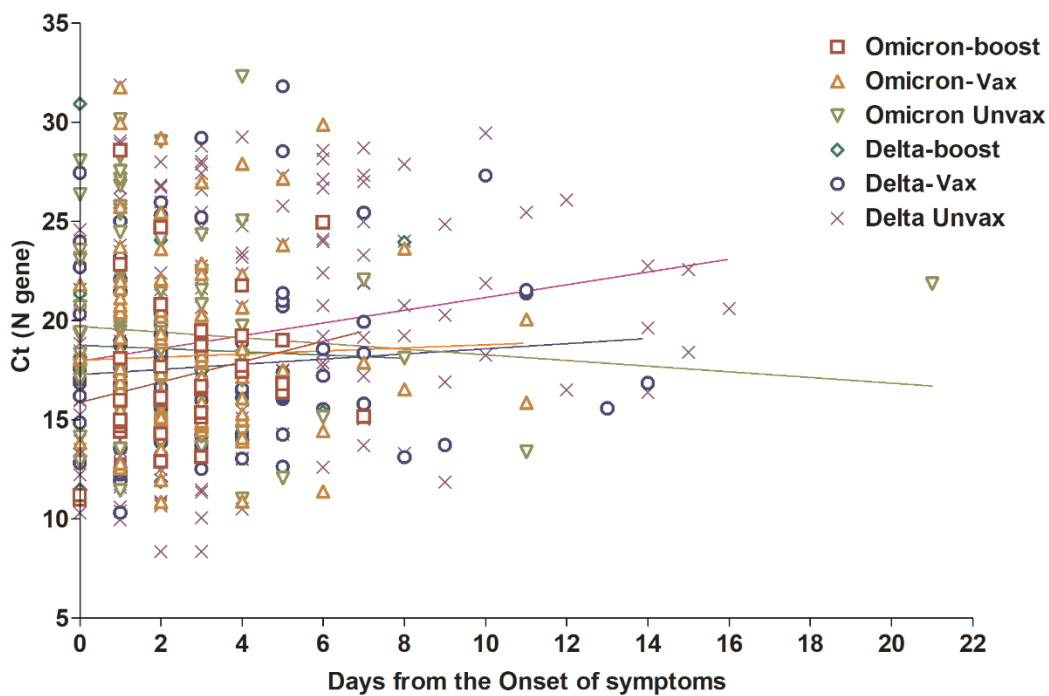


Figure 2. Omicron and Delta variants cycle threshold (Ct) values in upper respiratory samples.

A) Ct values of Omicron and Delta from all samples with available Ct values (N gene) stratified by vaccination status. B) Ct values of Omicron and Delta from samples collected 5 days or less from the onset of symptoms. For this analysis, samples from asymptomatic patients were not included. C) Ct values of Omicron and Delta variants broken down by vaccination status and associated with days after the onset of symptoms. Vax, fully vaccinated patients who didn't receive a booster dose (panel A only, Vax includes boosted patients); Unvax, unvaccinated; boost, patients with booster dose.

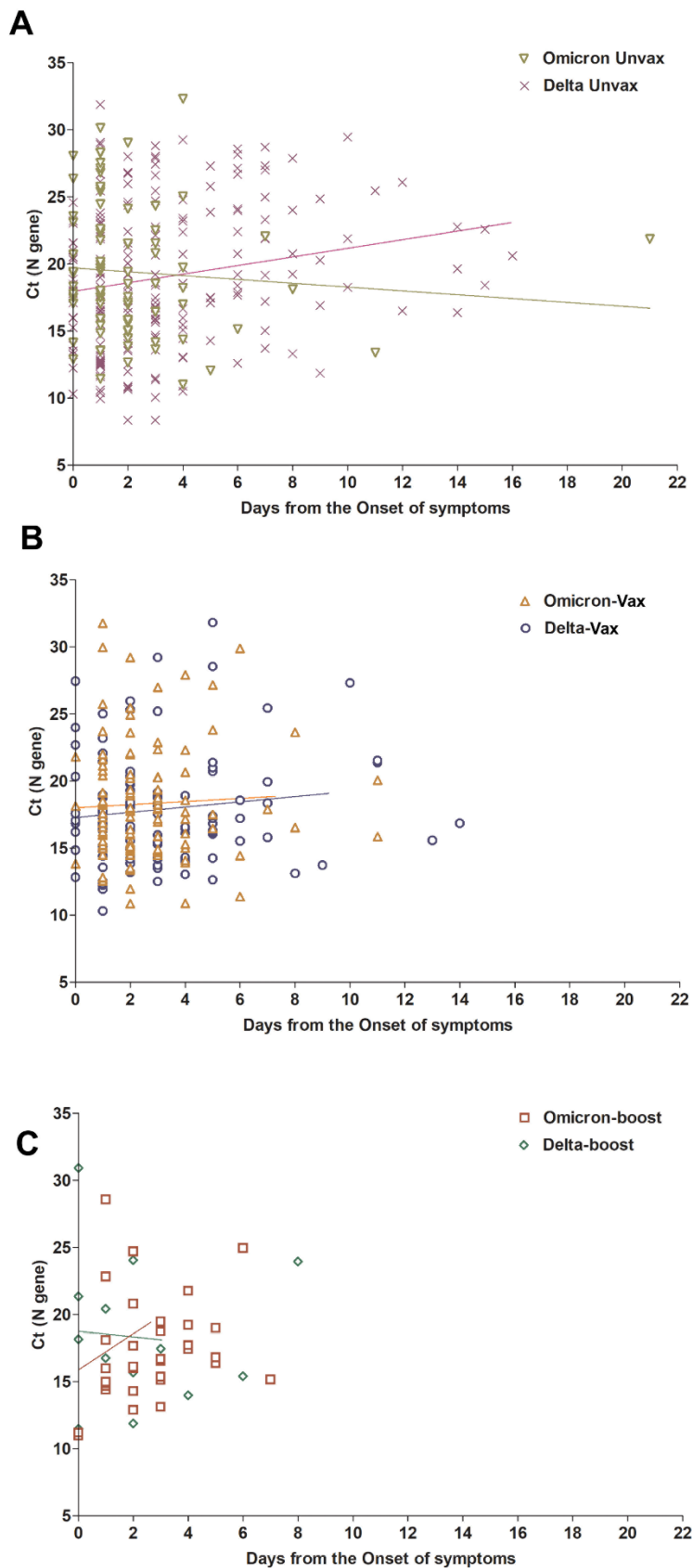


Figure S2. Omicron and Delta variants cycle threshold (Ct) values in upper respiratory samples.

Ct values of Omicron and Delta variants broken down by vaccination status and associated with days after the onset of symptoms in A) Unvax, unvaccinated, B) Vax, fully vaccinated patients who didn't receive a booster, and C) boost, patients with booster vaccinations.

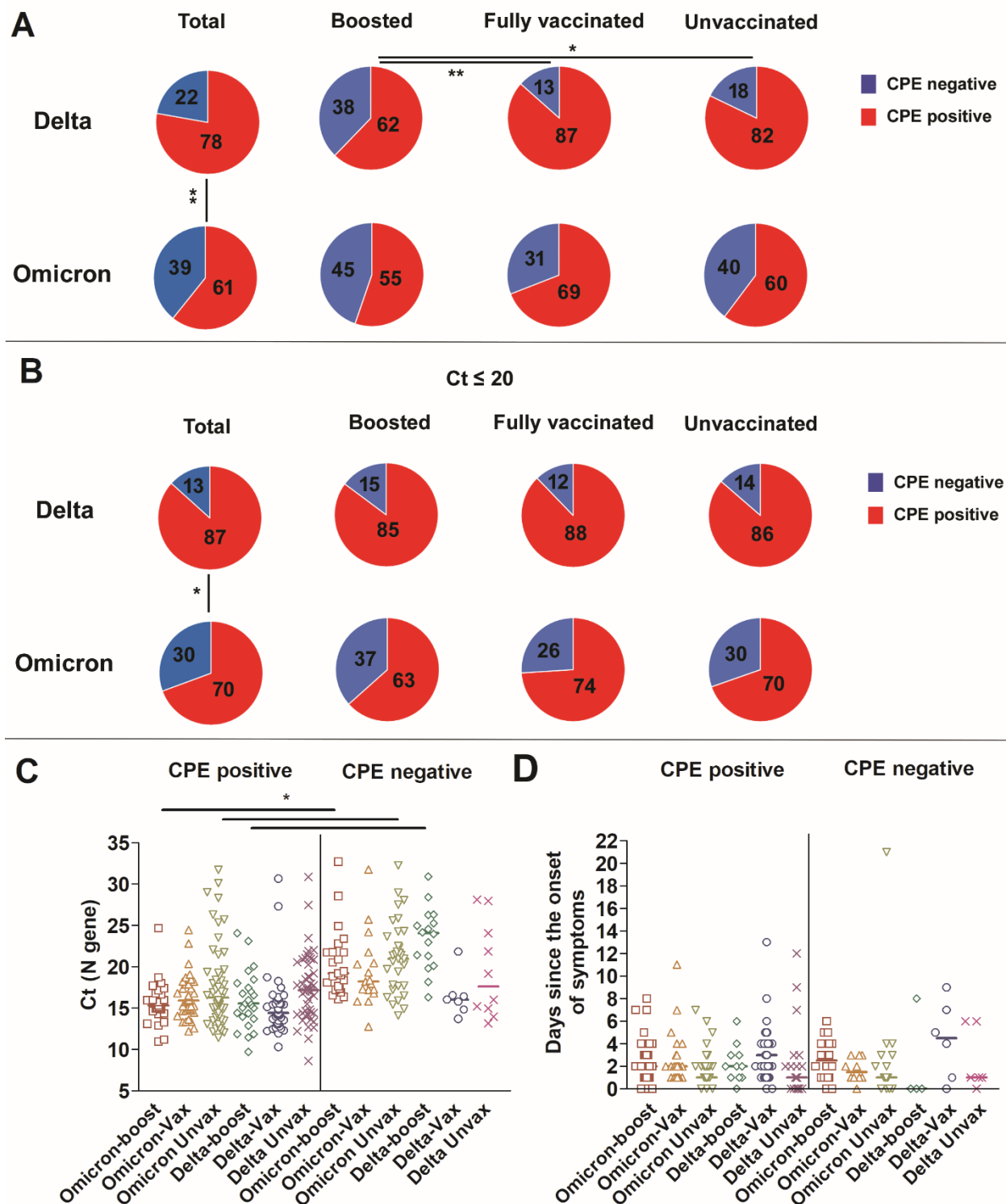


Figure 3. Recovery of infectious virus from respiratory samples of patients infected with Delta or Omicron. A) Percent CPE positives and negatives for Delta and Omicron; total, patients who received a booster, fully vaccinated, and unvaccinated groups. Chi-squared test * $p < 0.05$, ** $p < 0.001$. B) Percent

CPE positives and negatives for Delta and Omicron; total, patients who received a booster, fully vaccinated, and unvaccinated groups with Ct values less than 20. Chi-squared test * $p < 0.05$. C) Ct range and medians of Delta and Omicron samples CPE positive and negative. One-way ANOVA * $p < 0.05$ D) Distribution of sample collection time from each group in relation to days from the onset of symptoms.

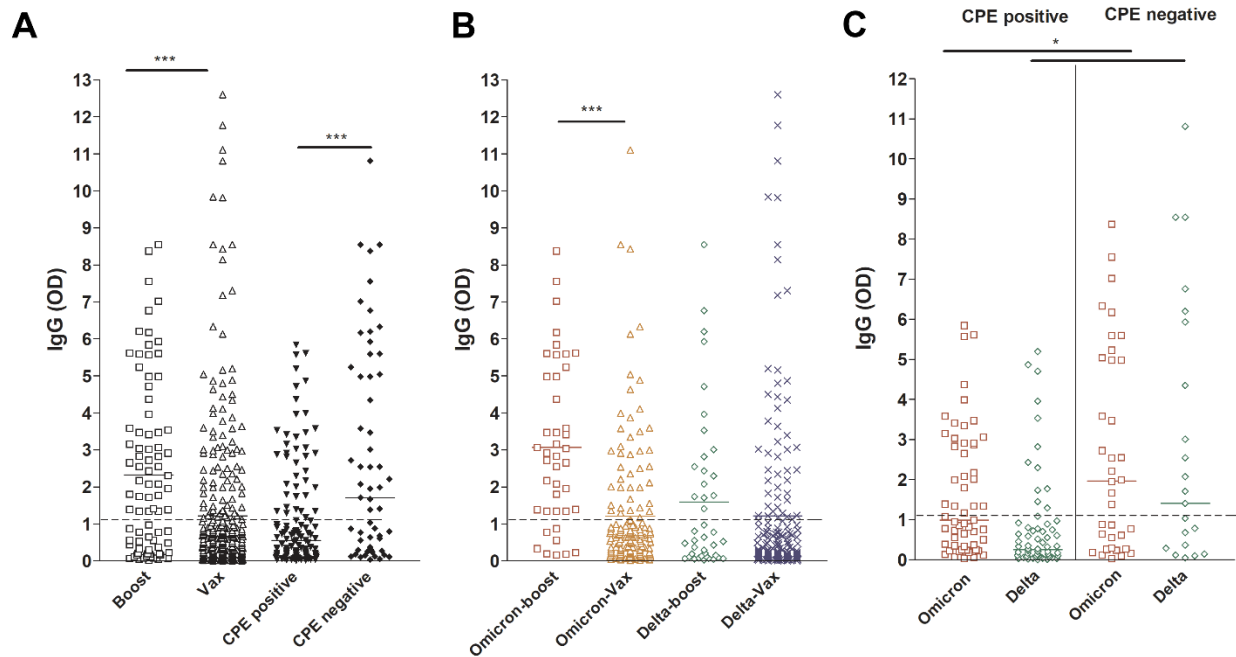


Figure 4. SARS-CoV-2 IgG levels in upper respiratory samples of infected vaccinated patients. Boost, patients with booster dose; Vax, fully vaccinated patients who didn't receive a booster dose. Dashed lines demarcate the limit of borderline and negative ELISA results as specified per assay's package insert.

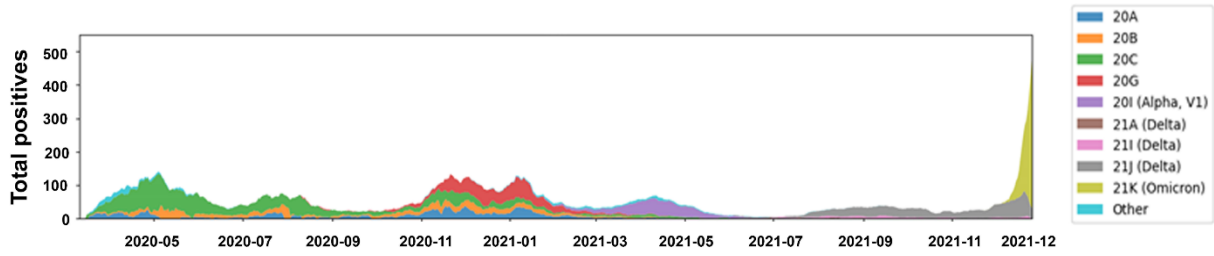


Figure S1. SARS-CoV-2 positivity and variants trends March 2020- December 2021. SARS-CoV-2 clade distribution between March 2020 and December 2021 relative to the 7 day rolling average positives from Johns Hopkins system.

	Omicron				Delta			
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21I (Delta)	24	36	4.35	1.65
21J (Delta)	468	791	84.78	36.30
21K (Omicron)	1	1208	0.18	55.44
Other or low QC	59	144	10.69	6.61

Table 2. SARS-CoV-2 testing and positivity in November and December 2021 and total sequenced. SARS-CoV-2 testing is performed by different molecular assays as detailed in the methods section and the total tested reflects testing by all assays. Sequence counts were up to the time of writing this manuscript.

	Omicron		Delta		
Total	1121		910		P
Collection range	11/25/2021 to 12/31/21	%	11/22/21 to 12/31/21	%	
Fully Vaccinated (No booster)	461	41.1	282	31.0	< 0.00001
Booster	134	12.0	83	9.1	0.04
Partially vaccinated	38	3.4	27	3.0	
Symptomatic	1037	92.5	837	92.0	
Asymptomatic	84	7.5	72	7.9	0.7
Age, median (stdev)	32 (18.5)		35 (23.4)		0.001
Comorbidities					
Hypertension	242	21.6	283	31.1	< 0.00001
Pregnancy	84	7.5	53	5.8	0.2
Lung Disease	225	20.1	206	22.6	0.2
Kidney Disease	69	6.2	126	13.8	< 0.00001
Immunosuppression	121	10.8	156	17.1	0
Diabetes	106	9.5	138	15.2	0.0001
Heart Failure	29	2.6	62	6.8	< 0.00001
Atrial Fibrillation	17	1.5	46	5.1	< 0.00001
Smoker	115	10.3	142	15.6	0.0004
Cerebrovascular Disease	46	4.1	69	7.6	0.001
Cancer	251	22.4	208	22.9	0.8
Coronary Artery Disease	85	7.6	151	16.6	< 0.00001
Outcome					
COVID Related Admission	34	3.0	126	13.8	< 0.00001
ICU Level Care	6	0.5	32	3.5	< 0.00001
Died	1	0.1	10	1.1	0.004

Table 3. Clinical and metadata of the Omicron and Delta infected patients. Statistics for ages were calculated by *t* test and all other statistics were calculated by Chi-squared test.

	Omicron				p	Delta				p
	Fully vaccinated including boosted		Unvaccinated			Fully vaccinated including boosted		Unvaccinated		
Total	595	%	488	%		365	%	518	%	
Symptomatic	547	91.93	454	93.03	0.60	329	90.14	385	74.32	0.3
Asymptomatic	48	8.07	34	6.97		36	9.86	32	6.18	
Age, median (stdev)	33 (18.2)		31.8 (18.8)		0.02	37 (23.2)		34 (23.6)		0.09
Comorbidities										
Hypertension	127	21.34	109	22.34	0.70	126	34.52	149	28.76	0.07
Pregnancy	50	8.40	30	6.15	0.20	24	6.58	27	5.21	0.50
Lung Disease	107	17.98	111	22.75	0.06	84	23.01	115	22.20	0.80
Kidney Disease	36	6.05	30	6.15	1.00	50	13.70	72	13.90	1.00
Immunosuppression	66	11.09	51	10.45	0.80	65	17.81	86	16.60	0.60
Diabetes	60	10.08	43	8.81	0.50	60	16.44	74	14.29	0.40
Heart Failure	15	2.52	13	2.66	1.00	30	8.22	30	5.79	0.20
Atrial Fibrillation	9	1.51	8	1.64	1.00	18	4.93	27	5.21	0.90
Smoker	64	10.76	45	9.22	0.40	53	14.52	84	16.22	0.50
Cerebrovascular Disease	24	4.03	20	4.10	1.00	26	7.12	41	7.92	0.70
Cancer	142	23.87	103	21.11	0.30	91	24.93	112	21.62	0.30
Coronary Artery Disease	51	8.57	31	6.35	0.20	60	16.44	88	16.99	0.80
Outcome										
COVID Related Admission	23	3.87	9	1.84	0.07	43	11.78	80	15.44	0.13
ICU Level Care	3	0.50	2	0.41	1.00	8	2.19	24	4.63	0.07
Died	0	0.00	1	0.20	0.40	4	1.10	6	1.16	1.00

Table 4. Clinical and metadata of Delta and Omicron vaccinated and unvaccinated patients. Statistics for ages were calculated by t test and all other statistics were calculated by Chi-squared test. stdev; standard deviation.

	Omicron						p (boosted to unvaccinated, Omicron)	Delta						p (boosted to unvaccinated, Delta)	P (boosted Omicron to boosted Delta)
	Fully Vaccinated		Boosted		Unvaccinated			Fully Vaccinated		Boosted		Unvaccinated			
Total	461	%	134	%	488	%		282	%	83	%	518	%		
Symptomatic	420	91.1	127	94.8	454	93.0		255	90.4	74	89.2	385	74.3		
Asymptomatic	41	8.9	7	5.2	34	7.0		27	9.6	9	10.8	32	6.2		
Age (median)	32 (17.9)		34 (19.3)		31.8 (18.8)			36 (22.6)		39 (25.1)		34 (23.6)			
Comorbidities															
Hypertension	95	20.6	32	23.9	109	22.3	0.7	98	34.8	28	33.7	149	28.8	0.3	0.12
Pregnancy	37	8.0	13	9.7	30	6.1	0.2	21	7.4	3	3.6	27	5.2	0.8	0.114
Lung Disease	85	18.4	22	16.4	111	22.7	0.1	62	22.0	22	26.5	115	22.2	0.4	0.08
Kidney Disease	30	6.5	6	4.5	30	6.1	0.5	34	12.1	16	19.3	72	13.9	0.2	0.0008
Immunosuppression	51	11.1	15	11.2	51	10.5	0.9	45	16.0	20	24.1	86	16.6	0.1	0.01
Diabetes	45	9.8	15	11.2	43	8.8	0.4	43	15.2	17	20.5	74	14.3	0.1	0.07
Heart Failure	14	3.0	1	0.7	13	2.7	0.3	24	8.5	6	7.2	30	5.8	0.6	0.01
Atrial Fibrillation	7	1.5	2	1.5	8	1.6	1.0	12	4.3	6	7.2	27	5.2	0.4	0.01
Smoker	54	11.7	10	7.5	45	9.2	0.6	42	14.9	11	13.3	84	16.2	0.6	0.0002
Cerebrovascular Disease	20	4.3	4	3.0	20	4.1	0.8	21	7.4	5	6.0	41	7.9	0.7	0.3
Cancer	104	22.6	38	28.4	103	21.1	0.1	65	23.0	26	31.3	112	21.6	0.1	0.6
Coronary Artery Disease	41	8.9	10	7.5	31	6.4	0.7	45	16.0	15	18.1	88	17.0	0.8	0.03
Outcome															
COVID Related Admission	20	4.3	3	2.2	9	1.8	0.7	30	10.6	13	15.7	80	15.4	1.0	0.0007
ICU Level Care	2	0.4	1	0.7	2	0.4	0.5	6	2.1	2	2.4	24	4.6	0.6	0.6
Died	0	0.0	0	0.0	1	0.2	1.0	2	0.7	2	2.4	6	1.2	0.3	0.1

Table S1. Clinical and metadata of Delta and Omicron vaccinated and unvaccinated patients. Statistics for ages were calculated by t test and all other statistics were calculated by Chi-squared test. stdev; standard deviation.

	Omicron admissions		Delta admissions		p value
		%		%	
Total	34		126		
Fully vaccinated	20	58.8	30	23.8	0.0001
Booster	3	8.8	13	10.3	1
Median age (stdev)	48 (25.8)		58 (21.1)		0.130
Comorbidities					
Hypertension	17	50.0	82	65.1	0.11
Pregnancy	4	11.8	4	3.2	0.06
Lung Disease	11	32.4	38	30.2	0.8
Kidney Disease	18	52.9	61	48.4	0.7
Immunosuppression	20	58.8	59	46.8	0.24
Diabetes	13	38.2	51	40.5	0.84
Heart Failure	10	29.4	25	19.8	0.24
Atrial Fibrillation	7	20.6	27	21.4	1
Smoker	8	23.5	32	25.4	1
Cerebrovascular Disease	7	20.6	26	20.6	1
Cancer	12	35.3	46	36.5	1
Coronary Artery Disease	15	44.1	64	50.8	0.56
Outcome					
Supplemental O2	23	67.6	92	73.0	0.5
ICU Level Care	6	17.6	32	25.4	0.5
Died	1	2.9	10	7.9	0.5

Table 5. Clinical and metadata of Delta and Omicron admitted patients. Statistics for ages were calculated by t test and all other statistics were calculated by Chi-squared test.