

Covid-19 vaccine-induced antibodies are attenuated and decay rapidly in infliximab treated patients

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Brief Communication

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1 **Covid-19 vaccine-induced antibodies are attenuated and decay rapidly in infliximab**
2 **treated patients**

3

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52 **Key words:** SARS-CoV-2, immune-mediated inflammatory diseases, inflammatory bowel disease,
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55

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57

58 **Abstract**

59 To inform healthcare policy for immunosuppressed patients there is a need to define SARS-CoV-2
60 vaccine responses. Here we report SARS-CoV-2 vaccine-induced antibody and T cell responses in
61 patients treated with anti-tumour necrosis factor (anti-TNF), a commonly used biologic in
62 inflammatory diseases, compared to patients treated with vedolizumab, a gut-specific antibody
63 targeting integrin $\alpha4\beta7$ that does not impair systemic immunity. In anti-TNF recipients, the
64 magnitude of anti-SARS-CoV2 antibodies was reduced five-fold, and rapidly decayed towards the
65 seroconversion threshold by 14 weeks after second dose of vaccine. In contrast, anti-SARS-CoV-2
66 antibodies were sustained up to 16 weeks in vedolizumab-treated patients. Anti-SARS-CoV2
67 antibody decay was not observed in vaccinated patients previously infected with SARS-CoV-2. T cell
68 responses were absent in one-fifth of anti-TNF and vedolizumab-treated patients after a second
69 dose of either vaccine. Our data have important implications for anti-TNF recipients, including the
70 need for vaccine prioritization, booster doses, and social distancing strategies.

71

72 **Main**

73 Vaccination programmes have reduced SARS-CoV-2 infections, transmission, hospitalisations and
74 deaths. Whether the durability of vaccine responses will stem further waves of disease, including the
75 spread of the delta variant is controversial. Public health bodies in the United Kingdom¹ and other
76 countries have committed to a booster dose of vaccines later this year; however, in the USA and
77 Europe, the Centre for Disease Control and prevention (CDC) and the European Medicines Agency
78 (EMA) are waiting for data on infection breakthrough in the vaccinated population before further
79 guidance is issued^{2,3}. Specific recommendations will be needed for the large minority of the
80 population who may for various reasons mount suboptimal immune responses.

81

82 Patients treated with immunosuppressive drugs were excluded from the registration trials of the
83 SARS-CoV-2 vaccines and real-world effectiveness data are limited. Drugs targeting tumor necrosis

84 factor (TNF), such as infliximab, are the most frequently prescribed biological therapies used in the
85 treatment of immune-mediated inflammatory disorders (IMIDs). Observational studies indicate that
86 most patients with inflammatory bowel disease (IBD), an archetypal IMID, mount serological
87 responses following SARS-CoV-2 vaccines; although most were underpowered to discern the impact
88 of specific drugs, including immunomodulators (azathioprine, mercaptopurine, and methotrexate)
89 and/or biologic therapies^{4,5}. We recently reported, however, that antibody responses following
90 SARS-CoV-2 infection or a single-dose of either the BNT162b2 or ChAdOx1 nCoV-19 SARS-CoV-2
91 vaccines are impaired in anti-TNF treated patients^{6,7}. We hypothesised here that antibody and T cell
92 responses following the second doses of BNT162b2 or ChAdOx1 nCoV-19 vaccines would be
93 attenuated and less durable in infliximab-treated patients.

94

95 CLARITY IBD is a 40-week prospective observational study investigating immune responses to SARS-
96 CoV2 infection and vaccination in IBD patients⁶. We measured anti-SARS-CoV-2 spike (S) receptor
97 binding domain (RBD) antibodies in patients with IBD treated with either infliximab, or vedolizumab,
98 a gut-specific antibody targeting integrin $\alpha 4\beta 7$, that does not impair systemic immunity. We report
99 data from 2052 infliximab- and 925 vedolizumab-treated participants without evidence of prior
100 SARS-CoV-2 infection, who had received uninterrupted biologic therapy since recruitment and had
101 an antibody test performed between 14 and 70 days after a second dose of either the BNT162b2 or
102 ChAdOx1 nCoV-19 SARS-CoV-2 vaccines. Participant characteristics are shown in Supplementary
103 Table 2. Secondary outcome analyses are also presented for 283 infliximab- and 137 vedolizumab-
104 treated patients who had had PCR-confirmed SARS-CoV-2 infection prior to vaccination.

105 Seroconversion was defined as an anti-S RBD antibody concentration ≥ 15 U/mL, a threshold
106 associated with viral neutralization of $\geq 20\%$ with a positive predictive value of 99.10 % (95% CI:
107 97.74-99.64)⁷. Anti-SARS-CoV-2 antibody non-persistence was defined as the time to a four-fold
108 decrease in anti-S RBD antibodies. Anti-S RBD antibody levels were compared with samples from 605
109 fully vaccinated adult participants from the Virus Watch study, a community cohort of 10,000

110 individuals representative of the UK population of England and Wales⁸. T cell responses to first and
111 second doses of either vaccine are reported in 225 infliximab- and 76 vedolizumab-treated patients
112 without prior infection. T cell responses were measured using interferon- γ ELISpot assays following
113 stimulation of peripheral blood mononuclear cells (PBMC) with a pool of SARS-CoV-2 spike peptides.
114
115 Geometric mean [geometric SD] anti-S RBD antibody concentrations were significantly lower in
116 patients treated with infliximab than vedolizumab, following a second dose of both the BNT162b2
117 (547.5 U/mL [6.3] vs 3980.4 U/mL [5.5], $p < 0.0001$) and ChAdOx1 nCoV-19 (189.3 U/mL [5.1] vs
118 781.5 U/mL [3.6], $p < 0.0001$) vaccines (Fig. 1a). Multivariable linear regression analyses in patients
119 without prior SARS-CoV-2 infection confirmed that antibody concentrations were attenuated
120 between four and five-fold in infliximab-, compared with vedolizumab-, treated patients in
121 participants who received either the BNT162b2 (fold change [FC] 0.17 [95% CI 0.13, 0.22], $p < 0.0001$)
122 or ChAdOx1 nCoV-19 ([FC] 0.25 [95% CI 0.21, 0.30], $p < 0.0001$) vaccines. Age ≥ 60 years, thiopurine or
123 methotrexate use in patients who received the BNT162b2, but not the ChAdOx1 nCoV-19 vaccine,
124 current smoking and Crohn's disease were also independently associated with lower anti-S RBD
125 antibody concentrations. Conversely, non-white ethnicity was associated with higher antibody
126 concentrations when data from both vaccines were taken together (Extended Data Fig. 1).
127
128 Seroconversion rates after the first vaccine dose were lower in infliximab- compared to
129 vedolizumab-treated patients. However, administration of a second dose of vaccine triggered a
130 >100 -fold increase in antibody concentrations with the BNT162b2 vaccine and >30 -fold with the
131 ChAdOx1 nCoV-19 vaccine in both treatment groups (Fig. 1a). More infliximab- than vedolizumab-
132 treated patients failed to seroconvert after their second vaccine dose (6.1% vs 1.3%, $p < 0.0001$).
133
134 Following two doses of either vaccine, anti-S RBD antibodies were sustained to more than 16 weeks
135 in patients treated with vedolizumab (Fig. 1b) and were not different to those observed in

136 participants in the Virus Watch community cohort (Extended Data Fig 2): however, in infliximab-
137 treated patients geometric mean concentrations decayed towards the seroconversion threshold,
138 defined as anti-S RBD ≥ 15 U/mL, by 18 and 14 weeks after a second dose of the BNT162b2 and
139 ChAdOx1 nCoV-19 vaccines, respectively (Fig. 1b). Cox proportional regression analysis
140 demonstrated that infliximab compared to vedolizumab treatment was independently associated
141 with anti-SARS-CoV-2 antibody non-persistence (hazard ratio (HR) 2.95 (95% CI 2.17 to 4.02), $p <$
142 0.0001) (Extended Data Fig. 3).

143

144 Amongst patients with SARS-CoV-2 infection prior to vaccination, geometric mean [SD] anti-S RBD
145 antibody concentrations were lower in infliximab- compared with vedolizumab-treated patients
146 after a second dose of BNT162b2 (1811.3 U/mL [3.5] vs 10079.6 U/mL [2.2], $p < 0.0001$) and
147 ChAdOx1 nCoV-19 (575.1 U/mL [5.2] vs 2595.1 [3.8] $p < 0.0001$) vaccines. In all patients, antibody
148 concentrations following vaccination were higher than those observed in patients without prior
149 infection (Fig. 1b). Irrespective of vaccine or biologic type, anti-S RBD antibodies were maintained to
150 more than 14 weeks.

151

152 There were no significant differences in the magnitude of anti-spike T cell responses observed in
153 infliximab- compared with vedolizumab-treated patients after one or two doses of either vaccine
154 (Fig. 2a). The proportion of patients failing to mount detectable T cell responses were similar in both
155 groups (infliximab 19.6% vs. vedolizumab 19.2%). For recipients of one and two doses of BNT162b2
156 vaccine there was a modest positive correlation between T cell responses and antibody
157 concentration. This association was not observed in recipients following either dose of the ChAdOx1
158 nCoV-19 vaccine (Fig. 2b). When T cell responses were ranked by magnitude of antibody responses,
159 most patients who did not mount an antibody response had a detectable T cell response (Extended
160 Data Fig. 4). In addition to the uncoupling of the T cell and antibody responses demonstrated, this
161 analysis emphasised that about one fifth made no T cell responses irrespective of vaccine used and a

162 minority of individuals carry neither detectable antibody nor T cell responses after 2 doses of vaccine
163 (Fig. 2b, Extended Data Fig. 4).

164

165 As many countries enter the third wave of COVID-19, our data have important implications for
166 millions of patients treated with anti-TNF drugs, who could remain susceptible to infection even
167 after vaccination. However, the sustained antibody responses observed in vaccinated patients with
168 prior infection indicates that a third antigen exposure significantly bolsters the serological response
169 and supports the rationale for providing booster doses to this patient population, who otherwise
170 may face further prolonged periods of restrictive social distancing. Early data from solid organ
171 transplant recipients reported that seroprevalence rates improved by about a third following a third
172 dose of the BNT162b2 vaccine after two months⁹. When starting a biologic, it would be reasonable
173 to consider differences in SARS-CoV-2 vaccine response as one of the factors when determining
174 which drug to use. For patients who need to start anti-TNF therapy, they and their families should
175 receive SARS-CoV-2 vaccines without an extended delay between doses. Whether the timing of
176 booster doses, the temporary discontinuation of immunomodulators¹⁰, the use of adjuvants
177 including the influenza vaccines (ComFluCOV)¹¹ and/or switching between vaccines with different
178 mechanisms of action¹² is more effective in immunosuppressed patients warrants further study.

179

180 The biology underpinning loss of durable antibody responses and uncoupling of the B cell and T cell
181 responses merit further research. TNF is a pleiotropic cytokine and its activities include maturation
182 of antigen presenting cells, modulation of T cell responses and stimulation of immunoglobulin
183 synthesis¹³⁻¹⁵. TNF neutralization, or genetic ablation, results in substantial loss of B-cells in primary
184 follicles in germinal centres, reduced numbers of memory B-cells in the periphery but preserved
185 numbers of T cells¹³. Uncoupling of humoral and T cell immunity to SARS-CoV-2 has been observed
186 in healthy individuals¹⁶, and although the relative contributions of memory B cell and T cell
187 responses have yet to be fully defined in SARS-CoV-2 immunity, the preservation of T cell immunity

188 reported here should provide some reassurance for anti-TNF treated patients although it is
189 noteworthy that one fifth made no anti-spike T cell response following two doses of either vaccine.
190 Chronic TNF exposure, a feature of many IMiDs, can render T cells anergic and can be reversed by
191 anti-TNF treatment¹⁷. This may in part explain why the magnitude of T cell responses observed in
192 anti-TNF-treated patients in this study did not differ significantly from patients treated with
193 vedolizumab.

194

195 We acknowledge some limitations in our study. Although our data show major differences in the
196 magnitude and durability of antibody responses, we have not assessed immunoglobulin classes, the
197 quality of antibody responses, or their effectiveness against the Wuhan and variants of concern.
198 However, there are strong positive associations between vaccine efficacy and viral neutralization
199 across the COVID-19 vaccine trials^{18,19}. Importantly, anti-RBD antibodies, such as the ones measured
200 in this study, strongly correlate with Wuhan Hu-1 live virus neutralization assays²⁰.

201

202 In conclusion, our data show that in infliximab-treated patients, anti-SARS-CoV-2 spike antibody
203 responses are attenuated and less durable following BNT162b2 and ChAdOx1 nCoV-19 SARS-CoV-2
204 vaccination. As early as 14 to 18 weeks after completing the vaccination course, many anti-TNF
205 treated patients have lost antibody-mediated protection from the virus, potentially leaving them
206 susceptible to infection. One fifth of both infliximab- and vedolizumab-treated patients did not
207 mount a T cell response and a small subset of patients had both poor antibody and T cell responses.
208 This could have important implications for health policy recommendations for patients taking anti-
209 TNF drugs, including vaccine prioritization, dosing intervals, booster requirements, and social
210 distancing strategies.

211

212 **Methods**

213 ***Anti-SARS-CoV2 Serology***

214 To determine antibody responses specific to vaccination we used the Roche Elecsys Anti-SARS-CoV-2
215 spike (S) immunoassay²¹ alongside the nucleocapsid (N) immunoassay²². This double sandwich
216 electrochemiluminescence immunoassay uses a recombinant protein of the receptor binding
217 domain on the spike protein as an antigen for the determination of antibodies against SARS-CoV-2.
218 Sample electrochemiluminescence signals are compared to an internal calibration curve and
219 quantitative values are reported as units (U)/mL. In-house assay validation experiments were
220 previously reported^{6,7}. Seroconversion was defined at a threshold of 15 U/mL. ElecSys Anti-SARS-
221 CoV-2 spike (S) RBD concentrations of greater than or equal to 15 U/ml are associated with
222 neutralization of $\geq 20\%$ with a positive predictive value of 99.10 % (95% CI: 97.74-99.64)⁷.

223 At entry to CLARITY IBD and at follow-up visits, all patients were tested for previous SARS-CoV-2
224 infection using the Roche Elecsys anti-SARS-CoV-2 (N) immunoassay. Because antibody responses
225 are impaired following PCR-confirmed natural infection we set a threshold of 0.25 times the cut-off
226 index (COI) at or above which patients were deemed to have had prior infection⁶. We defined a
227 second threshold of 0.12 times the COI, below which patients were deemed to have no evidence of
228 prior infection. Patients with a PCR test confirming SARS-CoV-2 infection at any time prior to
229 vaccination were deemed to have evidence of past infection irrespective of any antibody test result.

230 ***Peripheral blood mononuclear cell isolation***

231 Whole blood was collected in lithium heparin tubes and peripheral blood mononuclear cells (PBMCs)
232 were isolated by density-gradient centrifugation using LymphoprepTM (Stem Cell Technologies)
233 layered on to SepMateTM (Stem Cell Technologies) tubes. PBMC isolation was performed within 12
234 hours of venepuncture. Purified PBMCs were cryopreserved in 10% DMSO/50% FBS and stored in
235 liquid nitrogen pending batch analysis. For T cell assays blood was sampled 3-6 weeks after
236 vaccination.

237 ***Spike-peptide specific T cell responses***

238 IFN γ T cell ELISpot assays were performed using pre-coated plates (Mabtech 3420-2APT) and using
239 the protocol described previously^{16,20}. Two-hundred thousand cells were seeded per well and cells
240 were stimulated with a peptide pool, containing 18 peptides derived from SARS-CoV-2 spike
241 protein²³ at a concentration of 10 μ g/ml/peptide. Plates were cultured for 18-20 hours before
242 development and data collected using an AID classic ELISpot plate reader (Autoimmun Diagnostika
243 GMBH). Results are expressed as difference in (delta) spot forming cells (SFC) per 10⁶ PBMC between
244 peptide stimulation and a media only control. A response below 2 standard deviations of the media
245 only control wells was deemed to be a null response. Data was excluded if response to the positive
246 control anti-CD3 stimulation was <200 SFC per 10⁶ PBMCs.

247 ***Ethical consideration and role of funders***

248 CLARITY IBD is an investigator-led, UK National Institute for Health Research COVID-19 urgent public
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252 Celltrion Healthcare (South Korea), Takeda (UK), and Galapagos NV (Belgium). None of our funding
253 bodies had any role in study design, data collection or analysis, writing, or decision to submit for
254 publication. Patients were included after providing informed, written consent. The sponsor was the
255 Royal Devon and Exeter NHS Foundation Trust. The Surrey Borders Research Ethics committee
256 approved the study (REC reference: REC 20/HRA/3114) in September 2020. The protocol is available
257 online at <https://www.clarityibd.org>. The study was registered with the ISRCTN registry
258 (ISRCTN45176516).

259 ***Statistics***

260 The sample size for CLARITY IBD was based on the number of participants required to demonstrate a
261 difference in the impact of infliximab and vedolizumab on seroprevalence and seroconversion

262 following SARS-CoV-2 infection, with an estimated background seroprevalence of 0.05. We
263 calculated that a sample of 6970 patients would provide 80% power to detect differences in the
264 seroprevalence of SARS-CoV-2 antibodies in infliximab- compared with vedolizumab-treated
265 patients, whilst controlling for immunomodulator status at the 0.05 significance level.

266 Statistical analyses were undertaken in R 4.0.4 (R Foundation for Statistical Computing, Vienna,
267 Austria). All tests were two tailed and p-values reported without any correction for multiple testing.
268 P-values <0.05 were considered significant. We included patients with missing clinical data in
269 analyses for which they had data and have specified the denominator for each variable. Anti-S RBD
270 antibody concentrations are reported as geometric means and standard deviations. Other
271 continuous data are reported as median and interquartile range, and discrete data as numbers and
272 percentages, unless otherwise stated.

273 Univariable analyses, using t-tests of log-transformed anti-S RBD antibody concentration and
274 Spearman's rank correlation coefficients, were used to identify demographic, disease, vaccine, and
275 treatment-related factors associated with the concentration of anti-S RBD antibodies.

276 Mann-Whitney U test was used to compare the magnitude of T cell response (SFC/10⁶ PBMCs)
277 stratified by treatment and vaccine received, and Spearman's rank correlation coefficient was
278 calculated to determine correlation between antibody and T cell responses. Multivariable linear
279 regression models were used to identify factors independently associated with log anti-S RBD levels.
280 A priori, we included age, ethnicity, biological medication and immunomodulator use. No stepwise
281 regression was performed. Results are presented after exponentiation, so that the coefficients of the
282 model correspond to the fold change (FC) associated with each binary covariate. For age, a cut-off
283 was chosen based on graphical inspection of the relationship between age and anti-S RBD antibody
284 concentrations.

285 We compared the durability of antibody responses by calculating 15-day rolling geometric mean
286 anti-S RBD antibody concentrations. For this analysis we included participants who had an antibody

287 test carried out between 1 and 70 days after second vaccine dose. Time to a four-fold reduction in
288 detectable anti-S RBD antibodies were visualised using Kaplan-Meier curves. Cox proportional
289 hazard regression models were used to identify demographic, disease and treatment-related factors
290 associated with anti-S RBD antibody non-persistence.

291 We conducted sensitivity analyses to compare antibody responses stratified by participants with
292 serological or PCR evidence of SARS-CoV-2 infection at any time prior to vaccination.

293 **Data availability**

294 The study protocol including the statistical analysis plan is available at www.clarityibd.org. Individual
295 participant de-identified data that underlie the results reported in this article will be available
296 immediately after publication for a period of 5 years. The data will be made available to
297 investigators whose proposed use of the data has been approved by an independent review
298 committee. Analyses will be restricted to the aims in the approved proposal. Proposals should be
299 directed to tariq.ahmad1@nhs.net. To gain access data requestors will need to sign a data access
300 agreement.

301 **Code availability**

302 Code used for data analysis will be available upon request directed to nick.kennedy1@nhs.net.

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337 **Author Contributions**

338 NAK, JRG, CB, SS, NP, TA participated in the conception and design of this study. CB was the project
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341 DC, CB, MJ, SS, JLA, LC, JCL, CDM, ALH, PMI, GRJ, KBK, CAL, CWL, DMA, RJB, JRG, NP, TA were
342 involved in the acquisition, analysis, or interpretation of data. DMS, CJR, KML, DKB, and FFP
343 performed, analysed and interpreted T cell experiments. T cell experiments
344 were supervised, designed, analysed and interpreted by RJB and DMA. Data analysis was done by
345 NAK, DMS and RJB. Drafting of the manuscript was done by SL, NAK, NC, SS, CWL, DMA, RJB, JRG,
346 NP, TA. NP and TA obtained the funding for the study. All the authors contributed to the critical
347 review and final approval of the manuscript. NAK, NP and TA have verified the underlying data.

348

349 **Competing Interests**

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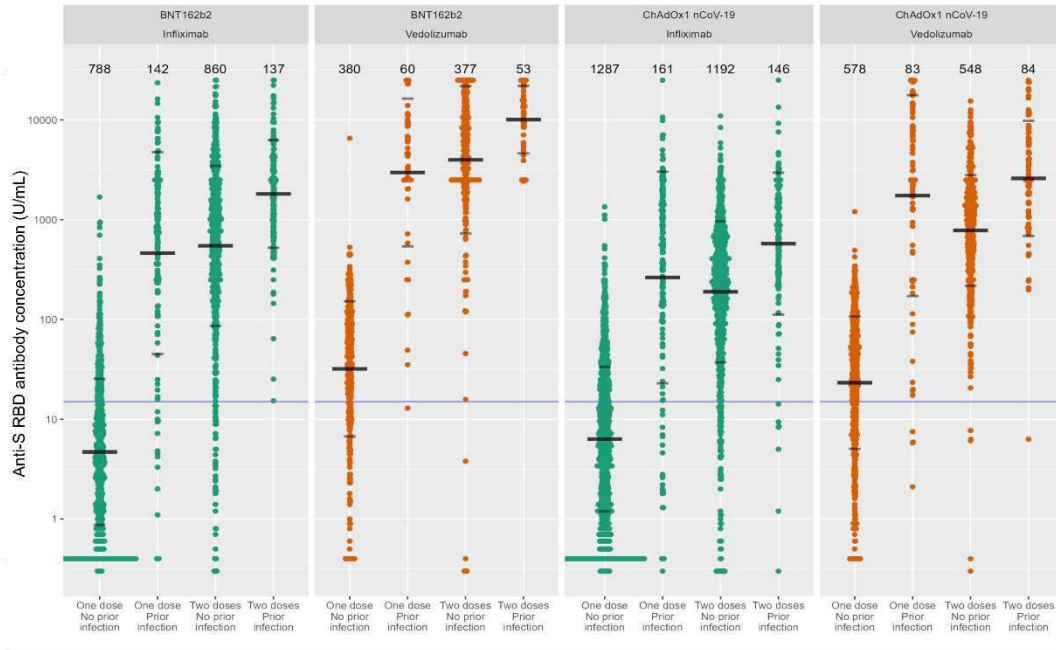
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463 convalescent individuals following COVID-19. *Nat. Immunol.* **21**, 1336 (2020).

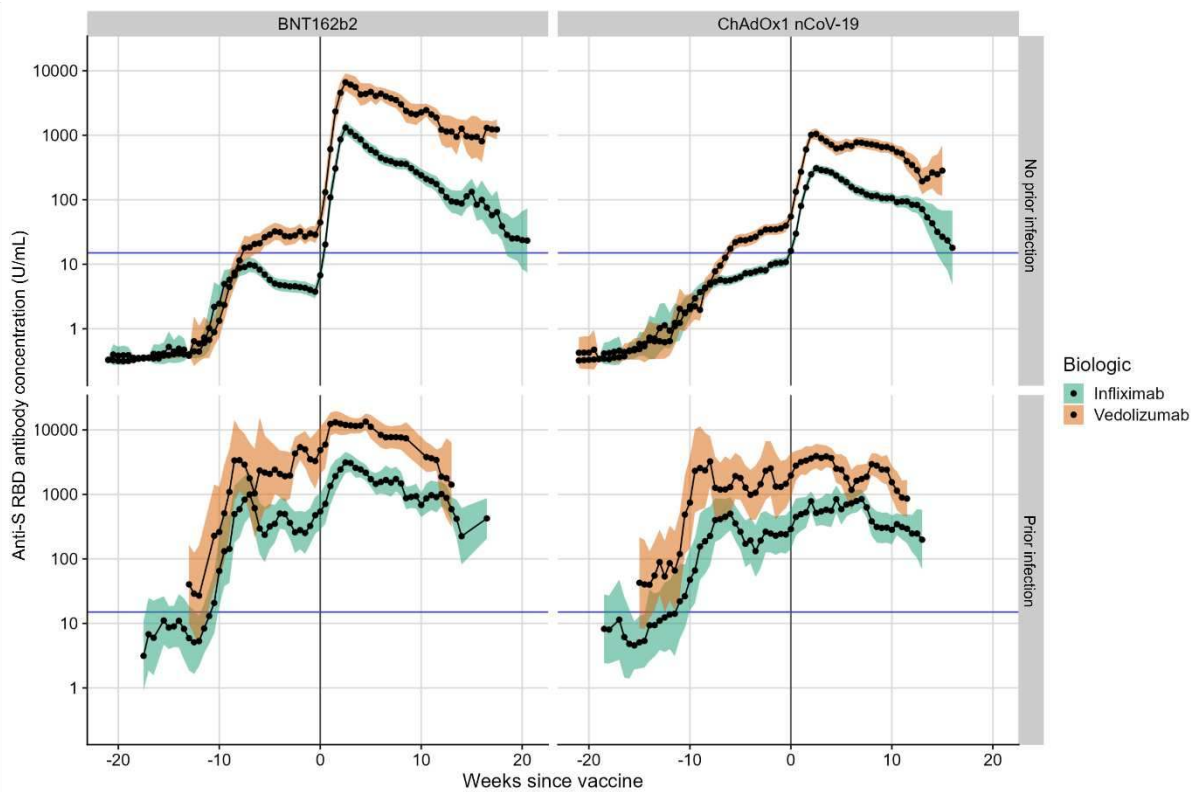
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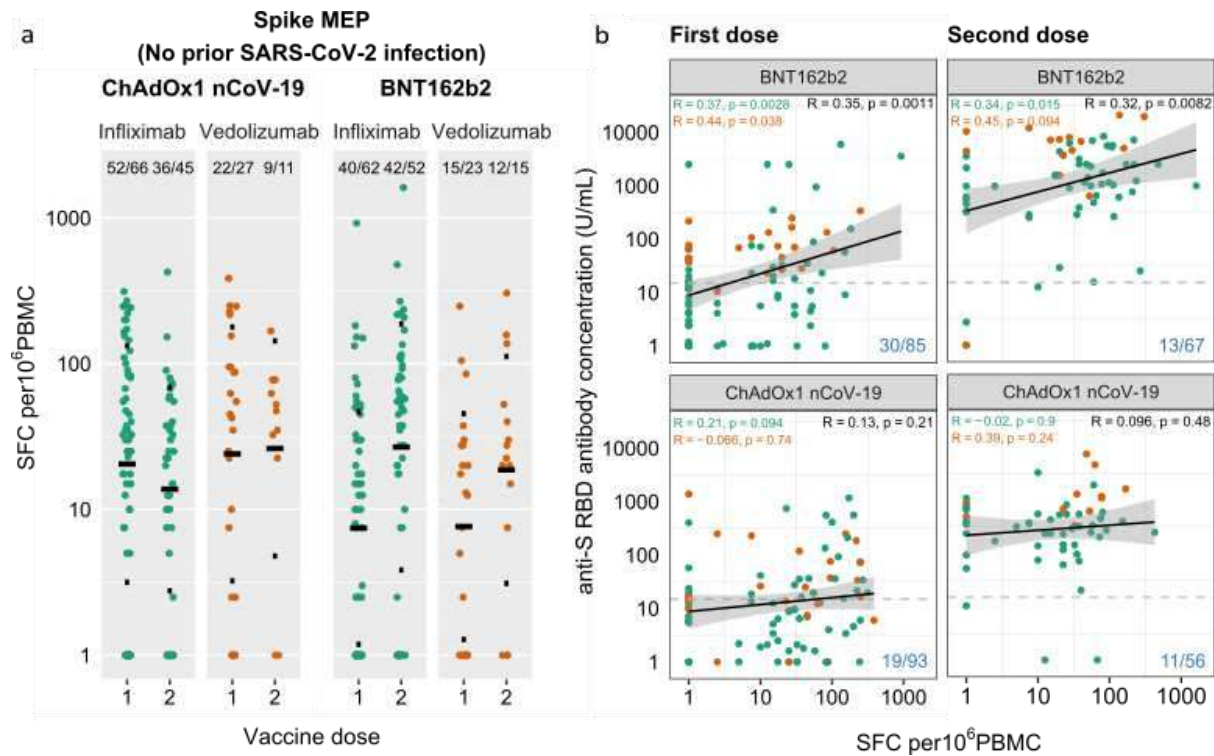
467 **Figure 1: Anti-SARS-CoV-2 spike (S) receptor binding domain (RBD) antibody response and**
 468 **durability**

469 **a.** Anti-SARS-CoV-2 spike RBD (anti-S RBD) antibody concentration stratified by biologic therapy
 470 (infliximab vs vedolizumab), type of vaccine, vaccine dose and prior infection. The wider bar
 471 represents the geometric mean, while the narrower bars are drawn one geometric standard

472 deviation either side of the geometric mean. Based on neutralization assays a threshold shown of 15
473 U/mL was used to determine seroconversion⁷. **b.** Rolling geometric mean antibody concentration
474 over time stratified by biologic therapy (infliximab vs vedolizumab), vaccine, and prior infection.
475 Geometric means are calculated using a rolling 15-day window (i.e. 7 days either side of the day
476 indicated). The shaded areas represent the 95% confidence intervals of the geometric means. The
477 blue line represents the seroconversion threshold (15 U/mL). Overall, data from 4470
478 participants with no prior infection (3029 on infliximab and 1441 on vedolizumab) and 683
479 participants with prior infection (459 on infliximab and 224 on vedolizumab) were included in this
480 graph between 22 weeks prior and 18 weeks post second dose vaccination.

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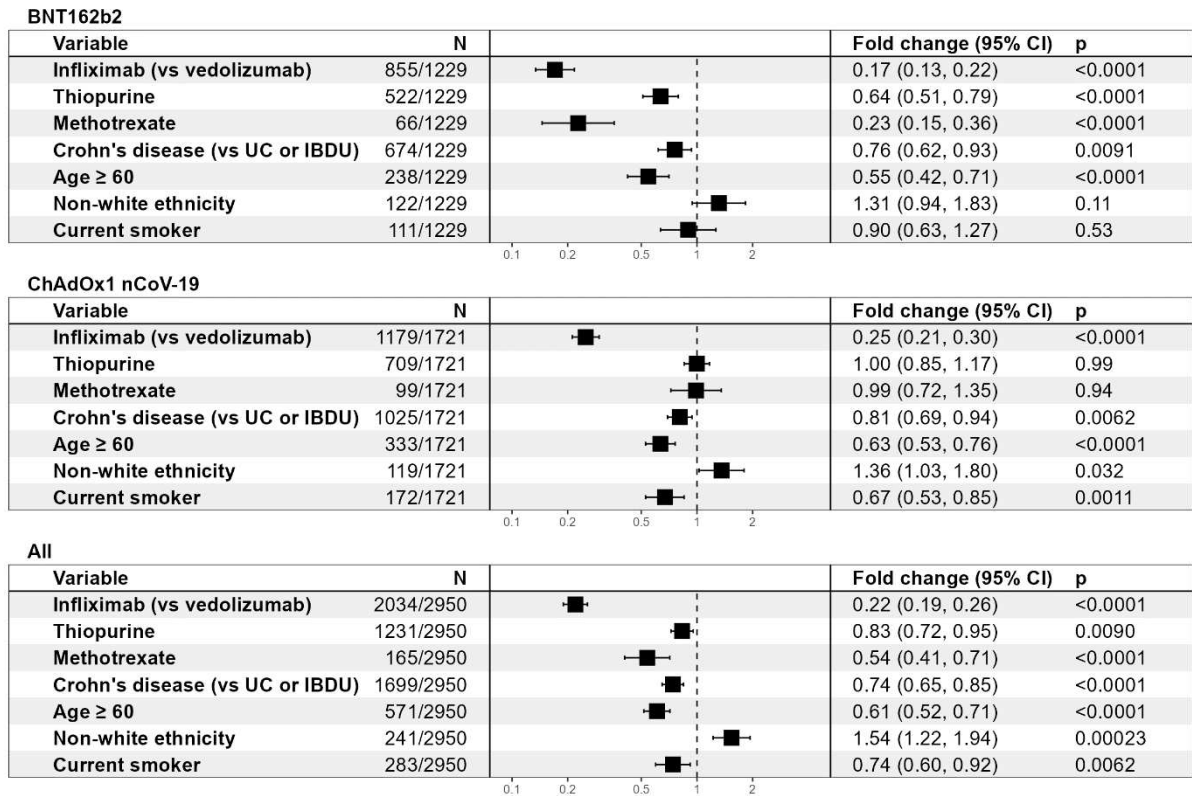
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485 **Figure 2. Anti-SARS-CoV-2 spike T cell responses stratified by biologic therapy (infliximab vs**
 486 **vedolizumab), vaccine type (BNT162b2 vs ChAdOx1 nCoV-19) and vaccine dose (one vs two).**

487 **a.** Spike MEP T cell responses SFC per 10^6 PBMC stratified by biologic therapy (infliximab vs
 488 vedolizumab), type of vaccine and vaccine dose. The horizontal bar represents the geometric mean,
 489 while the narrower bars represent one geometric standard deviation either side of the geometric
 490 mean. The number of T cell responders / total number of individuals tested are shown in black at the
 491 top of each panel. **b.** Scatterplot demonstrating the correlation between T cell responses against
 492 spike (SFC per 10^6 PBMC) and anti-SARS-CoV-2 spike antibody concentration after the first (LHS) and
 493 second (RHS) dose of BNT162B2 (top) and ChAdOx1 nCoV-19 (bottom). The number of non-T cell
 494 responders / total number of individuals tested is shown in blue on the bottom RHS of each panel.
 495 The horizontal dotted line in **b.** represents a threshold of 15 U/mL of anti-S1 SARS-CoV-2 antibody.
 496 The biologic infliximab is show in green and vedolizumab is shown in orange. R, Spearman's rank
 497 correlation. SFC, spot forming cells. MEP, mapped epitope peptide.

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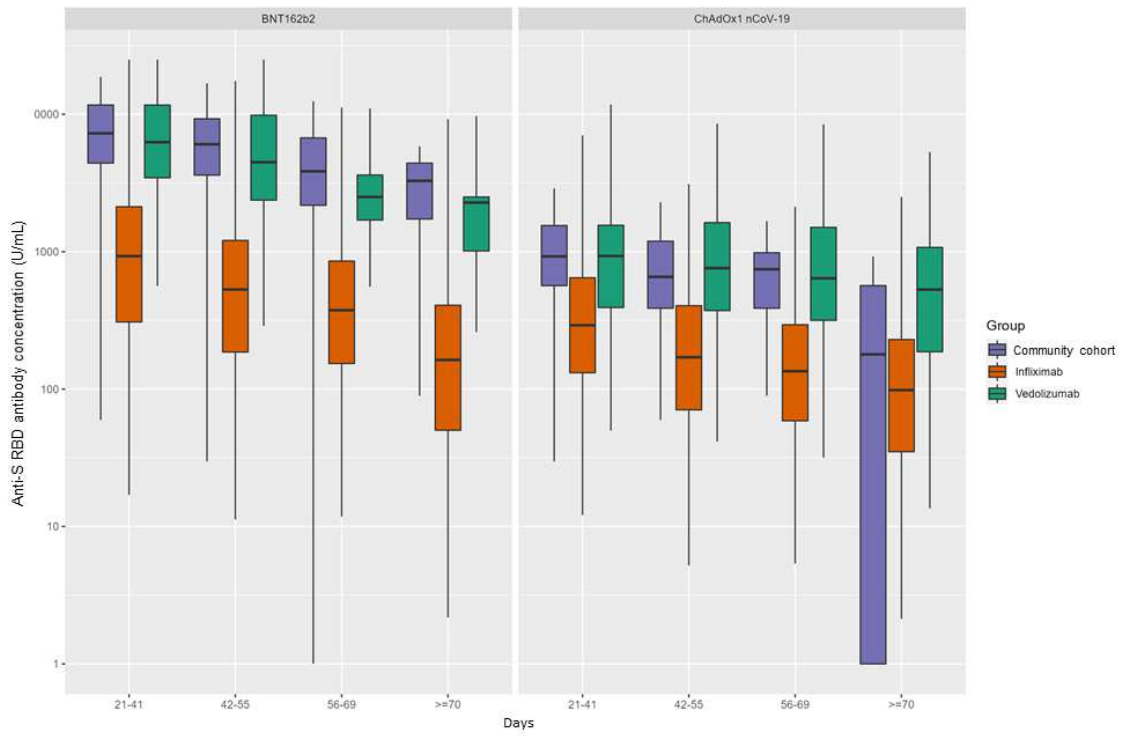
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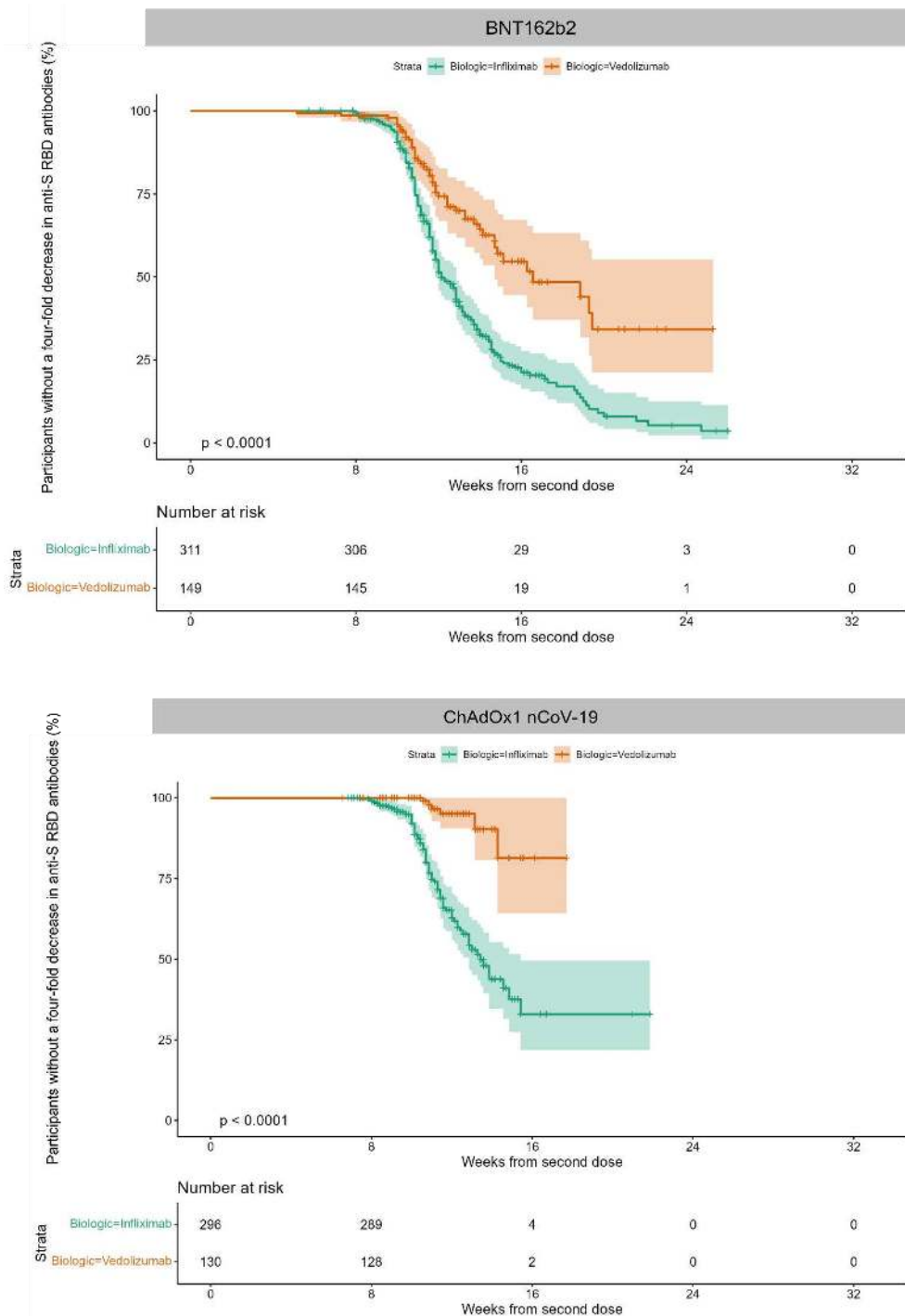
Extended Data Figure 1: Exponentiated coefficients of linear regression models of log(anti-S RBD antibody concentration)

The resultant values represent the fold change of antibody concentration associated with each variable. Each vaccine was modelled separately, and then a further model was created using all of the available data. Abbreviations: UC = ulcerative colitis, IBDU = IBD unclassified



508
 509 **Extended Data Figure 2: Anti-S RBD antibody levels at defined time points (days) after a second**
 510 **dose of vaccine in patients stratified by type of vaccine and biologic therapy (infliximab,**
 511 **vedolizumab) compared with individuals in the Virus Watch community cohort.**
 512

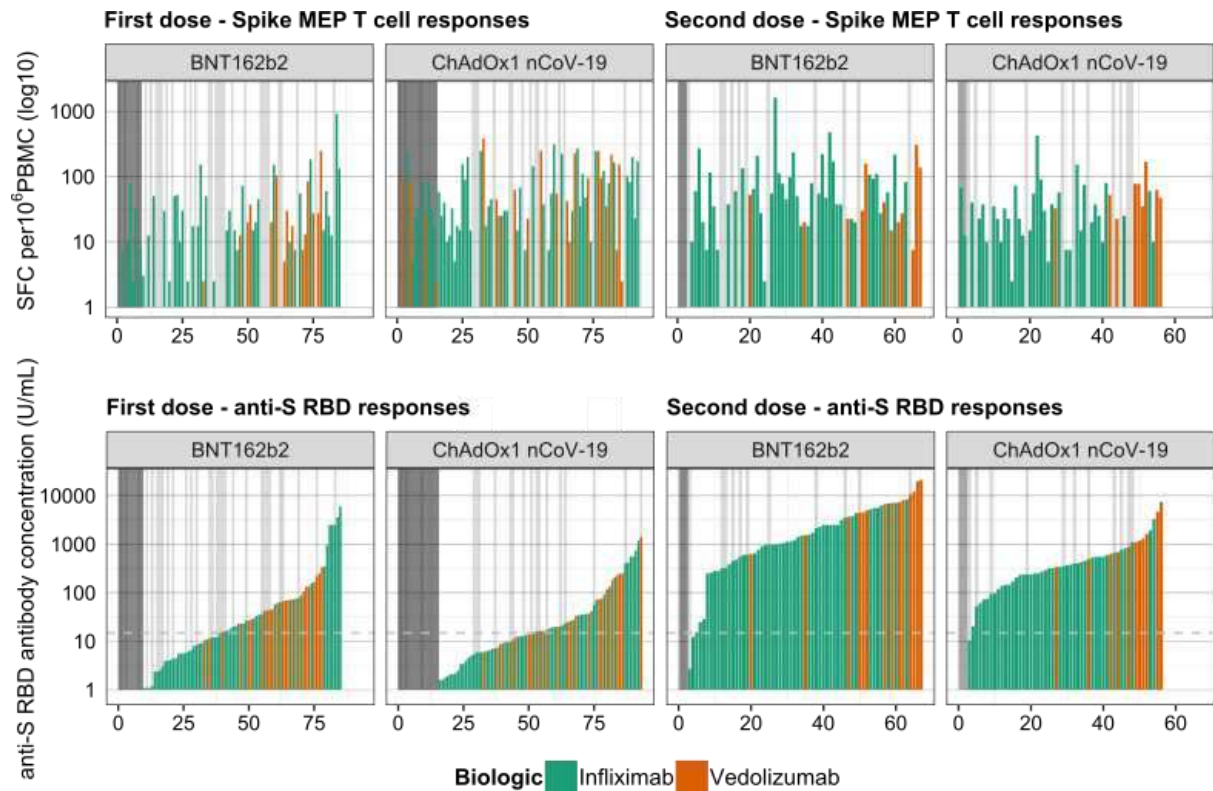
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527 **Extended Data Figure 3: Kaplan-Meier graphs showing the time to a four-fold drop in anti-S RBD**
 528 **antibody following the second dose of vaccination stratified by type of vaccine**
 529

530 Patients who had more than one anti-S RBD antibody measurement at least 2 weeks after a second
 531 dose of either vaccine were included in this analysis. Overall, data from 886 patients (311 infliximab-
 532 and 139 vedolizumab-treated patients who received the BNT162b2 vaccine and 296 infliximab- and
 533 130 vedolizumab-treated patients who received the ChAdOx1 nCoV-19 vaccine) were included in
 534 this analysis. P-value was defined using a log-rank test.

535



536

537 **Extended Data Figure 4: Anti-spike T cell responses ordered by cumulative magnitude of anti-S**
 538 **RBD following two doses of the BNT162b2 or ChAdOx1 nCoV-19 vaccine shows uncoupling of the T**
 539 **cell and antibody responses**

540

541 Top panel shows T cell responses to spike, and bottom panel shows anti-S RBD responses plotted for
 542 individual study participants ordered by increasing magnitude of anti-S RBD antibody concentration
 543 (U/mL). The vertical grey bars indicate individuals with no T cell response against spike. The
 544 horizontal dotted line represents a threshold shown of 15 U/mL of anti-S RBD.

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Supplementary Files

This is a list of supplementary files associated with this preprint. Click to download.

- [SupplementaryInformationCLARITY.pdf](#)