

Antibody decay, T cell immunity and breakthrough infections following two SARS-CoV-2 vaccine doses in inflammatory bowel disease patients treated with infliximab and vedolizumab

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Anti tumour necrosis factor (anti-TNF) drugs increase the risk of serious respiratory infection and impair protective immunity following pneumococcal and influenza vaccination. Here we report SARS-CoV-2 vaccine-induced immune responses and breakthrough infections in patients with inflammatory bowel disease, who are treated either with the anti-TNF antibody, infliximab, or with vedolizumab targeting a gut-specific anti-integrin that does not impair systemic immunity. Geometric mean [SD] anti-S RBD antibody concentrations are lower and half-lives shorter in patients treated with infliximab than vedolizumab, following two doses of BNT162b2 (566.7 U/mL [6.2] vs 4555.3 U/mL [5.4], $p < 0.0001$; 26.8 days [95% CI 26.2 – 27.5] vs 47.6 days [45.5 – 49.8], $p < 0.0001$); similar results are also observed with ChAdOx1 nCoV-19 vaccination (184.7 U/mL [5.0] vs 784.0 U/mL [3.5], $p < 0.0001$; 35.9 days [34.9 – 36.8] vs 58.0 days [55.0 – 61.3], p value < 0.0001). One fifth of patients fail to mount a T cell response in both treatment groups. Breakthrough SARS-CoV-2 infections are more frequent (5.8% (201/3441) vs 3.9% (66/1682), $p = 0.0039$) in patients treated with infliximab than vedolizumab, and the risk of breakthrough SARS-CoV-2 infection is predicted by peak anti-S RBD antibody concentration after two vaccine doses. Irrespective of the treatments, higher, more sustained antibody levels are observed in patients with a history of SARS-CoV-2 infection prior to vaccination. Our results thus suggest that adapted vaccination schedules may be required to induce immunity in at-risk, anti-TNF-treated patients.

Vaccination programmes have reduced SARS-CoV-2 transmission, hospitalisation and deaths¹. Patients treated with immunosuppressive drugs were excluded from the original trials for COVID-19 vaccines^{2,3}. Consequently, data relating to the magnitude and durability of immune responses and subsequent vaccine effectiveness in this population are limited⁴.

Drugs targeting tumour necrosis factor (TNF), such as infliximab, are the most frequently prescribed biologic therapies used in the treatment of immune-mediated inflammatory disorders (IMiDs). Observational studies indicate that most patients with inflammatory bowel disease (IBD), an archetypal IMiD, mount serological responses following SARS-CoV-2 vaccines, although most were underpowered to discern the impact of specific drugs, including immunomodulators (azathioprine, mercaptopurine and methotrexate) and/or biologic therapies^{5–8}. We reported that antibody responses following SARS-CoV-2 infection^{9,10} or a single dose of either the BNT162b2 or ChAdOx1 nCoV-19 SARS-CoV-2 vaccines were impaired in anti-TNF treated patients when compared to vedolizumab-treated patients¹¹. Vedolizumab, is a gut-selective anti-integrin $\alpha 4\beta 7$ monoclonal antibody that, unlike anti-TNF drugs, is not associated with increased susceptibility to systemic infection or attenuated serological responses to vaccination¹².

In this work, we show that anti-SARS-CoV-2 spike antibody responses are attenuated and less durable following two doses of the BNT162b2 and ChAdOx1 nCoV-19 SARS-CoV-2 vaccines in infliximab-treated compared with vedolizumab-treated patients with IBD. Irrespective of biologic drug type, one-fifth of all patients do not mount a T cell response and a minority mount neither antibody nor T cell responses. Breakthrough SARS-CoV-2 infections, which are associated with lower antibody levels after the second dose of vaccine, are more common and occur earlier in infliximab-treated patients. Higher and more sustained antibody levels are observed in patients with a history of SARS-CoV-2 infection. Further work to define immunity after third primary and booster vaccine doses is needed to inform the need for adapted vaccination schedules in at-risk anti-TNF treated patients.

Results

Patient characteristics. Between September 22, 2020 and December 23, 2020, 7226 patients were recruited to the CLARITY study from 92 UK hospitals¹⁰. In this analysis we included 2279 infliximab-treated and 1031 vedolizumab-treated participants without a history of prior SARS-CoV-2 infection, who had received uninterrupted biologic therapy since recruitment and had an antibody test between 14 and 70 days after the second dose of either the BNT162b2 and ChAdOx1 nCoV-19 SARS-CoV-2 vaccines. Participant characteristics are shown in Table 1.

Additional analyses are presented for a subset of 211 infliximab-treated and 71 vedolizumab-treated patients included in our T cell experiments (Supplementary Table 1), and a further 530 infliximab-treated and 224 vedolizumab-treated participants who had a history of SARS-CoV-2 infection before vaccination (Supplementary Table 2).

Anti-SARS-CoV-2-spike (S) antibody level following two doses of SARS-CoV-2 vaccine. Overall, the geometric mean [geometric SD] of anti-S receptor-binding domain (RBD) antibody concentration was higher in recipients of two doses of the BNT162b2 than ChAdOx1 nCoV-19 vaccines (1084.1 U/mL [7.6] vs 289.9 U/mL [5.2], $p < 0.0001$). Anti-S RBD antibody concentrations were lower in patients treated with infliximab than in those treated with vedolizumab, following a second dose of BNT162b2

(566.7 U/mL [6.2] vs 4555.3 U/mL [5.4], $p < 0.0001$) and ChAdOx1 nCoV-19 (184.7 U/mL [5.0] vs 784.0 U/mL [3.5], $p < 0.0001$) vaccines (Fig. 1).

Crude sensitivity analyses, excluding patients treated with a concomitant immunomodulator, confirmed lower anti-S RBD antibody concentrations in patients treated with infliximab alone versus vedolizumab alone (BNT162b2 809.1 U/mL [4.9] vs 4691.5 U/mL [5.9], $p < 0.0001$, ChAdOx1 nCoV-19 178.5 U/mL [4.6] vs 778.0 U/mL [3.5], $p < 0.0001$).

After propensity matching for immunomodulator use and the other factors associated with choice of biologic, we confirmed lower anti-S RBD antibody concentrations in infliximab-treated compared to vedolizumab-treated patients (BNT162b2 600.1 U/mL [6.0] vs 4674.1 U/mL [4.7], $p < 0.0001$, ChAdOx1 nCoV-19 195.2 U/mL [4.5] vs 779.2 U/mL [3.6], $p < 0.0001$) (Supplementary Table 3).

Multivariable linear regression analyses in patients without prior SARS-CoV-2 infection confirmed that antibody concentrations were reduced six and four-fold in infliximab-treated compared with vedolizumab-treated participants who received the BNT162b2 (fold change [FC] 0.15 [95% CI 0.12, 0.19], $p < 0.0001$) and ChAdOx1 nCoV-19 ([FC] 0.24 [95% CI 0.21, 0.28], $p < 0.0001$) vaccines (Fig. 2a, b respectively). Age ≥ 60 years and Crohn's disease were also independently associated with lower anti-S RBD antibody concentrations in vaccinated participants. Thiopurine or methotrexate use was independently associated with lower anti-S RBD antibody concentrations in participants who received the BNT162b2, but not the ChAdOx1 nCoV-19, vaccine. Current smoking, non-white ethnicity and steroid use were associated with lower anti-S RBD antibody concentrations in participants who received the ChAdOx1 nCoV-19 but not the BNT162b2 vaccine. To assess the effect of vaccine type on antibody responses, we combined our response data in a model that included vaccine type in addition to the significant factors above. Vaccination with the BNT162b2 vaccine compared to the ChAdOx1 nCoV-19 was independently associated with a 3.7 fold [95% CI 3.29–4.12] higher anti-S RBD antibody concentration ($p < 0.0001$) (Fig. 2c).

Seroconversion rates after the first vaccine dose were lower in infliximab-treated compared to vedolizumab-treated participants (Fig. 1). However, administration of a second vaccine dose resulted in a >100 -fold and >25 -fold increase in antibody concentrations in recipients of the BNT162b2 and ChAdOx1 nCoV-19 vaccines, respectively (Fig. 1). Overall, more infliximab-treated than vedolizumab-treated patients failed to seroconvert after their second vaccine dose (5.9% vs 1.3%, $p < 0.0001$). Seroconversion rates stratified by biologic therapy and vaccine type are reported in Supplementary Table 4.

Anti-spike T cell responses following two doses of BNT162b2 and ChAdOx1 nCoV-19 SARS-CoV-2 vaccines. There were no significant differences in the magnitude of anti-spike T cell responses observed in infliximab-treated compared with vedolizumab-treated patients after one or two doses of either vaccine (Fig. 3a). The proportion of patients failing to mount detectable T cell responses were similar in both groups (infliximab 19.6% vs. vedolizumab 19.2%). For recipients of one and two doses of the BNT162b2 vaccine, there was a modest positive correlation between T cell responses and antibody concentration. This association was not observed in recipients following either dose of the ChAdOx1 nCoV-19 vaccine (Fig. 3b). When T cell responses were ranked by magnitude of antibody responses, most patients who did not mount an antibody response after the first vaccine dose (indicated by the dark grey bar) had a detectable T cell response (Fig. 4). In addition to the uncoupling of the T cell

Table 1 Baseline characteristics of participants who had anti-S receptor-binding domain antibodies measured 2 to 10 weeks following two doses of SARS-CoV-2 vaccine.

Variable	Vedolizumab	Infliximab	<i>p</i>
Vaccine			
BNT162b2	40.1% (413/1031)	40.1% (914/2279)	1.0
ChAdOx1 nCoV-19	59.9% (618/1031)	59.9% (1365/2279)	
Age (years)	48.0 (35.2–61.6)	40.2 (30.1–53.1)	<0.0001
Sex			
Female	48.0% (492/1024)	45.9% (1040/2267)	0.15
Male	51.8% (530/1024)	54.1% (1226/2267)	
Intersex	0.0% (0/1024)	0.0% (0/2267)	
Prefer not to say	0.2% (2/1024)	0.0% (1/2267)	
Ethnicity			
White	89.9% (920/1023)	92.5% (2096/2266)	0.037
Asian	6.5% (66/1023)	4.6% (104/2266)	
Mixed	2.4% (25/1023)	1.4% (32/2266)	
Black	0.5% (5/1023)	0.8% (18/2266)	
Other	0.7% (7/1023)	0.7% (16/2266)	
Diagnosis			
Crohn's disease	36.9% (380/1031)	67.2% (1531/2279)	<0.0001
UC/IBDU	63.1% (651/1031)	32.8% (748/2279)	
Duration of IBD (years)	9.0 (5.0–17.0)	8.0 (3.0–16.0)	0.00017
Age at IBD diagnosis (years)	33.5 (22.8–47.3)	27.6 (20.1–39.6)	<0.0001
Immunomodulators at vaccine	20.4% (210/1031)	57.2% (1304/2279)	<0.0001
5-ASA	33.9% (349/1031)	20.6% (469/2279)	<0.0001
Steroids	5.9% (61/1031)	2.8% (64/2279)	<0.0001
BMI	26.0 (23.0–29.8)	25.9 (22.8–30.0)	0.74
Heart disease	4.9% (50/1023)	2.6% (59/2261)	0.0011
Diabetes	7.6% (78/1023)	3.6% (82/2261)	<0.0001
Lung disease	16.4% (168/1023)	13.1% (296/2261)	0.013
Kidney disease	1.9% (19/1023)	0.8% (17/2261)	0.065
Cancer	1.6% (16/1023)	0.3% (6/2261)	<0.0001
Smoker			
Yes	8.2% (84/1023)	10.8% (245/2261)	0.0010
Not currently	37.7% (386/1023)	30.2% (682/2261)	
Never	54.1% (553/1023)	59.0% (1334/2261)	
Exposure to documented cases of COVID-19	7.8% (80/1023)	8.6% (195/2261)	0.46
Income deprivation score	0.089 (0.055 - 0.146)	0.091 (0.052 - 0.153)	0.92
Active disease (PRO2)	10.4% (102/977)	5.2% (111/2153)	<0.0001
Time between vaccine doses (weeks)	10.9 (9.7–11.1)	11.0 (10.0–11.3)	0.0030
Time from second dose to serum sample (weeks)	5.7 (3.7– 7.7)	5.7 (3.7–7.7)	0.73

Values presented are median (interquartile range) or percentage (numerator/denominator).
P values represent the results of a Mann-Whitney *U*, Kruskal-Wallis or Fisher's exact test.
 UC ulcerative colitis, IBDU IBD unclassified, IBD inflammatory bowel disease, 5-ASA 5-aminosalicylic acid, BMI body mass index, PRO2 IBD disease activity.

and antibody responses demonstrated, this analysis emphasised that about one-fifth of participants made no T cell responses irrespective of vaccine used (indicated by the light grey bars). Moreover, a minority of individuals (3/67) 4.5% for BNT162b2 and (1/56) 1.8% for ChAdOx1 nCoV-19 vaccines carry neither detectable antibody nor T cell responses after two doses of vaccine (Figs. 3b, 4).

Durability of antibody responses following two doses of BNT162b2 and ChAdOx1 nCoV-19 SARS-CoV-2 vaccines. The estimated half-life of anti-S RBD antibodies was shorter in participants receiving the BNT162b2 compared to the ChAdOx1 nCoV-19 vaccines (30.8 days [95% CI 30.3–31.5] vs 40.5 days [95% CI 39.2–41.6], *p* value <0.0001). When stratified by biologic, half-life estimates were shorter in infliximab-treated than vedolizumab-treated patients following two doses of BNT162b2 (26.8 days [95% CI 26.2–27.5] vs 47.6 [95% CI 45.5–49.8], *p* value <0.0001) and ChAdOx1 nCoV-19 (35.9 days [95% CI 34.9–36.8] vs 58.0 days [95% CI 55.0–61.3], *p* value <0.0001) (Supplementary Fig. 1 and Supplementary Table 5).

Overall, following two doses of either vaccine, anti-S RBD antibodies showed minimal decay to the last follow-up in patients treated with vedolizumab (Fig. 5 and Supplementary Fig. 2) and were similar to those observed in participants in the Virus Watch community cohort (Supplementary Fig. 3). However, in infliximab-treated participants, the geometric mean concentrations dropped to the seroconversion threshold by about 25 weeks after the second dose irrespective of the vaccine administered (Fig. 5). Infliximab compared to vedolizumab treatment, current smoking and white ethnicity were associated with a faster fall in anti-S RBD antibody concentration below the seroconversion threshold. (Supplementary Figs. 4, 5).

Breakthrough SARS-CoV-2 infections following two doses of vaccine. Of 5123 participants without polymerase chain reaction (PCR)-positive or serological evidence of prior SARS-CoV-2 infection, 267 had a first positive SARS-CoV-2 PCR test 2 or more weeks after the second vaccine dose. Overall, 89.2% patients were symptomatic: the most commonly reported symptoms were fatigue (73.7%), anosmia/ageusia (71.4%), fever (57.1%), cough

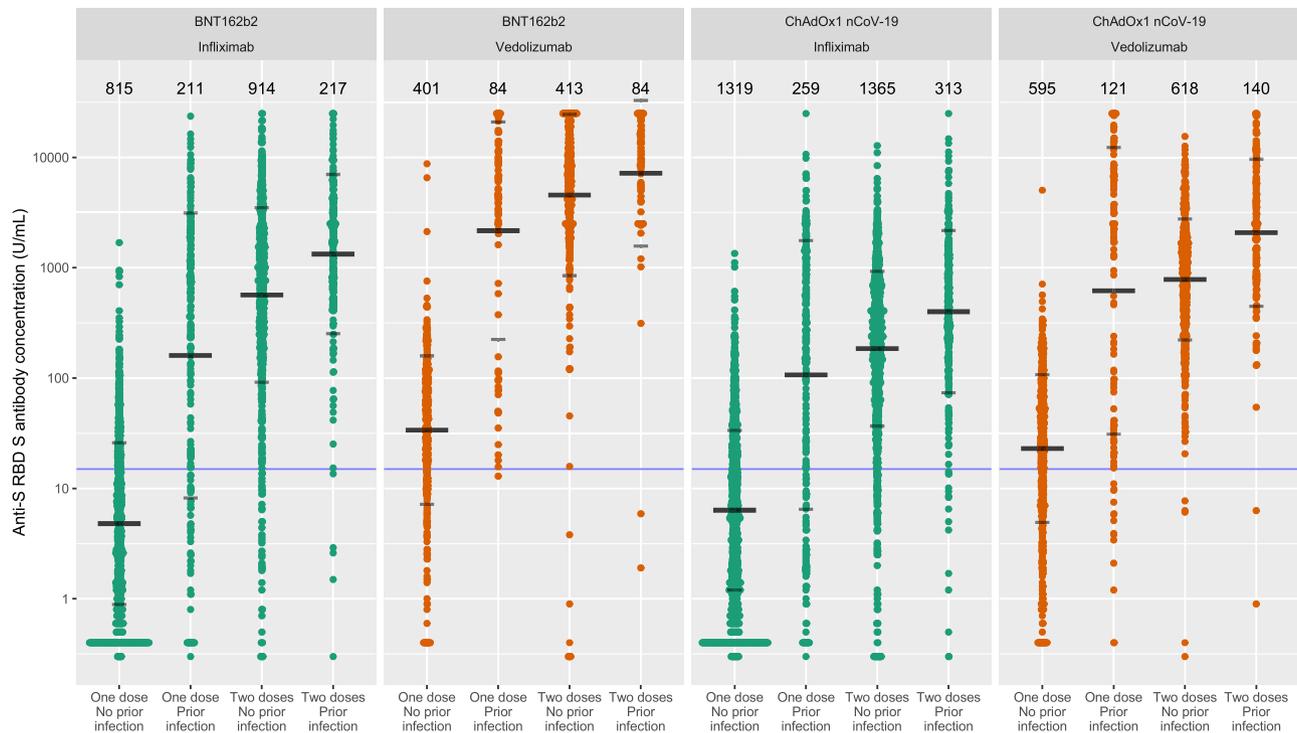


Fig. 1 Anti-S RBD antibody concentration stratified by biologic therapy (infliximab vs vedolizumab), type of vaccine, vaccine dose and history of prior SARS-CoV-2 infection. The wider bar represents the geometric mean, while the narrower bars are drawn one geometric standard deviation on either side of the geometric mean. Based on published data using neutralisation assays threshold shown of 15 U/mL was used to determine seroconversion¹¹. The biologic treatment infliximab is shown in green and vedolizumab in orange. The number of individuals tested for each group are shown in black at the top of each panel. Source data are provided as a Source Data file.

(54.9%), myalgia (45.9%), hoarse voice (30.8%), confusion (27.8%) and chest pains (23.3%). Overall, 1.2% (3/253) of participants with PCR-confirmed infection were hospitalised because of COVID-19.

Breakthrough SARS-CoV-2 infections were more frequent (5.8% (201/3441) vs 3.9% (66/1682), $p = 0.0039$) and the time to breakthrough shorter in patients treated with infliximab than vedolizumab ($p = 0.0027$) (Fig. 6b). In contrast biologic class did not impact on time to PCR-confirmed infection prior to vaccination ($p = 0.63$) (Fig. 6a). In a model that included biologic and vaccine type, shorter time to breakthrough infection was associated with infliximab (Hazard Ratio (HR) 1.52 [95% CI 1.15–2.01], $p = 0.003$) and having received the ChAdOx1 nCoV-19 (HR 1.49 [95% CI 1.15–1.92], $p = 0.0023$) vaccine. Geometric mean [geometric SD] anti-S RBD antibody concentrations measured 2 to 10 weeks after a second vaccine dose were significantly lower in participants who subsequently had a PCR-confirmed breakthrough SARS-CoV-2 infection: for every tenfold rise in anti-S RBD antibody concentration we observed a 0.8-fold reduction in odds of breakthrough infection ([95% CI 0.70–0.99], $p = 0.03$).

Antibody responses in patients with prior SARS-CoV-2 infection. Amongst patients with a history of SARS-CoV-2 infection before vaccination, geometric mean [SD] anti-S RBD antibody concentrations were lower in infliximab-treated compared with vedolizumab-treated patients after a second dose of BNT162b2 (1330.0 U/mL [5.3] vs 7169.5 U/mL [4.6], $p < 0.0001$) and ChAdOx1 nCoV-19 (399.7 U/mL [5.4] vs 2077.3 [4.6] $p < 0.0001$) vaccines. In all patients, antibody concentrations following vaccination were higher in patients without a history of SARS-CoV-2 infection (Fig. 1). Irrespective of vaccine or biologic type, minimal

decay of anti-S RBD antibodies were observed up to a follow-up of 21 weeks.

Discussion

We have shown that in infliximab-treated patients, anti-SARS-CoV-2 spike antibody responses are attenuated following two doses of the BNT162b2 and ChAdOx1 nCoV-19 SARS-CoV-2 vaccines. One-fifth of both infliximab-treated and vedolizumab-treated patients did not mount a T cell response and a small subset of patients had neither antibody nor T cell responses. Antibody half-lives were shorter in infliximab-treated patients. Breakthrough SARS-CoV-2 infections were more common and occurred earlier in infliximab-treated patients who received the ChAdOx1 nCoV-19 vaccine. The risk of breakthrough infection was predicted by lower antibody levels after the second dose of the vaccine. Irrespective of biologic treatment, higher and more sustained antibody levels were observed in patients with a history of SARS-CoV-2 infection.

Sustained antibody responses observed in vaccinated patients with a history of prior SARS-CoV-2 infection indicate that third antigen exposure enhances the serological response. This supports the rationale for prioritising a third dose of vaccine to clinically vulnerable patient populations^{13–16}, who otherwise may face further periods of social distancing or hospitalisation following infection. Whilst drawing direct comparisons between IBD patients and patients treated with more potent chemotherapies is limited by the degree to which patients are immunosuppressed, data from solid organ transplant recipients shows that a third dose of vaccine also leads to sustained immune responses¹⁷.

Irrespective of biologic or immunosuppressant use, and in keeping with the original trials^{2,18}, the highest antibody responses

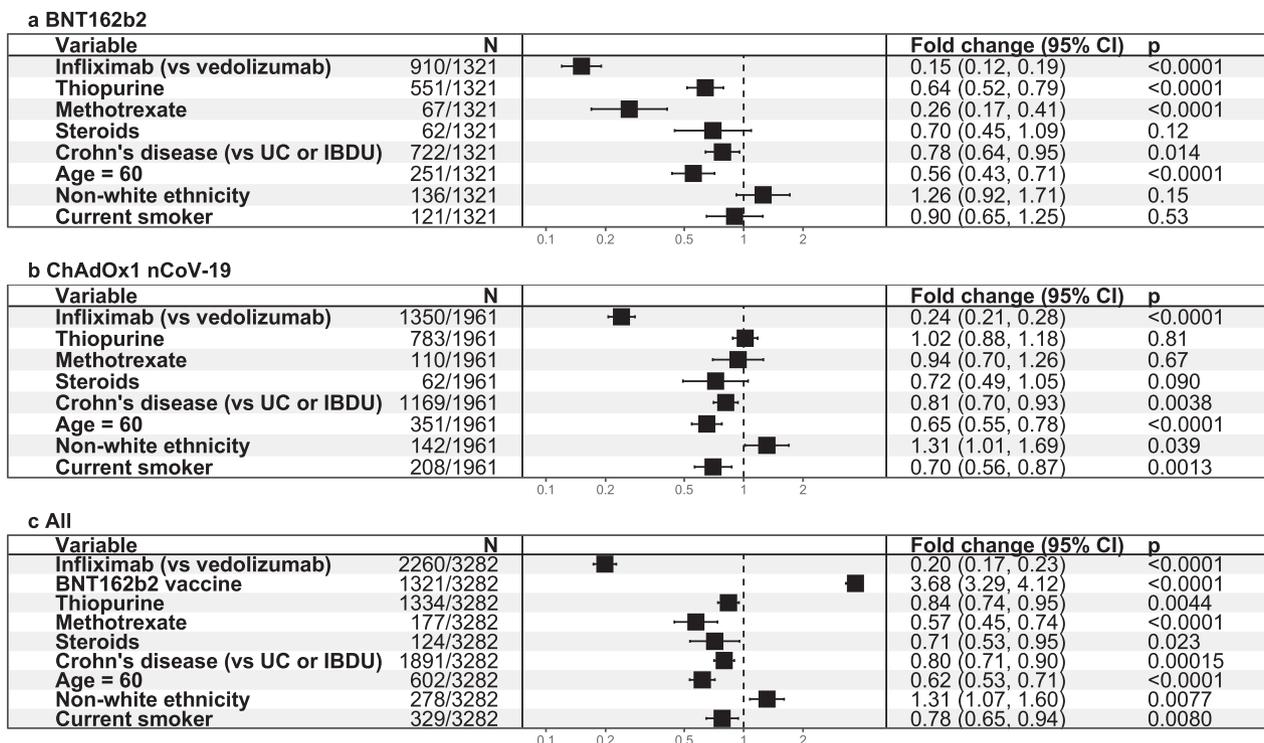


Fig. 2 Exponentiated coefficients of linear regression models of log anti-S RBD antibody concentration. Exponentiated coefficients of the linear regression model of log anti-S RBD antibody concentration in participants who received **a** BNT162b2 vaccine. **b** ChAdOx1 nCoV-19 vaccine. **c** either the BNT162b2 or ChAdOx1 nCoV-19 vaccine. The resultant values represent the fold change of antibody concentration associated with each variable (black square). The horizontal solid line through each square represents the 95% confidence interval. Each vaccine was modelled separately, and then a further model was created using all available data. The vertical dotted line represents a fold change of 1. Tests were two-tailed. *p* values were derived from linear regression using the *t*-test statistic and reported without correction for multiple testing. Source data are provided as a Source Data file. UC ulcerative colitis, IBDU IBD unclassified.

were seen in recipients of the BNT162b2 vaccine. Like in the general population, these responses waned more quickly than in the recipients of the ChAdOx1 nCoV-19 vaccine¹⁹. Unlike the general population²⁰, but similar to renal transplant recipients⁴, we did not observe differences in T cell ELISpot responses between recipients of the BNT162b2 and ChAdOx1 nCoV-19 vaccines. The differences observed in breakthrough infection by vaccine type reported here are consistent with the differences in efficacy reported in the respective clinical trials^{2,3,21}. The higher peak antibody levels and the lower rate of SARS-CoV-2 breakthrough infections suggest that the BNT162b2 rather than the ChAdOx1 nCoV-19 vaccine should be used for primary vaccination in infliximab-treated patients and, although untested, supports the use of BNT162b2 for third doses in all patients treated with an anti-TNF regardless of the primary vaccine type.

All patients treated with anti-TNF therapy should receive a third primary dose of the SARS-CoV-2 vaccine and our data support recent recommendations that this should occur about 4–8 weeks after the second dose^{13,14,16} during periods of high transmission in the population. Our data demonstrate that patients treated with vedolizumab and infliximab-treated patients with prior SARS-CoV-2 infection have sustained antibody levels beyond 6 months.

When starting a biologic, it would be reasonable to consider differences in SARS-CoV-2 vaccine response as one of the factors when determining which drug to use. For patients who need to start anti-TNF therapy, the benefits of combination immunomodulator therapy should be weighed against the risk of attenuated vaccine response, and whenever feasible, patients should first receive a SARS-CoV-2 vaccine dose. Further research to

determine whether timing third vaccine doses towards the end of anti-TNF treatment cycles when drug levels are lowest leads to greater immunogenicity⁹ is needed. Other strategies including the temporary discontinuation of immunomodulators²², the use of heterologous vaccines²³ and adjuvants including the influenza vaccines (ComFluCOV)²⁴ need to be studied in immunosuppressed patient groups.

The biology underpinning loss of durable antibody responses and uncoupling of the B cell and T cell responses merit further research. TNF is a pleiotropic cytokine and its activities include maturation of antigen-presenting cells, modulation of T cell responses and stimulation of immunoglobulin synthesis^{25–27}. TNF neutralisation, or genetic ablation, results in substantial loss of B-cells in primary follicles in germinal centres, reduced numbers of memory B-cells in the periphery but preserved numbers of T cells²⁵. Uncoupling of humoral and T cell immunity to SARS-CoV-2 has been observed in healthy individuals²⁸, and although the relative contributions of memory B cell and T cell responses have yet to be fully defined in SARS-CoV-2 immunity, the preservation of T cell immunity reported here should provide some reassurance for anti-TNF treated patients. However, it is noteworthy that one-fifth made no anti-spike T cell response following two doses of either vaccine. Chronic TNF exposure, a feature of many IMiDs, can render T cells anergic and can be reversed by anti-TNF treatment²⁹. This may in part explain why the magnitude of T cell responses observed in anti-TNF-treated patients in this study did not differ significantly from patients treated with vedolizumab.

Although our data show major differences in the magnitude and durability of antibody responses, we have not assessed the

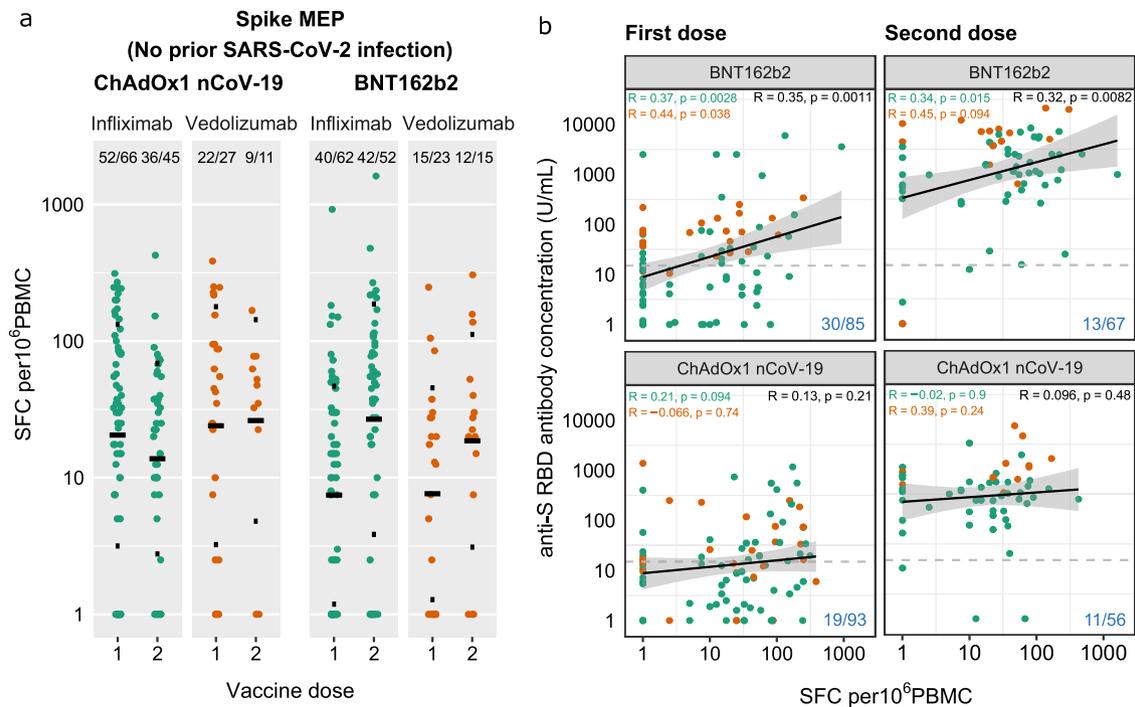


Fig. 3 Anti-SARS-CoV-2 spike T cell responses stratified by vaccine platform (BNT162b2 vs ChAdOx1 nCoV-19), biologic therapy (infliximab vs vedolizumab) and vaccine dose (one vs two). **a** Spike MEP T cell responses SFC per 10^6 PBMC stratified by vaccine platform, biologic therapy (infliximab vs vedolizumab) and the number of vaccine doses. The horizontal bar represents the geometric mean and the narrow bars represent one geometric standard deviation on either side of the geometric mean. The number of T cell responders / total number of individuals tested are shown in black at the top of each panel. **b** Scatterplot demonstrating the correlation between T cell responses against spike MEP pool (SFC per 10^6 PBMC) and anti-SARS-CoV-2 spike antibody concentration after the first (LHS) and second (RHS) dose of BNT162b2 (top) and ChAdOx1 nCoV-19 (bottom) vaccine. The number of non-T cell responders/total number of individuals tested is shown in blue on the bottom RHS of each panel. The shaded grey band represents the 95% confidence interval. The horizontal dotted line in **b** represents a threshold of 15 U/mL of anti-S1 SARS-CoV-2 antibody. The tests were two-tailed and *p* values were reported without correction for multiple testing. The biologic infliximab is shown in green and vedolizumab is shown in orange. Source data are provided as a Source Data file. MEP mapped epitope peptide, SFC spot forming cells, PBMC peripheral blood mononuclear cell, LHS left-hand side, RHS right-hand side, R Spearman's rank correlation.

impact of biologic therapy on specific immunoglobulin classes, antibody neutralisation or mucosal immune responses, which may be impaired, in particular, with anti-a4b7 therapy^{30,31}. However, previous studies have demonstrated that anti-RBD antibody levels such as the ones measured in this study, strongly correlate with Wuhan Hu-1 live virus and variant S RBD neutralisation assays^{32,33}, and we have demonstrated here that early antibody responses to vaccination correlates with the subsequent risk of breakthrough infection in immunosuppressed patients.

Infliximab was associated with attenuated, less durable vaccine-induced anti-SARS-CoV-2 spike antibody responses and a 50% increase in subsequent breakthrough SARS-CoV-2 infection. Further work to define immunity after third primary and booster vaccine doses is needed to inform the need for adapted vaccination schedules in at-risk anti-TNF treated patients.

Methods

Patient and settings. *impaCt of bioLogic therApy on saRS-cov-2 Infection and immuniTY (CLARITY) IBD* is a UK-wide, multicentre, prospective observational cohort study investigating the impact of infliximab and vedolizumab and/or concomitant immunomodulators (azathioprine, mercaptopurine and methotrexate) on SARS-CoV-2 acquisition, illness and immunity in patients with IBD.

Study methods have been previously described^{10,11}. Consecutive patients were recruited at the time of attendance at infusion units between 22 September 2020 and 23 December 2020 (Supplementary Table 1). Patients aged 5 years and over, with a diagnosis of IBD, treated with infliximab or vedolizumab were eligible for inclusion. Follow-up visits coincided with biologic infusions and occurred eight-weekly. Here, we report vaccine-induced antibody responses after the second dose of either the BNT162b2 or ChAdOx1 nCoV-19 vaccines. Participants were eligible for our primary immunogenicity analysis, if they had had an anti-S RBD antibody

test between 14 and 70 days after a second-dose vaccine, defined as the second dose of any of the licenced COVID-19 vaccines, 10-14 weeks after the first dose. Anti-S RBD antibody levels were compared with samples from 605 fully vaccinated adult participants from the Virus Watch study, a household community cohort of 10,000 individuals representative of the UK population of England and Wales recruited between 1 June 2020 to 31 August 2021¹⁹. Peripheral blood mononuclear cells (PBMC) for T cell experiments were collected from patients 4 to 6 weeks after the first and second dose of vaccine at the time of biologic infusions, at selected sites which could facilitate PBMC extraction within 12 h of venepuncture.

Outcome measures. Our primary outcome was anti-S RBD antibodies 2 to 10 weeks after the second dose of the BNT162b2 or ChAdOx1 nCoV-19 vaccines. Secondary outcomes were:

- the proportion of participants who seroconverted
- anti-spike T cell responses in patients following the first and second dose of vaccines
- the durability of vaccine responses
- risk of breakthrough infections two or more weeks after two doses of vaccine
- antibody concentrations and seroconversion rates in patients with PCR or serological evidence of past SARS-CoV-2 infection at, or prior, to the post-vaccination serum sample

Variables. Variables recorded by participants were demographics (age, sex, ethnicity, comorbidities, height and weight, smoking status, and postcode), IBD disease activity (PRO2), SARS-CoV-2 symptoms aligned to the COVID-19 symptoms study (symptoms, previous testing, and hospital admissions for COVID-19) and vaccine uptake (type and date of primary vaccination). Study sites completed data relating to IBD history (age at diagnosis, disease duration, and phenotype according to the Montreal classifications, previous surgeries, and duration of current biologic and immunomodulator therapy)¹⁰. We linked our data by NHS number or Community Health Index to Public Health England, Scotland, and Wales archive dates and results of all SARS-CoV-2 PCR tests undertaken and

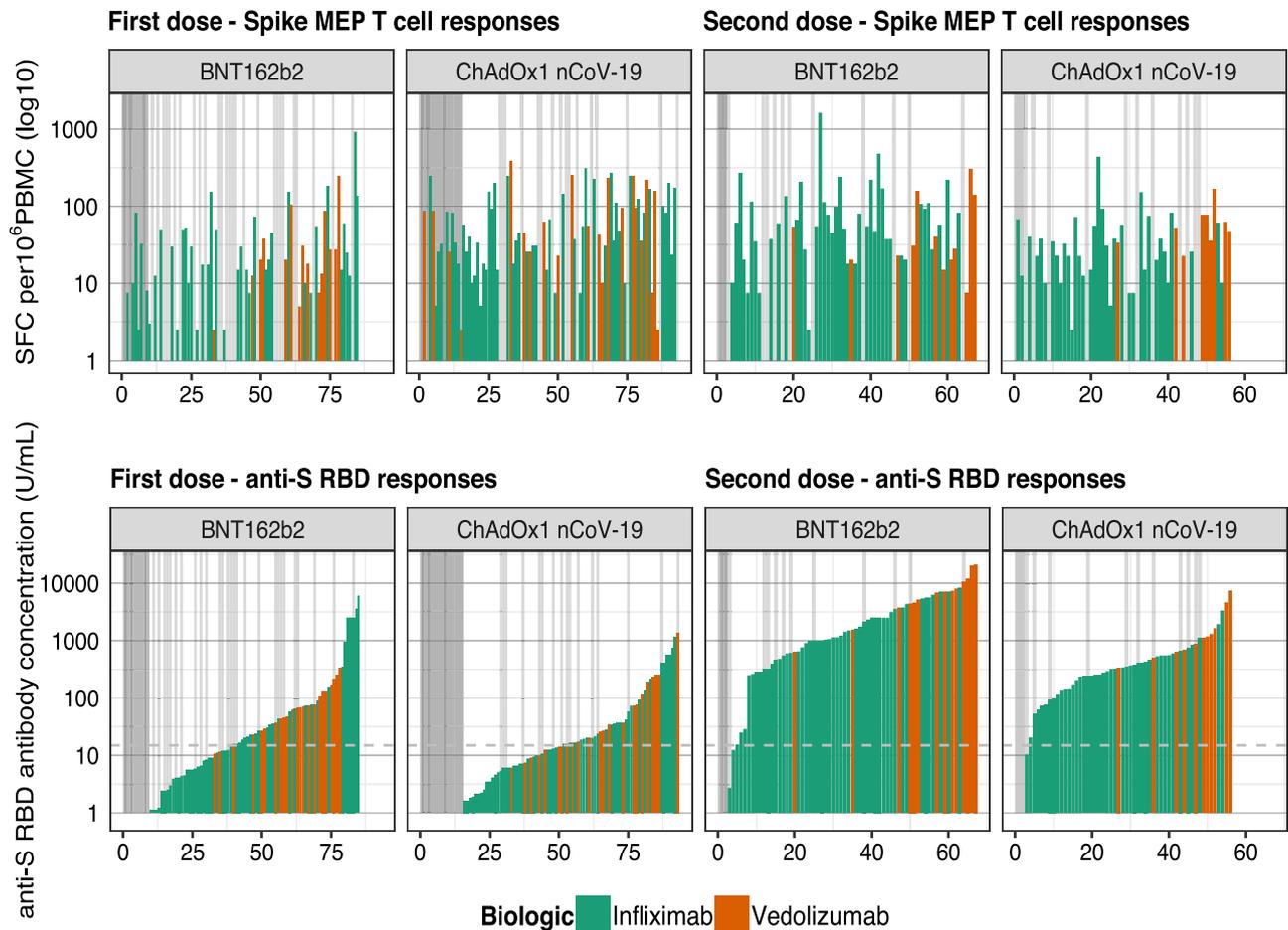


Fig. 4 Anti-spike T cell responses ordered by the cumulative magnitude of anti-S RBD following two doses of the BNT162b2 or ChAdOx1 nCoV-19 vaccine show uncoupling of the T cell and antibody responses. Top panel shows T cell responses to spike, and the bottom panel shows anti-S RBD responses plotted for individual study participants ordered by increasing magnitude of anti-S RBD antibody concentration (U/mL). The vertical dark grey bars at the LHS of the panels indicate individuals with no anti-S RBD response. The vertical light grey bars in the panels indicate individuals with no T cell response. The horizontal dotted line represents a threshold shown of 15 U/mL of anti-S RBD. Source data are provided as a Source Data file. LHS left-hand side, MEP mapped epitope peptide, SFC spot forming cells.

vaccines administered. Data were entered electronically into a purpose-designed REDCap database hosted at the Royal Devon and Exeter NHS Foundation Trust³⁴. Participants without access to the internet or electronic device completed their questionnaires on paper case record forms that were subsequently entered by local research teams.

Laboratory methods. To determine antibody responses specific to vaccination we used the Roche Elecsys Anti-SARS-CoV-2 spike (S) immunoassay³⁵ alongside the nucleocapsid (N) immunoassay³⁶. This double sandwich electrochemiluminescence immunoassay uses a recombinant protein of the receptor-binding domain on the spike protein as an antigen for the determination of antibodies against SARS-CoV-2. Sample electrochemiluminescence signals are compared to an internal calibration curve and quantitative values are reported as units (U)/mL. In-house assay validation experiments on the Roche Elecsys Anti-SARS-CoV-2 spike (S) immunoassay were performed on 20 samples from healthy individuals who have been vaccinated. This demonstrated:

- The intra-assay and inter-assay coefficient of variation were 1.3% and 5.6%, respectively
- Anti-SARS-CoV-2 (S) antibodies were stable in uncentrifuged blood and serum at ambient temperature for up to seven days permitting postal transport
- No effect was observed on recovery of anti-SARS-CoV-2 (S) antibodies following four freeze/thaw cycles
- No analytical interference was observed for the detection of anti-SARS-CoV-2 (S) with infliximab or vedolizumab up to 10,000 and 60,000 mg/L, respectively, or with anti-drug antibodies to infliximab or vedolizumab up to 400 and 38 AU/mL, respectively (data not shown).

Seroconversion was defined at a threshold of 15 U/mL. Elecsys Anti-SARS-CoV-2 spike (S) RBD concentrations of greater than or equal to 15 U/ml are

associated with neutralisation of $\geq 20\%$ with a positive predictive value of 99.10% (95% CI: 97.74–99.64)¹¹.

At the entry to CLARITY IBD and at follow-up visits, all patients were tested for previous SARS-CoV-2 infection using the Roche Elecsys anti-SARS-CoV-2 (N) immunoassay. We have previously reported that anti-N antibody responses following SARS-CoV-2 natural infection are impaired in patients treated with infliximab or vedolizumab¹¹. As such, a threshold 0.12 times above the cut-off index was set, using receiver operator characteristic curve and area under the curve analysis of anti-N antibody results from participants two weeks following a PCR-confirmed infection to maximise specificity, beyond which patients were deemed to have had prior SARS-CoV-2 infection (Supplementary Fig. 6). Patients with a PCR test confirming SARS-CoV-2 infection at any time prior to vaccination were deemed to have evidence of past infection irrespective of any antibody test result. Breakthrough infections were defined by a positive SARS-CoV-2 PCR test 2 or more weeks after the second vaccine dose.

Peripheral blood mononuclear cell isolation. Whole blood was collected in lithium heparin tubes and PBMCs were isolated by density-gradient centrifugation using LymphoprepTM (Stem Cell Technologies) layered onto SepMateTM (Stem Cell Technologies) tubes. PBMC isolation was performed within 12 h of venepuncture. Purified PBMCs were cryopreserved in 10% DMSO/50% FBS and stored in liquid nitrogen pending batch analysis.

Spike-peptide specific T cell responses. IFN- γ T cell ELISpot assays were performed using pre-coated plates (Mabtech 3420-2APT) and using the protocol described previously^{28,32}. Two-hundred thousand cells were seeded per well and cells were stimulated with a peptide pool, containing 18 peptides derived from SARS-CoV-2 spike protein³⁷ at a concentration of 10 $\mu\text{g}/\text{ml}/\text{peptide}$; the peptide pool utilises a mapped epitope pool (MEP) or 12–20mer peptides, mapped as

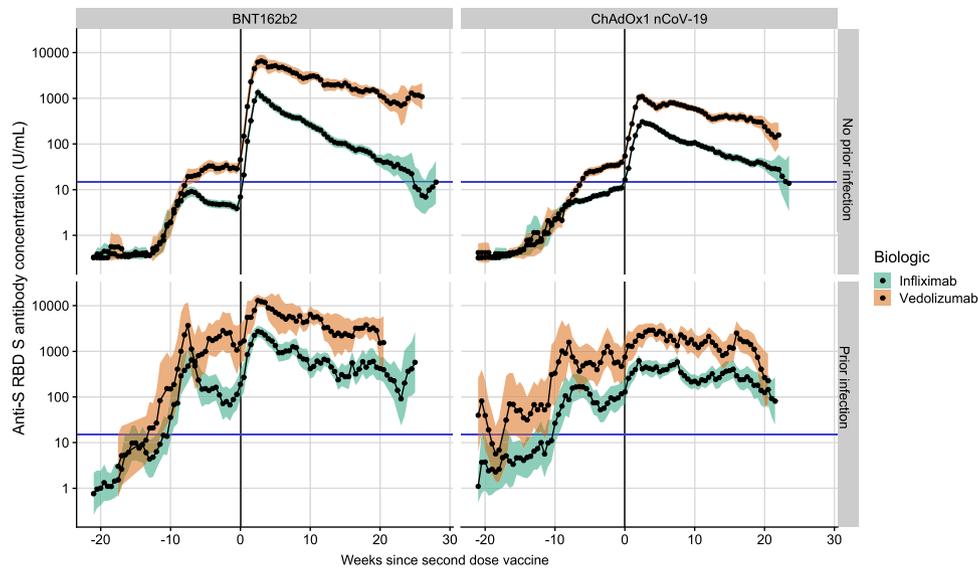


Fig. 5 Rolling geometric mean antibody concentration over time from the date of the second dose of the SARS-CoV-2 vaccine (week 0) stratified by biologic therapy (infliximab vs vedolizumab), vaccine and history of prior SARS-CoV-2 infection. Geometric means are calculated using a rolling 15-day window (i.e. 7 days on either side of the day indicated). The shaded areas represent the 95% confidence intervals of the geometric means. The horizontal blue line represents the seroconversion threshold (15 U/mL). The number of participants included at each time point is presented in Supplementary Fig. 2. Overall, data from 4474 participants with no history of prior infection (3029 on infliximab and 1445 on vedolizumab) and 1179 participants with a history of prior infection (833 on infliximab and 346 on vedolizumab) were included in this graph between 22 weeks before and 29 weeks after the second vaccine dose. The biologic treatment infliximab is shown in green and vedolizumab is shown in orange. Source data are provided as a Source Data file.

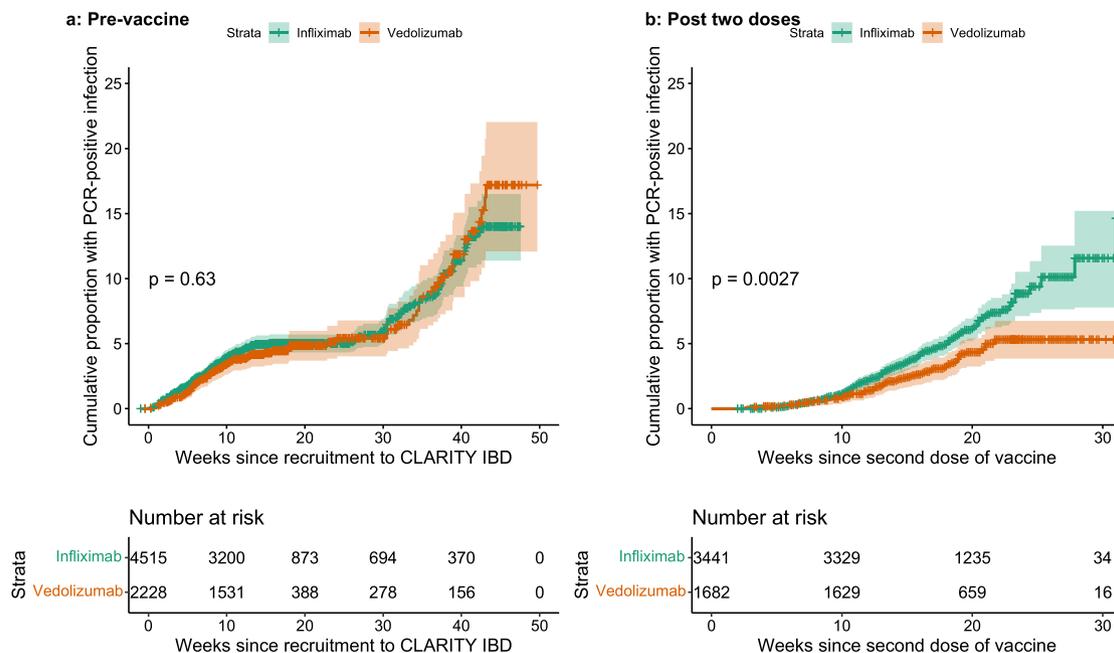


Fig. 6 Kaplan-Meier graphs comparing the time to PCR-confirmed SARS-CoV-2 infection stratified by biologic therapy (infliximab vs vedolizumab) in participants before vaccination and after receiving two doses of vaccine. **a** The time to PCR-confirmed SARS-CoV-2 infection in participants who have not received any dose of either vaccine stratified by biologic therapy (infliximab vs vedolizumab). **b** The time to a PCR-confirmed SARS-CoV-2 breakthrough infection in participants following two doses of either vaccine stratified by biologic therapy. The biologic treatment infliximab is shown in green and vedolizumab in orange. The number of participants at each time point are displayed in black at the bottom of each figure. *P* values are calculated using the log-rank test. Source data are provided as a Source Data file.

eliciting high-prevalence CD4 responses covering diverse HLA-II haplotypes^{28,32}. Use of this spike MEP in otherwise healthy SARS-CoV-2 seropositive individuals elicits a T cell response in 83% of individuals at 16–18 weeks after natural SARS-CoV-2 infection and 91% of healthy individuals 2–3 weeks after two-dose vaccination with seronegative individuals showing a level of response indistinguishable from pre-pandemic controls^{28,32}. Plates were cultured for 18–20 h before

development and data were collected using an AID classic ELISpot plate reader (Autoimmun Diagnostika GMBH). Results are expressed as differences in (delta) spot forming cells (SFC) per 10⁶ PBMC between peptide stimulation and a media-only control. A response below 2 standard deviations of the media-only control wells was deemed to be a null response. Data were excluded if the response to the positive control anti-CD3 stimulation was <200 SFC per 10⁶ PBMCs.

Sample size. The sample size for CLARITY IBD was based on the number of participants required to demonstrate a difference in the impact of infliximab and vedolizumab on seroprevalence and seroconversion following SARS-CoV-2 infection, with an estimated background seroprevalence of 0.05. We calculated that a sample of 6970 patients would provide 80% power to detect differences in the seroprevalence of SARS-CoV-2 antibodies in infliximab-treated compared with vedolizumab-treated patients, whilst controlling for immunomodulator status at the 0.05 significance level.

Statistical analyses. Analyses were undertaken using R 4.1.2 (R Foundation for Statistical Computing, Vienna, Austria). All tests were two-tailed and *p* values were reported without correction for multiple testing. *P* values <0.05 were considered significant. We included patients with missing clinical data in analyses for which they had data and have specified the denominator for each variable. Anti-S RBD antibody concentrations are reported as geometric means and standard deviations. Other continuous data are reported as a median and interquartile range, and discrete data as numbers and percentages, unless otherwise stated.

Univariable analyses, using Spearman's rank correlation coefficients, and *t*-tests of log-transformed anti-S RBD antibody concentration were used to identify demographic, disease, vaccine and treatment-related factors associated with the concentration of anti-S RBD antibodies across the cohort. Crude sensitivity analyses excluding patients treated without a concomitant immunomodulator were undertaken to control for the effect of immunomodulator use on anti-S RBD antibody concentrations. Propensity matching was used to account for the other significant differences in baseline variables between infliximab-treated and vedolizumab-treated patients using the MatchIt package³⁸. A priori, patients were matched exactly on diagnosis, immunomodulator use, and then using optimal matching, on age, the number of comorbidities, ethnicity, and presence of active disease. Multivariable linear regression models were used to identify factors independently associated with log anti-S RBD concentration. A priori, we included age, ethnicity, biologic medication and immunomodulator use. Results are presented after exponentiation so that the coefficients of the model correspond to the fold change (FC) associated with each binary covariate. For age, a cut-off was chosen based on a graphical inspection of the relationship between age and anti-S RBD antibody concentrations.

Mann-Whitney *U*-test was used to compare the magnitude of T cell response (SFC/10⁶ PBMCs) stratified by treatment and vaccine received, and Spearman's rank correlation coefficient was calculated to determine the correlation between antibody and T cell responses.

Anti-S RBD antibody half-lives were estimated using an exponential model of decay. Linear mixed models were fit using the lme4 and lmerTest package, with biologic treatment and vaccine type as fixed effects and each subject as a random effect. Each of these effects were estimated independently for gradient and intercept. 95% confidence intervals of fixed effects were calculated using likelihood ratios. *P* values for comparison of half-lives were estimated from the full linear mixed-effects model that incorporated vaccine, biologic drug and prior SARS-CoV-2 infection status.

We visualised the durability of antibody responses by calculating 15-day rolling geometric mean anti-S RBD antibody concentrations. For this analysis we included participants who had an antibody test carried out between 1 and 70 days after the second vaccine dose. Cox proportional hazard regression models were used to identify the demographic, disease and treatment-related factors associated with the time to fall in anti-S RBD antibody concentration below the seroconversion threshold.

Kaplan-Meier curves and Cox proportional hazard regression model was used to identify treatment-related factors associated with time to breakthrough infection. A linear regression model of log-transformed geometric mean anti-S RBD antibody concentration was used to determine the risk of breakthrough infections.

Where appropriate the same analyses were used to compare antibody responses in participants with PCR evidence of SARS-CoV-2 infection at any time prior to vaccination.

Ethical consideration. Patients were included after providing informed, written consent and compensation for participation was not provided. The sponsor was the Royal Devon and Exeter NHS Foundation Trust. The Surrey Borders Research Ethics committee approved the study (REC reference: REC 20/HRA/3114) in September 2020. The protocol is available online at <https://www.clarityibd.org>. The study was registered with the ISRCTN registry (<https://doi.org/10.1186/ISRCTN45176516>).

Reporting summary. Further information on research design is available in the Nature Research Reporting Summary linked to this article.

Data availability

The study protocol including the statistical analysis plan is available at <https://www.clarityibd.org/>. Individual participant de-identified data that underlie the results reported in this article will be available immediately after publication for a period of 5 years. Due to the sensitive nature of the data, this will be made available to investigators whose proposed use of the data has been approved by an independent review committee.

Analyses will be restricted to the aims in the approved proposal. Proposals should be directed to tariq.ahmad1@nhs.net. To gain access data requestors will need to sign a data access agreement. Data from the Virus Watch study is currently being archived on the Office of National Statistics Secure Research Service and will be available shortly. Source data are provided with this paper in the Source Data file. Source data are provided with this paper.

Code availability

Statistical analyses were undertaken in R 4.1.2 (R Foundation for Statistical Computing, Vienna, Austria). Code has been made available at: <https://github.com/exeteribd/clarityibd-public>.

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Author contributions

N.A.K., J.R.G., C.B., S.S., N.P. and T.A. participated in the conception and design of this study. C.B. was the project manager and coordinated patient recruitment. R.N. and T.J.M. coordinated all biochemical analyses and central laboratory aspects of the project. S.L., N.A.K., A.S., D.M.S., C.J.R., R.C.S., S.H.K., F.P.P., K.-M.L., D.K.B., N.C., D.C., C.B., M.J., S.S., J.L.A., L.C., J.C.L., C.D.M., A.L.H., P.M.I., G.-R.J., K.B.K., C.A.L., C.W.L., D.M.A., R.J.B., J.R.G., N.P. and T.A. were involved in the acquisition, analysis or interpretation of data. D.M.S., C.J.R., K.-M.L., D.K.B. and F.F.P. performed, analysed and interpreted T cell experiments. T cell experiments were supervised, designed, analysed and interpreted by R.J.B. and D.M.A. Data analysis was done by N.A.K., D.M.S. and R.J.B. Drafting of the manuscript was done by S.L., N.A.K., N.C., S.S., C.W.L., D.M.A., R.J.B., J.R.G., N.P. and T.A. R.J.B., N.P. and T.A. obtained the funding for the study. All the authors contributed to the critical review and final approval of the manuscript. N.A.K., N.P. and T.A. have verified the underlying data.

Competing interests

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Additional information

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⁹⁵Stockport NHS Foundation Trust, Stockport, UK. ⁹⁶Surrey and Sussex Healthcare NHS Trust, Surrey, UK. ⁹⁷Tameside and Glossop Integrated Care NHS Foundation Trust, Greater Manchester, UK. ⁹⁸Torbay and South Devon NHS Foundation Trust, Torquay, UK. ⁹⁹University Hospitals Birmingham NHS Foundation Trust, Birmingham, UK. ¹⁰⁰University Hospitals Bristol NHS Foundation Trust, Bristol, UK. ¹⁰¹University Hospitals of Derby and Burton NHS Foundation Trust, Derby, UK. ¹⁰²University Hospitals of Leicester NHS Trust, Leicester, UK. ¹⁰³University Hospitals Plymouth NHS Trust, Plymouth, UK. ¹⁰⁴United Lincolnshire Hospitals NHS Trust, Lincoln, UK. ¹⁰⁵University College London Hospitals NHS Foundation Trust, London, UK. ¹⁰⁶University Hospital Llandough, Penarth, UK. ¹⁰⁷University Hospital Southampton NHS Foundation Trust, Southampton, UK. ¹⁰⁸University Hospital of Wales (Paediatrics), Cardiff, UK. ¹⁰⁹Exeter NIHR Clinical Research Facility, University of Exeter, Exeter, UK. ¹¹⁰West Hertfordshire Hospitals NHS Trust, Hertfordshire, UK. ¹¹¹West Middlesex University Hospital, London, UK. ¹¹²West Suffolk NHS Foundation Trust, Ipswich, UK. ¹¹³Western General Hospital, Edinburgh, UK. ¹¹⁴Withybush General Hospital, Haverfordwest, UK. ¹¹⁵Yeovil District Hospital NHS Foundation Trust, Yeovil, UK. ¹¹⁶York Teaching Hospital NHS Foundation Trust, York, UK. ¹¹⁷Ysbyty Gwynedd, Bangor, UK.