# Phase I/II study of COVID-19 RNA vaccine BNT162b1 in adults

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In March 2020, the World Health Organization (WHO) declared coronavirus disease 2019 (COVID-19), which is caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2)<sup>1</sup>, a pandemic. With rapidly accumulating numbers of cases and deaths reported globally<sup>2</sup>, a vaccine is urgently needed. Here we report the available safety, tolerability and immunogenicity data from an ongoing placebo-controlled, observer-blinded dose-escalation study (Clinical Trials.gov identifier NCT04368728) among 45 healthy adults (18–55 years of age), who were randomized to receive 2 doses-separated by 21 days-of 10 μg, 30 μg or 100 μg of BNT162b1. BNT162b1 is a lipid-nanoparticle-formulated, nucleoside-modified mRNA vaccine that encodes the trimerized receptor-binding domain (RBD) of the spike glycoprotein of SARS-CoV-2. Local reactions and systemic events were dose-dependent, generally mild to moderate, and transient. A second vaccination with 100 µg was not administered because of the increased reactogenicity and a lack of meaningfully increased immunogenicity after a single dose compared with the 30-µg dose. RBD-binding IgG concentrations and SARS-CoV-2 neutralizing titres in sera increased with dose level and after a second dose. Geometric mean neutralizing titres reached 1.9-4.6-fold that of a panel of COVID-19 convalescent human sera, which were obtained at least 14 days after a positive SARS-CoV-2 PCR. These results support further evaluation of this mRNA vaccine candidate.

In December 2019, a pneumonia outbreak of unknown cause occurred in Wuhan, China. By January 2020, a new coronavirus was identified as the aetiological agent. Within a month, the genetic sequence of the virus became available (MN908947.3). Infections with SARS-CoV-2 and the resulting disease, COVID-19, have spread globally. On 11 March 2020, the WHO declared the COVID-19 outbreak a pandemic<sup>1</sup>. So far, the United States has reported the highest number of cases globally<sup>2,3</sup>. No vaccines are currently available to prevent SARS-CoV-2 infection or COVID-19.

The RNA vaccine platform has enabled rapid vaccine development in response to this pandemic. RNA vaccines provide flexibility in the design and expression of vaccine antigens that can mimic the structure and expression of the antigen during natural infection. RNA is required for protein synthesis, does not integrate into the genome, is transiently expressed, is metabolized and eliminated by the natural mechanisms of the body and is therefore considered safe<sup>4-7</sup>. RNA-based prophylactic infectious-disease vaccines and RNA therapeutic agents have been shown to be safe and well-tolerated in clinical trials. In general, vaccination with RNA elicits a robust innate immune response. RNA directs the expression of the vaccine antigen in host cells and has intrinsic adjuvant effects<sup>8</sup>. A strength of the RNA-vaccine manufacturing platform irrespective of the encoded pathogen antigen—is the ability to rapidly produce large quantities of vaccine doses against a new pathogen<sup>9,10</sup>.

Vaccine RNA can be modified by incorporating 1-methylpseudouridine, which dampens innate immune sensing and increases mRNA translation in vivo<sup>11</sup>. The BNT162b1 vaccine candidate that is currently investigated clinically incorporates such nucleoside-modified mRNA and encodes the RBD of the spike protein of SARS-CoV-2, a key target of virus-neutralizing antibodies<sup>12-14</sup>. The RBD antigen expressed by BNT162b1 is modified by the addition of a T4 fibritin-derived foldon trimerization domain to increase its immunogenicity<sup>15</sup> by multivalent display<sup>16</sup>. The proper folding of the RBDs in the resulting protein construct has been confirmed by high resolution structural analysis (A.B.V. et al., manuscript in preparation). The vaccine RNA is formulated in lipid nanoparticles for more-efficient delivery into cells after intramuscular injection<sup>17</sup>. BNT162b1 is one of several RNA-based SARS-CoV-2 vaccine candidates<sup>18</sup> that are studied in parallel for selection to advance

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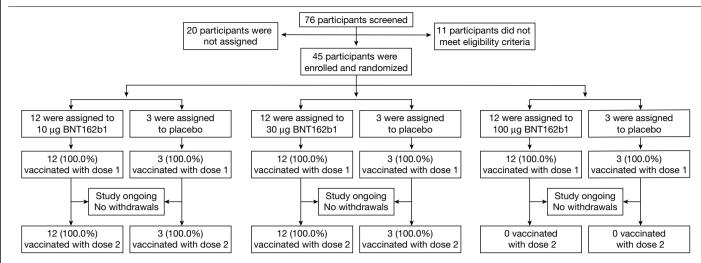


Fig. 1| Study design. Participants who were not assigned (n=20) were screened but not randomized because enrolment had closed.

to a safety and efficacy trial. Here, we present the available data, up to 14 days after a second dose in adults (18–55 years of age) from an ongoing phase I/II vaccine study with BNT162b1, which is also enrolling adults who are 65–85 years of age (ClinicalTrials.gov identifier, NCT04368728).

#### Study design and demographics

Between 4 May 2020 and 19 June 2020, 76 participants were screened, and 45 participants were randomized and vaccinated. Per dose level (10  $\mu g$  and 30  $\mu g$ ), 12 participants were vaccinated with BNT162b1 on days 1 and 21, 12 participants received a 100- $\mu g$  dose on day 1 and 9 participants received placebo (Fig. 1). The study population consisted of healthy male and female participants with a mean age of 35.4 years (range, 19–54 years); 51.1% were male and 48.9% were female. Most participants self-reported as white (82.2%) and non-Hispanic/non-Latinx (93.3%) (Extended Data Table 1).

#### Safety and tolerability

In the 7 days after vaccination doses 1 and 2, pain at the injection site was the most-frequent solicited local reaction, reported after the first dose by 58.3% (7 out of 12) in the 10-µg BNT162b1 group, 100.0% (12 out of 12 each) in the 30-µg and 100-µg BNT162b1 groups, and 22.2% (2 out of 9) in the placebo group. After the second dose, pain was reported by 83.3% (10 out of 12) and 100.0% of individuals who received 10 µg and 30 µg BNT162b1, respectively, and by 16.7% of individuals who received the placebo. All local reactions were mild or moderate in severity except for one report of severe pain after the first dose of 100 µg BNT162b1 (Fig. 2).

The most-common systemic events reported in the 7 days after each vaccination in both BNT162b1 and placebo groups were mild to moderate fatigue and headache. Reports of fatigue and headache were more common in the BNT162b1 groups than in the placebo group. In addition, chills, muscle pain and joint pain were reported by individuals who received BNT162b1 but not by individuals who received the placebo. Systemic events increased with dose level and were reported in a greater number of participants after the second dose (10-µg and 30-µg groups). After the first dose, fever (defined as  $\geq$ 38.0 °C) was reported by 8.3% (1 out of 12) of participants who received 10 µg and 30 µg BNT162b1 and by 50.0% (6 out of 12) of individuals who received 100 µg BNT162b1. After the second dose, 8.3% (1 out of 12) of participants who received 10 µg BNT162b1 and 75.0% (9 out of 12) of participants who received 30 µg BNT162b1 reported fever of  $\geq$ 38.0 °C. On the basis of the reactogenicity reported after the first dose of 100 µg and the second dose of

 $30 \, \mu g$ , participants who received an initial  $100 \, \mu g$  dose did not receive a second  $100 \, \mu g$  dose. Fevers generally resolved within 1 day of onset. No grade 4 systemic events or fever were reported (Fig. 3a, b). Most local reactions and systemic events peaked by day 2 after vaccination and resolved by day 7.

Adverse events (Extended Data Table 2) were reported by 50.0% (6 out of 12) of participants who received either 10 or  $30\,\mu g$  of BNT162b1, 58.3% (7 out of 12) of participants who received  $100\,\mu g$  of BNT162b1, and 11.1% (1 out of 9) of placebo recipients. Two participants reported a severe adverse event: grade 3 fever 2 days after vaccination in the  $30\,\mu g$  group, and sleep disturbance 1 day after vaccination in the  $100\,\mu g$  group. Related adverse events were reported by 25% (3 out of 12 in the  $10\,\mu g$  group) to 50% (6 out of 12 each in the  $30\,\mu g$  and  $100\,\mu g$  groups) of individuals who received BNT162b1 and by 11.1% (1 out of 9) of participants who received the placebo. No serious adverse events were reported.

No grade 1 or greater change in routine clinical laboratory values or laboratory abnormalities were observed for most participants after either of the BNT162b1 vaccinations. Of those with laboratory changes, the largest changes were decreases in the lymphocyte count after the first dose in 8.3% (1 out of 12), 45.5% (5 out of 11) and 50.0% (6 out of 12) of participants who received 10 µg, 30 µg and 100 µg BNT162b1, respectively. One participant each in the 10-µg (8.3% (1 out of 12)) and 30-μg (9.1% (1 out of 11)) groups and 4 participants in the 100-μg group (33.3% (4 out of 12)) had grade 3 decreases in the lymphocyte count. These decreases in lymphocyte count after the first dose were transient and returned to normal 6-8 days after vaccination (Extended Data Fig. 1). In addition, grade-2 neutropenia was noted 6–8 days after the second dose in 1 participant each in the 10-µg and 30-µg BNT162b1 groups. These two participants continue to be followed in the study, and no adverse events or clinical manifestations of neutropenia have been reported to date. None of the post-vaccination abnormalities observed were associated with clinical findings.

#### **Immunogenicity**

RBD-binding IgG concentrations and SARS-CoV-2-neutralizing titres were assessed at baseline, at 7 and 21 days after the first dose, at 7 days (day 28) and 14 days (day 35) after the second dose of BNT162b1. By 21 days after the first dose (for all three dose levels), geometric mean concentrations (GMCs) of RBD-binding IgG ranged from 534 to 1,778 U ml<sup>-1</sup> (Fig. 4a). In comparison, a panel of 38 SARS-CoV-2 infection and/or COVID-19 convalescent sera drawn at least 14 days after a PCR-confirmed diagnosis from patients with COVID-19 (18–83 years

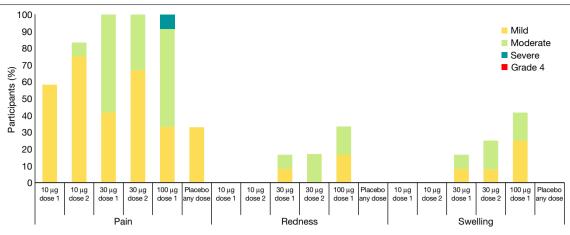


Fig. 2 | Local reactions reported within 7 days of vaccination for all dose levels. Solicited injection-site (local) reactions were: pain at injection site (mild, does not interfere with activity; moderate, interferes with activity; severe, prevents daily activity; grade 4, emergency room visit or hospitalization) and redness and swelling (mild, 2.0-5.0 cm in diameter;

moderate, >5.0-10.0 cm in diameter; severe, >10.0 cm in diameter; grade 4: necrosis or exfoliative dermatitis for redness, and necrosis for swelling). Data were collected with the use of electronic diaries for 7 days after each vaccination.

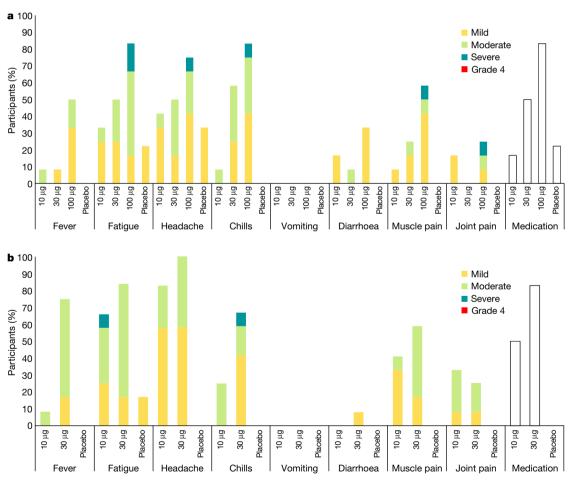


Fig. 3 | Systemic events and medication use reported within 7 days after vaccination. a, Systemic events and medication use reported within 7 days after vaccination 1 for all dose levels. b, Systemic events and medication use reported within 7 days after vaccination 2 for the 10-μg and 30-μg dose levels. Solicited systemic events were: fatigue, headache, chills, new or worsened  $muscle\ pain, new\ or\ worsened\ joint\ pain\ (mild, does\ not\ interfere\ with\ activity;$ moderate, some interference with activity; severe, prevents daily activity), vomiting (mild, 1-2 times in 24 h; moderate, >2 times in 24 h; severe, requires

intravenous hydration), diarrhoea (mild, 2-3 loose stools in 24 h; moderate, 4-5 loose stools in 24 h; severe: 6 or more loose stools in 24 h); grade 4 for all events: emergency room visit or hospitalization; and fever (mild, 38.0-38.4 °C; moderate, 38.5-38.9 °C; severe, 39.0-40.0 °C; grade 4, >40.0 °C). Medication indicates the proportion of participants who reported the use of antipyretic or pain medication. Data were collected with the use of electronic diaries for 7 days after each vaccination.

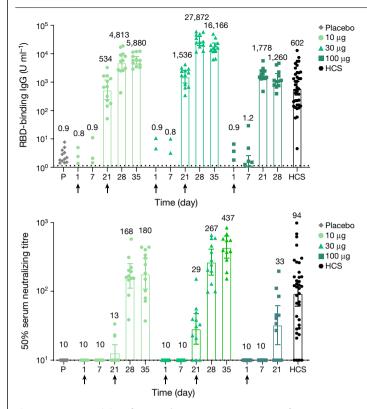


Fig. 4 | Immunogenicity of BNT162b1. Participants in groups of 15 were vaccinated with the indicated dose levels of BNT162b1 (n = 12) or with placebo (n=3) on days 1 (all dose levels and placebo) and 21 (10-µg and 30-µg dose levels and placebo). Reponses in individuals who received the placebo for each of the dosing groups are combined. The 28- and 35-day blood samples were obtained 7 and 14 days after the second vaccination. Sera were obtained before vaccination (day 1), and 7, 21, 28 and 35 days after the first vaccination. Human COVID-19 convalescent sera (HCS, n = 38) were obtained at least 14 days after PCR-confirmed diagnosis and at a time when the donors were asymptomatic. a, GMCs of recombinant RBD-binding IgG. Because the measured antibody concentrations using the Luminex assay are obtained in arbitrary units, they cannot be directly translated into concentrations on a molar or mass basis. The lower limit of quantitation is 1.15. b, The 50% SARS-CoV-2-neutralizing GMTs. Each data point represents a serum sample, and each vertical bar represents a geometric mean with 95% confidence interval. The number above the bars are either the GMC (a) or GMT (b) for the group. Arrows indicate the timing of vaccination (blood was obtained before vaccination on the vaccination days).

of age) had an RBD-binding IgG GMC of  $602~U~ml^{-1}$ . (Additional information on the convalescent serum panel is included in the Methods.) By 7 days after the second dose (for the 10- $\mu g$  and 30- $\mu g$  dose levels), RBD-binding IgG GMCs had increased to 4,813 and to  $27,872~U~ml^{-1}$ , respectively. RBD-binding antibody concentrations among participants who received one dose of  $100~\mu g$  BNT162b1 did not increase further at 21 days after the first vaccination. In the participants who received the 10- $\mu g$  and 30- $\mu g$  doses of BNT162b1, highly elevated RBD-binding antibody concentrations persisted to the last time point evaluated (day 35, 14 days after the second dose). These RBD-binding antibody concentrations were 5,880– $16,166~U~ml^{-1}$  compared to  $602~U~ml^{-1}$  in the panel of human convalescent sera.

For all doses, small increases in SARS-CoV-2-neutralizing geometric mean titres (GMTs) were observed 21 days after the first dose (Fig. 4b). Substantially greater serum neutralizing GMTs were achieved 7 days after the second 10- $\mu g$  and 30- $\mu g$  dose, reaching 168–267. Neutralizing GMTs further increased by 14 days after the second dose to 180 (10- $\mu g$  dose level) and 437 (30- $\mu g$  dose level), compared to 94 for the panel of human convalescent sera. The kinetics and durability of the neutralizing titres are being monitored.

#### Discussion

The RNA-based SARS-CoV-2 vaccine candidate BNT162b1, which was administered as 10-µg, 30-µg or 100-µg doses in healthy adults (18-55 years of age), exhibited a tolerability and safety profile consistent with those previously observed for mRNA-based vaccines<sup>5</sup>. A clear dose-level response in elicited neutralizing titres was observed after doses 1 and 2 in participants with a particularly steep dose response between the 10 µg and 30 µg dose levels.

On the basis of the tolerability profile of the first dose at  $100 \, \mu g$  and the second dose at  $30 \, \mu g$ , participants randomized to the  $100 \, \mu g$  group did not receive a second vaccination. Reactogenicity was generally greater after the second dose in the other two dosing levels; however, symptoms were transient and resolved within a few days. Transient decreases in lymphocyte counts (grades 1-3) were observed within a few days after vaccination, and returned to baseline within 6-8 days in all participants. These laboratory abnormalities were not associated with clinical findings. RNA vaccines are known to induce type-l interferon, which has been associated with transient migration of lymphocytes into tissues 10-22.

Robust immunogenicity was observed after vaccination with BNT162b1. RBD-binding lgG concentrations were detected at 21 days after the first dose, and these were substantially increased 7 days after the second dose given at day 21. After the first dose, the RBD-binding lgG GMCs (10- $\mu$ g dose) were similar to those observed in a panel of 38 convalescent human serum samples, obtained at least 14 days after a PCR-confirmed diagnosis of SARS-CoV-2 infection and/or COVID-19. After the first dose, GMCs were similar in the 30- $\mu$ g and 100- $\mu$ g groups and higher than those in the panel of human convalescent sera. After the second dose, with 10  $\mu$ g or 30  $\mu$ g BNT162b1, the RBD-binding lgG GMCs were around 8.0–50-fold that of the GMC of the convalescent serum panel.

The higher RBD-binding IgG GMC elicited by the vaccine relative to the GMC of the human convalescent serum panel may be attributed, in part, to antibodies that bind to epitopes that are exposed on the RNA-expressed RBD immunogen and the recombinant RBD target antigen of the binding assay but are buried and inaccessible to antibodies on the RBDs that are incorporated into the spikes of SARS-CoV-2 virions. Neutralization provides a measure of the vaccine-elicited antibody response that is more relevant to potential protection. Neutralization titres were measurable after a single vaccination at day 21 for all dose levels. At day 28 (7 days after the second dose), substantial SARS-CoV-2 neutralization titres were observed. The virus-neutralizing GMTs after the second dose of 10 µg and 30 µg were, respectively, 1.8-fold and 2.8-fold the GMT of the convalescent serum panel. By day 35 (14 days after the second dose)—despite a decrease in RBD-binding IgG titres since day 28-neutralizing GMTs continued to increase, to 1.9-fold and 4.6-fold the GMT of the convalescent panel for the 10  $\mu$ g and 30  $\mu$ g doses, respectively, which is consistent with affinity maturation.

Assuming that the neutralization titres that are induced by natural infection provide protection from COVID-19, comparing vaccine-induced SARS-CoV-2 neutralization titres to those from sera of convalescent humans provides a benchmark for the magnitude of the vaccine-elicited response and the potential of the vaccine to provide protection. Because the titre at which human neutralizing antibodies are protective remains unknown, these findings are not proof of vaccine efficacy. Efficacy will be determined in a pivotal phase III trial. Because the cohort that received the  $100~\mu g$  dose level did not receive the booster dose, no data for immunogenicity after a second vaccination at this dose level are available; however, there were no substantial differences in immunogenicity between the  $30~\mu g$  and  $100~\mu g$  dose levels after the first dose. This observation suggests that a well-tolerated and immunogenic dose level may be between  $10~\mu g$  and  $30~\mu g$  for this vaccine candidate.

Our study had several limitations. Although we used convalescent sera as a comparator, the kind of immunity (T cells versus B cells or both) and level of immunity needed to protect from COVID-19 are unknown. Furthermore, this analysis of available data did not assess

immune responses or safety beyond 2 weeks after the second dose of vaccine. Both are important to inform the public health use of this vaccine. Follow-up will continue for all participants and will include collection of serious adverse events for 6 months and COVID-19 infection and multiple additional immunogenicity measurements for up to 2 years. Although our population of healthy adults up to 55 years of age is appropriate for a phase I/II study, it does not accurately reflect the population at highest risk for COVID-19. Adults who are 65 years of age and over have already been enrolled in this study and results will be reported as they become available. Later phases of this study will prioritize enrolment of more diverse populations, including those with chronic underlying health conditions and from racial and ethnic groups that are adversely affected by COVID-19<sup>23</sup>.

The clinical testing of BNT162b1 described here has taken place in the context of a broader, ongoing COVID-19-vaccine-development program. That program includes the clinical testing of three additional vaccine candidates, including candidates that encode the full-length spike protein, and a parallel trial in Germany, in which additional immune responses, including neutralizing responses against variant strains and cell-mediated responses, are being assessed<sup>24</sup>. The resulting comparative data will allow us to address whether a full-length spike immunogen, which presents additional epitopes, is better able to elicit high virus-neutralizing titres that are robust to potential antigenic drift of SARS-CoV-2 than the relatively small RBD immunogen that is encoded by BNT162b1. The clinical findings for the BNT162b1 RNA-based vaccine candidate are encouraging and strongly support accelerated clinical development, including efficacy testing, and at-risk manufacturing to maximize the opportunity for the rapid production of a SARS-CoV-2 vaccine to prevent COVID-19.

#### **Online content**

Any methods, additional references, Nature Research reporting summaries, source data, extended data, supplementary information, acknowledgements, peer review information; details of author contributions and competing interests; and statements of data and code availability are available at https://doi.org/10.1038/s41586-020-2639-4.

- World Health Organization. WHO Director-General's Opening Remarks at the Media Briefing on COVID-19, https://www.who.int/dg/speeches/detail/who-director-general-sopening-remarks-at-the-media-briefing-on-covid-19---7-september-2020 (2020)
- Coronavirus Resource Center, COVID-19 Dashboard by the Center for Systems Science and Engineering (CSSE) at Johns Hopkins University (JHU). https://coronavirus.jhu.edu/ map.html (Johns Hopkins University & Medicine, 2020).

- World Health Organization, Coronavirus Disease 2019 (COVID-19) Situation Report 154. https://www.who.int/docs/default-source/coronaviruse/situation-reports/20200622covid-19-sitrep-154.pdf?sfvrsn=d0249d8d 2 (2020).
- Alberer, M. et al. Safety and immunogenicity of a mRNA rabies vaccine in healthy adults: an open-label, non-randomised, prospective, first-in-human phase 1 clinical trial. Lancet 390, 1511-1520 (2017).
- Feldman, R. A. et al. mRNA vaccines against H10N8 and H7N9 influenza viruses of pandemic potential are immunogenic and well tolerated in healthy adults in phase 1 randomized clinical trials. Vaccine 37, 3326-3334 (2019).
- Kranz, L. M. et al. Systemic RNA delivery to dendritic cells exploits antiviral defence for cancer immunotherapy. Nature 534, 396-401 (2016)
- Şahin, U. et al. Personalized RNA mutanome vaccines mobilize poly-specific therapeutic immunity against cancer. Nature 547, 222-226 (2017).
- Petsch, B. et al. Protective efficacy of in vitro synthesized, specific mRNA vaccines against influenza A virus infection. Nat. Biotechnol. 30, 1210-1216 (2012).
- Rauch, S., Jasny, E., Schmidt, K. E. & Petsch, B. New vaccine technologies to combat outbreak situations, Front, Immunol, 9, 1963 (2018).
- Sahin, U., Karikó, K. & Türeci, Ö. mRNA-based therapeutics—developing a new class of drugs, Nat. Rev. Drug Discov. 13, 759-780 (2014).
- Karikó, K. et al. Incorporation of pseudouridine into mRNA yields superior nonimmunogenic vector with increased translational capacity and biological stability. Mol. Ther. 16, 1833-1840 (2008).
- He, Y. et al. Receptor-binding domain of SARS-CoV spike protein induces highly potent neutralizing antibodies: implication for developing subunit vaccine. Biochem. Biophys. Res. Commun. 324, 773-781 (2004).
- Zost, S. J. et al. Rapid isolation and profiling of a diverse panel of human monoclonal antibodies targeting the SARS-CoV-2 spike protein. Nat. Med. 26, 1422-1427 (2020).
- Brouwer, P. J. M. et al. Potent neutralizing antibodies from COVID-19 patients define multiple targets of vulnerability. Science 369, 643-650 (2020)
- Güthe, S. et al. Very fast folding and association of a trimerization domain from bacteriophage T4 fibritin. J. Mol. Biol. 337, 905-915 (2004).
- Bachmann, M. F. & Zinkernagel, R. M. Neutralizing antiviral B cell responses. Annu. Rev. Immunol. 15, 235-270 (1997).
- Pardi, N. et al. Expression kinetics of nucleoside-modified mRNA delivered in lipid nanoparticles to mice by various routes. J. Control. Release 217, 345-351 (2015).
- Walsh, E. E. RNA-based COVID-19 vaccine BNT162b2 selected for a pivotal efficacy study. Preprint at https://doi.org/10.1101/2020.08.17.20176651 (2020).
- Foster, G. R. et al. IFN-a subtypes differentially affect human T cell motility. J. Immunol. **173**, 1663-1670 (2004).
- 20. Hopkins, R. J. et al. Randomized, double-blind, placebo-controlled, safety and immunogenicity study of 4 formulations of Anthrax Vaccine Adsorbed plus CPG 7909 (AV7909) in healthy adult volunteers, Vaccine 31, 3051-3058 (2013).
- Regules, J. A. et al. A recombinant vesicular stomatitis virus Ebola vaccine. N. Engl. J. Med. 376, 330-341 (2017).
- Lai, L. et al. Emergency postexposure vaccination with vesicular stomatitis virus-vectored Ebola vaccine after needlestick. J. Am. Med. Assoc. 313, 1249-1255 (2015).
- Stokes, F. K. et al. Coronavirus disease 2019 case surveillance United States, January 22-May 30, 2020. MMWR Morb. Mortal. Wkly. Rep. 69, 759-765 (2020).
- Şahin, U. et al. Concurrent human antibody and T<sub>H</sub>1 type T-cell responses elicited by a COVID-19 RNA vaccine. Preprint at https://doi.org/10.1101/2020.07.17.20140533 (2020).

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## **Methods**

#### Study design

This study was conducted in healthy men and women (who were not pregnant) who were 18–55 years of age to assess the safety, tolerability and immunogenicity of ascending dose levels of various BNT162 mRNA vaccine candidates. In the part of the study reported here, assessment of three dose levels (10  $\mu g$ , 30  $\mu g$  or 100  $\mu g$ ) of the BNT162b1 candidate was conducted at two sites in the USA. This study used a sentinel cohort design with progression and dose escalation taking place after review of data from the sentinel cohort at each dose level. The study is registered at ClinicalTrials.gov (NCT04368728). The phase I portion of this study was observer-blinded at the site level. Investigators were blinded to participant-level study intervention assignment; but investigators were not blinded to group-level assignment for the dataset included in this Article.

#### **Eligibility**

Key exclusion criteria included individuals with known infection with human immunodeficiency virus, hepatitis C virus or hepatitis B virus; immunocompromised individuals and those with a history of autoimmune disease; and those with increased risk for severe COVID-19, previous clinical or microbiological diagnosis of COVID-19, receipt of medications intended to prevent COVID-19, previous vaccination with any coronavirus vaccine, a positive serological test for SARS-CoV-2 IgM and/or IgG at the screening visit, and a SARS-CoV-2 nucleic acid amplification test-positive nasal swab within 24 h before study vaccination.

The final protocol and informed consent document were approved by institutional review boards for each of the participating investigational centres. This study was conducted in compliance with all International Council for Harmonisation good clinical practice guidelines and the ethical principles of the Declaration of Helsinki. A signed and dated informed consent form was required before any study-specific activity was performed.

#### **End points**

In this report, results from the following study primary end points are presented: the proportion of participants who reported solicited local reactions, systemic events and use of antipyretic and/or pain medication within 7 days after vaccination, adverse events and serious adverse events (available up to around 45 days after dose 1), and the proportion of participants with clinical laboratory abnormalities 1 and 7 days after vaccination and grading shifts in laboratory assessments between baseline and 1 and 7 days after dose 1, and between dose 2 and 7 days after dose 2. Secondary end points included: SARS-CoV-2-neutralizing GMTs and SARS-CoV-2 RBD-binding IgG GMCs 7 and 21 days after dose 1, and 7 and 14 days after dose 2.

#### **Procedures**

Study participants were randomly assigned to a vaccine group using an interactive web-based response technology system with each group comprising 15 participants (12 active vaccine recipients and 3 placebo recipients). Participants received two 0.5-ml doses of either BNT162b1 or placebo, administered by intramuscular injection into the deltoid muscle.

BNT162b1 incorporates a good manufacturing practice-grade mRNA drug substance that encodes the trimerized SARS-CoV-2 spike glycoprotein RBD antigen. The coding sequence for the antigen has been deposited with GenBank (accession number, MN908947.3). The mRNA is formulated with lipids as the mRNA–lipid nanoparticle drug product. The vaccine was supplied as a buffered-liquid solution for intramuscular injection and was stored at  $-80\,^{\circ}\text{C}$ . The placebo was a sterile saline solution for injection (0.9% sodium chloride injection, in a 0.5-ml dose).

#### Safety assessments

Safety assessments included a 4-h observation after vaccination (for the first 5 participants vaccinated in each group), or a 30-min observation (for the remainder of participants) for immediate adverse events. The safety assessments also included self-reporting of solicited local reactions (redness, swelling and pain at the injection site), systemic events (fever, fatigue, headache, chills, vomiting, diarrhoea, muscle pain and joint pain), the use of antipyretic and/or pain medication in an electronic diary for 7 days after vaccination, and the reporting of unsolicited adverse events and serious adverse events after vaccination. Haematology and chemistry assessments were conducted at screening, 1 and 7 days after the first dose, and 7 days after the second dose.

There were protocol-specified safety stopping rules for all sentinel cohort participants. Both an internal review committee and an external data monitoring committee reviewed all safety data. No stopping rules were met before the publication of this report.

#### Human convalescent serum panel

The 38 human SARS-CoV-2 infection and/or COVID-19 convalescent sera were drawn from participants, who were 18-83 years of age, at least 14 days after PCR-confirmed diagnosis, and at a time when participants were asymptomatic. The mean age of the donors was 45 years of age. Neutralizing GMTs in subgroups of the donors were as follows:  $\leq$ 55 years of age, 82 (n=29); >55 years of age, 142 (n=9); symptomatic infections, 90 (n=35); asymptomatic infections, 156 (n=3). The antibody titre for the one individual who was hospitalized was 618. The sera were obtained from Sanguine Biosciences, the MT Group and Pfizer Occupational Health and Wellness.

#### **Immunogenicity assessments**

For immunogenicity assessments, 50 ml of blood was collected before each study vaccination, at 7 and 21 days after the first dose, and at 7 and 14 days after the second dose. In the RBD-binding IgG assay, a recombinant SARS-CoV-2 RBD containing a C-terminal Avitag (Acro Biosystems, SPD-C82E9) and no foldon domain was bound to streptavidin-coated Luminex microspheres. In brief, 1.25 × 10<sup>7</sup> microspheres/ml were coated with streptavidin by 1-ethyl-3-[3-dimethylaminopropyl] carbodiimide hydrochloride reaction. Recombinant RBD Avitag was coupled to streptavidin beads by incubating for 90 min at room temperature with shaking (35 rpm). Beads were blocked in 1% BSA buffer for 30 min at room temperature. Heat-inactivated serum from participants was diluted 1:500, 1:5,000 and 1:50,000 in assay buffer (PBS with 0.5% BSA, 0.05% Tween-20 and 0.02% sodium azide). Following a 16–20-h incubation at 2–8 °C with shaking (300 rpm), plates were washed three times in a solution containing 0.05% Tween-20. An R-phycoerythrin-conjugated goat anti-human polyclonal antibody (Jackson Labs) was then added to plates for 90 min at room temperature with shaking (300 RPM). Plates were then washed a final time in a solution containing 0.05% Tween-20. Data were captured as median fluorescent intensities using a Luminex reader and converted to U/ml antibody concentrations using a reference standard curve with arbitrary assigned concentrations of 100 U/ ml and accounting for the serum dilution factor. The reference standard was composed of a pool of five COVID-19 convalescent serum samples (>14 days after PCR diagnosis). Three dilutions are used to increase the likelihood that at least one result for any sample will fall within the usable range of the standard curve. Assay results were reported in U/ ml of IgG. The final assay results are expressed as the GMC of all sample dilutions that produced a valid assay result within the assay range.

The SARS-CoV-2 neutralization assay used a previously described strain of SARS-CoV-2 (USA\_WA1/2020) that had been rescued by reverse genetics and engineered by the insertion of an mNeonGreen gene into open-reading frame 7 of the viral genome<sup>25</sup>. This reporter virus generates similar plaque morphologies and indistinguishable growth curves from the wild-type virus. Viral master stocks  $(2 \times 10^7)$ 

plaque-forming units per ml) used for the neutralization assay were grown in Vero E6 cells as previously described<sup>25</sup>. When testing patient convalescent serum specimens, the fluorescent neutralization assay produced comparable results as the conventional plaque reduction neutralization assay<sup>26</sup>. In brief, serial dilutions of heat-inactivated sera from participants were incubated with the reporter virus to yield an infection rate of approximately 10-30% of the Vero monolayer) for 1 h at 37 °C before inoculating Vero CCL81 cell monolayers (targeted to have 8,000–15,000 cells per well) in 96-well plates to enable the accurate quantification of infected cells. Total cell counts per well were enumerated by nuclear stain (Hoechst 33342) and fluorescent virally infected foci were detected 16-24 h after inoculation with a Cytation 7 Cell Imaging Multi-Mode Reader (BioTek) with Gen5 Image Prime v.3.09. Titres were calculated in GraphPad Prism v.8.4.2 by generating a four-parameter logistical fit of the percentage neutralization at each serial serum dilution. The 50% neutralization titre was reported as the interpolated reciprocal of the dilution that yielded a 50% reduction in fluorescent viral foci.

#### Statistical analysis

The sample size for the reported part of the study was not based on statistical hypothesis testing. The primary safety objective was evaluated by descriptive summary statistics for local reactions, systemic events, abnormal haematology and chemistry laboratory parameters, adverse events and serious adverse events after each vaccine dose for each vaccine group. The secondary immunogenicity objectives were descriptively summarized at the various time points. All participants with data available were included in the safety and immunogenicity analyses.

#### **Reporting summary**

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

#### **Data availability**

Upon request, and subject to review, Pfizer will provide the data that support the findings of this study. Subject to certain criteria, conditions and exceptions, Pfizer may also provide access to the related individual anonymized participant data. See https://www.pfizer.com/science/clinical-trials/trial-data-and-results for more information. These data

are interim data from an ongoing study for which the database is not locked. Data have not yet been source-verified or subjected to standard quality check procedures that would occur at the time of database lock and may therefore be subject to change.

- Xie, X. et al. An infectious cDNA clone of SARS-CoV-2. Cell Host Microbe 27, 841–848 (2020).
- Muruato, A. E. et al. A high-throughput neutralizing antibody assay for COVID-19 diagnosis and vaccine evaluation. Nat. Commun. 11, 4059 (2020).

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**Author contributions** K.U.J., P.R.D., W.C.G., N.K., S.L., A.G., R.B., O.T. and U.Ş. were involved in the design of the overall study and strategy. K.N., M.J.M., E.E.W., R.F. and A.R.F. provided feedback on the study design. W.K., D.C., K.A.S., K.R.T., C.F.-G. and P.Y.S. performed the immunological analyses. M.J.M., K.N., E.E.W., R.F., A.R.F., K.E.L. and V.R. collected data as study investigators. P.L. and K.K. developed the statistical design and oversaw the data analysis. J.A., K.U.J., P.R.D. and W.C.G. drafted the initial version of the manuscript. All authors reviewed and edited the manuscript and approved the final version.

Competing interests N.K., J.A., A.G., S.L., R.B., K.A.S., P.L., K.K., W.K., D.C., K.R.T., P.R.D., W.C.G. and K.U.J. are employees of Pfizer and may hold stock options. U.Ş. and Ö.T. are stock owners, management board members and employees at BioNTech and are inventors on patents and patent applications related to RNA technology. M.J.M., K.E.L., K.N., E.E.W., A.R.F., R.F. and V.R. received compensation from Pfizer for their role as study investigators. C.F.-G. and P.-Y.S. received compensation from Pfizer to perform the neutralization assay.

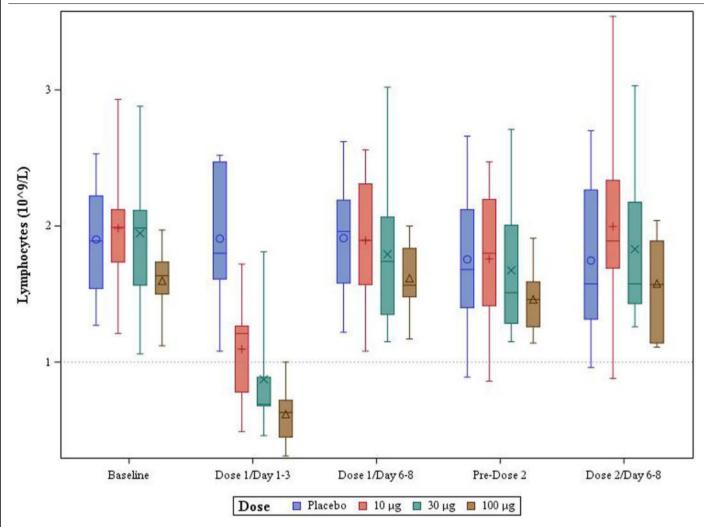
#### Additional information

**Supplementary information** is available for this paper at https://doi.org/10.1038/s41586-020-2639-4.

Correspondence and requests for materials should be addressed to J.A.

Peer review information Nature thanks Barbra Richardson and the other, anonymous, reviewer(s) for their contribution to the peer review of this work.

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**Extended Data Fig. 1** | **Post vaccination changes in lymphocyte count over time.** The following time points are shown: dose 1/day1-3, around 1 day after dose 1; dose 1/day6-8, around 7 days after dose 1; pre-dose 2, before dose 2; dose 2/day6-8, around 7 days after dose 2. Symbols denote group means;

circle, placebo; plus,  $10\,\mu g$ ; cross,  $30\,\mu g$ ; triangle,  $100\,\mu g$ . The box-and-whisker plots show the median (centre), first and third quartiles (lower and upper edges), and minimum and maximum values (lower and upper whiskers).

## Extended Data Table 1 | Demographic characteristics

	10 μg (N=12) n (%)	30 μg (N=12) n (%)	100 μg (N=12) n (%)	Placebo (N=9) n (%)	Total (N=45) n (%)
Sex					
Male	7 (58.3)	6 (50.0)	5 (41.7)	5 (55.6)	23 (51.1)
Female	5 (41.7)	6 (50.0)	7 (58.3)	4 (44.4)	22 (48.9)
Race					
White	8 (66.7)	10 (83.3)	11 (91.7)	8 (88.9)	37 (82.2)
Black or African American	1 (8.3)	0	0	0	1 (2.2)
Asian	3 (25.0)	2 (16.7)	1 (8.3)	1 (11.1)	7 (15.6)
Ethnicity					
Hispanic/Latino	1 (8.3)	1 (8.3)	0	0	2 (4.4)
Non-Hispanic/non-Latino	11 (91.7)	10 (83.3)	12 (100.0)	9 (100.0)	42 (93.3)
Not reported	0	1 (8.3)	0	0	1 (2.2)
Age at vaccination (years)					
Mean (SD)	29.4 (6.39)	35.8 (9.96)	38.3 (9.34)	39.0 (11.16)	35.4 (9.71)
Median	26.5	33.5	38.0	41.0	33.0
Min, max	(24, 42)	(23, 52)	(25, 53)	(19, 54)	(19, 54)

N, the number of participants in the specified group or the total sample. This value is the denominator for the percentage calculations. n, the number of participants with the specified characteristic.

## Extended Data Table 2 | Adverse events

	10 μg (N=12)	30 μg (N=12)	100 μg (N=12)	Placebo (N=9)
Adverse Event	n (%)	n (%)	n (%)	n (%)
Any event	6 (50.0)	6 (50.0)	7 (58.3)	1 (11.1)
Related	3 (25.0)	6 (50.0)	6 (50.0)	1 (11.1)
Severe	0	1 (8.3)	1 (8.3)	0
Life-threatening	0	0	0	0
Any serious adverse event	0	0	0	0
Related	0	0	0	0
Severe	0	0	0	0
Life-threatening	0	0	0	0
Any adverse event leading to withdrawal	0	0	0	0
Related	0	0	0	0
Severe	0	0	0	0
Life-threatening	0	0	0	0
Death	0	0	0	0

N, the number of participants in the specified group or the total sample. This value is the denominator for the percentage calculations. n, the number of participants who reported at least one occurrence of the specified adverse event category. For 'any event', n indicates the number of participants who reported at least one occurrence of any adverse event. Related, assessed by the investigator as related to the investigational product.

# nature research

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Last updated by author(s):	Jul 27, 2020

## **Reporting Summary**

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For	all statistical ar	nalyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.				
n/a	Confirmed					
	The exact	sample size $(n)$ for each experimental group/condition, given as a discrete number and unit of measurement				
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	The statis Only comm	itical test(s) used AND whether they are one- or two-sided non tests should be described solely by name; describe more complex techniques in the Methods section.				
$\boxtimes$	A descrip	tion of all covariates tested				
$\boxtimes$	A descrip	tion of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons				
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$\boxtimes$	For null hypothesis testing, the test statistic (e.g. <i>F</i> , <i>t</i> , <i>r</i> ) with confidence intervals, effect sizes, degrees of freedom and <i>P</i> value noted <i>Give P values as exact values whenever suitable.</i>					
$\boxtimes$	For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings					
$\boxtimes$	For hierar	rchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes				
$\boxtimes$	$\square$ Estimates of effect sizes (e.g. Cohen's $d$ , Pearson's $r$ ), indicating how they were calculated					
	Our web collection on <u>statistics for biologists</u> contains articles on many of the points above.					
So	ftware an	d code				
Poli	cy information	about availability of computer code				
Da	Pata collection (Inform (for data collected in the case report form) and electronic diary (Signant Health platform) for participant self reported reactogenicity					
Da	ata analysis	SAS 9.4				
For m		g custom algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors and				

## Data

Policy information about <u>availability of data</u>

All manuscripts must include a data availability statement. This statement should provide the following information, where applicable:

- Accession codes, unique identifiers, or web links for publicly available datasets
- A list of figures that have associated raw data
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Upon request, and subject to review, Pfizer will provide the data that support the findings of this study. Subject to certain criteria, conditions and exceptions, Pfizer may also provide access to the related individual anonymized participant data. See https://www.pfizer.com/science/clinical-trials/trial-data-and-results for more information

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Life scier	nces stu	udy design			
All studies must dis	close on these	points even when the disclosure is negative.			
Sample size	The sample size for this interim report was not based on statistical hypothesis testing. A total of 45 participants were enrolled in this part of the study. For the purposes of tolerability and dose escalation study a total of 15 participants (12 receiving vaccine and 3 receiving placebo) was deemed sufficient for a dosing finding phase study.				
Data exclusions		All safety and immunogenicity data that were available at the time of the data snapshot were included in the interim report. No data were excluded from the analyses.			
Replication	This is an interi	im report of an ongoing human clinical trial. There was no attempt at replication of study findings			
Randomization	This is an randomized controlled trial. Study participants were randomly assigned to a vaccine group using an interactive web-based response technology system with each group comprising 15 participants (12 active vaccine recipients and 3 placebo recipients).				
Blinding	This is an observer blinded study which is investigator blinded but Sponsor unblinded during Stage 1 (the stage from which data in the manuscript are presented). Investigators were unblinded to group level data but not subject level data for the purposes of interpretation and summary of the results included in this interim report.				
We require information	on from authors	pecific materials, systems and methods about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.			
Materials & exp	perimental s	systems Methods			
n/a Involved in th	ie study	n/a Involved in the study			
Antibodies		ChIP-seq			
Eukaryotic cell lines		Flow cytometry			
Palaeontology and archaeology MRI-based neuroimaging					
Animals and other organisms					
Human research participants					
Unical data  Dual use research of concern					
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Human rese	arch parti	icipants			
Policy information	about <u>studies i</u>	nvolving human research participants			
infection with huma those with a history microbiological diag		Study participants were healthy men or women 18-55 years of age. Key exclusion criteria included individuals with known infection with human immunodeficiency virus, hepatitis C virus, or hepatitis B virus; immunocompromised individuals and those with a history of autoimmune disease; those with increased risk for severe COVID-19; previous clinical or microbiological diagnosis of COVID 19; receipt of medications intended to prevent COVID 19; previous vaccination with any coronavirus vaccine; a positive serological test for SARS-CoV-2 IgM and/or IgG at the screening visit; and a SARS-CoV-2 NAAT-			

positive nasal swab within 24 hours before study vaccination.

Recruitment

Study participants were recruited at the two individual sites and recruitment strategies were at the discretion of individual sites and could include identification of interested individuals from the sites local database or through advertising in the local community. Once recruited participants were screened for eligibility based on pre-specified protocol criteria. Eligible participants were then randomized to vaccine or placebo in a blinded manner. These processes therefore did not led themselves to enrollment biases however participants who did not know about the study may have had less of an opportunity to participate.

Ethics oversight

The study protocol was approved by the western institutional review board for one site and by the Langone Health New York University Institutional IRB prior to enrollment of any participants

Note that full information on the approval of the study protocol must also be provided in the manuscript.

## Clinical data

Policy information about <u>clinical studies</u>

All manuscripts should comply with the ICMJE guidelines for publication of clinical research and a completed CONSORT checklist must be included with all submissions.

Clinical trial registration | ClinicalTrials.gov identifier: NCT04368728

Study protocol

Details of protocol elements can be accessed from clinicaltrials.gov

Data collection

Data were collected at screening (up to 14 days before vaccination) and for randomized participants at the investigative site at baseline, 1 day, 7 days and 21 days, after Dose 1, 7 days after dose 2 and up to 14 days after dose 2. Both safety and/or serum  $collection\ for\ immunogenicity\ assessments\ were\ collected\ for\ all\ stated\ time\ points.\ In\ addition,\ reactogenicity\ data\ were\ assessed$ through participant self reports via an electronic diary for 7 days after dose 1.

Outcomes

In this interim report, the following study primary endpoints are presented: the proportion of participants reporting prompted local reactions, systemic events, and use of antipyretic and/or pain medication within 7 days after vaccination, AEs and serious adverse events (SAEs) (available through up to ~45 days after Dose 1), and the proportion of participants with clinical laboratory abnormalities 1 and 7 days after vaccination and grading shifts in laboratory assessments between baseline and 1 and 7 days after Dose 1 and between Dose 2 and 7 days after Dose 2. Secondary endpoints included: SARS CoV 2 neutralizing geometric mean titers (GMTs); SARS CoV 2 RBD-binding IgG geometric mean concentrations (GMCs) 7 and 21 days after Dose 1 and 7 and 14 days after Dose 2