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Immunogenicity [of CO](https://jamanetwork.com/journals/jama/fullarticle/2780202#)VID-19 mRNA Vaccinos Pregnant and Lactating Women

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COVID-19 Resource Center

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Key Points

Question [What is the immun](https://jamanetwork.com/journals/jama/pages/coronavirus-alert)ogenicity of COVID-19 messenger RNA (mRNA) vaccines in and lactating women?

Findings [In this cohort study involving 103 women who received a COVID](#page-0-0)-19 mRNA vace

Findings In this cohort study involving 103 women who received a COVID-19 mRNA vaccine, 30 of whom were pregnant and 16 of whom were lactating, immunogenicity was demonstrated in all, and vaccine-elicited antibodies were found in infant cord blood and breast milk. Pregnant and nonpregnant vaccinated women developed cross-reactive immune responses against SARS-CoV-2 variants of concern.

Meaning In a small convenience sample, COVID-19 mRNA vaccines were immunogenic in pregnant and lactating women and induced immune responses against SARS-CoV-2 variants.

Abstract

Importance Pregnant women are at increased risk of morbidity and mortality from COVID-19 but have been excluded from the phase 3 COVID-19 vaccine trials. Data on vaccine safety and immunogenicity in these populations are therefore limited.

Objective To evaluate the immunogenicity of COVID-19 messenger RNA (mRNA) vaccines in pregnant and lactating women, including against emerging SARS-CoV-2 variants of concern.

Design, Setting, and Participants An exploratory, descriptive, prospective cohort study enrolled 103 women who received a COVID-19 vaccine from December 2020 through March 2021 and 28 women who had confirmed SARS-CoV-2 infection from April 2020 through March 2021 (the last follow-up date was March 26, 2021). This study enrolled 30 pregnant, 16 lactating, and 57 neither pregnant nor lactating women who received either the mRNA-1273 (Moderna) or BNT162b2 (Pfizer-BioNTech) COVID-19 vaccines and 22 pregnant and 6 nonpregnant unvaccinated women with SARS-CoV-2 infection.

Main Outcomes and Measures SARS-CoV-2 receptor binding domain binding, neutralizing, and functional nonneutralizing antibody responses from pregnant, lactating, and nonpregnant women were assessed following vaccination. Spike-specific T-cell responses were evaluated using IFN-γ enzyme-linked immunospot and multiparameter intracellular cytokine–staining assays. Humoral and cellular immune responses were determined against the original SARS-CoV-2 USA-WA1/2020 strain as well as against the B.1.1.7 and B.1.351 variants.

Results This study enrolled 103 women aged 18 to 45 years (66% non-Hispanic White) who received a COVID-19 mRNA vaccine. After the second vaccine dose, fever was reported in 4 pregnant women (14%; SD, 6%), 7 lactating women (44%; SD, 12%), and 27 nonpregnant women (52%; SD, 7%). Binding, neutralizing, and functional nonneutralizing antibody responses as well as CD4 and CD8 T-cell responses were present in pregnant, lactating, and nonpregnant women following vaccination. Binding and neutralizing antibodies were also observed in infant cord blood and breast milk. Binding and neutralizing antibody titers against the SARS-CoV-2 B.1.1.7 and B.1.351 variants of concern were reduced, but T-cell responses were preserved against viral variants.

Introduction

Pregnant women with symptomatic COVID-19 have a higher risk of intensive care unit admechanical ventilation, and death compared with other women in their reproductive years. creases in preterm birth and stillbirth also have been observed in pregnancies complicated COVID-19.² Maternal-fetal virus transmission in utero is rare,² and it appears that newbo ceive passive immunity through antibody transfer via the placenta and from breast milk natural infection.^{3,4} Vaccination during pregnancy has reduced maternal morbidity and r from influenza and neonatal morbidity from pertussis through passive immunity.^{5,6}

The theoretical risks of COVID-19 vaccination in pregnancy and during lactation are limit the current vaccines have a favorable safety profile and high efficacy in nonpregnant ind The Cente[rs](#page-8-0) for Disease Control and Prevention⁷ recommen[de](#page-8-0)d that pregnant and lactat women have access to the available COVID-19 vaccines. In the month following Emerger Authorization of [2](#page-12-0) COVID-19 messenger RNA (mRNA) vaccines in December 2020, 11087 nant women received a COVID-19 vaccine in the United States.⁸ However, pregnant and women were excluded from phase 3 vaccine efficacy trials⁹⁻¹¹; thus, data on vaccine safe immunogenicity in these populations remain limited.

New genetic variants have evolved from the initial SARS-CoV-2 sequence. The D614G va associated with enhanced infectivity,¹² the B.1.1.7 variant is associated with greater trans ty,¹³ and the B.1.351 variant appears to evade natural immunity from prior infection^{14,15} and tially escapes from neutralizing antibodies. The objective of this study was to assess the genicity of the current COVID-19 mRNA vaccines in pregnant and lactating women again the original SARS-CoV-2 USA-WA1/2020 strain as well as against the B.1.1.7 and B.1.351 v of concern.

Methods

Study Population

The Beth Israel Deaconess Medical Center institutional review board approved this study parent biorepository study; participants provided written informed consent. We conduct

ferred from flyers posted in the hospital. All participants provided blood, some provided infant cord blood at delivery, and some provided breast milk. Samples were collected close to cine dose and 2 to 8 weeks after the second dose for the mRNA-1273 (Moderna) or BNT1 (Pfizer-BioNTech) COVID-19 vaccine. The analysis presented herein includes pregnant, la and nonpregnant women aged 18 to 45 years who were vaccinated or infected (Table 1). ther characterize the study population, participants were asked to provide their race and ty based on specified categories for each; they could select multiple race categories. Participants also reported if they had fever symptoms following either vaccine dose.

Enzyme-Linked Immunosorbent Assay

SARS-CoV-2 spike receptor binding domain (RBD)-specific binding antibodies in serum a were assessed by enzyme-linked immunosorbent assay (ELISA) (Table 2). The 96-well pl were coated with 2 μg/mL of wild-type SARS-CoV-2, variant B.1.1.7 (containing mutation (A.G. Schmidt),¹⁶ or B.1.351 (containing mutations K417N, E484K, N501Y) RBD protein in becco phosphate-buffered saline (DPBS) and incubated at 4 °C overnight.

After incubation, plates were washed once with wash buffer (0.05% Tween 20 in 1× DPE blocked with 350 μL of casein block solution per well for 2 to 3 hours at room temperature. lowing incubation, block solution was discarded and plates were blotted dry. Serial dilut heat-inactivated serum or breast milk diluted in Casein block were added to wells, and p were incubated for 1 hour at room temperature, prior to 3 more washes and a 1-hour incu with a 1:4000 dilution of anti-human IgG horseradish peroxidase (HRP) (Invitrogen, The er Scientific) or a 1:1000 dilution of anti-human IgA HRP (Bethyl Laboratories Inc) at roo perature in the dark. Plates were washed 3 times, and 100 μL of SeraCare KPL TMB Sure Start solution was added to each well; plate development was halted by adding 100 µL of Care KPL TMB Stop solution per well. The absorbance at 450 nm, with a reference at 65 was recorded with a VersaMax microplate reader (Molecular Devices). For each sample, the ELISA end point titer was calculated using a 4-parameter logistic curve fit to calculate the recip serum dilution that yields a corrected absorbance value (450 nm-650 nm) of 0.2. Interp end point titers were reported.

Pseudovirus Neutralizing Antibody Assay

cession number, EPI_ISL_712096). The supernatants containing the pseudotype viruses were collected 48 h[o](#page-12-0)urs after transfection; pseudotype viruses were purified by filtration with a filter.

To determine the neutralization activity of human adult and infant cord blood serum and breast milk, HEK293T-hACE2 cells were seeded in 96-well tissue culture plates at a dens 1.75×10^4 cells per well overnight. Three-fold serial dilutions of heat-inactivated serum s were prepared and mixed with 50 µL of pseudovirus. The mixture was incubated at 37 °C hour before adding to HEK293T-hACE2 cells. After 48 hours, cells were lysed in Steadyciferase Assay (Promega Corp) according to the manufacturer's instructions. SARS-CoV-2 ization titers (NT50) were defined as the sample dilution at which a 50% reduction in re light units was observed relative to the average of the virus control wells.

Systems Serology

For the functional analysis of sera samples, bead-based assays were used to quantify anti dependent cellular phagocytosis (ADCP), antibody-dependent neutrophil phagocytosis (and antibody-dependent complement deposition (ADCD), as previously described.¹⁹ Flu streptavidin beads (ThermoFisher) were coupled to biotinylated SARS-CoV-2 Spike trime Pharma) and incubated with diluted serum (ADCP and ADNP, 1:100; ADCD, 1:10). For AD cells (ATCC), derived from a human monocytic cell line, were added to the immune comp and incubated for 16 hours at 37 °C. For ADNP, primary neutrophils were isolated using a nium chloride potassium lysis buffer from whole blood.

After a 1-hour incubation at 37 °C, neutrophils were stained with an anti-CD66b PacBlue tion antibody (Biolegend). For the ADCD assay, lyophilized guinea pig complement comp (C3b) (Sigma) was resuspended according to manufacturer's instructions and diluted in a veronal buffer with calcium and magnesium (Boston BioProducts). After incubation, C3 v tected with fluorescein-conjugated goat IgG fraction to guinea pig complement C3 (MPb). ADCP, events were gated on bead-positive cells, whereas neutrophils were defined as CI positive followed by gating on bead-positive neutrophils for ADNP. ADCP and ADNP data reported as the phagocytic score, calculated using the following formula: phagocytic score {[percentage of bead-positive cells] × [geometric mean MFI (mean fluorescence index) for beadpositive cells]}/1000. ADCD was reported as the MFI of C3 deposition.

positive cells]}/1000. ADCD was reported as the MFI of C3 deposition.

IFN-**γ** Enzyme-Linked Immunospot Assay

Enzyme-linked immunospot (ELISPOT) assay plates were coated with mouse anti–human IFN-γ monoclonal antibody (MabTech) at 1 μg per well and incubated overnight at 4 °C. Plates were washed with DPBS and blocked with R10 media (RPMI with 10% heat-inactivated fetal bovine serum [FBS] with 1% of 100× penicillin-streptomycin, 1 M of HEPES buffer, 100 mM of sodium pyruvate, 200 mM of L-glutamine, and 0.1% of 55 mM of 2-mercaptoethanol) for 2 to 4 hours at 37 °C. Peptides from wild-type, B.1.1.7, and B.1.351 variant spike (21st Century Biochemicals) were prepared and plated at a concentration of 2 μg per well, and 100000 cells per well were added to the plate.

The peptides and cells were incubated for 15 to 20 hours at 37 °C. All steps following this incubation were performed at room temperature. The plates were washed with an ELISPOT wash buffer and incubated for 2 to 4 hours with 1 μg/mL of biotinylated mouse anti–human IFN-γ monoclonal antibody (MabTech). The plates were washed again and incubated for 2 to 3 hours with 1.33 μg/mL of conjugated goat antibiotin alkaline phosphatase (Rockland Inc). The final wash was followed by adding nitor-blue tetrazolium and 5-bromo-4-chloro-3-indolyphosphate *p*-toludine salt (NBT/BCIP chromogen) substrate solution for 7 minutes. The chromogen was discarded, and the plates were washed with water and dried in a dim place for 24 hours. Plates were scanned and counted on an immunospot analyzer (Cellular Technologies Ltd).

Intracellular Cytokine Staining Assay

Peripheral blood mononuclear cells were resuspended at a concentration of 10⁶ cells in 100 μL of R10 media supplemented with a CD49d monoclonal antibody (1 μg/mL) and a CD28 monoclonal antibody (1 μg/mL). Each sample was assessed with mock (100 μL of R10 plus 0.5% dimethyl sulfoxide; background control), peptides (2 μg/mL), and/or 10 pg/mL of phorbol myristate acetate and 1 μg/mL of ionomycin (Sigma-Aldrich) (100 μL; positive control) and incubated at 37 °C for 1 hour. After incubation, 0.25 μL of GolgiStop (BD Bioscience), which contains monensin, and 0.25 μL of GolgiPlug (BD Bioscience), which contains brefeldin A, in 50 μL of R10 was added to each well and incubated at 37 °C for 8 hours and then held at 4 °C overnight.

The next day, the cells were washed twice with DPBS, stained with aqua live-or-dead dye for 10 minutes, and then stained with predetermined titers of monoclonal antibodies (mAbs) against CD279 (clone EH12.1, BB700), CD4 (clone L200, BV711), CD27 (clone M-T271, BUV563), CD8 (clone SK1, BUV805), and CD45RA (clone 5H9, APC H7) for 30 minutes. Cells were then washed twice with a 2% FBS-DPBS buffer and incubated for 15 minutes with 200 μL of BD CytoFix/Cyto-Perm fixation/permeabilization solution. Cells were washed twice with 1× Perm Wash buffer (BD Biosciences Perm/Wash Buffer 10× in the CytoFix/CytoPerm Fixation/ Permeabilization kit diluted with MilliQ water and passed through 0.22-μm filter) and stained intracellularly with mAbs

Statistical Analysis

Descriptive statistics were calculated using SAS 9.4 (SAS Institute Inc) and GraphPad Pris (GraphPad Software). Data are presented as median with interquartile range (IQR) or pro with standard deviation (SD).

Results

Enrollment

The hospital-wide biorepository enrolled 103 women aged 18 to 45 years who received and management of the mann COVID-19 vaccine and had serum available for analysis; an additional 4 individuals declin participate. Among these 103 participants, 30 were pregnant; 16 were lactating; and 57 ther pregnant nor lactating (Table 1). Samples were obtained a median of 21 days (IQR, 1 days) after the second vaccine dose from nonpregnant women, 21 days (IQR, 14-36 days pregnant women, and 26 days (IQR, 19-31 days) from lactating women. Nine pregnant women delivered during the study and contributed infant cord blood. Fifty-six participants (54%) ceived BNT162b2; 47 (46%) received mRNA-1273. Prior SARS-CoV-2 infection was diagno $(4%)$ of the vaccinated participants. Among pregnant participants, 5 (17%) received their vaccine dose in the first trimester, 15 (50%) in the second, and 10 (33%) in the third.

The hospital-wide biorepository also enrolled 70 women aged 18 to 45 years who were to SARS-CoV-2, including 60 pregnant women; an additional 76 individuals declined to par The analysis presented herein includes 22 pregnant and 6 nonpregnant unvaccinated wo with SARS-CoV-2 infection as comparators who had serum available for analysis. These participants were more likely to self-identify as Black or Hispanic than were the vaccinated wo ble 1). Among women who had not been vaccinated but had been infected, the median to symptom onset (or positive polymerase chain reaction [PCR] test result among those wh asymptomatic) to sample collection was 12 days (IQR, 10-20 days) for nonpregnant won 41 days (IQR, 15-140 days) for pregnant women. Among participants infected but not vac 1 nonpregnant (17%) and 3 pregnant (14%) women experienced severe disease.

Reactogenicity

 A fter the second dose, fever was reported in 27 nonpregnant (520), SD; 70(), 4 pregnant

tating (23497) women after the second vaccine dose were higher than their baseline pre tion titers (28) (**Figure 1**A). Among pregnant women, median binding antibody titer was following vaccination and was 1321 after infection. The median binding antibody titer wa following vaccination and was 771 after infection in nonpregnant women. Similarly, the r pseudovirus NT50 in vaccinated nonpregnant (901), pregnant (910), and lactating indivi (783) were higher than the prevaccination titers (20) ($Figure 1B$). Among those who were vaccinated but were infected, the median NT50 values were 148 among those who were nant and 193 among those who were not pregnant. Among vaccinated individuals, ADNF was quantified [with med](#page-3-0)ian phagocytic score of 58 in nonpregnant, 27 in pregnant, and tating individuals (**Figure 1**C). The median MFI for ADCD among vaccinated nonpregnant was 376; pregnant women, 402; and lactating women, 333 (Figure 1D). The median phago score for ADCP among vaccinated nonpregnant women was 277; for pregnant women, 28 for lactating women, 249 (**Figure 1**E).

Binding and Neutralizing Antibodies in Cord Blood

Nine paired maternal and infant cord blood samples were used to evaluate transplacental fer of vaccine-elici[ted bind](#page-3-0)ing and neutralizing antibodies. Median maternal serum RBD ing antibody titers at delivery were 14953 compared with 1[9873 in c](#page-3-0)ord blood (Figure 2. median maternal NT50 at delivery was 1016 compared with 324 in cord blood (Figure 2B). unvaccinated infected mat[ernal and](#page-3-0) infant dyads, the median RBD IqG binding antibody delivery were 1342 in maternal sera and 635 in cord blood (Figure 2A), and the median N 151 in maternal sera compared with 164 in cord blood (**Figure 2**B).

Binding and Neutralizing Antibodies in Breast milk

RBD IgG and IgA binding antibodies and neutralizing antibodies were assessed in breast lowing vaccination and infection. The median serum IgG binding antibody titer was 2505 vaccination and 1593 following natural infection. Median breast milk IgG titer was 97 in ed and 203 in infected individuals (**Figure 2**C). The median serum IgA-binding antibodies 820 after vaccination and 152 after infection. The median breast milk IgA binding antibo 25 after vaccination and 1940 after infection (**Figure 2**D). The median NT50 in breast mil following vaccination and 153 following infection (**Figure 2**E).

cells, and CD8 central memory T cells were comparable in pregnant, lactating, and nonpregnant women (Figure 3).

Humoral and Cellular Immune Responses to SARS-CoV-2 Variants of Concern

Serum RBD IgG binding antibodies and neutralizing antibodies to the B.1.1.7 and B.1.351 v of concern were evaluated.²⁰ Binding antibody responses were comparable against wild USA-WA1/2020 and B.1.1.7 RBD proteins in nonpregnant, pregnant, and lactating womer infant cord samples but were lower for the B.1.351 RBD protein ($Figure 4A$). The median izing antibody titer in nonpregnant, pregnant, and lactating women was lower by 3.5-fo B.1.1.7 variant and 6-fold lower for the B.1.351 variant than for the USA-WA1/2020 variant 4B).

Spike-specific T-cell responses were also compared with the wildtype USA-WA1/2020, B B.1.351 peptides by ELISPOT and ICS assays following vaccination. There were no difference ELISPOT responses, CD4 T-cell responses, CD4 central memory T-cell responses, CD8 Tsponses, or CD8 central memory T-cell responses across these variants.

Discussion

COVID-19 mRNA vaccines were immunogenic, as quantified by both humoral and cellular responses, in pregnant, lactating, and nonpregnant, nonlactating women. Following the dose of the mRNA vaccines, 13% of pregnant women and 47% of nonpregnant women re fever. These findings need to be confirmed using the national v-safe Centers for Disease and Prevention registry.²¹ Moreover, similar to prior studies,²² this study validates that v tion elicits higher antibody responses than does infection.

The detection of binding and neutralizing antibodies in infant cord blood suggests efficient transplacental transfer of maternal antibodies. As with the recommendation for diphthe tetanus toxoids and acellular pertussis vaccination in pregnancy to protect vulnerable ne against pertussis, maternal COVID-19 vaccination in pregnancy may confer similar benefit newborns who may be ineligible for vaccination. Vaccination also elicited binding and ne ing antibodies in breast milk, although IgA responses were low in breast milk, with the e of early breast milk from participants receiving a vaccine during pregnancy. Differential breast Consistent with recent reports,^{15,25,26} reduced serum neutralizing antibody titers were e against the B.1.1.7 variant that was originally identified in the UK and particularly against the B.1.351 variant that was originally identified in South Africa. Both vaccinated pregnant w and infant cord blood showed reductions in neutralizing antibody titers against these variant contrast, minimal reductions were observed against these variants for nonneutralizing an binding and for CD4 and CD8 T-cell responses in both pregnant and nonpregnant women ing vaccination. These data suggest that there may be greater cross-reactivity for function neutralizing antibodies and cellular immune responses than for neutralizing antibodies a SARS-CoV-2 variants of concern. The mechanistic roles of these different immune respor protecting against COVID-19 infection and disease remain to be determined, but data from human primates suggest that both humoral and cellular immune responses may contribute protection.17

Limitations

This study has several limitations. First, the study size is small, and thus conclusions abo cine safety and tolerability could not be made. Second, the correlates of immunogenicity and protection against COVID-19 infection and disease have not yet been determined. Third, hort study rather than a randomized clinical trial, any differences in the findings among to groups cannot be assumed to be causal. Fourth, given the reliance on a convenience sam women who were willing to be vaccinated, the generalizability of the findings may be line Fifth, immune responses were evaluated at a short interval after vaccination; thus, concl regarding durability cannot be drawn from these results.

Conclusions

In this exploratory analysis of a convenience sample, receipt of a COVID-19 messenger R cine was immunogenic in pregnant women, and vaccine-elicited antibodies were transport infant cord blood and breast milk. Pregnant and nonpregnant women who were vaccinat veloped cross-reactive antibody responses and T-cell responses against SARS-CoV-2 vari concern.

Back to top

Author Contributions: Drs Collier and Barouch had full access to all of the data in the sti take responsibility for the integrity of the data and the accuracy of the data analysis.

Concept and design: Collier, Alter, Barouch.

Acquisition, analysis, or interpretation of data: Collier, McMahan, Yu, Tostanoski, Aguayo, Chandrashekar, Patel, Apraku Bondz[ie, Sellers, Barrett, Sanbo](http://jamanetwork.com/article.aspx?doi=10.1001/jama.2021.7563)rn, Wan, Chang, Anioke, NI Bradshaw, Jacob-Dolan, Feldman, Gebre, Borducchi, Liu, Schmidt, Suscovich, Linde, Hacker, Barouch.

Drafting of the manuscript: Collier, McMahan, Yu, Chang, Anioke, Barouch.

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Statistical analysis: Collier, Yu, Tostanoski, Chang, Hacker.

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Administrative, technical, or material support: Collier, McMahan, Yu, Aguayo, Ansel, Chandrashekar, Patel, Apraku Bondzie, Sellers, Barrett, Sanborn, Chang, Anioke, Bradshaw, Fe Gebre, Liu, Suscovich, Linde, Barouch.

Supervision: Collier, McMahan, Ansel, Nkolola, Schmidt, Alter, Barouch.

Other - Performing assays: Wan.

Other - methodology, resources, data curation: Alter.

Conflict of Interest Disclosures: Dr Suscovich reported that he is an employee at and own of SeromYx Systems Inc. Dr Linde reported that she is an employee of SeromYx Systems Alter reported cofounding and serving as a consultant to, and having a patent pending t **Funding/Support:** This study was funded by grant CA260476 from the National Institute Health (NIH), and grants from the Ragon Institute of Massachusetts General Hospital, M achusetts Institute of Technology, Harvard, the Massachusetts Consortium for Pathogen ness, and the Musk Foundation (DHB); AI146779 from the NIH (AGS); HD000849 from t productive Scientist Development Program from the Eunice Kennedy Shriver National In Child Health & Human Development and from Burroughs Wellcome Fund (AYC), AI00738 the Multidisciplinary AIDS Training Program (LHT), and TR002541 from the Harvard Clini Translational Science Center (MRH).

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