Letters

RESEARCH LETTER

COVID-19 Cases and Disease Severity in Pregnancy and Neonatal Positivity Associated With Delta (B.1.617.2) and Omicron (B.1.1.529) Variant Predominance

The Omicron (B.1.1.529) variant of SARS-CoV-2 has spread rapidly but appears to cause less severe disease than the Delta (B.1.617.2) variant.¹ During pregnancy, Delta was associated with increased COVID-19 severity,² but infections and sever-

+

Related articles

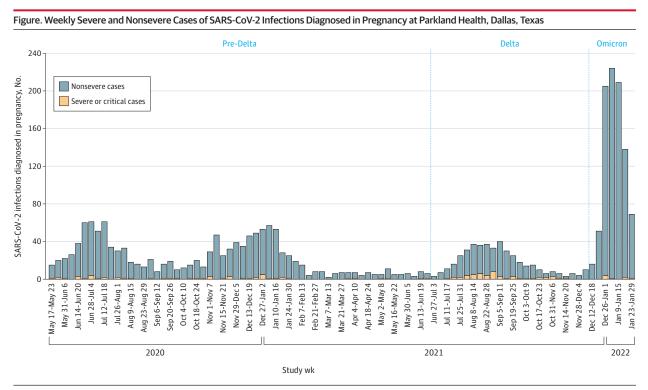
ity have not been examined during Omicron. We examined infections, illness sever-

ity, vaccinations, and early neonatal infections among obstetric patients during the pre-Delta, Delta, and Omicron epochs.

Methods | We prospectively studied pregnant patients diagnosed with SARS-CoV-2 infection at Parkland Health, an urban prenatal system encompassing a centralized acute care hospital and 10 community-based prenatal clinics around Dallas, Texas. Infections were defined based on a positive result of SARS-CoV-2 reverse transcriptase-polymerase chain reaction (RT-PCR) tests administered to all inpatients, symptom-based RT-PCR or antigen tests in outpatient and emergency departments, and patient-reported positive results of PCR or antigen tests conducted elsewhere. Positive tests were grouped by week of diagnosis. Severe or critical illness was defined as requiring supplemental oxygen, high-flow nasal cannula, mechanical ventilation, or extracorporeal membrane oxygenation.³ COVID-19 vaccines were offered beginning December 2020. Following delivery, neonatal nasal RT-PCR was performed at 24 and 48 hours for infants born within 4 weeks of maternal diagnosis or when clinically indicated until late 2021, when 1-time testing was permitted.

Local SARS-CoV-2 variant surveillance was conducted beginning in early 2021 using a combination of whole genome sequencing and genotyping using a validated multiplex, non-sequencing-based capillary electrophoresis assay in a Clinical Laboratory Improvement Amendments-certified laboratory.⁴

Poisson regression was used to model new cases each week. Covariates included dominant variant, week (to account for differing lengths of the 3 periods), and complete vaccination status. Dominant variant was assigned based on 50% or more of surveillance samples having Delta or Omicron predominance (pre-Delta epoch, May 17, 2020, through June 26, 2021; Delta epoch, June 27 through December 11, 2021; and Omicron epoch, December 12, 2021, through January 29, 2022). Complete vaccination was defined as at least 2 weeks elapsed following a complete primary series. Estimates are presented



The pre-Delta epoch is May 17, 2020, through June 26, 2021; Delta epoch, June 27 through December 11, 2021; and Omicron epoch, December 12, 2021, through January 29, 2022. Variant epoch was assigned based on greater than 50% of local surveillance samples having Delta or Omicron predominance.

jama.com

Table. COVID-19 Infections and liness Seventy in Pregnancy During Pre-Deita, Deita, and Omicron Epochs					
Variant epoch	No. of weeks	No. of infections	Incidence rate ratio (95% CI) ^a	Severe or critical illness, No. (%)	Odds ratio (95% CI) ^a
Pre-Delta	58	1298	1 [Reference]	53 (4.1)	1 [Reference]
Delta	24	431	3.07 (2.46-3.82)	51 (11.8)	2.93 (1.18-7.69)
Omicron	7	912	10.09 (7.42-13.69)	8 (0.9)	0.20 (0.05-0.83)

Table. COVID-19 Infections and Illness Severity in Pregnancy During Pre-Delta, Delta, and Omicron Epochs

^a The IRRs and ORs are adjusted for variant, week, and complete vaccination status, defined as 2 weeks or longer after a complete primary series.

as incidence rate ratios. Logistic regression was used to model case severity, with dominant variant, week, and complete vaccination status as covariates. Statistical analysis was conducted in R version 3.6.1; a 2-sided *P* < .05 was considered statistically significant. This study was approved by the University of Texas Southwestern institutional review board with a waiver of informed consent.

Results | The median number of deliveries per week was similar in the 3 periods (216 during pre-Delta, 239 during Delta, and 222 during Omicron epochs). There were 2641 SARS-CoV-2 infections diagnosed in pregnancy (Figure), 1298 (median, 17 cases per week) during the pre-Delta epoch, 431 (median, 14 cases per week) during Delta, and 912 (median, 138 cases per week) during Omicron. Two infected individuals (0.15%) were fully vaccinated pre-Delta, 49 (11.4%) during Delta, and 256 (28.1%) during Omicron. Of the 2641 cases, 112 (4.2%) were severe or critical. These included 53 (4.1%) severe or critical cases pre-Delta (0 vaccinated), 51 (11.8%) during Delta (2 vaccinated), and 8 (0.9%) during Omicron (2 vaccinated). Compared with the pre-Delta epoch, periods of Delta and Omicron predominance were associated with increased infections (incidence rate ratios, 3.07 [95% CI, 2.46-3.82] and 10.09 [95% CI, 7.42-13.69], respectively) (Table). Delta predominance was associated with increased (odds ratio, 2.93 [95% CI, 1.18-7.69]) and Omicron associated with decreased (odds ratio, 0.20 [95% CI, 0.05-0.83]) severe or critical illness in pregnancy compared with pre-Delta infections after adjusting for vaccination.

Of 1919 infants delivered during the study period, 1015 were tested and 32 (3.1%) were positive for SARS-CoV-2 (13 during pre-Delta, 8 during Delta, and 11 during Omicron epochs); none had significant illness. There was no difference in early neonatal positivity in the pre-Delta, Delta, or Omicron epochs (P = .39). Twenty-nine infants (90.6%) were born to individuals with nonsevere illness, and 18 (90%) of 20 infected neonates were born after December 2020 to vaccine-eligible but unvaccinated individuals.

Discussion As in nonpregnant people, Delta and Omicron variant predominance were associated with increased SARS-CoV-2 infections in pregnancy, with the majority occurring in unvaccinated individuals. Delta variant predominance was associated with increased illness severity and Omicron with decreased illness severity after adjusting for prior vaccination. The majority of early neonatal SARS-CoV-2 infections occurred among unvaccinated mothers with nonsevere COVID-19. Long-term risks of early neonatal SARS-CoV-2 infection are unknown, but maternal vaccination may be protective.⁵

Limitations include the cases from a single institution and potentially missing data on vaccination or positive results of tests conducted outside the health care system. Rates of SARS-CoV-2 exposure and vaccinations among uninfected individuals were not available. Whether the decreased illness severity during Omicron is related to greater numbers of pregnant people previously infected or vaccinated or to intrinsic virological properties cannot be determined.⁶

Emily H. Adhikari, MD Lorre MacDonald, CNM Jeffrey A. SoRelle, MD Jessica Morse, MD Jessica Pruszynski, PhD Catherine Y. Spong, MD

Author Affiliations: Department of Obstetrics and Gynecology, University of Texas Southwestern Medical Center, Dallas (Adhikari, Pruszynski, Spong); Parkland Health, Dallas, Texas (MacDonald); Department of Pathology, University of Texas Southwestern Medical Center, Dallas (SoRelle); Department of Pediatrics, University of Texas Southwestern Medical Center, Dallas (Morse).

Accepted for Publication: March 8, 2022.

Published Online: March 24, 2022. doi:10.1001/jama.2022.4356

Corresponding Author: Emily H. Adhikari, MD, Department of Obstetrics and Gynecology, University of Texas Southwestern Medical Center, 5323 Harry Hines Blvd, Dallas, TX 75390 (emily.adhikari@utsouthwestern.edu).

Author Contributions: Drs Adhikari and Pruszynski had full access to all the data in the study and take responsibility for the integrity of the data and the accuracy of the data analysis.

Concept and design: Adhikari, Spong.

Acquisition, analysis, or interpretation of data: All authors.

Drafting of the manuscript: Adhikari, MacDonald, Spong.

Critical revision of the manuscript for important intellectual content: Adhikari, SoRelle, Morse, Pruszynski, Spong.

Statistical analysis: Pruszynski.

Administrative, technical, or material support: Adhikari, MacDonald, SoRelle, Morse, Spong.

Supervision: Adhikari, Spong.

Conflict of Interest Disclosures: Dr SoRelle reported having a patent pending for genotyping PCR test for variants. No other disclosures were reported.

Funding/Support: Support for investigator and statistical effort, infrastructure, and sequencing effort was provided by the American Heart Association, the Doris Duke Charitable Foundation, the Harry S. Moss Heart Trust, the Once Upon a Time Foundation Next Generation Sequencing Lab, and the Centers for Disease Control and Prevention (grant NU5OCKO00501).

Role of the Funder/Sponsor: The above entities had no role in the design or conduct of the study; collection, management, analysis, and interpretation of the data; preparation, review, or approval of the manuscript, or decision to submit the manuscript for publication.

Additional Contributions: We acknowledge Mohammed Kanchwala, MS, McDermott Center Bioinformatics Core at University of Texas Southwestern Medical Center (not compensated specifically for this work), who performed all bioinformatic analysis and SARS-CoV-2 lineage assignment, and Ellen Araj, MD, Department of Pathology, University of Texas Southwestern Medical Center (not compensated specifically for this work), who created the clinical informatics platform to review and analyze sequencing data.

1. Iuliano AD, Brunkard JM, Boehmer TK, et al. Trends in disease severity and health care utilization during the early Omicron variant period compared with previous SARS-CoV-2 high transmission periods—United States, December 2020-January 2022. *MMWR Morb Mortal Wkly Rep.* 2022;71(4):146-152. doi:10. 15585/mmwr.mm7104e4

2. Adhikari EH, SoRelle JA, McIntire DD, Spong CY. Increasing severity of COVID-19 in pregnancy with Delta (B.1.617.2) variant surge. *Am J Obstet Gynecol*. 2022;226(1):149-151. doi:10.1016/j.ajog.2021.09.008

3. Society for Maternal-Fetal Medicine. Management considerations for pregnant patients with COVID-19. Accessed February 16, 2022. https://s3. amazonaws.com/cdn.smfm.org/media/2734/SMFM_COVID_Management_of_COVID_pos_preg_patients_2-2-21_(final).pdf

4. Addetia A, Lin MJ, Peddu V, Roychoudhury P, Jerome KR, Greninger AL. Sensitive recovery of complete SARS-CoV-2 genomes from clinical samples by use of Swift Biosciences' SARS-CoV-2 multiplex amplicon sequencing panel. *J Clin Microbiol*. 2020;59(1):e02226-e20. doi:10.1128/JCM.02226-20

5. Gale C, Quigley MA, Placzek A, et al. Characteristics and outcomes of neonatal SARS-CoV-2 infection in the UK: a prospective national cohort study using active surveillance. *Lancet Child Adolesc Health*. 2021;5(2):113-121. doi:10. 1016/S2352-4642(20)30342-4

6. Hui KPY, Ho JCW, Cheung MC, et al. SARS-CoV-2 Omicron variant replication in human bronchus and lung ex vivo. *Nature*. Published online February 1, 2022. doi:10.1038/s41586-022-04479-6