treatment follow-up visits with blood testing of malaria patients can be challenging in Haiti, healthcare professionals should strive to implement these goals. Implementation would enable continuous in vivo monitoring of drug susceptibility of parasites and provide real-time data to public health authorities to formulate evidence-based policy.

Acknowledgments
We thank Masami Nakatsu and Moritoshi Iwagami for providing advice on analysis of the k13 gene and the Eiken Chemical Co., Ltd. (Tokyo, Japan), for providing technical advice on loop-mediated isothermal amplification.

S.K. was partly supported by the Research Program on the Challenges of Global Health Issues: US–Japan Cooperative Medical Sciences Program (grant 16jk0210006h0001).

About the Author
Dr. Vincent is a doctoral candidate in the Department of Tropical Medicine and Malaria of the Research Institute National Center for Global Health and Medicine, Tokyo, Japan. Her research interests include Plasmodium biology, malaria diagnosis, field studies for controlling malaria, and other public health issues.

References

Address for correspondence: Shigeyuki Kano, Department of Tropical Medicine and Malaria, Research Institute, National Center for Global Health and Medicine, 1-21-1 Toyama, Shinjuku, Tokyo 162-8655, Japan; email: kano@ri.ncgm.go.jp


Scott W. Olesen, Yonatan H. Grad

Author affiliations: Harvard T.H. Chan School of Public Health, Boston, Massachusetts, USA (S.W. Olesen, Y.H. Grad); Brigham and Women’s Hospital, Boston (Y.H. Grad)

DOI: https://doi.org/10.3201/eid2411.180762

Using a US nationwide survey, we measured disparities in antimicrobial drug acquisition by race/ethnicity for 2014–2015. White persons reported twice as many antimicrobial drug prescription fills per capita as persons of other race/ethnicities. Characterizing antimicrobial drug use by demographic might improve antimicrobial drug stewardship and help address antimicrobial drug resistance.

Antimicrobial drug use varies by sex, age, and geography (1), and antimicrobial drug prescribing practice for specific medical conditions and age cohorts varies by patients’ race/ethnicity (2–4). Many studies on the role of patient race/ethnicity in antimicrobial drug prescribing practice focus on acute respiratory illnesses because antimicrobial drugs are often inappropriately prescribed for these conditions. The subjective diagnostic criteria for respiratory illnesses might result in race/ethnicity influencing prescribing practice more for these illnesses than for other illnesses (4). Despite our increasing knowledge of the role of patient race/ethnicity in drug prescribing practice for specific
conditions, how or whether these specific effects translate into overall antimicrobial drug use by race/ethnicity remains unclear. In this report, we address this gap in knowledge by describing the extent of racial/ethnic disparities in overall antimicrobial drug prescription fill rates in the United States.

We used the nationwide Medical Expenditure Panel Survey (MEPS) to acquire data about race/ethnicity and outpatient antimicrobial drug use for 2014–2015, the latest years with data available. MEPS contains data on members of a nationally representative sample of households (5,6). A person’s race/ethnicity is reported by the respondent and imputed in <0.1% of cases. Information about race and Hispanic ethnicity are collected in separate questions. Data regarding prescriptions filled at outpatient pharmacies are collected from the respondent and, if the respondent approves, verified with the filling pharmacy. Data are afterward cross-referenced and cleaned by survey preparers (5).

We used 2 exposure variables. The first was a categorical race/ethnicity variable with 5 values: Hispanic, non-Hispanic white only, non-Hispanic black only, non-Hispanic Asian only, and non-Hispanic other or multiple race. The second exposure variable was dichotomous and indicated whether white was the race or 1 of the races of the respondent. This exposure variable included all persons from the non-Hispanic white category, some from the Hispanic category, and some from the other or multiple race category. The main outcome was reported outpatient antimicrobial drug prescription fills per 1,000 persons per year stratified by major antimicrobial drug class (penicillins, macrolides, quinolones, sulfonamides, other). The complex survey design was accounted for when computing rates and CIs with survey package version 3.32 in R version 3.4.1 (online Technical Appendix, https://wwwnc.cdc.gov/EID/article/24/11/18-0762-Techapp1.pdf).

The reported annual outpatient prescription fill rate for all antimicrobial drugs was 373 (95% CI 358–388) fills/1,000 persons. This rate varied by race/ethnicity; non-Hispanic whites reported the highest rate, followed by persons of other or multiple race/ethnicity, Hispanics, non-Hispanic blacks, and non-Hispanic Asians (Figure, panel A). White persons reported 2.0 (95% CI 1.9–2.2)–fold more fills per capita than nonwhite persons (Figure, panel B). The reported fill rate disparity was similar for penicillins (2.0 [95% CI 1.8–2.4]–fold higher), sulfonamides (2.2 [95% CI 1.8–2.7]–fold higher), and quinolones (2.3 [95% CI 1.9–2.8]–fold higher) but smaller for macrolides (64% [95% CI 48%–82%] higher).

We found a large disparity in antimicrobial drug fill rates by race/ethnicity: white persons reported making twice as many antimicrobial drug prescription fills as persons who were not white. Disparities were apparent for each major antimicrobial drug class, rather than different drug classes being used more predominantly by persons of particular race/ethnicities.

This study is subject to several limitations. First, the survey is not a perfect measure of antimicrobial drug use. Survey respondents report medications they remember filling, subjecting results to systematic differences in respondents’ abilities to recall medications (6). Survey preparers then obtain information about those medications from the pharmacies the respondents visited, which themselves might have systematic differences in completeness of records (5). Also, respondents might not have complete information about other household members’ antimicrobial drug use or might choose not to disclose all antimicrobial drug use. Second, nonprescription antimicrobial drug use might be higher among minority groups (7), perhaps mitigating the observed disparity. Last, the survey measures reported antimicrobial drug fills and not actual use (8); the fill rates we report are substantially lower than those measured by others using sales data (1) or other national surveys (9).

Whether differences in antimicrobial drug use and race/ethnicity lead to disparities in antimicrobial drug resistance is unclear. We expect that disparities in use, regardless of their cause, will lead to disparities in the proportions of carried bacteria that are antimicrobial drug–resistant. Absolute rates of antimicrobial drug–resistant
infections, however, might follow different patterns (10). For example, higher macrolide use among white persons might lead to macrolide resistance in a greater proportion of *Streptococcus pneumoniae* bacteria carried by whites, but if white persons have fewer *S. pneumoniae* infections, then they would incur a lower absolute rate of macrolide-resistant *S. pneumoniae* infections. Further studies comparing antimicrobial drug use, antimicrobial drug resistance, and disease prevalence by race/ethnicity will be critical for addressing this question and improving antimicrobial drug stewardship.

S.W.O. was supported by cooperative agreement U54GM088558 from the National Institute of General Medical Sciences.

About the Author
Dr. Olesen is a postdoctoral fellow at the Harvard T.H. Chan School of Public Health, Boston, Massachusetts, USA. His research interests include antimicrobial drug use and resistance. Dr. Grad is the Melvin J. and Geraldine L. Glimcher Assistant Professor of Immunology and Infectious Diseases at the Harvard T.H. Chan School of Public Health. His research interests include antimicrobial drug use and resistance and the evolution and spread of pathogens.

References

Address for correspondence: Yonatan H. Grad, Harvard T. H. Chan School of Public Health, 665 Huntington Ave, Bldg 1, Rm 715, Boston, MA 02115, USA; email: ygrad@hsph.harvard.edu

**Congenital Zika Virus Infection with Normal Neurodevelopmental Outcome, Brazil**

**Alessandra Lemos de Carvalho, Carlos Brites, Tânia Barreto Taguchi, Suely Fernandes Pinho, Gúbio Campos, Rita Lucena**

Author affiliations: SARAH Network of Rehabilitation Hospitals, Salvador, Brazil (A.L. de Carvalho, T.B. Taguchi, S.F. Pinho); Federal University of Bahia, Salvador (C. Brites, G. Campos, R. Lucena)

DOI: https://doi.org/10.3201/eid2411.180883

We describe a case of a 20-month-old girl with probable congenital Zika virus infection and normal neurodevelopment, despite microcephaly and abnormal neuroimaging. This case raises questions about early prognostic markers and draws attention to the need for investigation in suspected Zika cases, even if the child’s early neurodevelopment is normal.

**Zika virus** is a mosquito-borne RNA virus (genus *Flavivirus*, family Flaviviridae) that was first isolated in 1947 from monkeys in the Zika Forest in Uganda (1). In November 2015, there was an outbreak of congenital microcephaly in the northeast states of Brazil (2). Further confirmation of this syndrome’s relationship with Zika virus infection during pregnancy was then possible (3). Congenital Zika syndrome has been recognized as a new clinical entity (4,5). Most published case series focus on the picture of severely affected infants (6,7). We describe a case of a child with probable congenital Zika virus infection whose neurodevelopment was normal, despite