

## State Legislation, Regulations, and Hospital Guidelines for Newborn Screening for Critical Congenital Heart Defects — United States, 2011–2014

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Critical congenital heart defects (CCHD) occur in approximately two of every 1,000 live births (*1*). Newborn screening provides an opportunity for reducing infant morbidity and mortality (*2,3*). In September 2011, the U.S. Department of Health and Human Services (HHS) Secretary endorsed the recommendation that critical congenital heart defects be added to the Recommended Uniform Screening Panel (RUSP) for all newborns (*4*). In 2014, CDC collaborated with the American Academy of Pediatrics (AAP) Division of State Government Affairs and the Newborn Screening Technical Assistance and Evaluation Program (NewSTEPs) to assess states' actions for adopting newborn screening for CCHD. Forty-three states have taken action toward newborn screening for CCHD through legislation, regulations, or hospital guidelines. Among those 43, 32 (74%) are collecting or planning to collect CCHD screening data; however, the type of data collected by CCHD newborn screening programs varies by state. State mandates for newborn screening for CCHD will likely increase the number of newborns screened, allowing for the possibility of early identification and prevention of morbidity and mortality. Data collection at the state level is important for surveillance, monitoring of outcomes, and evaluation of state CCHD newborn screening programs.

Congenital heart defects occur in approximately eight of every 1,000 live births, one fourth of which are considered to be CCHD (*1*). CCHD are defined as those requiring surgery or catheterization before age 1 year. In the absence of early detection, infants with CCHD are at risk for serious complications or death within the first few days or weeks of life (*1*). Newborn screening for CCHD uses pulse oximetry, a noninvasive technology to measure blood oxygen saturation. Low oxygen saturation indicates hypoxemia, an early clinical sign of CCHD. Additional testing (e.g., repeat screening, echocardiogram) is

needed following an abnormal pulse oximetry screen (*1*) to determine whether CCHD are present (or to determine the cause of the abnormal result). Thus, unlike most newborn screening conditions, screening for CCHD is not based on performing a blood test. In addition, hypoxemia detected by screening could indicate a medical problem, and requires immediate follow-up before discharge from the hospital.

When accompanied by early identification and treatment, newborn screening provides an opportunity to reduce infant morbidity and mortality (*2,3*). The Secretary's Advisory Committee on Heritable Disorders in Newborns and Children has provided national guidelines and recommendations on

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newborn screening, known as the RUSP, and this panel is reviewed and endorsed by the HHS Secretary (3). As of March 2015, 32 conditions were included in the RUSP. States use the RUSP as guidance when considering adopting conditions for their own screening panels (3). State decisions might differ depending on method of screening required or the legislative authority of the newborn screening program. When states add conditions to their state-specific screening panels, they do so by state legislation, or rules and regulations (5). In 2010, the Secretary's Advisory Committee on Heritable Disorders in Newborns and Children recommended adding CCHD to the RUSP for all newborns (4). In September 2011, the HHS Secretary endorsed the recommendation.

To assess states' actions for adopting newborn screening for CCHD, CDC collaborated with the AAP Division of State Government Affairs and NewSTEPs. AAP obtained primary information through direct contact and partnership with AAP state chapters. AAP monitored state legislation by use of tracking software; regulations and hospital guidelines were researched on state websites.

NewSTEPs is a program of the Association of Public Health Laboratories in collaboration with the Colorado School of Public Health, funded through a cooperative agreement from the Health Resources and Services Administration (6). NewSTEPs maintains a data repository of state newborn screening program metrics and provides education and technical assistance to newborn screening programs. In January 2014, NewSTEPs distributed a survey on CCHD newborn screening

adoption and data collection practices to state CCHD newborn screening programs. The survey requested the status of CCHD mandates and requirements for data collection. If data collection was required at the state level, additional information was requested on the type of data collected. All 50 states and the District of Columbia participated.

The survey findings indicated that 43 states have legislation, regulations, or hospital guidelines in place supporting CCHD newborn screening; 35 states have legislation, and 13 have regulations related to CCHD screening (Table). Among the 43, three states (Indiana, Maryland, and New Jersey) enacted legislation before the Secretary's approval of adding CCHD to the RUSP in 2011 (Table). State adoption of CCHD screening peaked in 2013 with 25 states adopting screening (Figure 1).

The manner in which these 43 states developed universal screening varied substantially (Figure 2), and for some was a multistage process (Table). For example, California passed legislation requiring that CCHD screening be offered to parents of newborns. In 2013, Pennsylvania issued a regulation requiring reporting of results and diagnoses of screened newborns. However, the regulation did not mandate screening. In 2014, Pennsylvania enacted a law requiring screening. In 2012, Tennessee initially passed legislation that required the state's genetic advisory committee to develop a program for addition of CCHD to its screening panel. In 2013, Tennessee added CCHD to its panel via regulation. In 2012, Virginia's governor issued an executive order establishing a work group to develop a CCHD screening implementation plan, and

The *MMWR* series of publications is published by the Center for Surveillance, Epidemiology, and Laboratory Services, Centers for Disease Control and Prevention (CDC), U.S. Department of Health and Human Services, Atlanta, GA 30329-4027.

**Suggested citation:** [Author names; first three, then et al., if more than six.] [Report title]. *MMWR Morb Mortal Wkly Rep* 2015;64:[inclusive page numbers].

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**TABLE. State approvals of legislation, regulation, and hospital guidelines for newborn screening for critical congenital heart defects (CCHD) — United States, 2011–2014**

State	Mechanism of current approval for CCHD screening				Screening supported as standard of care	Data collection system at state level	Type of data reported (current or proposed)
	Enacted date	Effective date	Legislation*	Regulation/Guidelines†			
Alabama	May 2013	June 2013		X <sup>§</sup>		Planned	All oxygen saturations/times on all failed screens
Alaska	September 2013	January 2014 (January 2016 for providers who attend fewer than 20 births a year)	X <sup>§</sup>			Yes	Aggregate data
Arizona	April 2014	July 2015	X <sup>§</sup>			Planned	All oxygen saturations/times
Arkansas	April 2013	August 2013	X <sup>§</sup>			Planned	Pass/Fail on all newborns
California*	October 2012	July 2013	X (Screening is required to be offered)			Yes	Pass/Fail on all newborns
Colorado					X	Planned	All oxygen saturations/times on diagnosed cases
Connecticut	May 2012	January 2013		X <sup>§</sup>		No	All oxygen saturations/times
Delaware	May 2013	May 2013		X <sup>§</sup>		Yes	Pass/Fail on all newborns
District of Columbia					X <sup>¶</sup>	Yes	All oxygen saturations/times (one hospital)
Florida	October 2014	October 2014		X <sup>§</sup>		Yes	Final oxygen saturations/times
Georgia	May 2014	June 2014		X <sup>§</sup>		Planned	All oxygen saturations/times
Hawaii**					X	Planned	All oxygen saturations/times
Idaho					X	No	
Illinois	August 2013	August 2013	X <sup>§</sup>			No	
Indiana	May 2011	January 2012	X <sup>§</sup>			Yes	All oxygen saturations/times
Iowa (guidelines)	August 2012	August 2012	X <sup>§</sup>	X		No	
Iowa (legislation)	April 2013	July 2013					
Kansas					X	Yes	All oxygen saturations/times (four hospitals); Aggregate data (other hospitals)
Kentucky	March 2013	January 2014	X <sup>§</sup>			Yes	All oxygen saturations/times; Echocardiogram results <sup>††</sup>
Louisiana	June 2013	August 2013	X <sup>§</sup>			No	
Maine	July 2013	July 2013	X <sup>§</sup>			Planned	All oxygen saturations/times
Maryland	May 2011	July 2011	X <sup>§</sup>			Yes	Pass/Fail on all newborns; Option to enter all oxygen saturations/times
Massachusetts (guidelines)	May 2013	May 2013	X <sup>§</sup>	X		Yes	Aggregate data only
Massachusetts (legislation)	March 2014	January 2015					
Michigan	October 2013	April 2014		X <sup>§</sup>		Yes	All oxygen saturations/times; Echocardiogram results
Minnesota	May 2013	August 2013	X <sup>§</sup>			Yes	All oxygen saturations/times
Mississippi	October 2014	November 2014		X <sup>§</sup>		Planned	Aggregate data
Missouri	July 2013	January 2014	X <sup>§</sup>			Yes	Aggregate data; Plan to include newborn data with all oxygen saturations/times
Montana	June 2014	July 2014		X <sup>§</sup>		Planned	Pass/Fail on all newborns
Nebraska	June 2013	September 2013	X <sup>§</sup>			No	
Nevada	June 2013	July 2015	X <sup>§</sup>			Yes	Aggregate data only (hospitals participating in a pilot program)
New Hampshire	June 2012	August 2012	X <sup>§</sup>			No	
New Jersey	June 2011	September 2011	X <sup>§</sup>			Yes	Aggregate data; Plan to collect all oxygen saturations/times

See table footnotes on next page.

**TABLE. (Continued) State approvals of legislation, regulation, and hospital guidelines for newborn screening for critical congenital heart defects (CCHD) — United States, 2011–2014**

State	Mechanism of current approval for CCHD screening			Regulation/ Guidelines <sup>†</sup>	Screening supported as standard of care	Data collection system at state level	Type of data reported (current or proposed)
	Enacted date	Effective date	Legislation*				
New Mexico	March 2014	May 2014	X <sup>§</sup>			Planned	All oxygen saturations/times
New York	July 2013	January 2014	X <sup>§</sup>			No	
North Carolina	May 2013	May 2013	X <sup>§</sup>			Yes	Aggregate data
North Dakota	April 2013	August 2013	X <sup>§</sup>			No	
Ohio	June 2013	September 2013	X <sup>§</sup>			Planned	All oxygen saturations/times
Oklahoma	April 2013	July 2013	X <sup>§</sup>			Yes	Pass/Fail on all newborns
Oregon	June 2013	June 2013	X <sup>§</sup>			No	
Pennsylvania (regulation) <sup>†</sup>	December 2012	March 2013	X <sup>§</sup>	X		Yes	Aggregate data only; Oxygen saturations/time for confirmed cases only
Pennsylvania (legislation)	(regulation) July 2014	(regulation) September 2014					
Rhode Island	August 2014	July 2015		X <sup>§</sup>		Yes	Pass/Fail on newborns (some hospitals)
South Carolina	June 2013	June 2013	X <sup>§</sup>			No	
South Dakota	March 2013	July 2013	X <sup>§</sup>			No	
Tennessee (legislation)*	March 2012	January 2013	X	X <sup>§</sup>		Yes	Pass/Fail and date/time of screen on all newborns
Tennessee (regulation) <sup>†</sup>	(legislation) May 2013	(legislation) May 2013					
Texas	(regulation) June 2013	(regulation) September 2013	X <sup>§</sup>			Yes	All oxygen saturations on diagnosed cases only
Utah	March 2013	October 2014	X <sup>§</sup>			Yes	Pass/Fail on all newborns Planned: All oxygen saturations/times
Vermont					X <sup>¶</sup>	Planned	Aggregate data only on all newborns; Oxygen saturations/times on failed screens
Virginia (executive order) <sup>§§</sup>	June 2012	June 2012	X <sup>§</sup>			Planned	Oxygen saturations/times on failed screens
Virginia (legislation)	February 2014	July 2014					
Washington					X <sup>¶</sup>	No	
West Virginia	March 2012	June 2012	X <sup>§</sup>			Yes	Pass/Fail on all newborns
Wisconsin*	March 2014	March 2014	X*	X <sup>§</sup>		Yes	Pass/Fail on all newborns; All oxygen saturations/times from some hospitals
(legislation)	(legislation)	(legislation)					
Wisconsin (regulation)	June 2014	July 2014					
Wyoming	(regulation)	(regulation)			X	Planned	All oxygen saturations/times

\* A total of 35 states have enacted legislation related to newborn screening for CCHD; 32 of those state laws require screening. California's law requires the screen to be offered to parents of newborns before discharge. Tennessee's law requires the state to develop a program for CCHD screening. Wisconsin's law allows the state department of health to add conditions or diseases to the state's newborn screening panel.

† A total of 13 states issued regulations or hospital guidelines related to newborn screening; 10 of those states issued regulations requiring screening. Iowa and Massachusetts issued guidelines to hospitals and birthing centers on screening, but the guidelines do not require screening. Pennsylvania issued a regulation requiring reporting of results and diagnoses of screened newborns, but the regulation does not require screening. Tennessee issued a regulation, after enacting legislation, adding CCHD to the state's newborn screening panel.

§ Mandates CCHD screening of newborns.

¶ State reports that all hospitals are performing CCHD screening.

\*\* Legislation in Hawaii to require screening failed in 2014.

†† Echocardiogram is the diagnostic test that follows a failed pulse oximetry screen.

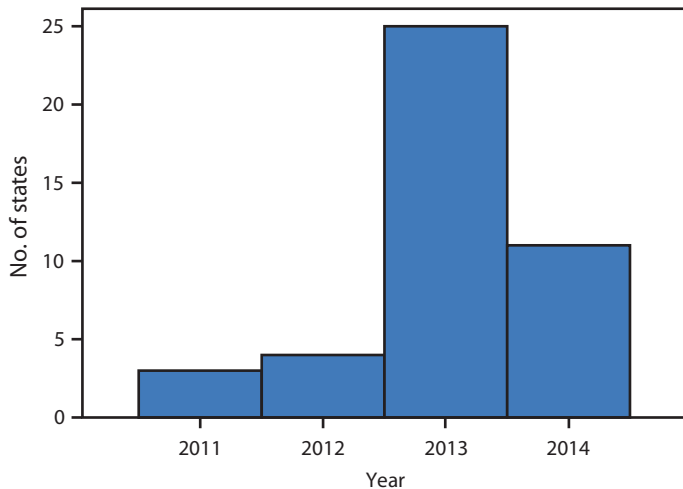
§§ Virginia's former governor issued a directive in 2012 that established a workgroup to develop a plan for implementing screening.

legislation for mandatory screening was passed in 2014. In 2013, Massachusetts issued guidelines that recommended hospitals screen newborns and passed mandatory screening legislation in 2014. In 2014, Wisconsin enacted a law that allows the state department of health to add conditions to its state panel via regulation. Soon after enactment, regulations were issued adding CCHD to its panel.

Seven states and the District of Columbia support CCHD newborn screening as the standard of care with no mandate in place. Two states and the District of Columbia report that all hospitals are screening for CCHD (Table).

By December 2014, among the 50 states and the District of Columbia, data collection within each newborn screening program varied from no data collection to collection of all

**FIGURE 1. Number of states (N = 43) adopting legislation, regulation, or hospital guidelines for universal newborn screening for critical congenital heart defects, by year — United States, 2011–2014**

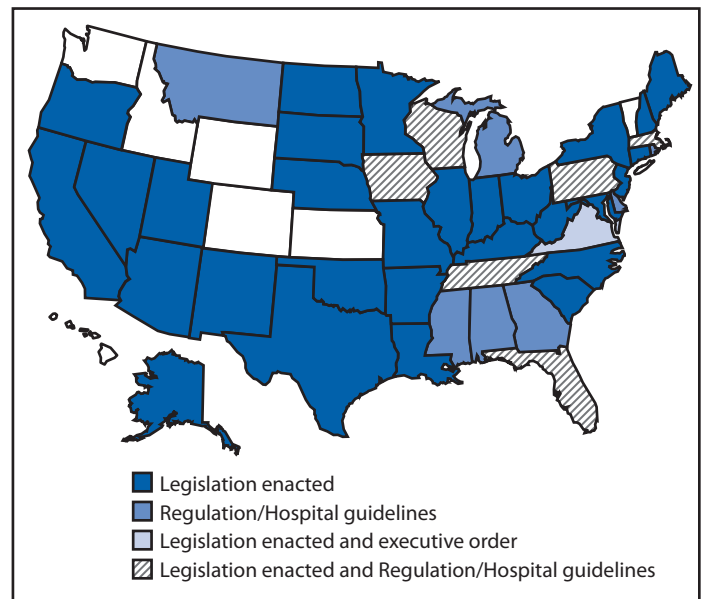


screening results for every newborn. Of the states that have implemented, or are planning to implement CCHD screening, 24 reported current data collection, 14 reported planning future data collection, and 13 reported no plans for data collection (Table). The types of data collection vary from aggregate data collection only, collection of pass/fail results on all newborns, oxygen saturation results on all newborns, oxygen saturation results on failed newborns only, or a combination of these (Table).

### Discussion

The increasing number of states mandating newborn screening for CCHD will likely increase the number of newborns screened, allowing for early identification and the potential for the prevention of morbidity and mortality. Most newborn screening conditions are tested through a heel stick test, with bloodspot analysis at public health or contracted laboratories. Screening for CCHD is a point-of-care test that occurs in hospitals before a newborn is discharged, with results entered into the medical record. Therefore, the role of public health is different than that for newborn bloodspot screening (7). This role might present challenges in data collection and surveillance for evaluating CCHD screening, because uniform reporting systems might not be established between public health programs, birthing centers, and hospitals (8). States have previously reported barriers to involvement with CCHD screening, such as the lack of legislative authority, staffing, funding, and informatics infrastructure (9). This report represents the first assessment of state legislative activities, requirements for collection of screening data, and progress made with screening activities, despite previously reported barriers.

**FIGURE 2. Actions taken by states to adopt newborn screening for critical congenital heart defects — United States, 2011–2014\***



\* Actions taken as of December 2014.

State-level data collection is vital for surveillance, monitoring of outcomes, and evaluation of state CCHD newborn screening programs. Although all types of screening data can be valuable, individual-level data are important for surveillance and evaluation. Collecting data related to factors associated with false-positive and false-negative results could help refine the recommended CCHD screening algorithm and screening activities (7). As states evaluate the implementation of CCHD screening, they are encouraged to consider programmatic changes that would improve their screening program, such as the inclusion of individual-level data reporting.

Enactment of a state law or regulation does not translate into immediate and universal change in clinical practice. In addition to policy changes, the proper public health infrastructure, including infrastructure needs for data collection and reporting of CCHD screening results, is vital to ensure a successful CCHD newborn screening program.

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## Summary

### What is already known on this topic?

Congenital heart defects occur in approximately eight in every 1,000 live births, one fourth of which are considered to be critical congenital heart defects (CCHD). Newborn screening using pulse oximetry can detect hypoxemia, a clinical sign of CCHD.

### What is added by this report?

This report represents the first assessment of state's actions to adopt newborn screening for CCHD and requirements for collection of CCHD screening data. Forty-three states have taken action toward newborn screening for CCHD through statute, regulations, or hospital guidelines. Among the 43 states, 32 (74%) are collecting or planning to collect CCHD screening data.

### What are the implications for public health practice?

State mandates for newborn screening for CCHD might increase the number of newborns screened, allowing for early identification and prevention of morbidity and mortality. Data collection and reporting are essential to evaluate the effect of this public health program.

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## Opioid Overdose Prevention Programs Providing Naloxone to Laypersons — United States, 2014

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Drug overdose deaths in the United States have more than doubled since 1999 (1). During 2013, 43,982 drug overdose deaths (unintentional, intentional [suicide or homicide], or undetermined intent) were reported (1). Among these, 16,235 (37%) were associated with prescription opioid analgesics (e.g., oxycodone and hydrocodone) and 8,257 (19%) with heroin (2). For many years, community-based programs have offered opioid overdose prevention services to laypersons who might witness an overdose, including persons who use drugs, their families and friends, and service providers. Since 1996, an increasing number of programs provide laypersons with training and kits containing the opioid antagonist naloxone hydrochloride (naloxone) to reverse the potentially fatal respiratory depression caused by heroin and other opioids (3). In July 2014, the Harm Reduction Coalition (HRC), a national advocacy and capacity-building organization, surveyed 140 managers of organizations in the United States known to provide naloxone kits to laypersons. Managers at 136 organizations completed the survey, reporting on the amount of naloxone distributed, overdose reversals by bystanders, and other program data for 644 sites that were providing naloxone kits to laypersons as of June 2014. From 1996 through June 2014, surveyed organizations provided naloxone kits to 152,283 laypersons and received reports of 26,463 overdose reversals. Providing opioid overdose training and naloxone kits to laypersons who might witness an opioid overdose can help reduce opioid overdose mortality.

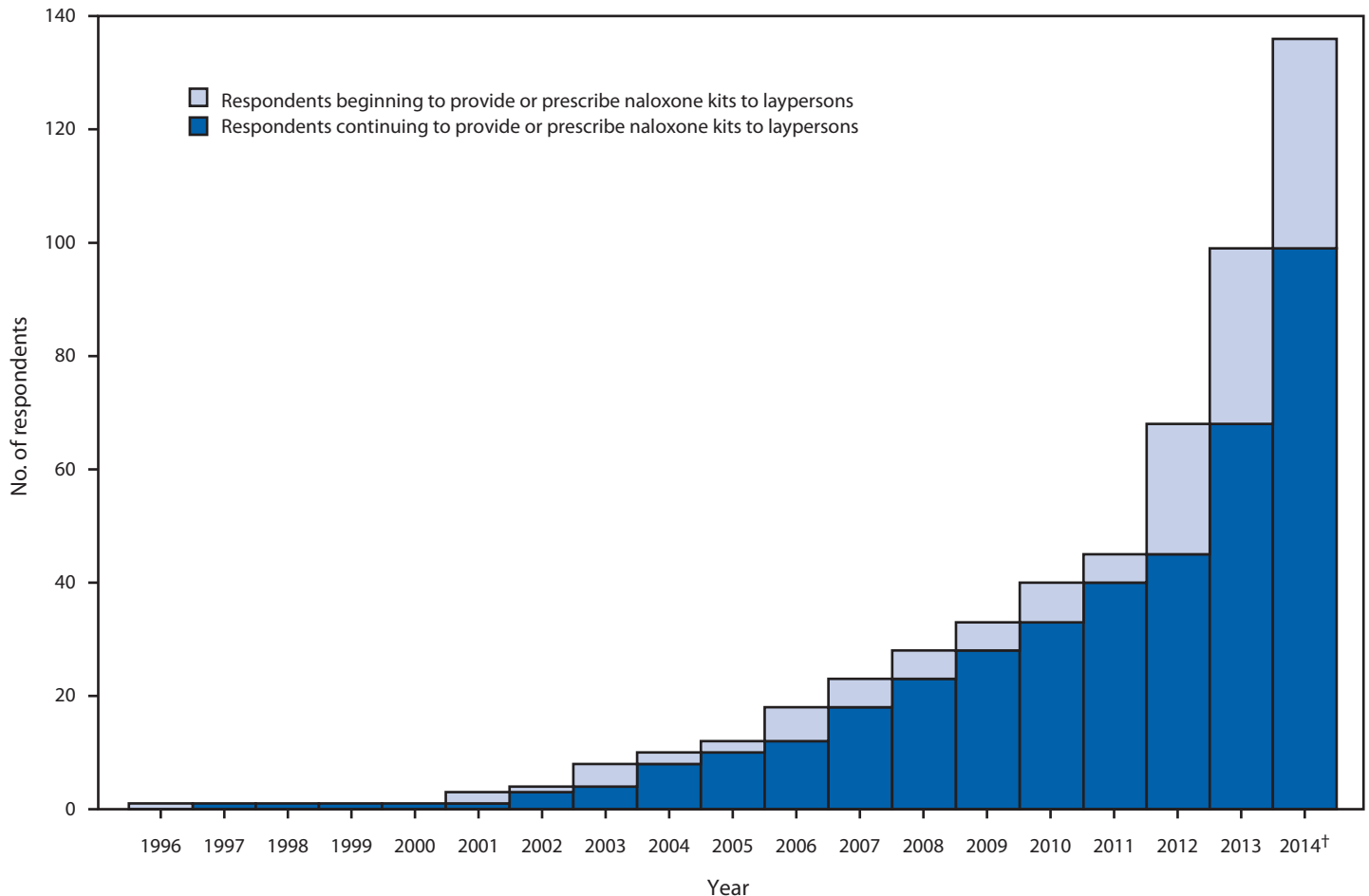
Since 2008, HRC has maintained a database of organizations providing naloxone kits to laypersons. The Opioid Safety and Naloxone Network is a national network of naloxone experts, program administrators, and advocates. Before the survey, HRC staff polled network participants for information on any new organizations providing naloxone kits to laypersons that should be included in the survey. In July 2014, HRC e-mailed a link to an online survey to managers of 140 organizations known to provide naloxone kits to laypersons. These organizations included public health departments, pharmacies, health care facilities, substance use treatment facilities, and community-based organizations providing services to persons who use drugs, including current or former opioid (heroin or pharmaceutical) users, and other potential witnesses to overdoses. Law enforcement organizations, emergency medical

services, and other professional first responders using naloxone were not included in this survey.

The survey included questions about the year the organization began operating; the numbers of sites or local programs providing naloxone kits; the number of persons trained in overdose prevention and provided naloxone kits; and the number of reports of overdose reversals (administration of naloxone by a trained layperson in the event of an overdose) (4), as well as whether the reports were based on program data or were estimates. The survey also asked about the naloxone formulations currently provided in kits, models for training and providing naloxone kits, funding sources, and any difficulties obtaining naloxone. To obtain data for a recent full calendar year, organizations providing naloxone kits during calendar year 2013 were asked to provide specific data for that year, including numbers of persons provided naloxone kits, reversals reported, and naloxone vials provided; characteristics of persons who received naloxone kits (e.g., persons who use drugs, friends and family members, service providers); characteristics of persons reporting overdose reversals; and the drugs involved in reported overdose reversals. HRC staff used follow-up e-mails and telephone calls to encourage participation and clarify responses.

Managers from 136 (97.1%) organizations completed the survey, including those from 84 community-based organizations, 18 health care facilities, 10 Veterans Administration health care systems, 18 state or local health departments, and six pharmacies. Half of the responding organizations began operating during January 2013–June 2014 (Figure 1). Respondents provided reports for 644 local opioid overdose prevention sites that provide naloxone kits, located in 30 states and the District of Columbia (DC) (Figure 2). Thirty-eight respondents provided consolidated data for multiple local sites providing naloxone kits. Some organizations estimated responses; for example, one health department estimated the number of laypersons receiving naloxone kits on the basis of the number of kits distributed to local sites. Three state health departments (Massachusetts, New Mexico, and New York) oversee operations of statewide naloxone programs, with 334 local sites (51.9% of the 644 local sites).

From 1996, when the first organization began providing naloxone, through June 2014, the 136 responding organizations reported providing training and naloxone kits to 152,283

**FIGURE 1. Number of survey respondents reporting beginning or continuing to provide naloxone kits to laypersons, by year — United States, 1996–June 2014\*†**

\* Results of a survey conducted in July 2014 by the Harm Reduction Coalition, in which 136 organizations reported 644 local sites where laypersons were trained to recognize an opioid drug overdose and provided or prescribed naloxone kits.

† As of June 2014.

laypersons (range = 1–36,450; median = 100; mean = 1,120).<sup>\*</sup> The 109 organizations that collect reports of reversals documented 26,463 overdose reversals (range = 0–5,430; median = 9; mean = 243).<sup>†</sup>

During 2013, 93 organizations reported distributing or prescribing naloxone to 37,920 laypersons (range = 0–9,000; median = 75; mean = 407.7).<sup>§</sup> The 68 (50%) organizations that collect reports of reversals documented 8,032 overdose reversals (range = 0–2,079; median = 10; mean = 118.1).<sup>¶</sup>

<sup>\*</sup> Estimated by 57 respondents (55,201 [36.2%] laypersons) and based on program data for 79 (97,082 [36.8%]).

<sup>†</sup> Estimated by 28 respondents (5,245 [19.8%] reversals) and based on program data for 81 (21,218 [80.2%]).

<sup>§</sup> Estimated by 36 respondents (6,483 [17.1%] laypersons) and based on program data for 57 (31,437 [82.9%]).

<sup>¶</sup> Estimated by 13 respondents (659 [8.2%] reversals) and based on program data for 55 (7,373 [91.8%]).

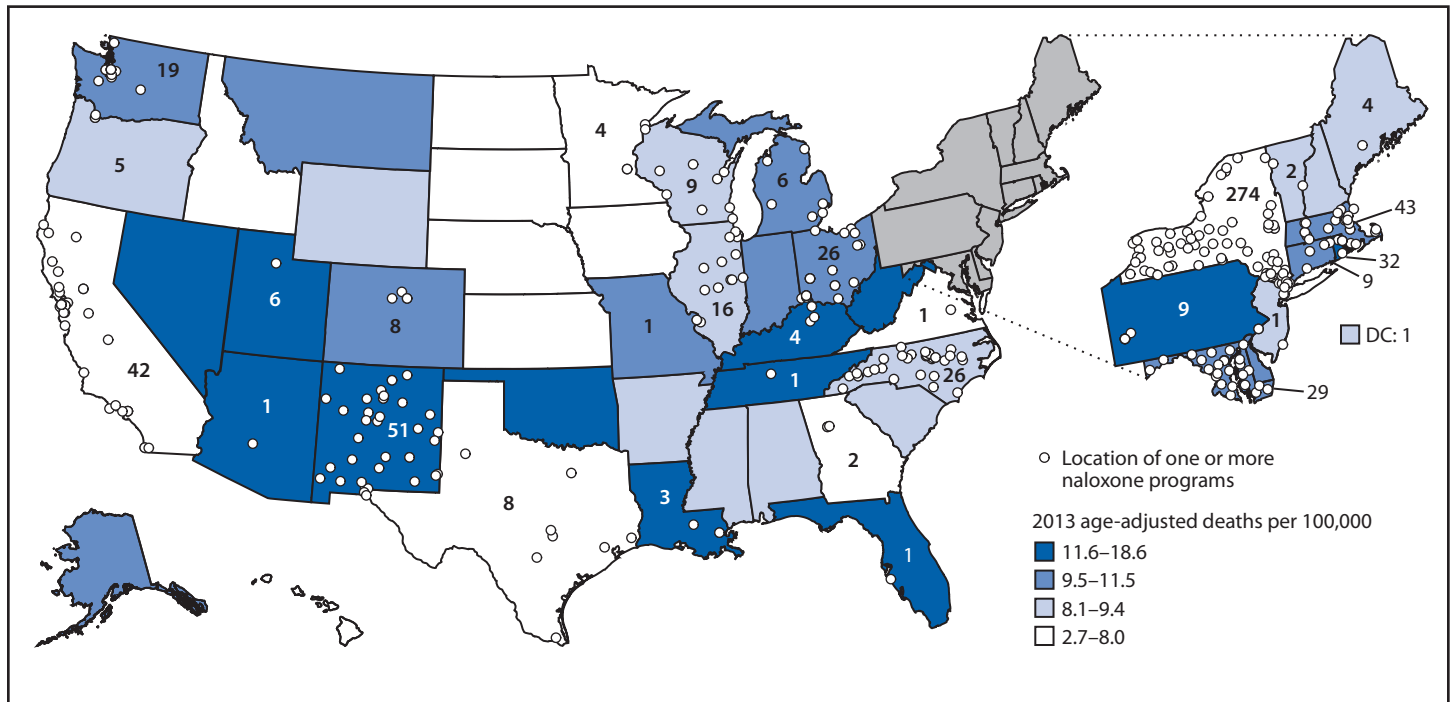
Ninety-three organizations collected information on the characteristics of laypersons who were provided naloxone kits. Laypersons who received naloxone kits were characterized as persons who use drugs (81.6%); friends and family members (11.7%); service providers (3.3%); or unknown (3.4%).<sup>\*\*</sup> Sixty-eight organizations provided information about laypersons who reported administering naloxone, characterizing them as persons who use drugs (82.8%); friends and family members (9.6%); service providers (0.2%); or unknown (7.4%).<sup>††</sup> Forty-two organizations collected information from laypersons about the drugs that appeared to be involved

<sup>\*\*</sup> Estimated by 48 (51.6%) respondents and based on program data for 45 (48.4%).

<sup>††</sup> Estimated by 26 (38.2%) respondents and based on program data for 42 (61.8%).



**FIGURE 2. Number\* and location of local drug overdose prevention programs providing naloxone to laypersons, as of June 2014, and age-adjusted rates† of drug overdose deaths§ in 2013 — United States**



\* Total N = 644; numbers on map indicate the total number of programs within each state.

† Per 100,000 population.

§ CDC, National Center for Health Statistics; Compressed Mortality File 1999–2013 on CDC WONDER Online Database, released January 2015.

in the reversed overdoses; heroin was involved in 81.6% and prescription opioids in 14.1%.<sup>§§</sup>

Various program models were used by organizations to provide naloxone to laypersons, including distribution of naloxone kits by trained nonmedical staff or volunteers under a standing order (60 [44.1%]), by medical staff (49 [36.0%]), prescriptions written by a medical provider and filled at a pharmacy (39 [28.7%]), pharmacists dispensing directly via collaborative practice agreements and other mechanisms (12 [8.8%]), and other protocols (19 [14.0%]). Thirty-three organizations used more than one model.

During 2013, 90 (66.2%) of the 136 organizations reported distributing 140,053 naloxone vials, including refills (range = 1–53,200; median = 179.5; mean = 1,556.1).<sup>¶¶</sup> Three respondents whose organizations were operational in 2013 did not report on the number of vials because they furnished prescriptions to be filled at a pharmacy. The remaining 43 organizations indicated that they were not yet providing naloxone kits during 2013. Sixty-nine respondents (50.7%)

reported their organization provided only injectable naloxone, 51 (37.5%) provided only intranasal naloxone, and 16 (11.8%) provided both injectable and intranasal naloxone.<sup>\*\*\*</sup> A total of 111,602 vials (79.7%) of injectable naloxone (21.4% 10 mL and 58.1% 1 mL) and 28,446 (20.3%) vials of intranasal naloxone were provided to laypersons. Organizations were characterized as small, medium, large, or very large, on the basis of the number of naloxone vials distributed during 2013. The 11 large and very large organizations provided naloxone to 28,604 laypersons, representing 75.4% of all 2013 recipients (Table). Forty (29.4%) organizations reported difficulties maintaining an adequate supply of naloxone, and 73 (53.7%) reported inadequate resources to sustain or expand their organization's efforts to disseminate naloxone kits.

## Discussion

Organizations that provide naloxone kits to laypersons have expanded substantially since a similar survey in 2010 (5), reflecting a 183% (from 48 to 136) increase in the number of

<sup>§§</sup> Estimated by 18 (42.9%) respondents and based on program data for 24 (57.1%).

<sup>¶¶</sup> Estimated by 37 survey respondents (31,838 [22.7%] vials) and based on program data for 53 (108,215 [77.2%]).

<sup>\*\*\*</sup> Organizations provide laypersons with naloxone for injection and/or for intranasal administration. Injectable naloxone is distributed in multi-dose (10 mL) and single-dose (1 mL) vials with concentrations of 0.4 mg/mL. Intranasal naloxone is distributed in single-dose 2 mL vials with concentration of 1 mg/mL that are adapted for intranasal use with a mucosal atomizer.

**TABLE. Reported number of laypersons receiving or prescribed naloxone kits, overdose reversals, and opioid overdose prevention programs, by survey respondent program size — United States, 1996–June 2014**

Category (by size)*	Respondents		Sites		Calendar year 2013				1996–June 2014			
					Laypersons received/ prescribed kits <sup>†</sup>		Opioid overdose reversals <sup>§</sup>		Laypersons received/ prescribed kits <sup>¶</sup>		Opioid overdose reversals <sup>**</sup>	
	No.	(%)	No.	(%)	No.	(%)	No.	(%)	No.	(%)	No.	(%)
Small (<100)	84	(61.8)	154	(23.9)	1,709	(4.5)	134	(1.7)	7,867	(5.2)	641	(2.4)
Medium (101–1,000)	41	(30.1)	129	(20.0)	7,607	(20.1)	1,351	(16.8)	19,239	(12.6)	4,414	(16.7)
Large (1,001–10,000)	7	(5.1)	62	(9.6)	6,117	(16.1)	4,329	(53.9)	29,099	(19.1)	11,807	(44.6)
Very large (>10,000)	4	(2.9)	299	(46.4)	22,487	(59.3)	2,218	(27.6)	96,078	(63.1)	9,601	(36.3)
<b>Total</b>	<b>136</b>	<b>(100.0)</b>	<b>644</b>	<b>(100.0)</b>	<b>37,920</b>	<b>(100.0)</b>	<b>8,032</b>	<b>(100.0)</b>	<b>152,283</b>	<b>(100.0)</b>	<b>26,463</b>	<b>(100.0)</b>

\* Based on reported number of vials of naloxone provided during 2013.

<sup>†</sup> Calendar year 2013 information provided by 93 survey respondents distributing kits/prescribing naloxone during that year, with 36 estimating (6,483 [17.1%] persons) and 57 based on program data (31,437 [82.9%]).

<sup>§</sup> Sixty-eight of 93 respondents distributing kits/prescribing naloxone in 2013 provided information on reported reversals, with 13 estimating (659 [8.2%] reversals) and 55 based on program data (7,373 [91.8%]).

<sup>¶</sup> Estimated by 57 survey respondents (55,201 [36.2%] persons) and 79 based on program data (97,082 [63.8%]).

<sup>\*\*</sup> Program began in 1996; as of June 2014, 109 respondents distributing kits/prescribing naloxone provided information on reported reversals, with 28 estimating (5,245 [19.8%] reversals) and 81 based on program data (21,218 [80.2%]).

responding organizations; a 243% (from 188 to 644) increase in the number of local sites providing naloxone; a 187% (from 53,032 to 152,283) increase in the number of laypersons provided naloxone kits; a 160% (from 10,171 to 26,463) increase in the number of reversals reported; and a 94% (from 16 to 30) increase in states (including DC) with at least one organization providing naloxone. Half of the responding organizations began operating during January 2013–June 2014. Although early adopters of naloxone kit provision were mainly syringe exchanges, other programs, including substance use treatment facilities, Veterans Administration health care systems, primary care clinics, and pharmacies have started providing naloxone to laypersons.

Providing naloxone kits to laypersons reduces overdose deaths (4), is safe (3), and is cost-effective (6). U.S. and international health organizations recommend providing naloxone kits to laypersons who might witness an opioid overdose (3,7); to patients in substance use treatment programs (3,7,8); to persons leaving prison and jail (3,7,8); and as a component of responsible opioid prescribing (8).

Although the number of organizations providing naloxone kits to laypersons is increasing, in 2013, 20 states had no such organization, and nine had less than one layperson per 100,000 population who had received a naloxone kit. Among these 29 states with minimal or no access to naloxone kits for laypersons, 11 had age-adjusted 2013 drug overdose death rates higher than the national median (2).

Some organizations reported information on the laypersons receiving naloxone kits (N = 99 organizations), using naloxone in overdose reversals (N = 68), and the drugs that appeared to have caused the overdose (N = 42). Persons who use drugs accounted for 81.6% of laypersons who received naloxone kits; they also performed the majority (82.8%) of reported

overdose reversals. A majority (81.6%) of the overdoses that were reversed involved heroin, indicating that organizations are reaching laypersons who witness heroin overdoses. A study of a community-based naloxone program in San Francisco also found that persons who use drugs play a major role in reversing heroin overdoses (9). Additional interventions are needed to reach persons who may witness prescription opioid analgesic overdoses, which account for nearly twice as many deaths as heroin overdoses.

Forty (29.4%) respondents reported that their organization has experienced problems obtaining naloxone. Prices of intranasal naloxone more than doubled in the second half of 2014 (10) and Opioid Safety and Naloxone Network members report that cost increases are reducing the quantity of naloxone purchased and provided to laypersons (Matt Curtis, VOCAL NY, personal communication, 2015).

The findings in this report are subject to at least four limitations. First, despite extensive knowledge of naloxone distribution programs by the Harm Reduction Coalition and Opioid Safety and Naloxone Network, organizations providing naloxone kits are increasing rapidly and some might not yet be known to HRC and therefore, might not be included in the survey, which may underestimate the impact of these programs. Second, survey responses are based on unconfirmed reports from organizations providing naloxone kits. Third, some reports provided by organizations are based on estimates. These three limitations could result in either under or over-reporting of persons provided naloxone kits. Finally, the numbers of overdose reversals likely were under-reported, because some sites, such as pharmacies, do not collect reversal reports.

Organizations providing naloxone kits to laypersons receive many reports of overdose reversals and can reach large numbers of potential overdose bystanders. Comprehensive prevention

## Summary

### What is already known on this topic?

Drug overdose deaths in the United States have more than doubled since 1999, reaching a total of 43,982 in 2013. Heroin and prescription opioids are major causes of drug overdose deaths. Naloxone is the standard medication used for reversal of the potentially fatal respiratory depression caused by opioid overdose.

### What is added by this report?

From 1996 through June 2014, a total of 644 local sites in 30 states and the District of Columbia reported providing naloxone kits to 152,283 laypersons and receiving reports of 26,463 drug overdose reversals using naloxone from 1996 through June 2014. Most laypersons who reported using the kits to reverse an overdose were persons who use drugs, and many of the reported reversals involved heroin overdoses. Medical clinics and pharmacies have started providing naloxone kits to laypersons, and the reported number of organizations providing kits almost doubled from January 2013 through June 2014.

### What are the implications for public health practice?

Organizations training and providing naloxone kits to laypersons can reach large numbers of potential overdose witnesses and result in many reported overdose reversals. Comprehensive prevention measures that include teaching laypersons how to respond to overdoses and administer naloxone prevent opioid-related drug overdose deaths. Additional methods are needed to provide naloxone kits to persons who might witness prescription opioid analgesic overdoses.

measures that include teaching laypersons how to respond to overdoses and administer naloxone might help prevent opioid drug overdose deaths. This report suggests that many programs reach persons who witness heroin-related overdoses; additional methods are needed to provide naloxone kits to persons who might witness prescription opioid analgesic overdoses.

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## Coccidioidomycosis in a State Where It Is Not Known To Be Endemic — Missouri, 2004–2013

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During 1998–2012, coccidioidomycosis cases increased nationally nearly eightfold (1). To describe the epidemiology of coccidioidomycosis in Missouri, a state without endemic coccidioidomycosis, coccidioidomycosis surveillance data during 2004–2013 at the Missouri Department of Health and Senior Services were retrospectively reviewed. The incidence of reported coccidioidomycosis increased from 0.05 per 100,000 population in 2004 to 0.28 per 100,000 in 2013, with cases distributed throughout all regions of Missouri. Persons aged >60 years were most affected. In cases in which patients had disease manifestations, the most common were pneumonia (37%) and influenza-like illness (31%). Nearly half (48%) of patients had traveled to an area where coccidioidomycosis is endemic, whereas approximately one-quarter (26%) of patients did not report such travel. Those with history of travel to endemic areas were significantly more likely to receive a diagnosis by positive culture or polymerase chain reaction (PCR) testing, compared with those without a history of travel to endemic areas, who were more likely to receive a diagnosis by serological tests. Additional studies will be required to ascertain whether truly endemic cases exist in Missouri.

Coccidioidomycosis, or Valley Fever, is a systemic disease caused by the fungus *Coccidioides*, which is endemic to the southwestern United States, Mexico, and Central and South America (2). This fungus normally resides in soil, but airborne spores can cause infection if inhaled. Sixty percent of infections are asymptomatic and do not come to medical attention (3). Manifestations range from influenza-like illness to pneumonia, lung nodules, and disseminated infections. Laboratory tests typically used for coccidioidomycosis diagnosis include complement fixation, immunodiffusion, enzyme immunoassay, culture, histopathology, and PCR. In most patients, infection is mild and resolves without specific antifungal treatment. Azole antifungals are the most commonly used drugs in patients who require treatment (4).

During 1998–2011, among all coccidioidomycosis cases reported to CDC from 28 states and the District of Columbia, 97% were from Arizona and California, 1% from states where coccidioidomycosis is uncommon, and <1% from states where it is not endemic (4). In areas where coccidioidomycosis is endemic (excluding Texas), the age-adjusted incidence of reported cases increased sevenfold from 5.3 cases per 100,000 population in 1998 to 42.6 per 100,000 in 2011 (5). In states

where it is not endemic, in 2011, 240 coccidioidomycosis cases were reported, compared with only six in 1998 (1).

Coccidioidomycosis surveillance data during 2004–2013 at the Missouri Department of Health and Senior Services were retrospectively reviewed. Only cases meeting the definition of a confirmed case of coccidioidomycosis as defined by CDC's National Notifiable Diseases Surveillance System and the Council of State and Territorial Epidemiologists were included (6).

Patients with known travel history were categorized into three groups: 1) those who had not traveled to an area where coccidioidomycosis is known to be endemic, 2) those who had traveled at any time to an area where it is endemic, and 3) those who had recently traveled to an area where the disease is endemic. Recent travel was defined as travel associated with the experience of symptoms either during travel or within 21 days of leaving the endemic area. Poisson regression analysis was used to model the incidence of coccidioidomycosis reported in Missouri.

A total of 93 confirmed coccidioidomycosis cases were reported during the study period (Table). Disease incidence increased from 0.05 per 100,000 population in 2004 to 0.28 per 100,000 in 2013 ( $p < 0.001$ ) (Figure 1). The median age of patients was 58 years (range = 19–94 years). Among 51 (55%) patients with a known symptom onset date, median time to diagnosis was 25 days (range = 3–304). Fungal culture (31%) and complement fixation (30%) were the most common diagnostic tests. Forty-three (46%) patients required hospitalization (five in intensive care). Among 29 patients who received antifungal drugs, 14 were treated as outpatients and 15 were inpatients. Fluconazole was the most used antifungal drug (20% of patients). Eight (8.6%) of the 93 patients died: three deaths were attributed to coccidioidomycosis, three to other illnesses, and the cause of death in the remaining two patients was not reported.

Mapping of cases by residence at the time of diagnosis revealed that patients, with or without travel to an area where coccidioidomycosis is endemic, were distributed throughout all regions of the state (Figure 2). Forty-five patients (48%) traveled to an area where the disease is endemic, 24 (26%) did not, and the travel history for the remaining 24 (26%) was unknown. Among the 45 patients with travel to an area with endemic disease, 19 had recent travel, 20 had travel that was



not recent or had a history of residence in an area with the disease, and six had travel timelines that could not be exactly established. The proportion of patients receiving a diagnosis by positive coccidioidomycosis culture or PCR was significantly higher in patients who had traveled (21 of 45) compared with those who had not (4 of 24) ( $p = 0.018$ ). Overall, among 24 patients without a history of travel, 11 received a diagnosis only on the basis of positive immunoglobulin M or qualitative enzyme immunoassay or immunodiffusion tests, and four received a diagnosis on the basis of positive coccidioidomycosis cultures (all of the latter were immunocompetent). Seventeen (18%) patients were immunocompromised, eight of whom had a history of travel to an area where coccidioidomycosis is endemic.

### Discussion

The increase in the incidence of reported coccidioidomycosis in Missouri from 2004 through 2013 was statistically significant and substantial. The increase is consistent with the national trend of increasing incidence of coccidioidomycosis that includes states with and without endemic disease (1). Mapping of cases with and without a history of travel to areas with endemic disease revealed that cases were occurring in all regions of Missouri.

One explanation for the increase in reported cases could be that coccidioidomycosis became a reportable condition in Missouri in 2003. In comparison, after coccidioidomycosis became reportable in Arizona in 1997, the reported incidence increased from 21 per 100,000 in 1997 to 91 per 100,000 population in 2006 (7). An additional contributing factor could be an increased awareness among health care providers and the public of coccidioidomycosis, leading to more testing, as well as better availability of diagnostic tests offered by commercial laboratories.

A false-positive immunoglobulin M test result might lead to incorrect diagnosis of coccidioidomycosis if diagnosis is confirmed solely by this serological test (8). The positive predictive value of the enzyme immunoassay for coccidioidomycosis has been shown to vary depending on the circumstances under which the assay was used (9). In this study, those with history of travel to areas where coccidioidomycosis is endemic were significantly more likely to receive a diagnosis on the basis of positive culture, PCR testing, or both, compared with those without such travel, who were more likely to receive a diagnosis with serological tests. Because culture and PCR are more accurate tests for diagnosing recent coccidioidomycosis compared with the serological tests, whether all patients with no travel history were experiencing current infection is unknown. The coccidioidomycosis surveillance case definition makes no distinction between those with travel or residence in an area

**TABLE. Demographic and clinical characteristics of coccidioidomycosis cases — Missouri, 2004–2013**

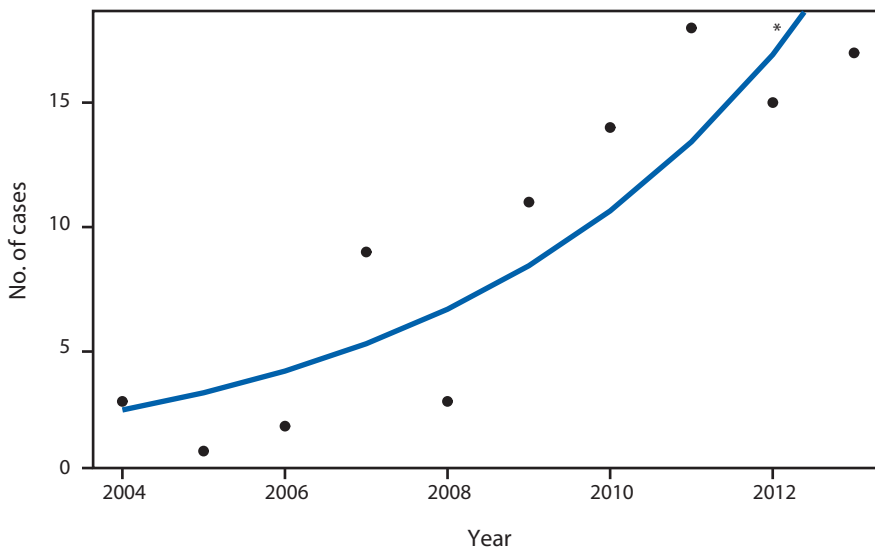
Characteristic	No.	(%)
<b>Total</b>	<b>93</b>	<b>(100)</b>
<b>Sex</b>		
Male	67	(72)
Female	26	(28)
<b>Age (yrs)</b>		
≥70	22	(24)
60–69	21	(23)
50–59	16	(17)
40–49	15	(16)
30–39	12	(13)
20–29	6	(6)
10–19	1	(1)
<b>Race</b>		
White	50	(54)
Black	7	(8)
Pacific Islander	1	(1)
Asian	1	(1)
Unknown	34	(37)
<b>Manifestations</b>		
Symptomatic lung lesions/Pneumonia	37	(40)
Flu-like illness	31	(33)
Hemoptysis	5	(5)
Headache/Confusion	3	(3)
Skin lesions	3	(3)
Sepsis/Disseminated	2	(2)
Asymptomatic lung lesions	2	(2)
Arthritis/Arthralgia	2	(2)
Meningitis	1	(1)
Unknown	7	(8)
<b>Laboratory tests</b>		
Culture	29	(31)
CF	28	(30)
Immunodiffusion	23	(25)
EIA/ELISA	13	(14)
PCR	4	(4)
Histopathology	2	(2)
Unknown serology	9	(10)

**Abbreviations:** CF = complement fixation; EIS/ELISA = enzyme immune assay/enzyme linked immunosorbent assay; PCR = polymerase chain reaction.

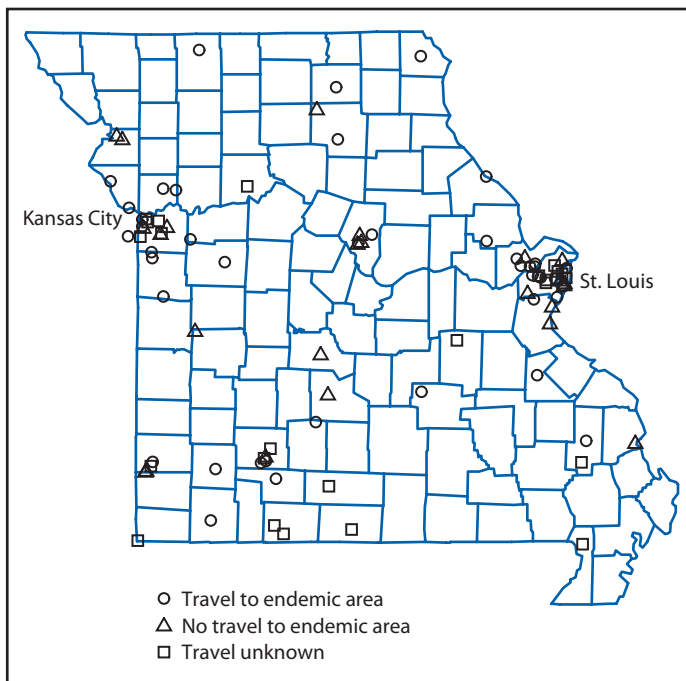
where coccidioidomycosis is known to be endemic and those without such a history, even though the history affects the positive predictive value of current diagnostic tests. Because persons living in areas without endemic disease have a much lower risk for having coccidioidomycosis, more stringent requirements for laboratory diagnosis of these cases might be prudent.

Four patients in the group that did not report travel received a diagnosis on the basis of positive coccidioidomycosis culture, raising the possibility that the disease was locally acquired. Soil analysis for *Coccidioides* spores in the area where those patients resided could have been helpful for clarification of whether the cases were truly locally acquired. Recently, *Coccidioides* was found in soil in south-central Washington, a state where coccidioidomycosis was not believed to be endemic; three acute coccidioidomycosis cases have been reported in Washington (10). No reports documenting the presence of *Coccidioides*



**FIGURE 1. Incidence of coccidioidomycosis, by year — Missouri, 2004–2013**

\* Line represents estimated Poisson Regression model  $\hat{y} = e^{-463.29 + 0.23 \times \text{year}}$ ;  $p < 0.001$ .

**FIGURE 2. Coccidioidomycosis cases, by location and travel status — Missouri, 2004–2013**

spores in Missouri soil have been published in the indexed literature. Cluster analyses of a larger sample of coccidioidomycosis cases using software that analyzes spatial, temporal, and space-time data using spatial, temporal, or space-time scan statistics might be helpful for more accurate estimation of the possibility of endemic cases in Missouri.

The findings in this report are subject to one main limitation. The retrospective analysis was conducted on routine public health surveillance data, and no medical chart review or direct patient interviews were conducted. The surveillance data were not sufficiently complete in some cases with respect to demographics, travel history, medical history, clinical symptoms, diagnosis, treatment, and follow-up. In some cases, the exact diagnostic tests used for serology (e.g., immunoglobulin G or immunoglobulin M) or the exact titer for those tested by complement fixation were not known.

Follow-up of patients with coccidioidomycosis to ensure that no alternative diagnoses emerged often was not available.

Epidemiology of coccidioidomycosis has been well described in states where it is known to be endemic, such as Arizona and California, but little information exists about it in other states. This research is a first attempt to study the epidemiology of coccidioidomycosis in a state without known endemic disease. Sustained surveillance for coccidioidomycosis in non-endemic states is important to ascertain whether locally acquired cases are occurring.

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## References

## Summary

## What is already known on this topic?

The incidence of reported coccidioidomycosis is increasing nationally, both in states where the disease is known to be endemic and those where it is not.

## What is added by this report?

This is the first study of the epidemiology of coccidioidomycosis in a state without endemic disease. In Missouri, during 2004–2013, reported coccidioidomycosis incidence per 100,000 population significantly increased from 0.05 to 0.28. Nearly half of the patients with known travel history had visited areas where coccidioidomycosis is endemic, and were more likely to receive a diagnosis of the disease by fungal culture and polymerase chain reaction, rather than serological assays.

## What are the implications for public health practice?

Surveillance for coccidioidomycosis is needed in non-endemic states to discover if locally acquired cases are occurring. For persons living in areas where coccidioidomycosis is not believed to be endemic, more stringent requirements for laboratory diagnosis of coccidioidomycosis might be appropriate.

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## Update on Vaccine-Derived Polioviruses — Worldwide, January 2014–March 2015

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Since the World Health Assembly's 1988 resolution to eradicate poliomyelitis (1), one of the main tools of the World Health Organization (WHO) Global Polio Eradication Initiative (GPEI) has been the live, attenuated oral poliovirus vaccine (OPV) (2). OPV might require several doses to induce immunity but provides long-term protection against paralytic disease. Through effective use of OPV, GPEI has brought polio to the threshold of eradication. Wild poliovirus type 2 (WPV2) was eliminated in 1999, WPV3 has not been detected since November 2012, and WPV1 circulation appears to be restricted to parts of Pakistan and Afghanistan (1). However, continued use of OPV carries two key risks. The first, vaccine-associated paralytic poliomyelitis (VAPP) has been recognized since the early 1960s (2,3). VAPP is a very rare event that occurs sporadically when an administered dose of OPV reverts to neurovirulence and causes paralysis in the vaccine recipient or a nonimmune contact. VAPP can occur among immunologically normal vaccine recipients and their contacts as well as among persons who have primary immunodeficiencies (PIDs) manifested by defects in antibody production; it is not associated with outbreaks. The second, the emergence of genetically divergent, neurovirulent vaccine-derived polioviruses (VDPVs) was recognized more recently (4). Circulating VDPVs (cVDPVs) resemble WPVs and, in areas with low OPV coverage, can cause polio outbreaks. Immunodeficiency-associated VDPVs (iVDPVs) can replicate and be excreted for years in some persons with PIDs; GPEI maintains a registry of iVDPV cases. Ambiguous VDPVs (aVDPVs) are isolates that cannot be classified definitively (4,5). This report updates previous surveillance summaries (5) and describes VDPVs detected worldwide during January 2014–March 2015. Those include new cVDPV outbreaks in Madagascar and South Sudan, and sharply reduced type 2 cVDPV (cVDPV2) circulation in Nigeria and Pakistan during the latter half of 2014. Eight newly identified persons in six countries were found to excrete iVDPVs, and a patient in the United Kingdom was still excreting iVDPV2 in 2014 after more than 28 years. Ambiguous VDPVs were found among immunocompetent persons and environmental samples in 16 countries. Because the large majority of VDPV case-isolates are type 2, WHO has developed a plan for coordinated worldwide withdrawal of trivalent (types 1, 2, and 3) OPV (tOPV) and replacement with bivalent (types 1 and 3) OPV (bOPV)

in April 2016, preceded by introduction of at least 1 dose of injectable inactivated poliovirus vaccine (IPV) into routine immunization schedules worldwide to maintain immunity to type 2 viruses (6).

### Properties of VDPVs

VDPVs are polioviruses whose genetic divergence from the parental OPV strains indicates prolonged replication or circulation (4,5). Poliovirus isolates are grouped into three categories: 1) WPVs; 2) vaccine-related polioviruses (VRPVs); and 3) VDPVs. Current WPVs are genetically unrelated to any vaccine strain. The demarcation between VRPVs and VDPVs is based on the known poliovirus evolution rate. Nucleotide substitutions accumulate in poliovirus genomes at an overall rate of approximately 1% per year and are routinely monitored by sequencing the ~900-nucleotide region encoding VP1, the major poliovirus surface protein. Although nucleotide substitutions might accumulate more rapidly in the early phases of OPV replication, fewer than five VP1 substitutions typically accumulate in the vaccine virus during the normal period of replication in an immunocompetent OPV recipient (4–6 weeks). Based on this rate of nucleotide substitution, type 1 and type 3 isolates that are <1.0% divergent and type 2 isolates that are <0.6% divergent in VP1 sequences from the corresponding vaccine strain are classified as VRPVs. Type 1 and type 3 isolates that are >1.0% divergent or type 2 isolates that are >0.6% divergent in VP1 sequences from the corresponding OPV strain are classified as VDPVs (5). VDPVs are further categorized as 1) cVDPVs when evidence of person-to-person transmission in the community exists; 2) iVDPVs, which are isolated from persons with PIDs; and 3) aVDPVs, which are either clinical isolates from persons with no known immunodeficiency and no evidence of transmission, or sewage isolates that are unrelated to known cVDPVs or iVDPVs and whose source is unknown (5).

### Virologic Testing for VDPVs

All poliovirus isolates are characterized by laboratories of the Global Polio Laboratory Network (5) using a real-time reverse transcription–polymerase chain reaction (rRT-PCR) nucleic acid amplification, targeted to nucleotide substitutions that typically revert to the parental WPV sequence during replication of OPV in the human intestine (7). The rRT-PCR

methods are used in 88 of 146 Global Polio Laboratory Network laboratories (5). Candidate VDPVs identified by rRT-PCR screening are sequenced in the VP1 region for definitive analysis; the complete genome is sequenced if required for higher-resolution analysis.

### cVDPVs

The number of countries with circulation of indigenously emergent cVDPVs decreased from seven during July 2012–December 2013 (5) to four (Pakistan, Nigeria, Madagascar, and South Sudan) during January 2014–March 2015. Outbreaks associated with indigenous cVDPV2 (Afghanistan, Chad, China, Somalia, and Yemen) and with imported cVDPV2 (Cameroon, Kenya, and Niger) (5) have been interrupted. Although the cVDPV2 outbreak in Pakistan has continued (5), the large outbreak in Nigeria has nearly stopped (5,8), and the two new outbreaks in Madagascar (cVDPV1) and South Sudan (cVDPV2) are small (Table, Figure 1). The most prevalent cVDPVs are type 2 (88.2%), followed by type 1 (10.3%) and type 3 (1.6%). Among the 686 cVDPV cases reported since 2006, >97% were associated with cVDPV2 (Figure 2).

**Madagascar.** One cVDPV1 was isolated from an acute flaccid paralysis (AFP) patient in Analalava, Mahjanga Province, on the northwest coast. Circulation is suspected because of the extent of VP1 nucleotide sequence divergence (2.2%) from the parental OPV strain, the absence of immunodeficiency in the AFP patient, and the infection of two nonhousehold contacts with closely related cVDPV1 viruses, as well as the history of repeated cVDPV emergence in Madagascar (5).

**Nigeria.** The large indigenous cVDPV2 outbreaks in northern Nigeria, associated with >20 independent cVDPV2 emergences, peaked in 2009 (8), but low-level circulation continued (5). Virus from the major cVDPV2 lineage group that first emerged in 2005 (8) was isolated from 11 AFP patients (most recent onset date: October 14, 2014) and 61 sewage samples (most recent positive sample: March 4, 2015) during the reporting period. Virus from an independent cVDPV2 emergence, apparently originating in Chad in 2012 (5), was isolated from 18 AFP patients (most recent onset date: November 3, 2014) and 32 sewage samples (most recent positive sample: June 18, 2014) in 2014. In addition, four Kaduna State sewage isolates from samples collected from August 2014 through January 2015 had shared nucleotide substitutions at six VP1 positions and the accumulation of VP1 substitutions (0.8%–1.4%) over time (8), both characteristics consistent with cVDPV2s. Circulating VDPV2s were found only in the northern states during the reporting period.

**Pakistan.** At least five independent cVDPV2 emergences have occurred in Pakistan since 2012. The emergence associated with most reported cases (71 in Pakistan and four in

Afghanistan) was first detected in Killa Abdullah, Balochistan, in August 2012 (5), spread to the insecure North Waziristan Agency in 2013, causing a large outbreak; to parts of Karachi in 2012–2013; and to neighboring Tribal Agencies and Khyber Pakhtunkhwa in 2014. Four cases in Kandahar, Afghanistan, in 2012–2013 were associated with this emergence. The last case from this emergence was reported in June 2014, and the most divergent isolate differed from OPV type 2 at 3.7% of VP1 nucleotide positions. Three additional independent emergences were detected in North Waziristan Agency, associated with five cases in 2013 (0.8%–1.1% VP1 divergence), three during 2013–2014 (0.8%–1.2% VP1 divergence), and two in 2014 (1.1% VP1 divergence), respectively. A fifth cVDPV2 emergence, associated with one AFP case (December 13, 2014), and 29 closely related but nonidentical 2014–2015 sewage isolates (0.8%–2.1% VP1 divergence), has been detected in an insecure part of Karachi, with subsequent introduction into Quetta, Balochistan.

**South Sudan.** In September 2014, two cVDPV2 isolates (1.0% VP1 divergence) were identified from patients with AFP in Rubkona, Unity State. The isolates shared three VP1 nucleotide substitutions, consistent with epidemiologic linkage.

### iVDPVs

Since the introduction of OPV in 1961, approximately 100 persons with PIDs worldwide have been found to be excreting iVDPVs, indicating prolonged infection; the majority of these immunodeficiencies were detected only after onset of paralysis. After implementation of intensified surveillance for VDPVs and special studies of iVDPV excretion among persons with PIDs in developing and middle-income countries (9), detection of new iVDPV infections increased from two during January 2008–June 2009, to nine during July 2009–June 2011, and to 12 during April 2011–June 2012, but decreased to 10 during July 2012–December 2013 (5), and to eight during the current reporting period (Table). Like cVDPVs, type 2 iVDPVs are the most prevalent (65%), followed by type 1 (18%) and type 3 (17%). Some patients have heterotypic (i.e., types 1 and 2 or types 2 and 3) iVDPV infections, with the extent of sequence divergence in each isolate of the heterotypic mixture consistent with derivation from a single tOPV source dose (4). Eight new patients with iVDPV infections were reported during January 2014–March 2015 (in addition to the patient with the longest known iVDPV infection, whose infection continued during the reporting period) are described as follows.

**Albania.** A boy aged 5 months with X-linked agammaglobulinemia, who first received OPV in March 2014 and developed paralysis in June 2014, his iVDPV3 infection cleared after September 2014.

TABLE. Vaccine-derived polioviruses (VDPVs) detected worldwide, January 2014–March 2015

Category	Country	Year(s) detected*	Source of isolates (total cases or specimens) <sup>†</sup>	Serotype	No. of isolates <sup>§</sup>		Non-AFP source	VP1 divergence from Sabin OPV strain (%)	Routine coverage with 3 doses of polio vaccine (%) <sup>¶</sup>	Estimated duration of VDPV replication**	Current status (date of last outbreak case, last patient isolate, or last environmental sample)
					Cases	Contacts					
cVDPV	Madagascar	2014	AFP patient	1	1	2	—	2.2	73	2 yrs	September 29, 2014
	Nigeria	2005–2015	Outbreaks (394 total cases) <sup>††</sup>	2	11	—	61	0.7–8.4	67	10 yrs	March 4, 2015 <sup>§§</sup>
	Nigeria	2013–2014	Importation <sup>¶¶</sup> (22 total cases)	2	18	—	32	1.2–3.9	67	3 yrs	November 3, 2014
	Pakistan	2012–2015	Outbreaks (82 total cases)	2	18	3	26	0.7–3.7	72	3 yrs	March 28, 2015***
iVDPV	South Sudan	2014	2 cases	2	2	—	—	1.0	50	~1 yr	September 12, 2014
	Albania	2014	AFP patient XLA	3	1	—	—	0.7–1.0	99	6 mos	September 12, 2014
	China	2014	AFP patient	3	1	—	—	1.4	99	~1 yr	November 26, 2014
	Iran	2014	Non-AFP SCID	1	—	—	1	2.4	98	10 mos	April 15, 2014
		2014	AFP patient XLA	1	1	—	—	1.8		1.5 yrs	August 2, 2014
		2014	AFP patient PID	2	1	—	—	0.7		<1 yr	September 13, 2014
	Libya <sup>†††</sup>	2014	Non-AFP SCID	2	—	—	1	0.7–1.0	95	4 mos	February 7, 2014
	Tunisia	2014	Non-AFP SCID	2	—	—	1	1.0	98	~1 yr	May 2014
	Turkey	2014	Non-AFP SCID	3	—	—	1	1.2	98	1.4 yrs	February 17, 2015
	UK	2014	Non-AFP CVID	2	—	—	1	17.9	96	>28 yrs	June 22, 2014
aVDPV	Brazil	2014	Environment	2	—	—	1	8.6	99	8 yrs	January 2014
	Chad	2015	AFP patient	2	1	—	—	0.8	50	<1 yr	January 8, 2015
	China	2014–2015	AFP patient	1	1	—	—	1.1	99	~1 yr	March 20, 2015
		2014–2015	AFP patients	2	4	—	—	0.7–2.4		<1 yr; 2 yrs	March 21, 2015
		2014	Non-AFP patient	1	—	—	1	1.1		~1 yr	October 2014
	DRC	2014	AFP patient	2	1	—	—	1.1	70	1 yr	January 15, 2015
	Egypt	2014	AFP patient	2	1	—	—	1.0	97	~1 yr	April 19, 2014
		2014	Environment	1	—	—	2	1.1; 2.7		1 yr; 2.5 yrs	April 20, 2014
		2014–2015	Environment	2	—	—	2	0.7		<1 yr	February 4, 2015
	Ethiopia	2014–2015	AFP patient	2	1	—	—	0.7–0.9	70	<1 yr	March 5, 2015
	Guinea	2014	AFP patient	2	1	—	—	1.3	64	~1 yr	August 30, 2014
	India	2014–2015	AFP patients	2	4	—	—	0.7–1.0	70	~1 yr	February 26, 2015
	Israel	1998–2014	Environment	2	—	—	2	>15%	94 <sup>§§§</sup>	>15 yrs	September 22, 2014
		2014	Environment	2	—	—	1	0.7		<1 yr	January 26, 2014
	Madagascar	2015	AFP patient	1	1	—	—	3.9	73	2 yrs	January 31, 2015
	Nigeria	2014	AFP patients	2	2	—	—	0.7	67	<1 yr	April 5, 2014
		2014–2015	Environment	2	—	—	8	0.7–1.4		≤1 yr	March 9, 2015
	Pakistan	2014–2015	AFP patients	2	9	1	—	0.8–2.3	72	≤1 yr; 2 yrs	February 9, 2015
		2014–2015	Environment	2	—	—	6	0.8–1.4		≤1 yr	January 2015
	Philippines	2015	AFP patient	2	1	—	—	0.8	88	<1 yr	December 18, 2014
	Russia	2014	AFP patient	3	1	3	—	1.1	98	~1 yr	July 10, 2014
	Turkey	2014	AFP contact	1	—	1	—	1.0	98	~1 yr	May 8, 2014
	Uganda	2014	AFP patients	2	2	—	—	0.7	82	<1 yr	August 13, 2014

**Abbreviations:** cVDPV = circulating VDPV; iVDPV = immunodeficiency-associated VDPV; aVDPV = ambiguous VDPV; OPV = oral poliovirus vaccine; IPV = inactivated poliovirus vaccine; AFP = acute flaccid paralysis; PID = primary immunodeficiency; SCID = severe combined immunodeficiency; XLA = X-linked agammaglobulinemia; CVID = common variable immunodeficiency; DRC = Democratic Republic of the Congo.

\* Total years detected and cumulative totals for previously reported cVDPV outbreaks (Nigeria and Pakistan).

<sup>†</sup> Outbreaks list total cases clearly associated with cVDPVs. Some VDPV case isolates from outbreak periods might be listed as aVDPVs.

<sup>§</sup> Total cases for VDPV-positive specimens from AFP cases and total VDPV-positive samples for environmental (sewage) samples.

<sup>¶</sup> Based on 2013 data from the World Health Organization (WHO) Vaccine Preventable Diseases Monitoring System (2014 global summary) and WHO–United Nations Children's Fund (UNICEF) coverage estimates, available at [http://www.who.int/immunization/monitoring\\_surveillance](http://www.who.int/immunization/monitoring_surveillance). National data might not reflect weaknesses at subnational levels.

\*\* Duration of cVDPV circulation was estimated from extent of VP1 nucleotide divergence from the corresponding Sabin OPV strain; duration of immunodeficiency-associated VDPV replication was estimated from clinical record by assuming that exposure was from initial receipt of OPV; duration of ambiguous VDPV replication was estimated from sequence data.

<sup>††</sup> Count does not include 29 cases with <10 substitutions in VP1 detected before 2010.

<sup>§§</sup> The most recent isolate was from an environmental sample.

<sup>¶¶</sup> Importation from Chad.

\*\*\* The most recent isolate was from an environmental sample.

<sup>†††</sup> The VDPV was detected and characterized in Germany where the patient had gone for treatment.

<sup>§§§</sup> Value for routine IPV immunization in 2013. Israel conducted two rounds with bivalent OPV in response to detection of imported wild poliovirus type 1 from environmental samples.

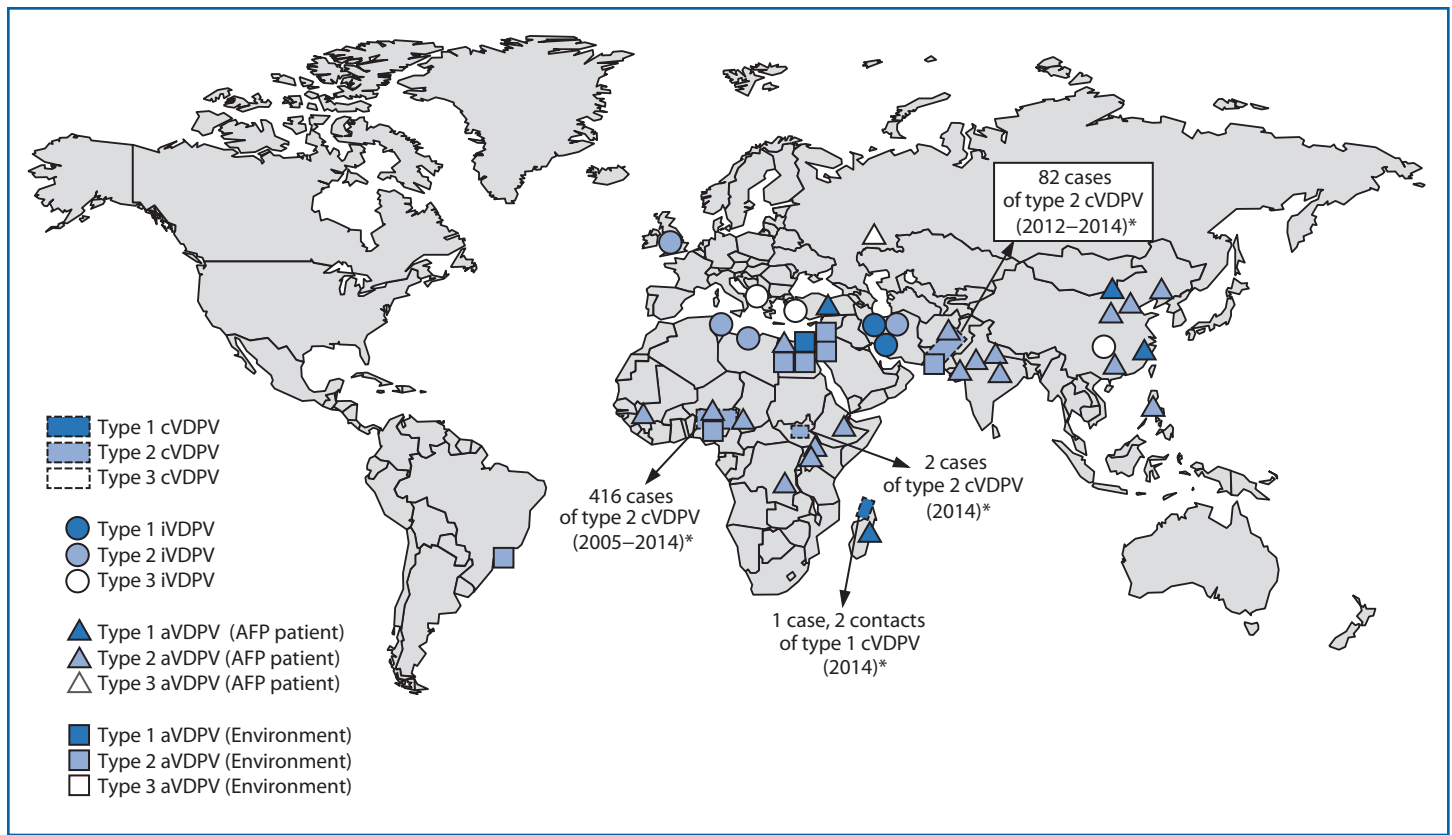
**China.** A boy aged 1 year with PID, who received his third OPV dose in February 2014, his iVDPV3 infection cleared soon after onset of AFP in November 2014.

**Iran.** Iran has maintained sensitive clinical and laboratory surveillance to screen persons with PIDs for poliovirus infections. During this reporting period, three patients (two with AFP) were found to be excreting iVDPVs. One was a nonparalyzed child aged 10 months with severe combined

immunodeficiency infected with an iVDPV1. Another boy, aged 10 months, with X-linked agammaglobulinemia received OPV in March 2014 and developed paralysis in May 2014; his iVDPV1 infection cleared after August 2014. A boy aged 9 months with PID and infected with iVDPV2 developed paralysis in June 2014 but stopped excreting iVDPVs after September 2014.



FIGURE 1. Vaccine-derived polioviruses (VDPVs) detected worldwide, January 2014–March 2015



**Abbreviations:** cVDPV = circulating VDPV; iVDPV = immunodeficiency-associated VDPV; aVDPV = ambiguous VDPV; AFP = acute flaccid paralysis.

\* Spread of cVDPVs followed the elimination of the corresponding serotype of indigenous wild poliovirus, but with continued introduction of oral poliovirus vaccine into communities with growing immunity gaps. All of the cVDPV outbreaks were detected first by the laboratory, using sequence data and evolutionary analyses.

**Libya.** A nonparalyzed girl aged 1 month with severe combined immunodeficiency traveled to Germany for treatment and was found to be infected with iVDPV2 during November 2013–February 2014; excretion stopped following bone marrow transplantation.

**Tunisia.** A nonparalyzed boy aged 11 years with severe combined immunodeficiency was infected with an iVDPV2. He stopped excreting iVDPVs after May 2014.

**Turkey.** A nonparalyzed girl aged 1 year with severe combined immunodeficiency was infected with an iVDPV3, which she continued to excrete into December 2014.

**United Kingdom.** A man aged 44 years with common variable immunodeficiency was found to be excreting iVDPV2 since 1995. He has no AFP, but the sequence properties of the isolates obtained from serial specimens are consistent with chronic iVDPV2 infection since his last OPV dose at age 7 years.

## aVDPVs

During January 2014–March 2015, aVDPVs were isolated in 16 countries (Table). Detection of aVDPVs in settings with <60% vaccination coverage with 3 doses of polio vaccine might signal cVDPV emergence and potential gaps in surveillance. Some aVDPVs, especially those with limited divergence detected in areas with high vaccination coverage and in patients with no known immunodeficiency, might represent limited spread of OPV or the upper limit of OPV sequence divergence in a single normal vaccine recipient or contact. The most divergent aVDPV was from Brazil, a country with >90% vaccination coverage with 3 doses of polio vaccine. Selected aVDPVs from the reporting period are described as follows.

**Brazil.** An aVDPV2 (8.6% VP1 divergence) was isolated from sewage in the Port of São Sebastião, São Paulo, in January 2014. The isolate resembles an iVDPV but is classified as an aVDPV because no immunodeficient source patient has been identified.

**China.** Sporadic aVDPVs were isolated in six different provinces during January 2014–March 2015; one aVDPV1

and four aVDPV2s were isolated from AFP patients, and one aVDPV1 was isolated from a healthy child.

**India.** Four aVDPV2s (0.7%–1.0% VP1 divergence) were isolated from AFP patients in four different states during January 2014–March 2015.

**Israel.** Two isolates of the highly divergent, neurovirulent aVDPV2 first detected in 1998 were detected in sewage samples collected on May 4 and September 22, 2014, and an independent aVDPV2 was found in sewage collected on January 26, 2014.

**Madagascar.** An aVDPV1 (3.9% VP1 divergence) was isolated from a patient in Nosy-Varika, Fianarantsoa Province (central east coast), with onset of paralysis on January 31, 2015. Despite a small number of VP1 substitutions shared with the 2014 cVDPV1 isolates from Analalava, this aVDPV1 appears to be an independent emergence.

**Nigeria.** Ten aVDPV2s (two from AFP patients, eight from sewage samples, and all with 0.6%–0.7% VP1 divergence) were isolated in the northern states and the Federal Capital Territory during the reporting period.

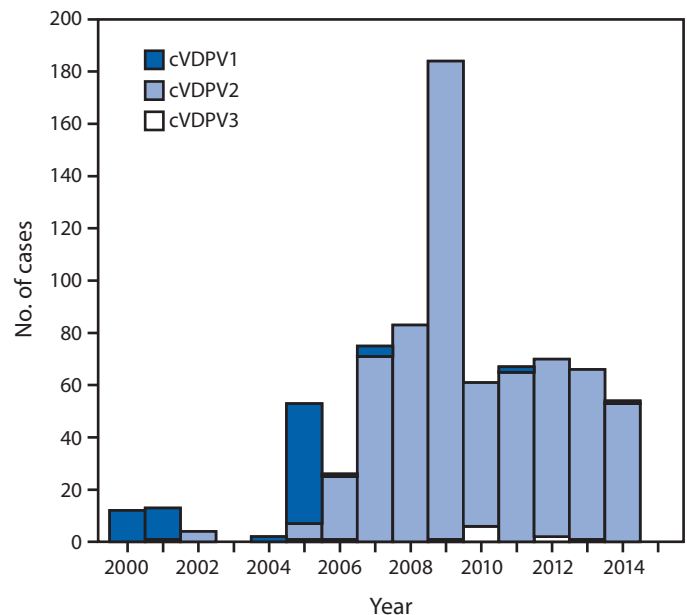
**Pakistan.** Fifteen aVDPV2s isolates (10 from AFP cases/contacts, four from sewage samples, and all with 0.8%–2.3% VP1 divergence) were isolated during January 2014–March 2015. The most recent aVDPV2 isolates were from the Khyber Agency (two AFP cases in February 2015 and 0.8% VP1 divergence), and Peshawar, Khyber Pakhtunkhwa (from a January 2015 sewage sample, with 0.8% VP1 divergence).

### Discussion

During January 2014–March 2015, the size and geographic distribution of cVDPV outbreaks further declined since July 2012–December 2013. However, new cVDPV2 lineages have emerged in both Nigeria and Pakistan in settings of insecurity and widening immunity gaps to cVDPV2. Inclusion of more tOPV rounds in the steadily improving supplementary immunization activities (SIAs)\* and ensuring increased access to unimmunized children are important factors in controlling cVDPV2 outbreaks. The new outbreaks in Madagascar and South Sudan underscore the importance of maintaining high population immunity to all polioviruses and of sensitive AFP surveillance.

Although expanded testing of sewage samples for the presence of poliovirus (environmental surveillance) in Nigeria and Pakistan has increased the sensitivity of poliovirus detection, especially cVDPV2, which has a 10-fold lower case-to-infection

**FIGURE 2. Circulating vaccine-derived poliovirus (cVDPV) cases detected worldwide, by serotype and year, January 2000–March 2015\***



\* Data through March 2015, as available by June 15, 2015.

ratio than WPV1 (4), it also presents new logistical and technical challenges to the Global Polio Laboratory Network, because VDPVs must be detected within the complex mixtures of polioviruses and other enteric viruses present in sewage. The rRT-PCR screening for VDPVs must be capable of recognizing the small number of genetic differences distinguishing VDPVs from the closely related VRPVs, which are currently of little public health interest, while striking a balance between sensitivity and specificity not required for the identification of WPVs, which are readily distinguishable from VRPVs and VDPVs (7); the requirement for high specificity has resulted in an increased need for nucleotide sequencing.

Interpreting the virologic data presents additional challenges: one VDPV isolate from an AFP patient might signal hundreds to thousands of inapparent cVDPV infections, whereas multiple VDPV sewage isolates might derive from a single iVDPV infection. Environmental cVDPV isolates can be recognized by their close genetic relationship with known cVDPVs from one or more AFP patients or by local detection of closely related VDPVs over several months that show progressive divergence from the parental OPV strain. These latter environmental cVDPVs are distinguishable by their sequence properties from those environmental aVDPVs (4) which very likely signal the presence of a chronic iVDPV excretor in the community. Indeed, highly divergent environmental aVDPVs that are probably iVDPVs from unidentified chronic excretors have been detected in five countries, most recently in Brazil.

\* SIAs are mass vaccination campaigns conducted in a short period (days to weeks) during which a dose of OPV is administered to all children aged <5 years, regardless of previous vaccination history. Campaigns can be conducted nationally or in portions of a country.

## Summary

### What is already known on this topic?

Genetically divergent vaccine-derived polioviruses (VDPVs) are detected by poliovirus surveillance and have biologic properties indistinguishable from wild polioviruses. High polio vaccination coverage can prevent circulating VDPV (cVDPV) outbreaks, but prolonged immunodeficiency-associated VDPV (iVDPV) infections will occur as long as oral poliovirus vaccine (OPV) is used.

### What is added by this report?

The intensity of cVDPV transmission fell after mid-2014. Recent cVDPV outbreaks in Afghanistan, Chad, Somalia, and Yemen have apparently stopped, and the large outbreak in Nigeria has nearly stopped. Virus of the major cVDPV emergence in Pakistan was last detected in June 2014, but low-level circulation of a new emergence was detected into 2015. New, possibly small, outbreaks were detected in Madagascar and South Sudan. Nine new prolonged iVDPV infections in seven countries were detected, either by characterization of isolates from patients with acute flaccid paralysis (AFP) or by intensified search for iVDPV excretion among persons with primary B-cell immunodeficiencies. Since 2006, >97% of cVDPVs detected have been type 2.

### What are the implications for public health practice?

Circulating VDPV outbreaks can be prevented and controlled by high OPV coverage. By contrast, only cessation of OPV use will prevent prolonged iVDPV infections. WHO has responded to the continued global type 2 VDPV risk by incorporating the following into its new strategic plan: 1) shifting from trivalent OPV to bivalent OPV (types 1 and 3) by April 2016; 2) including ≥1 dose of inactivated poliovirus vaccine into routine immunization schedules worldwide; 3) maintaining strategic stockpiles of monovalent OPV; 4) developing a robust acute flaccid paralysis and poliovirus surveillance and response capacity; and 5) encouraging development of antiviral drugs to clear prolonged iVDPV infections.

Detection of less divergent environmental VDPVs (especially VDPV2s) without linkage to known infected persons presents the greatest challenges for epidemiologic interpretation.

Special studies in several countries to search for VDPV infections among patients with PIDs have increased the number of known iVDPV excretors, while documenting the infrequency of iVDPV infections, even among persons with PIDs (9). Global AFP surveillance and environmental surveillance have proven most sensitive in detecting prolonged iVDPV excretion.

In view of the rising incidence of cVDPV2 outbreaks more than a decade after the last known WPV2 case, GPEI has incorporated the coordinated worldwide withdrawal of tOPV and replacement with bOPV into its new strategic plan, with the ultimate goal of stopping all OPV use (6). The switch from tOPV to bOPV, planned for April 2016, is predicated on the

absence of any known cVDPV2 transmission (6). The absence of any reported cases associated with cVDPV2 in 2015 (all cVDPV2 isolates through March 2015 were from environment samples) and the current low frequency of cVDPV2 detection worldwide is encouraging. To ensure that VDPV emergence is minimized and that any VDPV infections are detected, it will be essential to continue efforts to strengthen routine immunization services and to strengthen AFP and poliovirus surveillance during 2015. Most countries will incorporate at least 1 dose of IPV into routine childhood immunization schedules in 2015 (6).

Replacement of tOPV with bOPV will greatly reduce the risk for cVDPV2 outbreaks, and global cessation of OPV use will ultimately prevent all cVDPV outbreaks and all new iVDPV infections (6). However, a small number of persons with chronic iVDPV infections, as exemplified by the non-AFP common variable immunodeficiency patient from the United Kingdom, might continue to excrete poliovirus for a decade or more after receipt of the last OPV dose. Therefore, maintenance of high levels of population immunity through comprehensive IPV coverage will be necessary to protect against iVDPV becoming a source of spread in the community. Detection of chronic iVDPV excretors in all countries (9) and development of antivirals to clear chronic iVDPV infections are also important (10).

## Acknowledgments

Global Polio Laboratory Network. Humayun Asghar, MD, WHO Regional Office, Cairo, Egypt. Sunil Bahl, MD, National Polio Surveillance Program, New Delhi, India. Eugene Gavrillin, PhD, WHO Regional Office, Copenhagen, Denmark. Nicky Gumedé, PhD, WHO Regional Office, Brazzaville, Republic of the Congo. Gloria Rey, MSc, WHO Regional Office, Washington, DC. W. William Schluter, MD, WHO Regional Office, Manila, Philippines. Shohreh Shahmahmoodi, PhD, Tehran University of Medical Sciences, Tehran, Iran. Salmaan Sharif, PhD, Sohail Zaidi, MSc, National Institute of Health of Pakistan, Islamabad, Pakistan. Lester Shulman, PhD, Central Virology Laboratory, Tel Hashomer, Israel. Wenbo Xu, PhD, China CDC, Beijing, China. Eric Mast, MD, Global Immunization Division, Center for Global Health, CDC; Jane Iber, MSc, Qi Chen, MSc, David Kilpatrick, PhD, Steven Oberste, PhD, Division of Viral Diseases, National Center for Immunization and Respiratory Diseases, CDC.

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# Yellow Fever Vaccine Booster Doses: Recommendations of the Advisory Committee on Immunization Practices, 2015

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On February 26, 2015, the Advisory Committee on Immunization Practices (ACIP) voted that a single primary dose of yellow fever vaccine provides long-lasting protection and is adequate for most travelers (1). ACIP also approved recommendations for at-risk laboratory personnel and certain travelers to receive additional doses of yellow fever vaccine (Box). The ACIP Japanese Encephalitis and Yellow Fever Vaccines Workgroup evaluated published and unpublished data on yellow fever vaccine immunogenicity and safety. The evidence for benefits and risks associated with yellow fever vaccine booster doses was evaluated using the Grading of Recommendations, Assessment, Development, and Evaluation (GRADE) framework (2,3). This report summarizes the evidence considered by ACIP and provides the updated recommendations for yellow fever vaccine booster doses.

## Yellow Fever Epidemiology and Risk for Disease in Travelers

Yellow fever is a mosquito-borne viral disease that is endemic to sub-Saharan Africa and tropical South America. Worldwide, yellow fever virus causes an estimated 200,000 cases of clinical disease and 30,000 deaths annually (4). Clinical disease ranges

from a mild, nonspecific febrile illness to severe disease with jaundice and hemorrhage. The case-fatality ratio for severe yellow fever is 20%–50% (5). Because no specific treatment exists, prevention through vaccination is critical to reduce morbidity and mortality from yellow fever virus infection.

The risk of a traveler acquiring yellow fever varies based on season, location, activities, and duration of their travel. For a 2-week stay, the estimated risk for illness attributed to yellow fever for an unvaccinated traveler to West Africa is 50 cases per 100,000 population; for South America, the risk for illness is five cases per 100,000 population (6).

## Yellow Fever Vaccine Recommendations and International Health Regulations Requirements

Yellow fever vaccine is recommended for persons aged ≥9 months who are traveling to or living in areas with risk for yellow fever virus transmission (7). International Health Regulations allow countries to require proof of yellow fever vaccination from travelers entering their country (8). These requirements are intended to minimize the potential importation and spread of yellow fever virus. Currently, International Health Regulations specify that a dose of yellow fever vaccine is valid for 10 years. Therefore, at present, travelers to countries with a yellow fever vaccination entry requirement must have received a dose of yellow fever vaccine within the past 10 years.

**Recent changes to yellow fever vaccine recommendations.** In April 2013, the World Health Organization Strategic Advisory Group of Experts on Immunization concluded that a single primary dose of yellow fever vaccine is sufficient to confer sustained immunity and lifelong protection against yellow fever disease, and that a booster dose is not needed (9). This conclusion was based on a systematic review of published studies on the duration of immunity after a single dose of yellow fever vaccine, and on data that suggest vaccine failures are extremely rare and do not increase in frequency with time since vaccination (10). The advisory group noted that future studies and surveillance data should be used to identify specific risk groups, such as persons infected with human immunodeficiency virus (HIV) or infants, who might benefit from a booster dose. In May 2014, the World Health Assembly adopted the recommendation to remove the 10-year booster dose requirement from the International Health Regulations by June 2016 (11).

*Recommendations for routine use of vaccines in children, adolescents, and adults are developed by the Advisory Committee on Immunization Practices (ACIP). ACIP is chartered as a federal advisory committee to provide expert external advice and guidance to the Director of the Centers for Disease Control and Prevention (CDC) on use of vaccines and related agents for the control of vaccine-preventable diseases in the civilian population of the United States. Recommendations for routine use of vaccines in children and adolescents are harmonized to the greatest extent possible with recommendations made by the American Academy of Pediatrics (AAP), the American Academy of Family Physicians (AAFP), and the American College of Obstetricians and Gynecologists (ACOG). Recommendations for routine use of vaccines in adults are harmonized with recommendations of AAFP, ACOG, and the American College of Physicians (ACP). ACIP recommendations approved by the CDC Director become agency guidelines on the date published in the Morbidity and Mortality Weekly Report (MMWR). Additional information regarding ACIP is available at <http://www.cdc.gov/vaccines/acip>.*



## Yellow Fever Vaccine Long-term Immunogenicity Data

No data are available on vaccine efficacy or protective antibody titers (i.e., seroprotection) related to long-term immunogenicity after yellow fever vaccination. Benefits considered critical in assessing the need for booster doses of yellow fever vaccine for U.S. travelers or laboratory workers included vaccine effectiveness (i.e., a lack of vaccine failures) and evidence of seropositivity (i.e., yellow fever virus-specific antibodies detected in a blood sample) (3).

**Vaccine effectiveness.** A total of 23 vaccine failures were identified after the administration of >540 million doses of yellow fever vaccine (3). Of the 23 cases, five occurred <10 days after vaccination and were excluded because most persons are not expected to develop protective titers in that timeframe (5). Of the remaining 18 cases, 16 (89%) occurred in persons who reported receiving a dose of the vaccine within the previous 10 years (3). One vaccine failure occurred at 20 years and one at 27 years post-vaccination.

**Seropositivity.** Thirteen observational studies provided immunogenicity data on 1,137 persons vaccinated  $\geq 10$  years previously (3). Using a random effects model, the estimated seropositivity rate for persons vaccinated  $\geq 10$  years previously was 92% (95% confidence interval [CI] = 85%–96%). Of the 164 persons vaccinated  $\geq 20$  years previously, the estimated seropositivity rate was 80% (CI = 74%–86%).

## Yellow Fever Vaccine Booster Dose Safety Data

Serious adverse events, yellow fever vaccine–associated viscerotropic disease (a severe illness similar to wild-type disease), and yellow fever vaccine-associated neurologic disease were considered critical risks to assess the need for yellow fever vaccine booster doses (7).

**Serious adverse events.** Nine observational studies provided data on serious adverse events for 333 million distributed doses of yellow fever vaccine (3). Overall, 1,255 persons were reported to have a serious adverse event after yellow fever vaccination. For most (84%) persons, it was unknown if the adverse event occurred after a primary or booster dose of the vaccine. Of the 201 persons with a serious adverse event where dose type was known, 14 (7%) of the adverse events occurred after a booster dose of vaccine.

**Viscerotropic disease.** Eight observational studies provided data on viscerotropic disease for 437 million distributed doses of yellow fever vaccine (3). A total of 72 persons had yellow fever vaccine–associated viscerotropic disease. Of the 31 persons where dose type was known, one (3%) had viscerotropic disease after receiving a booster dose of the vaccine; no

### BOX. Recommendations for use of yellow fever vaccine booster doses\*

- A single primary dose of yellow fever vaccine provides long-lasting protection and is adequate for most travelers [Category A].
- Additional doses of yellow fever vaccine are recommended for certain travelers:
  - Women who were pregnant (regardless of trimester) when they received their initial dose of yellow fever vaccine should receive 1 additional dose of yellow fever vaccine before their next travel that puts them at risk for yellow fever virus infection [Category A];
  - Persons who received a hematopoietic stem cell transplant after receiving a dose of yellow fever vaccine and who are sufficiently immunocompetent to be safely vaccinated should be revaccinated before their next travel that puts them at risk for yellow fever virus infection [Category A];
  - Persons who were infected with human immunodeficiency virus when they received their last dose of yellow fever vaccine should receive a dose every 10 years if they continue to be at risk for yellow fever virus infection [Category A].
- A booster dose may be given to travelers who received their last dose of yellow fever vaccine at least 10 years previously and who will be in a higher-risk setting based on season, location, activities, and duration of their travel [Category B]. This would include travelers who plan to spend a prolonged period in endemic areas or those traveling to highly endemic areas such as rural West Africa during peak transmission season or an area with an ongoing outbreak.
- Laboratory workers who routinely handle wild-type yellow fever virus should have yellow fever virus-specific neutralizing antibody titers measured at least every 10 years to determine if they should receive additional doses of the vaccine. For laboratory workers who are unable to have neutralizing antibody titers measured, yellow fever vaccine should be given every 10 years as long as they remain at risk [Category A].

\* Persons being considered for additional doses of yellow fever vaccine should be assessed for contraindications or precautions in accordance with the current yellow fever vaccine ACIP recommendations (7).

laboratory testing to assess vaccine causality was performed for that case.

**Neurologic disease.** Eight observational studies provided neurologic disease data for approximately 462 million

distributed doses of yellow fever vaccine (3). A total of 218 persons had yellow fever vaccine–associated neurologic disease. Of the 110 persons where dose type was known, three (3%) persons reported neurologic disease after receiving a booster dose of the vaccine.

## Other relevant evidence

**Pregnant women.** The proportion of women who develop yellow fever virus antibodies is variable and might be related to the trimester in which they received the vaccine. Among pregnant women who received yellow fever vaccine primarily in their third trimester, 39% (32 of 83) had evidence of seroconversion to yellow fever virus at 2–4 weeks post-vaccination, compared with 94% (89 of 95) in the general population (12). Of 433 women vaccinated primarily in the first trimester (mean gestational age = 5.7 weeks; CI = 5.2–6.2), 425 (98%) developed yellow fever virus–specific neutralizing antibodies at 6 weeks post-vaccination (13).

**Hematopoietic stem cell transplant recipients.** Data are limited on safety and immunogenicity for yellow fever vaccine in hematopoietic stem cell transplant recipients. However, data suggest most recipients become seronegative to live viral vaccine antigens after transplantation (14). Infectious Diseases Society of America guidelines recommend re-administering live viral vaccines, such as measles, mumps, and rubella vaccine and varicella vaccine, to post-transplant patients if the recipient is seronegative and is no longer immunosuppressed (15).

**HIV-infected persons.** Two published studies provide immunogenicity data for yellow fever vaccines in HIV-infected persons (16,17). Both studies found lower rates of yellow fever virus–specific neutralizing antibodies among HIV-infected persons compared with uninfected controls at 10 to 12 months post-vaccination. Although the mechanisms for the diminished immune response in HIV-infected persons are uncertain, an inverse correlation exists between immune response and HIV RNA levels and a positive correlation with CD4+ cell counts (18).

**Young children.** Twelve studies provided data on the initial immune response to yellow fever vaccine in children aged 4 months–10 years (3). All studies included children who resided in endemic areas, and 10 studies included children who received at least one other vaccine at the same time as yellow fever vaccine. Based on a random effects model, the estimated seroconversion rate in 4,675 children was 93% (CI = 88%–96%). No difference was observed in the seroconversion rates between children aged <9 months and those aged ≥9 months (3).

**Other higher-risk groups.** Over the preceding 20 years, 90% of all yellow fever cases were reported from countries in West Africa, and epidemiologic data suggest that travelers to West Africa are at the highest risk for travel-associated yellow

## Summary

### What is currently recommended?

In 2009, the Advisory Committee on Immunization Practices (ACIP) approved yellow fever vaccine recommendations that noted International Health Regulations require revaccination at intervals of 10 years to boost antibody titer. Evidence from multiple studies demonstrates that yellow fever vaccine immunity persists for many decades and might provide life-long protection.

### Why are the recommendations being modified now?

The World Health Organization Strategic Advisory Group of Experts in Immunization concluded in April 2013 that a single primary dose of yellow fever vaccine is sufficient to confer sustained immunity and lifelong protection against yellow fever disease, and a booster dose of the vaccine is not needed. In May 2014, the World Health Assembly adopted the recommendation to remove the 10-year booster dose requirement from the International Health Regulations by June 2016. Once the International Health Regulations are updated, the current statement in the ACIP recommendation will no longer be relevant.

### What are the new recommendations?

A single primary dose of yellow fever vaccine provides long-lasting protection and is adequate for most travelers. The recommendations also provide considerations and recommendations for at-risk laboratory personnel and certain travelers to receive additional doses of yellow fever vaccine.

fever (5). Persons traveling to an area with an ongoing outbreak, persons traveling for a prolonged period in an endemic area, and laboratory workers who routinely handle wild-type yellow fever virus are also considered to be at higher risk for yellow fever virus exposure and disease than other persons for whom yellow fever vaccine is recommended.

## Rationale for Yellow Fever Vaccine Booster Dose Recommendations

The GRADE evaluation found that there are few vaccine failures documented after a primary dose of yellow fever vaccine, most (92%) primary vaccine recipients maintain detectable levels of neutralizing antibodies ≥10 years post-vaccination, and few serious adverse events have been reported after a booster dose of yellow fever vaccine (3). Based on the available data, ACIP voted to no longer recommend booster dose of yellow fever vaccine for most travelers, because a single dose of yellow fever vaccine provides long-lasting protection (Box). However, additional doses of yellow fever vaccine are recommended for certain populations (i.e., pregnant women, hematopoietic stem cell transplant recipients, and HIV-infected persons) who might not have as robust or sustained immune response to yellow fever vaccine compared with other recipients. Furthermore,

additional doses may be given to certain groups believed to be at increased risk for yellow fever disease either because of their location and duration of travel or because of more consistent exposure to virulent virus (i.e., laboratory workers). ACIP meeting minutes are available at <http://www.cdc.gov/vaccines/acip/meetings/meetings-info.html>.

### Acknowledgments

Members of the Advisory Committee on Immunization Practices (ACIP). ACIP Japanese Encephalitis and Yellow Fever Vaccines Workgroup. Bradley Biggerstaff, National Center for Emerging and Zoonotic Diseases, CDC. ACIP member roster for July 2014–June 2015 available at <http://www.cdc.gov/vaccines/acip/committee/members.html>.

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## Notes from the Field

### Tickborne Relapsing Fever Outbreak at an Outdoor Education Camp — Arizona, 2014

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Tickborne relapsing fever (TBRF) is a bacterial infection characterized by recurring episodes of fever, headache, muscle and joint aches, and nausea. In North America, TBRF primarily is caused by *Borrelia hermsii* spirochetes transmitted by *Ornithodoros hermsii* ticks (1). Once infected, these soft ticks are infectious for life (2) and transmit the spirochete to sleeping humans quickly (possibly within 30 seconds) during short feeds (15–90 minutes) (1–3). On August 10, 2014, the Coconino County Public Health Services District in Arizona was notified by a local hospital that five high school students who attended the same outdoor education camp had been hospitalized with fever, headache, and myalgias. Hantavirus infection initially was suspected because of reported exposure to rodent droppings, but after detecting spirochetes on peripheral blood smears from all five hospitalized students, TBRF was diagnosed. The camp was instructed to close immediately, and the health department, in collaboration with local university experts, investigated to identify additional cases, determine the cause, and prevent further infections. A total of 11 cases (six confirmed and five probable) were identified.

Camp staff members and attendees were interviewed during August 11–14. Medical records of the five hospitalized patients were reviewed, and the campsite was inspected for evidence of rodent or tick infestation. Consistent with the Arizona Department of Health Services case definition, a probable case was defined as an illness with at least three of the four major TBRF signs and symptoms (fever, chills, myalgias, and headache) without laboratory testing in a person attending the camp during August 1–3, 2014. A case was confirmed by visualization of spirochetes in an attendee's blood smear or by *Borrelia hermsii* isolation by culture.

During August 1–3, a total of 45 persons (39 high school football players and six adult coaches) attended a school-run outdoor education camp located in a wooded area in Coconino County. Thirty-one (69%) of the 45 persons at the camp were interviewed. Six confirmed cases (four by visualization of spirochetes on blood smear and isolation and two by visualization of spirochetes alone) and five probable cases were

identified (attack rate: 24%). Ten patients were students aged 15–17 years, and one was a coach aged 33 years.

All six persons with confirmed TBRF and four of the five persons with probable TBRF had slept in the camp's main cabin. Using the earliest date when common exposure might have occurred (August 1), the median incubation period was 6 days (range = 2–10 days). All six of the persons with confirmed TBRF had fever, headache, myalgias, and arthralgias; all five of those with probable TBRF had fever, headache, and myalgias (Table). Among the six with confirmed TBRF and known laboratory values, five had thrombocytopenia (platelets <150/ $\mu$ L); four had decreased albumin; and three had elevated transaminases. Eight of the 11 patients were treated with doxycycline and had no known major complications; the six patients with confirmed TBRF were treated with 100 mg doxycycline twice daily for 7–10 days. Attempts to obtain clinical and laboratory information on patients with probable TBRF were unsuccessful.

The investigation revealed that, during July 17–24, professional pest controllers had performed rodent-proofing activities at the main cabin; however, no acaricides (pesticides that kill ticks and mites) were applied. On August 12 and 28, the public health team inspected the cabin and found evidence of rodents and soft tick infestation, including rodent nesting material in a woodpile in a crawl space beneath the cabin, squirrel droppings in a chimney crevasse, and one live and one desiccated *Ornithodoros hermsii* tick. Among four chipmunks (*Tamias dosalis*) trapped on August 12, two were documented with *B. hermsii* by positive quantitative polymerase chain reaction. Testing of one soft tick for *B. hermsii* was negative. Camp management was provided written instructions regarding rodent-proofing and acaricide application. The camp reopened after recommendations were implemented; no additional cases have been identified.

During 1982–2013, a total of 22 TBRF cases (0–3 cases annually) were reported in Arizona residents. This 2014 outbreak of TBRF with 11 confirmed and probable cases is the largest recorded in Arizona since 1990. Health care providers and public health professionals should be aware that TBRF is a possible cause of febrile illness among patients with a travel history to areas where TBRF is endemic, particularly if they have slept in a rustic cabin (1). These findings suggest that pest control companies and cabin owners might benefit from education regarding prevention of tickborne diseases, including sleeping off the floor and away from walls, applying insect repellent on skin and clothing, and rodent-proofing.



**TABLE. Number of signs and symptoms of patients with confirmed or probable tickborne relapsing fever in an outbreak associated with an outdoor education camp — Arizona, 2014**

Sign or symptom	Total no. (N = 11)	Confirmed no. (n = 6)	Probable no. (n = 5)
Fever	11	6	5
Headache	11	6	5
Myalgias	11	6	5
Arthralgias	10	6	4
Abdominal pain	7	5	2
Fatigue	6	2	4
Vomiting	5	4	1
Cough	4	2	2
Dizziness	3	3	0
Syncope	2	2	0
Rash	2	2	0

To eliminate tick populations in a building, it is important to consider acaricide spraying concurrently with rodent-proofing because removing rodents from buildings can result in ticks losing their primary source of food and feeding on humans as an alternative (2).

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## Notes from the Field

### Update: Silicosis Mortality — United States, 1999–2013

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Silicosis is a potentially fatal but preventable occupational lung disease caused by inhaling respirable crystalline silica (silica) (*1*). Chronic silicosis, the most common form, occurs after exposure to relatively low silica concentrations for >10 years. Accelerated silicosis occurs after 5–10 years of exposure to higher silica levels, and acute silicosis can occur after only weeks or months of exposure to extremely high silica concentrations (*1*). New national mortality data for silicosis have become available since a previous report on silicosis surveillance was published earlier this year (*2*). CDC reviewed multiple cause-of-death mortality files from the National Center for Health Statistics to analyze deaths from silicosis (*International Classification of Diseases, 10th Revision* diagnosis code J62: a pneumoconiosis due to dust containing silica) reported during 1999–2013. Each record lists one underlying cause of death (the disease or injury that initiated the chain of events that led directly and inevitably to death), and up to 20 contributing causes of death (other significant conditions contributing to death but not resulting in underlying cause). Available death certificates from 35 states were reviewed for the period 2004–2006 to identify occupations associated with silicosis among decedents aged 15–44 years. Results indicate that despite substantial progress in eliminating silicosis, silicosis deaths continue to occur. Of particular concern are silicosis deaths in young adults (aged 15–44 years). These young deaths likely reflect higher exposures than those causing chronic silicosis mortality in older persons, some of sufficient magnitude to cause severe disease and death after relatively short periods of exposure. A total of 12 such deaths occurred during 2011–2013, with nine that had silicosis listed as the underlying cause of death.

During 1999–2013, a total of 2,065 decedents had silicosis listed as the underlying or as a contributing cause of death (1,122 [54.3%] decedents had silicosis listed as the underlying cause of death) (Table). The annual number of silicosis deaths declined 40% from 185 in 1999 to 111 in 2013 (p-value for trend <0.001), but the decline appears to have leveled off during 2010–2013. The lowest number of silicosis deaths (88) occurred in 2011. Higher numbers of deaths occurred in 2012 (103) and 2013 (111), but remained within the 95% confidence interval predicted by the first-order autoregressive

linear regression model used to evaluate trends for 1999–2013. Among all silicosis deaths, 47 (2.3%) decedents were aged 15–44 years; of these, 34 (72.3%) had silicosis coded as the underlying cause of death (Table). The annual number of silicosis deaths in persons aged 15–44 years varied and was 4, 0, and 8 in 2011, 2012, and 2013, respectively.

Death certificate review identified 62 silicosis deaths, accounting for 13.7% of the 451 reported silicosis deaths during 2004–2006. Of 39 (62.9%) decedents with silicosis listed as the underlying cause of death, three were aged 15–44 years. Entries on death certificates of these young decedents related to industry and occupation were classified\* as miscellaneous nonmetallic mineral product manufacturing (stationary engineers and boiler operators), construction (brickmasons and blockmasons), and cut stone and stone product manufacturing (crushing, grinding, and polishing machine setters, operators, and tenders). These industries and occupations are well-known for their association with exposure to crystalline silica (*1*).

Silicosis mortality in the United States has declined over time (*2,3*). The continuing occurrence of silicosis deaths in young adults and reports of new occupations and tasks that place workers at risk for silicosis, including fabricators and installers of quartz-containing engineered stone products and workers employed to extract natural gas by hydraulic fracturing (*4–7*), underscore the need for strengthening efforts to limit workplace exposure to crystalline silica. Effective silicosis prevention strategies for employers are available from the Occupational Safety and Health Administration† and CDC's National Institute for Occupational Safety and Health.§ State health departments can strengthen silicosis prevention efforts by identifying silicosis cases through review of state morbidity and mortality data and by investigating the circumstances surrounding silicosis cases.

\*The National Institute for Occupational Safety and Health Industry and Occupation Computerized Coding System was used to classify and code the industry and occupation according to the U.S. Census Bureau, North American Industry Classification System, and the U.S. Bureau of Labor Statistics, Standard Occupational Classification System, respectively. Additional information available at <http://wwwn.cdc.gov/niosh-nioccs>.

† Additional information available at [https://www.osha.gov/pls/oshaweb/owadisp.show\\_document?p\\_table=standards&p\\_id=12716](https://www.osha.gov/pls/oshaweb/owadisp.show_document?p_table=standards&p_id=12716).

§ Additional information available at <http://www.cdc.gov/niosh/topics/silica>.

### Acknowledgments

David N. Weissman, MD, Brent Doney, PhD, Division of Respiratory Disease Studies, National Institute for Occupational Safety and Health, CDC.

**TABLE. Number of silicosis deaths, by age group, other selected characteristics, and year — United States, 1999–2013**

Characteristic	Age group					
	15–44 yrs		≥45 yrs		Overall	
	Deaths	Underlying cause	Deaths	Underlying cause	Deaths	Underlying cause
<b>Total</b>	<b>47</b>	<b>34</b>	<b>2,018</b>	<b>1,088</b>	<b>2,065</b>	<b>1,122</b>
<b>Sex</b>						
Male	39	30	1,933	1,031	1,972	1,061
Female	8	4	85	57	93	61
<b>Race</b>						
White	37	27	1,727	927	1,764	954
Black	8	6	265	142	273	148
Other	2	1	26	19	28	20
<b>Ethnicity</b>						
Hispanic	9	7	131	83	140	90
Non-Hispanic	38	27	1,883	1,002	1,921	1,029
Unknown	0	0	4	3	4	3
<b>Year</b>						
1999	3	2	182	100	185	102
2000	5	5	146	66	151	71
2001	1	1	162	81	163	82
2002	5	4	141	85	146	89
2003	6	5	171	97	177	102
2004	2	0	163	76	165	76
2005	2	1	158	73	160	74
2006	6	3	120	64	126	67
2007	1	1	121	71	122	72
2008	2	2	144	83	146	85
2009	1	1	120	65	121	66
2010	1	0	100	52	101	52
2011	4	3	84	53	88	56
2012	0	0	103	58	103	58
2013	8	6	103	64	111	70
<b>P-value*</b>	<b>0.45</b>	<b>0.33</b>	<b>&lt;0.001</b>	<b>0.003</b>	<b>&lt;0.001</b>	<b>0.004</b>

\* For trend during 1999–2013. Trends examined using a first-order autoregressive linear regression model.

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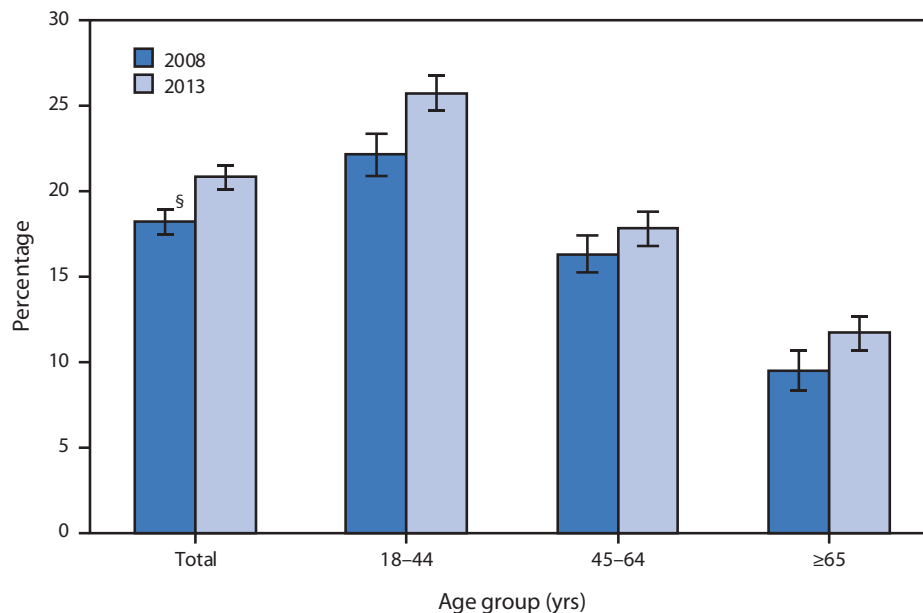
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## QuickStats

FROM THE NATIONAL CENTER FOR HEALTH STATISTICS

### Percentage of Adults Aged $\geq 18$ Years Who Met National Guidelines for Aerobic Activity and Muscle Strengthening,\* by Age Group — National Health Interview Survey, United States, 2008 and 2013<sup>†</sup>



\* Per U.S. Department of Health and Human Services *2008 Physical Activity Guidelines for Americans*. Available at <http://www.health.gov/paguidelines/guidelines/default.aspx>. Respondents defined as meeting both aerobic-activity and muscle-strengthening guidelines reported moderate-intensity physical activity for  $\geq 150$  minutes per week, vigorous-intensity physical activity for  $\geq 75$  minutes per week, or an equivalent combination of moderate- and vigorous-intensity activity, and engaging in physical activities specifically designed to strengthen muscles at least twice per week.

<sup>†</sup> Estimates are based on household interviews of a sample of the civilian, noninstitutionalized U.S. population and are derived from the National Health Interview Survey sample adult component.

<sup>§</sup> 95% confidence interval.

The percentage of adults aged  $\geq 18$  years who met the aerobic-activity and muscle-strengthening guidelines increased from 18.2% in 2008 to 20.8% in 2013. Adults aged 18–44 years were the most likely to meet the aerobic-activity and muscle-strengthening guidelines, and those aged  $\geq 65$  years were the least likely in both 2008 and 2013. For all age groups, the percentage meeting the guidelines increased from 2008 to 2013.

**Source:** CDC. National Health Interview Survey data, 2008 and 2013. Available at <http://www.cdc.gov/nchs/nhis.htm>.

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ISSN: 0149-2195