This study extends the known geographic range of *R. africae* in *A. variegatum* ticks in sub-Saharan Africa. The number of potentially infective ticks recorded in Uganda and Nigeria suggests that persons in rural areas of northern Uganda and central Nigeria might be at risk for African tick-bite fever. Awareness of this rickettsiosis should be raised, particularly among persons who handle cattle (e.g., herders and paraveterinary and veterinary personnel). Physicians in these areas as well as those who care for returning travelers, should consider African tick-bite fever in their differential diagnosis for patients with malaria and influenza-like illnesses.

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References

1. Parola P, Paddock CD, Raoult D. Tick-borne rickettsioses around the world: emerging diseases challenging old concepts. Clin Microbiol Rev. 2005;18:719–56. http://dx.doi.org/10.1128/CMR.18.4.719-756.2005

2. Socolovschi C, Huynh TP, Davoust B, Gomez J, Raoult D, Parola P. Transovarial and trans-stadial transmission of *Rickettsia africae* in *Amblyomma variegatum* ticks. Clin Microbiol Infect. 2009;15(Suppl. 2):317–8. http://dx.doi.org/10.1111/j.1469-0691.2008.02278.x

3. Ogo NI, de Mera IG, Galindo RC, Okubanjo OO, Inuwa HM, Agbede RI, et al. Molecular identification of tick-borne pathogens in Nigerian ticks. Vet Parasitol. 2012;187:572–7. http://dx.doi.org/10.1016/j.vetpar.2012.01.029

4. Reye AL, Arinola OG, Hübschen JM, Muller CP. Pathogen prevalence in ticks collected from the vegetation and livestock in Nigeria. Appl Environ Microbiol. 2012;78:2562–8. http://dx.doi.org/10.1128/AEM.06686-11

5. Walker AR, Bouaoutour A, Camicas JL, Estrada-Peña A, Horak IG, Latif A, et al. Ticks of domestic animals in Africa. A guide to identification of species. Edinburgh (UK): Bioscience Reports; 2003.

6. Regnery RL, Spruill CL, Plikayitsi BD. Genotypic identification of rickettsiae and estimation of intraspecies sequence divergence for portions of two rickettsial genes. J Bacteriol. 1991;173:1576–89.

7. Fournier PE, Roux V, Raoult D. Phylogenetic analysis of spotted fever group rickettsiae by study of the outer surface protein OmpA. Int J Syst Bacteriol. 1998;48:839–49. http://dx.doi.org/10.1099/00207713-48-3-839

8. Alberdi MP, Dalby MJ, Rodriguez-Andres J, Farzakerley JK, Kohl A, Bell-Sakyi L. Detection and identification of putative bacterial endosymbionts and endogenous viruses in tick cell lines. Ticks Tick Borne Dis. 2012;3:137–46. http://dx.doi.org/10.1016/j.ttbdis.2012.05.002

9. Robinson JB, Eremeanova ME, Olsen PE, Thornton SA, Medina MJ, Sumner JW, et al. New approaches to detection and identification of *Rickettsia africae* and *Ehrlichia ruminantium* in *Amblyomma variegatum* (Acari: Ixodidae) ticks from the Caribbean. J Med Entomol. 2009;46:942–51. http://dx.doi.org/10.1603/033.046.0429

10. Kelly PJ, Mason PR, Manning T, Slater S. Role of cattle in the epidemiology of tick-bite fever in Zimbabwe. J Clin Microbiol. 1991;29:256–9.
Jewish community; residency in the London boroughs of Barnet, Hackney, or Haringey; and notification during December 20, 2012–March 19, 2013.

After serologic confirmation of measles in the index case-patient, an unvaccinated Orthodox Jewish 4-year-old from Hackney, the case was reported to the Health Protection Team (HPT) on December 20, 2012. The family could not recall having contact with someone with measles. The child attended nursery while infectious; subsequently, cases in 3 secondary patients in the nursery were reported to the HPT. Transmission was observed within households, extended family groups, nurseries, schools, and a camp for Orthodox Jewish teenagers attended by 80 girls (mainly from Hackney) with staff from Italy. Five secondary cases from this camp were reported (in 3 residents of London, 1 resident of Sheffield, and 1 resident of Hertfordshire).

During December 20, 2012–March 19, 2013, a total of 62 notifications of measles cases meeting the case definition were received in residents of Barnet (8 cases), Hackney (47), and Haringey (7). Patients’ ages ranged from 7 months to 27 years (median 7 years). Thirty-four (55%) were female. Fifty-four (87%) had never received an MMR vaccine, and 8 (13%) had received only 1. Three were admitted to the hospital, and 5 were clinically assessed in accident and emergency departments (patients’ ages ranged from 7 months to 4 years).

All case-patients were assessed for risk by the local HPT for vulnerable contacts and source of infection. The HPT provided infection control guidance and an oral fluid testing kit. Sixteen (26%) case-patients could not recall any contact with a person with measles; the remainder stated various epidemiologic links to a case-patient (Figure).

Forty-two cases have been confirmed (measles IgM detected) by serologic testing (4 cases) or oral fluid (38). One notified case-patient did not have measles IgM on oral fluid testing but had an epidemiologic link to a case-patient and clinical symptoms. They are included in this analysis. Seventeen IgM-positive oral fluid samples were genotyped, and all were D8, currently the most common genotype in the United Kingdom.

One confirmed case was detected in an unvaccinated child from Haringey who was not Orthodox Jewish but was known to have had contact with a case-patient from the community. The child’s illness did not meet the case definition and is not included in this analysis.

In response to the outbreak, active case finding and awareness-raising have been undertaken by the HPT, National Health Service (NHS) public health departments, and community NHS services focused on health and education services and Orthodox Jewish communities. Information letters were sent to the 38 Orthodox Jewish schools and nurseries in Hackney and to attendees of the youth camp. Community NHS vaccination clinics have been maintained to complement standard immunization services offered in general practice surgeries. This includes a Sunday vaccination clinic. Furthermore, community NHS staff provided a vaccination clinic in a secondary school that had an attack rate of 7% (9 cases) at which 9 pupils received 1 MMR vaccine after parental consent. This was the only on-site school vaccination clinic offered; thus,
no comparative uptake data are available to supplement our evaluation of the intervention.

Information relating to the outbreak was placed in 2 Orthodox Jewish newspapers and targeted information for families (in English, Yiddish, and Hebrew) has been disseminated. Finally, all 25 HPTs were alerted to this outbreak and the national Public Health England database (HPZone) has been enhanced to capture notifications from Orthodox Jewish communities.

This ongoing outbreak highlights continued health risks in communities with low vaccination coverage. The outbreak has been largely contained within London’s Orthodox Jewish communities, with limited spread outside of the city and to just 1 local non–Orthodox Jewish child. Given the mobility of members, the risk for transmission outside of London is relatively high. The outbreak underscores the need for ongoing evidence-based and culturally appropriate health interventions that seek to improve vaccination coverage.

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References

1. Lernout T, Kissling E, Hutse V, Schrijver KD, Top G. An outbreak of measles in Orthodox Jewish communities in Antwerp, Belgium, 2007–2008: different reasons for accumulation of susceptibles. Euro Surveill. 2009;14:19087.

2. Stein-Zamir C, Abramson N, Shoob H, Zentner G. An outbreak of measles in an ultra-Orthodox Jewish community in Jerusalem, Israel, 2007—an in-depth report. Euro Surveill. 2008;13:8045.

3. Cohen BJ, McCann R, van den Bosch C, White J. Outbreak of measles in an Orthodox Jewish community. Euro Surveill. 2000;4:1675.

4. Ashmore J, Addiman S, Cordery R, Maguire H. Measles in North East and North Central London, England: a situation report. Euro Surveill. 2007;12:3271.

5. Muscat M. Who gets measles in Europe? J Infect Dis. 2011;204:S353–65. http://dx.doi.org/10.1093/infdis/jir067

6. Mayhew L, Harper G, Waples S, Mayhew Harper Associates. Counting Hackney’s population using administrative data—an analysis of change between 2007 and 2011 [cited 2013 Aug 2]. http://www.hackney.gov.uk/Assets/Documents/estimating-and-profiling-the-population-of-hackney/pdf

7. Public Health England. Quarterly vaccine coverage data tables [cited 2013 Aug 2]. http://www.hpa.org.uk/webc/HPAwebHPAwebStandard/HPAweb_C/1211441442288

8. Henderson L, Millet C, Thorogood N. Perceptions of childhood immunization in a minority community: qualitative study. J R Soc Med. 2008;101:244–51. http://dx.doi.org/10.1258/jrsm.2008.070363

9. Purdy S, Jones KP, Sherratt M, Fallon PV. Demographic characteristics and primary health care utilization patterns of strictly Orthodox Jewish and non-Jewish patients. Fam Pract. 2000;17:233–5.

10. Health Protection Agency. HPA national measles guidelines: local and regional services. 2010 [cited 2013 Aug 2]. http://www.hpa.org.uk/webc/HPAweb&HPAwebStandard/HPAweb_C/1274088429847

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Human Infection with Eurasian Avian-like Influenza A(H1N1) Virus, China

To the Editor: We report a human infection with avian-like swine A(H1N1) influenza virus first identified through a surveillance system for influenza like illness (ILI) in mainland China. An influenza virus, isolated from a patient with ILI, was originally subtyped as influenza A(H1N1)pdm09 virus with a hemagglutination inhibition (HI) test, but it was identified as a Eurasian avian-like influenza A(H1N1) virus (EA-H1N1) by full genome sequencing on January 30, 2013. The virus was named A/Hebei-Yuhua/SWL1250/2012 (H1N1v) (HB/1250/12), according to the definition of the World Health Organization (J).

The case-patient was a 3-year-old boy who had symptoms of fever and sore throat; his highest body temperature was 38°C on December 9, 2012. He was brought for medical treatment to an influenza sentinel hospital in the city of Shijiazhuang in Hebei Province, China, on December 12. He recovered within a week without hospitalization and oseltamivir treatment. A throat swab specimen was collected and sent to the local Chinese Center for Disease Control and Prevention for virus isolation and characterization, according to the Guidelines of the Chinese National Influenza Surveillance Network. A retrospective investigation was conducted to identify the potential infection source and any other possible cases. The case-patient was previously healthy and had no history of close contact with animals (live or dead wild birds, poultry, and swine) within 2 weeks before the onset of symptoms, nor a history of travel. He lived with his sister and parents; all other family members did not