

## References

1. Solomon T, Lewthwaite P, Perera D, Cardoso MJ, McMinn P, Ooi MH. Virology, epidemiology, pathogenesis, and control of enterovirus 71. *Lancet Infect Dis*. 2010;10:778–90. [http://dx.doi.org/10.1016/S1473-3099\(10\)70194-8](http://dx.doi.org/10.1016/S1473-3099(10)70194-8)
2. Bessaud M, Razadindratsimandresy R, Nougairède A, Joffret ML, Deshpande JM, Dubot-Pères A, et al. Molecular comparison and evolutionary analyses of VP1 nucleotide sequences of new African human enterovirus 71 isolates reveal a wide genetic diversity. *PLoS ONE*. 2014;9:e90624. <http://dx.doi.org/10.1371/journal.pone.0090624>
3. van der Sanden S, van der Avoort, Lemey P, Uslu G, Koopmans M. Evolutionary trajectory of the VP1 gene of human enterovirus 71 genogroup B and C viruses. *J Gen Virol*. 2010;91:1949–58. <http://dx.doi.org/10.1099/vir.0.019695-0>
4. Mirand A, Schuffenecker I, Henquell C, Billaud G, Jugie G, Falcon D, et al. Phylogenetic evidence of a recent spread of two populations of human enterovirus 71 in European countries. *J Gen Virol*. 2010;91:2263–77. <http://dx.doi.org/10.1099/vir.0.021741-0>
5. Schuffenecker I, Henquell C, Mirand A, Coste-Burel M, Marque-Juillet S, Desbois D, et al. New introductions of enterovirus 71 subgenogroup C4, France, 2012. *Emerg Infect Dis*. 2014;20:1343–6.
6. Fischer TK, Nielsen AY, Sydenham TV, Andersen PH, Andersen B, Midgley SE. Emergence of enterovirus 71 C4a in Denmark, 2009 to 2013. *Euro Surveill*. 2014;19:20911.
7. Badran SA, Midgley S, Andersen P, Böttlinger B. Clinical and virological features of enterovirus 71 infections in Denmark, 2005 to 2008. *Scand J Infect Dis*. 2011;43:642–8. <http://dx.doi.org/10.3109/00365548.2011.577094>
8. Wu WH, Kuo TC, Lin YT, Huang SW, Liu HF, Wang J, et al. Molecular epidemiology of enterovirus 71 infection in the central region of Taiwan from 2002 to 2012. *PLoS ONE*. 2013;8:e83711. <http://dx.doi.org/10.1371/journal.pone.0083711>
9. Linsuwanon P, Puenja J, Huang SW, Wang YF, Mauleekoonphairoj J, Wang JR, et al. Epidemiology and seroepidemiology of human enterovirus 71 among Thai populations. *J Biomed Sci*. 2014;21:16. <http://dx.doi.org/10.1186/1423-0127-21-16>

Address for correspondence: Audrey Mirand, CHU Clermont-Ferrand, Laboratoire de Virologie, Centre National de Référence des Entérovirus-Parechovirus–Laboratoire Associé, Clermont-Ferrand, France; email: [amirand@chu-clermontferrand.fr](mailto:amirand@chu-clermontferrand.fr)

## Avian Influenza A(H7N9) Virus Antibodies in Close Contacts of Infected Persons, China, 2013–2014

Mai-Juan Ma,<sup>1</sup> Guang-Yuan Ma,<sup>1</sup> Xiao-Xian Yang,<sup>1</sup> Shan-Hui Chen,<sup>1</sup> Gregory C. Gray, Teng Zhao, Jing Bao, Jing-Jing Zhou, Yan-Hua Qian, Bin Lu, Xia Ling,<sup>2</sup> Wu-Chun Cao<sup>2</sup>

Author affiliations: Beijing Institute of Microbiology and Epidemiology, Beijing, China (M.-J. Ma, W.-C. Cao); Wuxi Center for Disease Control and Prevention, Wuxi, China (G.-Y. Ma, S.-H. Chen, J. Bao, Y.-H. Qian, B. Lu, X. Ling); State Key

Laboratory of Pathogen and Biosecurity, Beijing (X.-X. Yang, T. Zhao, J.-J. Zhou); School of Public Health, Central South University, Changsha, China (X.-X. Yang); Duke University Medical Center, Durham, North Carolina, USA (G.C. Gray)

DOI: <http://dx.doi.org/10.3201/eid2104.141442>

**To the Editor:** From early 2013 (1) through November 2014, >460 human cases of laboratory-confirmed avian influenza A(H7N9) virus infection occurred in China. Although human-to-human transmission of subtype H7N9 virus is not common, evidence has been reported of probable transmission among several family clusters (2), between 2 household contacts (3), and between a doctor and an infected patient (4). Taken together, these observations suggest that family members, health care providers, and other close contacts (hereafter called contacts) of H7N9-infected persons may be at risk for infection.

In China, national guidelines regarding H7N9-infected patients call for observation of contacts for 7 days after exposure for signs and symptoms of infection and, if any occur, collection of throat swab specimens for testing by molecular assays (5). The guidelines do not call for serologic testing. Because human avian influenza infections may be mild or asymptomatic, we sought to determine whether serologic testing would show evidence of H7N9 virus infection among contacts of infected persons during the 2013–2014 epidemic in China. Contacts were defined in accordance with China's guidelines for prevention and control of human H7N9 virus infection (5,6). The institutional review board of Wuxi Center for Disease Control and Prevention, Wuxi, Jiangsu Province, China, reviewed and approved this study.

During the epidemic, we recruited contacts of patients in Wuxi and collected throat swab specimens when signs or symptoms of infection developed; serum samples were collected 2–3 weeks later. Swab specimens were tested for H7N9 virus by using real-time reverse transcription PCR (7). Serum samples were tested for antibodies against hemagglutinin antigens of 3 avian influenza viruses (A/Anhui/1/2013 [H7N9], A/Anhui/1/2005 [H5N1]-RG5, and A/chicken/Jiangsu/1/00 [H9N2]) (8) by using a horse erythrocyte hemagglutination inhibition (HI) assay and against the hemagglutinin antigens of 2 seasonal influenza viruses (A/California/07/2009 [H1N1] and A/Victoria/210/2009 [H3N2]) by using a turkey erythrocyte HI assay. Serum samples with HI titers  $\geq 1:40$  against H7N9 virus were confirmed positive by microneutralization assay.

Ten laboratory-confirmed human infections with H7N9 virus occurred in Wuxi during March 29, 2013–May 15, 2014. In total, 225 contacts of 7 H7N9-infected patients

<sup>1</sup>These authors contributed equally to this article.

<sup>2</sup>These senior authors contributed equally to this article.

were enrolled in the study (Table); contacts included 30 family members; 177 health care workers (54 physicians, 119 nurses who provided patient care with standard precautions, 2 hospital attendants, and 2 nurse assistants who provided services related to patient care, safety, and comfort, including anxiety relief, and medical observation); and 18 other contacts (8 friends who visited the patient in the hospital, 2 patients who shared the same room, and 8 patients who shared the same hospital area). The contacts of 3 other H7N9-infected patients declined to participate in the study.

Serologic assay results showed that, 14–28 days after their earliest exposure to an H7N9-infected patient, 22 (9.8%) contacts had elevated HI antibody titers ( $\geq 1:40$ ) against H7N9 virus; titers were 1:40 for 17 contacts and 1:80 for 5 contacts. Positive results for all 22 serum samples were validated by microneutralization assay; 15 (68.2%) samples had microneutralization antibody titers of  $\geq 1:10$  against H7N9 virus antigen (Table). Of the contacts with an HI titer of  $\geq 1:80$  and microneutralization titer of  $\geq 1:40$ , 3 were nurses, 1 was a nurse assistant, and 1 was a family member (a patient's daughter). All 5 of these contacts had antibody titers of  $< 1:40$  to influenza subtype H1N1, H5N1, and H9N2 viruses, and 2 of the nurses had HI antibody

titers of 1:80 against subtype H3N2 virus. All contacts denied having influenza-like respiratory symptoms during the 28 days of follow-up and also denied recent exposure to poultry or pigs or their environments. Of contacts with an HI titer of  $\geq 1:80$  to seasonal H1N1 virus, 3 had titer of 1:80, and 1 each had titer of 1:160 or 1:640. Of the 225 contacts, 108 had HI titers  $\geq 1:80$  against seasonal H3N2 virus (1:80 for 63 contacts, 1:160 for 27 contacts, 1:320 for 9 contacts, and  $\geq 1:640$  for 8 contacts). All contacts had influenza subtype H5N1 and H9N2 antibody titers of  $< 1:80$ .

A previous epidemiologic study (2) reported the medical monitoring of 2,657 contacts of H7N9-infected patients in mainland China and found that, for 28 of the contacts, respiratory symptoms developed within 7 days after monitoring began. Results of molecular assay testing of throat swab specimens for H7N9 virus were negative for all 28 contacts; the study did not include serologic testing. However, small serologic survey studies in Taiwan (9) and household contacts in mainland China (10) showed no evidence of human-to-human transmission among contacts.

A limitation of our study is that we did not collect serum samples from all contacts of infected persons or from controls; therefore, we could not assess the possibility of

**Table.** Demographic characteristics and HI antibody titers against influenza subtype H7N9, H5N1, H9N2, H1N1, and H3N2 viruses among close contacts of avian influenza A(H7N9)-infected persons, China, 2013–2014\*

Characteristics	Close contacts, N = 225		
	Family members, n = 30	Health care workers, n = 177	Others, n = 18
Mean age, y $\pm$ SD	48.03 $\pm$ 17.79	33.71 $\pm$ 7.97	68.50 $\pm$ 14.89
Sex			
F	18 (60.0)	135 (76.3)	4 (22.2)
M	12 (40.0)	42 (23.7)	14 (77.8)
Exposure duration, mean days $\pm$ SD	7.38 $\pm$ 4.70	4.42 $\pm$ 3.67	3 $\pm$ 1.48
Virus subtype and HI titer†			
H7N9			
$< 1:80$	29 (96.7)	173 (97.7)	18 (100.0)
$\geq 1:80$	1 (3.3)	4 (2.3)	0
H5N1			
$< 1:80$	30 (100.0)	177 (100.0)	20 (100.0)
$\geq 1:80$	0	0	0
H9N2			
$< 1:80$	30 (100.0)	177 (100.0)	20 (100.0)
$\geq 1:80$	0	0	0
H1N1			
$< 1:80$	30 (100.0)	172 (97.1)	18 (100.0)
$\geq 1:80$	0	5 (2.9)	0
H3N2			
$< 1:80$	20 (66.7)	89 (50.3)	9 (50.0)
$\geq 1:80$	10 (33.3)	88 (49.7)	9 (50.0)
MN titer, H7N9‡			
$< 1:10$	0	6 (35.3)	1 (100.0)
1:10	0	4 (23.5)	0
1:20	3 (75.0)	3 (17.6)	0
1:40	0	3 (17.6)	0
1:80	1 (25.0)	1 (5.9)	0

\*Data are no. (%) unless otherwise indicated. A comparison of HI titers for control serum samples against reference influenza virus strains used in this study is shown in the online Technical Appendix (<http://wwwnc.cdc.gov/EID/article/21/4/14-1442-Techapp1.pdf>). HI, hemagglutination inhibition; MN, microneutralization assay.

†HI titer cut points were selected conservatively at  $\geq 1:80$  on the basis of World Health Organization recommendations for human infection with influenza A(H5N1) virus (<http://www.who.int/influenza/resources/documents/RecAllabtestsAug07.pdf>).

‡Results for 22 close contacts (17 health care workers, 4 family members, and 1 other close contact) with an HI titer of  $\geq 1:40$ .

false-positive results or asymptomatic infections. However, our findings of elevated levels of subtype H7N9 antibody among 6.7% of contacts during this epidemic in China offer evidence that human-to-human transmission of H7N9 virus may occur among contacts of infected persons.

### Acknowledgments

We thank all subjects for participating in the study, the staff of the Wuxi Center for Disease Control and Prevention for collecting samples, and Yuelong Shu and Tian Bai for providing influenza subtype H7N9 and H5N1 virus samples.

This work was supported by the grants from the Program of International Science and Technology Cooperation (2013DFA30800) of the Ministry of Science and Technology of China; the National Natural Science Foundation of China (no. 81402730); the Key Project of Jiangsu Provincial Department of Health (H201448); and the Major Project of Wuxi Bureau of Health (G201201, Z201404).

### References

- Gao R, Cao B, Hu Y, Feng Z, Wang D, Hu W, et al. Human infection with a novel avian-origin influenza A (H7N9) virus. *N Engl J Med*. 2013;368:1888–97. <http://dx.doi.org/10.1056/NEJMoa1304459>
- Li Q, Zhou L, Zhou M, Chen Z, Li F, Wu H, et al. Epidemiology of human infections with avian influenza A(H7N9) virus in China. *N Engl J Med*. 2014;370:520–32. <http://dx.doi.org/10.1056/NEJMoa1304617>
- Qi X, Qian YH, Bao CJ, Guo XL, Cui LB, Tang FY, et al. Probable person to person transmission of novel avian influenza A (H7N9) virus in Eastern China, 2013: epidemiological investigation. *BMJ*. 2013;347:f4752. <http://dx.doi.org/10.1136/bmj.f4752>
- Shanghai Health and Family Planning Commission of China. A health care worker was infected with H7N9 avian influenza virus [in Chinese] [cited 2014 Jan 20]. <http://www.wsjsw.gov.cn/wsj/n422/n422/u1ai132455.html>
- China National Health and Family Planning Commission. Chinese guideline for prevention and control for human infection with A(H7N9) avian influenza; 2014 ed. [in Chinese] [cited 2014 Jan 27]. <http://www.moh.gov.cn/jkj/s3577/201401/8c1828375a7949cd85454a76bb84f23a.shtml>
- Editorial Team Journal of Thoracic Disease. Guideline on prevention and control of H7N9 avian influenza human infection. *J Thorac Dis*. 2013;Suppl 2:S168–72.
- World Health Organization. Real-time RT-PCR protocol for the detection of avian influenza A(H7N9) virus [cited 2013 Apr 15]. [http://www.who.int/influenza/gisrs\\_laboratory/cnic\\_realtime\\_rt\\_per\\_protocol\\_a\\_h7n9.pdf](http://www.who.int/influenza/gisrs_laboratory/cnic_realtime_rt_per_protocol_a_h7n9.pdf)
- World Health Organization. Serological detection of avian influenza A(H7N9) virus infections by modified horse red blood cells haemagglutination-inhibition assay [cited 2013 Dec 20]. [http://www.who.int/influenza/gisrs\\_laboratory/cnic\\_serological\\_diagnosis\\_hai\\_a\\_h7n9\\_20131220.pdf](http://www.who.int/influenza/gisrs_laboratory/cnic_serological_diagnosis_hai_a_h7n9_20131220.pdf)
- Hsieh SM, Huang YS, Chang SY, Lin PH, Chang SC. Serological survey in close contacts with a confirmed case of H7N9 influenza in Taiwan. *J Infect*. 2013;67:494–5. <http://dx.doi.org/10.1016/j.jinf.2013.08.003>
- Qiu C, Yuan S, Tian D, Yang Y, Zhang A, Chen Q, et al. Epidemiologic report and serologic findings for household contacts of three cases of influenza A (H7N9) virus infection. *J Clin Virol*. 2014;59:129–31. <http://dx.doi.org/10.1016/j.jcv.2013.12.004>

Address for correspondence: Wu-Chun Cao, State Key Laboratory of Pathogen and Biosecurity, Beijing Institute of Microbiology and Epidemiology, 20 Dong-Da St, Fengtai District, Beijing 100071, China; email: caowc@bmi.ac.cn

## Hepatitis E Epidemic, Biratnagar, Nepal, 2014

**Ananta Shrestha, Thupten K. Lama, Sneha Karki, Deepak R. Sigdel, Utsav Rai, Shyam K. Rauniyar, Mamun Al-Mahtab, Kazuaki Takahashi, Masahiro Arai, Sheikh M.F. Akbar, Shunji Mishiro**

Author affiliations: Liver Foundation Nepal, Kathmandu, Nepal (A. Shrestha); Civil Service Hospital, Kathmandu (T.K. Lama, S. Karki); Koshi Zonal Hospital, Biratnagar, Nepal (D.R. Sigdel, U. Rai, S.K. Rauniyar); Bangbandhu Sheikh Mujib Medical University, Dhaka, Bangladesh (M. Al-Mahtab); Toshiba General Hospital, Tokyo, Japan (K. Takahashi, M. Arai, S.M.F. Akbar, S. Mishiro)

DOI: <http://dx.doi.org/10.3201/eid2104.141512>

**To the Editor:** We report a recent epidemic of hepatitis E in Biratnagar, Nepal. During the third week of April 2014, a total of 11 patients with acute jaundice came to hospitals in Biratnagar. IgM against hepatitis E virus (HEV) was detected in serum samples from all 11 patients. During the next 7 weeks, 1,861 patients with acute jaundice came to the outpatient departments of 2 of 5 large hospitals in Biratnagar; 123 patients were admitted to these 2 hospitals.

Registries at these 2 hospitals indicated that 2 patients with acute jaundice came to these hospitals on April 14. On April 28; May 5, 12, 19, and 26; and June 2, 9, 16, 23, and 30, the number of patients with acute jaundice who came to these 2 hospitals were 42, 67, 58, 69, 48, 21, 5, 3, 1, and 0, respectively. Registries showed that this increased frequency of acute jaundice lasted until the end of May 2014, when it began to decrease and reached near zero by the first week of July. In addition, unusually large numbers of patients with acute jaundice came to 25 smaller private health care facilities in Biratnagar during April–May 2014.

The Private and Boarding Schools' Organization of Nepal closed 80 schools in Biratnagar and surrounding areas during the second week of May 2014 because of risk for disease transmission (1). The Biratnagar Zonal Health Authority and National Health Authority of Nepal issued special alerts by mass media regarding jaundice after the third week of April and advised using boiled water for consumption (2).

Registries of major hospitals, smaller health clinics, and private physicians indicated that ≈7,000 patients were