

were not part of the study, none were diagnosed with pandemic (H1N1) 2009 during the study period.

Our study began during the week in September 2009 in which the overall rate of incidence of pandemic (H1N1) 2009 in Spain reached 77.8 cases per 100,000 inhabitants (4), a level that was above the threshold established for the previous influenza season, and ended during the week in which influenza activity fell below this threshold level (5). Therefore, the study spanned the full cycle of the epidemic. The national peak, with an overall rate of incidence of 372.7 cases per 100,000 inhabitants, occurred in week 10 of our study.

This series included 1 asymptomatic carrier. We do not know if that finding could reflect a false-positive test or a low-virulence viral presence.

Notably, among the population of health care workers taking part in the study, only 4 (11%) had been vaccinated against the novel form of the influenza A virus, and none of them had positive PCR results for pandemic (H1N1) 2009 virus. On the other hand, 5 (15%) of workers not vaccinated had a positive PCR result. This finding suggests that, despite the climate of uncertainty concerning the evolution of the influenza outbreak, hospital workers had a greater fear of possible side effects of the vaccine than of the disease itself.

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## Pandemic (H1N1) 2009 and HIV Infection

**To the Editor:** In the United States during spring and fall of 2009, pandemic (H1N1) 2009 influenza A virus resulted in 2 major outbreaks of disease. Initial reports identified immunosuppression, including HIV infection, as a risk factor for the development of severe influenza (1–5). Subsequent reports did not confirm this association, but the number of HIV-infected patients in these studies was small (6,7). We describe the clinical course of pandemic (H1N1) 2009 in HIV-infected persons in a US hospital.

During 2009, 23 cases of laboratory-confirmed pandemic (H1N1) 2009 in HIV-infected persons were identified at Harborview Medical Center (Seattle, WA, USA) by querying the University of Washington HIV Information System (a database that enables complete capture of all HIV testing results at Harborview Medical Center) and by querying the Harborview Infection Control Registry for influenza subtype H1N1 infections. Most cases occurred during October and November. Baseline patient characteristics are noted in the Table. Most patients who sought care had fever and cough; median duration of symptoms before seeking care was 4 days. Overall mortality rate for the entire cohort was 8.7%.

Of the 23 patients, only 2 were not treated for influenza; each had mild signs and symptoms and neither required hospital admission. Each of the remaining 13 outpatients received a 5-day course of treatment with oseltamivir. The 8 patients who required hospitalization received therapy for a median of 6 (range 1–22) days.

Overall mortality rate among HIV-infected patients hospitalized for pandemic (H1N1) 2009 infection was 25% (2 of 8 patients). The 2



Table. Baseline characteristics for HIV-infected patients with pandemic (H1N1) 2009, Seattle, Washington, USA, 2009\*

Patient characteristics	All patients, n = 23	Inpatients, n = 8	Outpatients, n = 15	p value†
<b>Demographics</b>				
Male sex	19 (83)	5 (63)	14 (93)	0.1
Median age, y (range)	43 (22–72)	46 (34–72)	42 (22–64)	0.5
Race/ethnicity				1.0
White	14 (61)	5 (63)	9 (60)	
Black	5 (22)	2 (25)	3 (20)	
Hispanic	4 (17)	0	4 (27)	1.0
Other/refused to answer	4 (17)	1 (13)	3 (20)	
<b>History</b>				
Received 2009–10 seasonal influenza vaccine before illness	14 (61)	6 (75)	8 (53)	0.1
Ever smoked	15 (65)	7 (88)	8 (53)	0.2
<b>HIV-associated factors</b>				
CD4 count, cells/μL, median (range)	308 (32–1,024)	147 (32–1,024)	438 (119–833)	0.06
Undetectable HIV-1 RNA	19 (83)	6 (75)	13 (87)	0.6
Receiving antiretroviral therapy	21 (91)	8 (100)	13 (87)	0.5
<b>Predisposing risk factors</b>				
Prior lung disease	15 (65)	6 (75)	9 (60)	0.5
Prior lung disease	8 (35)	5 (63)	3 (20)	0.07
Cardiovascular disease	4 (17)	3 (38)	1 (6.7)	0.1
Obesity, body mass index >30	4 (17)	1 (13)	3 (20)	1.0
Neutropenia	3 (13)	2 (25)	1 (6.7)	0.3
Receiving immunosuppressive agent	3 (13)	3 (38)	0	0.03
Malignancy	4 (17)	2 (25)	2 (13)	0.6
Diabetes	2 (8.7)	0	2 (13)	0.5
<b>Signs and symptoms</b>				
Fever	18 (78)	7 (88)	11 (73)	0.6
Fatigue	5 (22)	2 (25)	3 (20)	1.0
Malaise	11 (48)	7 (88)	4 (27)	0.009
Myalgia	10 (43)	3 (38)	7 (47)	1.0
Sore throat	5 (22)	2 (25)	3 (20)	1.0
Cough	21 (91)	8 (100)	13 (87)	0.5
Dyspnea	8 (35)	4 (50)	4 (27)	0.4
Nausea/vomiting	10 (43)	2 (25)	8 (53)	0.4
Median duration of symptoms before seeking care, d (range)	4 (0–30)	4.5 (0–10)	3 (1–30)	0.6
<b>Physical examination findings</b>				
Median temperature, °C (range)	38.0 (35.7–40.2)	39.1 (37.3–40.2)	37.7 (35.7–38.5)	0.001
Median heart rate, beats/min (range)	96 (69–129)	109 (80–127)	94 (69–129)	0.1
Mean arterial blood pressure, mm Hg (range)	94 (66–116)	89 (66–98)	97 (75–116)	0.05
Median respiratory rate, breaths/min (range)	20 (14–40)	22 (18–40)	19 (14–36)	0.04
Abnormal lung sounds	11 (48)	8 (100)	3 (23)	0.001
<b>Laboratory findings</b>				
Leukocyte count, cells × 10 <sup>9</sup> /L (range)	4.53 (0.53–10.8)	3.2 (0.53–10.8)	5.4 (2.8–9.4)	0.4
Leukopenia, <5,000 cells/μL	9 (39)	5 (63)	4 (40)	0.6
Chest radiograph findings, new infiltrate	6 (26)	5 (63)	1 (13)	0.1
<b>Care received</b>				
Antiviral treatment for influenza	21 (91)	8 (100)	13 (87)	0.5
Intensive care unit admission	3 (13)	3 (38)	NA	
Mechanical ventilation	2 (8.7)	2 (25)	NA	
Vasopressors	2 (8.7)	2 (25)	NA	
<b>Outcomes</b>				
Secondary pneumonia	3 (13)	3 (38)	0	0.03
Thrombotic complications	1 (4.6)	1 (13)	0	0.4
Died	2 (8.7)	2 (25)	0	0.1

\*Values are no. (%) patients except as indicated. Among the 15 outpatients, 13 had a lung examination documented, 10 had a leukocyte count performed, and 8 had a chest radiograph taken. NA, not applicable.

†For comparison of characteristics between inpatients and outpatients.

inpatients who died had each received  $\geq 14$  days of therapy with oseltamivir. Three inpatients were admitted to the intensive care unit (ICU); of these, 2 had hypoxemic respiratory failure and bilateral infiltrates at the time of admission and a later diagnosis of acute respiratory distress syndrome, and 1 was hospitalized with fever and hemodynamic instability. Each patient with acute respiratory distress syndrome subsequently died; 1 had methicillin-resistant *Staphylococcus aureus* pneumonia at the time of admission, and 1 had severe hypoxemic respiratory failure requiring the use of rescue therapies (e.g., prone positioning and inhaled nitric oxide) and later treatment for ventilator-associated pneumonia. Of the 2 patients who died, 1 had concurrent conditions, including preexisting interstitial lung disease (believed to be associated with crack cocaine use) and a low CD4 cell count of 127 cells/ $\mu\text{L}$ , and 1 had a preserved CD4 cell count  $>1,000$  cells/ $\mu\text{L}$ , but 8 days passed before anti-influenza therapy was started, and thrombotic complications developed before death. The lengths of ICU stay for the patients who died were 13 and 29 days.

Our findings are similar to those reported by others, suggesting that HIV infection alone does not appear to be a risk factor for severe pandemic (H1N1) 2009, provided that patients are not severely immunocompromised, do not have other risk factors associated with poor outcomes, and are treated for influenza soon after signs and symptoms develop (6–9). Most of the 23 patients described here had mild disease and were treated as outpatients. Only 3 required ICU admission, and 2 of these died. Although the mortality rate reported here is higher than that reported in other studies, our sample size was relatively small, and the patients who died had additional risk factors for poor outcomes.

Our study has several limitations. It is a retrospective study, and HIV-infected patients at Harborview Medical Center were not all prospectively tested for pandemic (H1N1) 2009. Most pandemic (H1N1) 2009 virus was detected by reverse transcription PCR of nasal swab specimens; this testing was only available after October 2009, during the second wave of influenza. Infections occurring during the spring were diagnosed by insensitive testing with fluorescent antibody and culture, diagnosed by clinical criteria alone and not included in this analysis, or missed altogether.

Because of differences in pandemic (H1N1) 2009 virus testing, we were unable to compare the incidence of pandemic (H1N1) 2009 virus infection and outcomes between HIV-infected and HIV-uninfected patients. A total of 189 persons received a diagnosis of pandemic (H1N1) 2009 at Harborview Medical Center in 2009, and 79 were hospitalized. A total of 8 (10%) of 79 patients with pandemic (H1N1) 2009 died, including the 2 HIV-infected patients reported here. However, during the peak of the epidemic, many HIV-infected outpatients, who were receiving antiretroviral therapy and had preserved CD4 cell counts, were advised to remain at home if they had mild influenza-like symptoms and were therefore not tested for influenza. This circumstance could have produced a bias toward diagnosing and reporting only more severe disease. Outpatients who had influenza-like symptoms were tested and treated empirically pending test results. Our case series of HIV-infected patients with pandemic (H1N1) 2009 at a single institution in the United States suggests that HIV itself does not appear to be as major a risk factor for severe disease as are other previously reported concurrent conditions, delays in treatment, and development of secondary bacterial pneumonia.

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## Swine Influenza Virus A (H3N2) Infection in Human, Kansas, USA, 2009

**To the Editor:** Triple-reassortant swine influenza viruses (SIVs), which contain genes from human, swine, and avian influenza A viruses, have been enzootic among swine herds in the United States since the late 1990s (1). Although uncommon, occasional transmission of triple-reassortant SIVs from swine to humans has occurred (2–4). Before April 2009, only limited, nonsustained human-to-human transmission of SIVs had been reported (5–7). Although an animal source for pandemic (H1N1) 2009 virus has yet to be identified, the pandemic strain resulted from the reassortment of 2 different lineages of SIV (8).

On July 28, 2009, a 12-year-old Kansas boy sought treatment for fever, cough, and sore throat. Results of an influenza rapid antigen test were positive, and a specimen was sent to the Kansas Department of Health and Environment for further testing. Real-time reverse transcription PCR (rRT-PCR) testing determined the virus contained the surface hemagglutinin (HA) gene of influenza A (H3) and the internal nucleoprotein gene common to all triple-reassortant SIVs (9). The

specimen was sent to the Centers for Disease Control and Prevention (Atlanta, GA, USA) on August 3 and identified as swine-origin influenza virus A (H3N2) by rRT-PCR and sequence analysis.

The patient reported that during July 23–25, 2009, he touched healthy-appearing swine multiple times while attending a county fair. The boy received a standard treatment course of oseltamivir and recovered completely. None of his 3 household contacts attended the fair, and none reported signs or symptoms of illness in the weeks afterward.

The Kansas Department of Health and Environment and the local health department collaborated with the county extension office to identify and interview swine exhibitors at the county fair, focusing on influenza symptoms among exhibitors and household contacts during the week before and after the fair. Twenty-seven (79%) of 34 exhibitors participated in the survey; none reported signs or symptoms of influenza-like illness, defined as fever (temperature  $\geq 100^{\circ}\text{F}$ ) accompanied by either cough or sore throat. Two household contacts of separate exhibitors each reported a low-grade fever ( $< 100^{\circ}\text{F}$ ) and sore throat in the week after the fair. Both touched swine while attending the fair. Both visited a physician's office; neither was tested for influenza; and symptoms of both resolved without treatment. The veterinarian overseeing the swine barn reported no signs of respiratory illness among the swine during the fair. Most swine exhibited were slaughtered at the fair's conclusion.

On August 7, the Kansas Animal Health Department collected nasal swab specimens and blood samples from 13 swine belonging to 7 exhibitors. All samples were delivered to the Kansas State Veterinary Diagnostic Laboratory for analysis by influenza matrix rRT-PCR and virus isolation on nasal swab samples

and hemagglutination inhibition (HI) assays against classical swine influenza virus (H1N1) (A/swine/Iowa/73) and the prototype swine influenza virus (H3N2) (A/swine/Texas/98) on serum samples. In addition, the influenza virus (H3N2) “county fair” isolate, A/Kansas/13/2009 (H3N2), was sent from the Centers for Disease Control and Prevention, amplified, and used to develop a second HI assay that included the original swine serum samples as well as paired convalescent-phase samples from 3 of the swine (Table).

Influenza matrix RT-PCR and virus isolation on nasal swab samples were negative. HI assays demonstrated little or no antibody against the influenza (H1N1) indicator virus and low-level antibody reaction against the prototype swine influenza virus (H3N2). However, HI titers against the “county fair” influenza virus (H3N2) showed consistently elevated titers, which suggested that the animals might have been exposed to the virus 2 weeks earlier, during the time of the fair. The swine may have cleared the virus by the time the nasal swabs were collected, but without positive RT-PCR or virus isolation results, the situation remain inconclusive.

We compared the HA gene segment of A/Kansas/13/2009 (H3N2) with recent animal and human influenza (H3N2) viruses by using the neighbor-joining method, and it clustered with the HA from recent triple-reassortant SIV (H3N2) isolates (online Appendix Figure, [www.cdc.gov/EID/content/17/6/1143-appF.htm](http://www.cdc.gov/EID/content/17/6/1143-appF.htm)) (10). A/Kansas/13/2009 (H3N2) shares  $>97\%$  nucleotide identity with 2 swine viruses reported to have caused human infections, A/Ontario/RV1273/2005 and A/Ontario/1252/2007, and  $>90\%$  nucleotide identity with currently circulating seasonal (H3N2) viruses, such as A/Perth/16/2009. Sequence analysis for the remaining 7 gene segments confirmed A/