

Influenza A(H5N2) Virus Antibodies in Humans after Contact with Infected Poultry, Taiwan, 2012

Ho-Sheng Wu, Ji-Rong Yang, Ming-Tsan Liu,
Chin-Hui Yang, Ming-Chu Cheng,
and Feng-Yee Chang

Six persons in Taiwan who had contact with poultry infected with influenza A(H5N2) showed seroconversion for the virus by hemagglutinin inhibition or microneutralization testing. We developed an ELISA based on nonstructural protein 1 of the virus to differentiate natural infection from cross-reactivity after vaccination; 2 persons also showed seroconversion by this test.

Since 1959, highly pathogenic avian influenza A (HPAI) subtypes H5 and H7 have caused outbreaks in poultry resulting in high mortality rates and have also caused sporadic infections in humans (1–3). Some low pathogenicity avian influenza (LPAI) viruses can mutate to become HPAI virus by acquiring basic amino acid residues in the hemagglutinin (HA) cleavage site after multiple passages in chickens (4). In Taiwan, poultry infected by LPAI (H5N2) virus were reported during 2003–2004 and 2008–2011 (5–7), whereas HPAI (H5N2) viruses were first isolated in 2012 and caused subsequent outbreaks in poultry. Although >20 differences have been found in amino acids in the HA protein between the 2012 HPAI (H5N2) viruses and the 2003 LPAI (H5N2) virus (A/chicken/Taiwan/1209/2003), these viruses are antigenically similar (M.-C. Cheng, unpub. data) and related to those that circulated in Mexico in 1994 but unrelated to the subtype H5N1 viruses that re-emerged in 2003 and the subtype H5N1 vaccine strain (A/Vietnam/1194/2004) (5).

As of December 23, 2013, influenza A(H5N2) virus had not been isolated from humans, but previous studies have provided serologic evidence for subclinical infections

in persons who had frequent contacts with infected animals (3,8–11). We therefore investigated the possibility of infection among persons who were exposed to HPAI (H5N2) virus during outbreaks in chicken farms in Taiwan during January–March 2012.

The Study

For our study, we enrolled 141 persons who had close contact with poultry at 5 chicken farms that had influenza A(H5N2) outbreaks in chickens during January–March 2012. These contacts were 15 farm workers, 90 animal health officials, and 36 temporary employees who participated in culling of infected chickens; no symptoms of influenza-like illness occurred in these persons within 1 week after culling. All 15 poultry workers had been working at their poultry farms for >6 years, and most of the animal health officials had experience in stamping out infected poultry. However, for the 36 temporary employees, previous contact histories with infected chickens were unknown.

Throat swab specimens were collected from all contacts for virus detection within 7 days from the beginning of exposure to the virus, and paired serum samples were collected 21 days apart for serologic testing. Participants were offered an inactivated influenza A(H5N1) vaccine on a voluntary basis on the day the first serum specimens were collected, and vaccination histories within 1 year before the specimen collection date were recorded through oral questionnaires. A total of 102 (72.3%) of the 141 participants were vaccinated: 22 (15.6%) received influenza A(H5N1) only; 39 (27.7%) received seasonal influenza vaccine only; 41 (29.0%) received both vaccines; and 39 (27.7%) received neither (Table 1).

We found all swab specimens were negative for influenza viruses by real-time reverse transcription PCR. However, hemagglutination inhibition (HI) and/or microneutralization (MN) test results showed 7 persons had antibody titers ≥ 80 for subtype H5N2 virus; 6 of these persons showed seroconversion for the virus (Table 2). Elevated antibodies against subtype H3N2 or H5N1 viruses were detected in some of the 6 persons who showed seroconversion (Table 2), which suggests that positive results for subtype H5N2 could be the result of cross-reactive antibodies from previous influenza vaccinations or infections. All 6 persons who showed seroconversion for influenza A(H5N2) virus had received vaccinations for influenza A(H5N1) and seasonal influenza (Table 2). Further, persons who received both influenza vaccinations showed a significant ($p = 0.001$) geometric mean titer increase in HI antibody against influenza A(H5N2) virus in the second samples, whereas those who did not receive both vaccinations did not show a similar increase. This finding indicates these heterologous vaccinations may influence HI antibody titers against influenza A(H5N2) virus.

Author affiliations: Centers for Disease Control, Taipei, Taiwan (H.-S. Wu, J.-R. Yang, M.-T. Liu, C.-H. Yang, F.-Y. Chang); Taipei Medical University, Taipei (H.-S. Wu); Animal Health Research Institute, Taipei (M.-C. Cheng); and National Defence Medical Center, Taipei (F.-Y. Chang)

DOI: <http://dx.doi.org/10.3201/eid2005.131393>

Table 1. HI antibody titers for influenza A(H5N2) virus in paired serum samples of 141 persons who had contact with infected poultry*

HI titers	Total	Second sample						Influenza vaccination history during previous 12 mo			
		<10	10	20	40	80	≥160	A(H5N1) only	Seasonal only	Both	None
First sample											
<10	13	0	6	7	0	0	0	0	4 (30.8)	7 (53.8)	2 (15.4)
10	57	2	4	48	3	0	0	16 (28.1)	5 (8.8)	22 (38.6)	14 (24.5)
20	32	0	5	11	15	1*	0	3 (9.4)	17 (53.1)	3 (9.4)	9 (28.1)
40	39	0	1	10	22	4*	2*	3 (7.7)	13 (33.3)	9 (23.1)	14 (35.9)
80	0	0	0	0	0	0	0	0	0	0	0
≥160	0	0	0	0	0	0	0	0	0	0	0
Total	141	2	16	76	40	5	2	22 (15.6)	39 (27.7)	41 (29.0)	39 (27.7)

*Values are no. persons or no. (%) persons. Serum samples with hemagglutination inhibition (HI) titer ≥80 underwent microneutralization testing. Criteria for determining seroconversion in HI antibodies with regard to influenza A(H5N2) viruses: 1) a ≥4-fold increase in antibody titers in the paired serum samples; and 2) HI titer ≥80 for the second sample.

To investigate whether the influenza A(H5N2) antibodies were elevated as a result of exposure to that virus or because of vaccination with heterologous influenza viruses, we determined antibody levels to influenza A(H5N2) nonstructural protein 1 (NS1) (12–15). The NS1 protein is not readily incorporated into virions used to make inactivated influenza vaccine, so a response to NS1 protein would indicate active influenza A(H5N2) infection. Paired serum samples were analyzed by ELISA plates coated with 2 peptides, NS1^{36–48} (LRRDQKSLRGRGS, NS1-pA) and NS1^{204–225} (RSSNENGGPPLTPKQKREMAR, NS1-pB), synthesized on the basis of the NS1 protein sequences of influenza A(H5N2) virus (NS1-pA, A/chicken/Taiwan/1209/2003) and influenza A(H3N2) virus (NS1-pB, A/Taiwan/4055/2009), respectively. The NS1-pA of the 2012 influenza A(H5N2) outbreak strain has an S48N substitution that is not found in the 2003 strain.

For controls, we simultaneously analyzed 3 groups of paired serum samples with seroconversion (data not shown): 1) samples from 7 ferrets infected with different

influenza virus strains (H1N1, n = 3; H3N2, n = 1; H5N1 [A/Vietnam/1194/2004], n = 1; and A[H1N1]pdm09, n = 2); 2) samples from 8 persons infected with influenza A(H1N1)pdm09 virus; and 3) samples from 9 persons who received vaccinations against influenza A(H5N1) virus. The resulting NS1 antibody responses were plotted (Figure). Five (71.4%) of 7 ferrets showed positive NS1 response against NS1-pA and all against NS1-pB (Figure, panel A), which indicates that influenza virus infection can cause a measurable anti-NS1 response after virus challenge. For influenza virus-infected persons, 3 (37.5%) of 8 showed responses against NS1-pA and NS1-pB (Figure, panel B), but for vaccinated persons, 1 (11.1%) of 9 showed responses against NS1-pA and none against NS1-pB (Figure, panel C). These patterns suggest that anti-NS1 response elicited by natural infection is stronger than that induced by vaccination.

For the group of 7 contacts we identified who had elevated influenza A(H5N2) antibodies, 2 (contacts 1 and 3) had positive NS1 antibody response against both peptides;

Table 2. Serologic test results and vaccination and occupational histories for persons who had high antibody titers against influenza A(H5N2) virus and contact with infected poultry, Taiwan, 2012*†

Cont. no.	Sample date	Tested antigens, by influenza subtype					NS1-ELISA titer		Date and type of influenza vaccination			Occupation
		H5N2	H5N1	H1N1 _{pdm09}	H3N2	HI	NS1-pA	NS1-pB	A(H5N1)	Seasonal	Occupation	
1	Mar 3	40	<20	80	<10	10	1.68	1.4	2012 Mar	NA	Poultry worker	
	Mar 30	80	230	80	<10	10			3, 30			
2	Mar 5	40	160	40	20	80	0.89	0.86	2012 Mar	2011 Oct 8	Poultry worker	
	Mar 28	160	450	80	20	80			6, 28			
3	Mar 5	40	20	80	20	40	1.70	1.34	2012 Mar	2012 Jan	Animal health official	
	Mar 27	80	80	80	20	40			5	20		
4	Mar 5	40	<20	40	10	40	1.13	1.09	2012 Mar	2012 Jan	Animal health official	
	Apr 2	80	80	40	10	40			5, Apr 2	20		
5	Mar 6	40	20	40	<10	320	1.21	1.02	2012 Mar	2011 Nov	Temp. employee	
	Apr 2	80	30	40	<10	160			6, Apr 2	26		
6	Mar 6	20	<20	40	<10	160	1.17	1.08	2012 Mar	2012 Mar	Temp. employee	
	Apr 2	80	160	80	<10	640			6, Apr 2	5		
7	Mar 5	40	<20	40	<10	40	0.98	1.05	2012 Mar	2012 Mar	Temp. employee	
	Apr 2	320	1,280	160	10	640			5, Apr 2	5		

*Cont., contact; HI, hemagglutination inhibition; MN, microneutralization; NS1, nonstructural protein 1; H1N1pdm09, pandemic influenza A(H1N1); NA, not applicable (did not receive vaccination); temp., temporary.

†Boldface indicates a ≥4-fold rise in antibody titer by HI or MN tests and a positive anti-NS1 antibody response by NS1-ELISA test between the first and second samples of paired serum samples. A positive anti-NS1 antibody response by NS1-ELISA assay was defined as each paired serum sample having a test absorbance of the second sample 30% higher than that of its first sample (ratio >1.30).

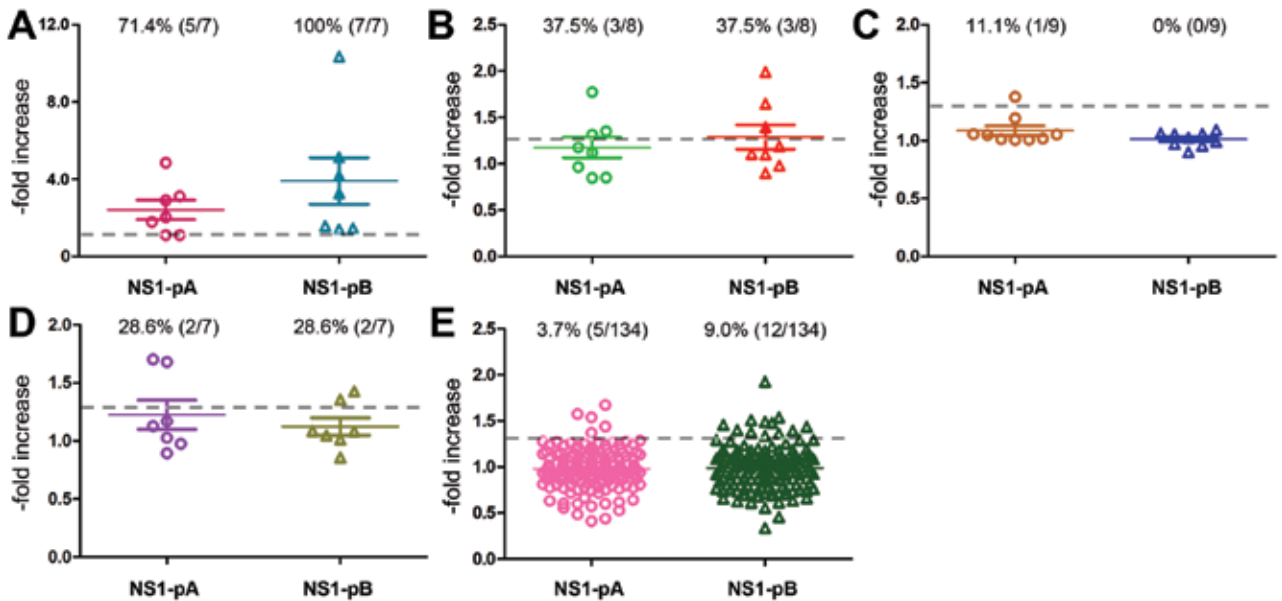


Figure. Antibody responses to 2 influenza A(H5N2) nonstructural protein 1 (NS1) peptides for paired serum samples from: A) influenza virus–infected ferrets; B) influenza virus–infected persons; C) influenza virus–vaccinated persons; D) persons in Taiwan who had contact with infected poultry during January–March 2012 and who showed seroconversion for influenza A(H5N2) virus exposure; and E) persons in Taiwan who had contact with infected poultry during January–March 2012 and who did not show seroconversion. Responses for each group were plotted by -fold increase from first to second sample against NS1-pA (circles) and NS1-pB (triangles); numbers and percentages of positive responses for each sample set are indicated above each plot. Dashed lines indicate cutoff value for defining a positive response: results for the second sample in each pair 30% higher than those for first sample.

the remaining 5 did not (Figure, panel D). These results suggest that contact 1, a poultry worker, and contact 3, an animal health official, may have experienced recent influenza infections.

To better establish the validity of using NS1 to distinguish infected from vaccinated persons, we analyzed paired serum samples for the 134 persons who did not show seroconversion for influenza A(H5N2) virus. Of these, 5 (3.7%) showed positive NS1 antibody response against NS1-pA and 12 (9.0%) against NS1-pB (Figure, panel E). This result suggests that an NS1-ELISA should not be used alone to determine influenza infection but can provide additional data to validate the results of protein-based serologic assays.

Conclusions

In this study, we sampled 141 persons exposed to poultry infected with influenza A(H5N2) virus to assess virus shedding and used multiple serologic assays (including a novel NS1 ELISA) to determine seroconversion status. We found that 6 (4.3%) persons had elevated HA antibodies detected by HI and/or MN assays; a lower percentage (1.4%, 2/141) of subclinical infections was suspected after validation by NS1 antibody assays. The NS1-peptide B was designed on the basis of influenza A(H3N2) virus; however, it also reacted with antibodies

elicited by viruses of different subtypes, which suggests that consensus residues may play an essential role in forming the epitope of NS1 protein.

Our study has limitations. Patient histories of exposure to avian influenza viruses and influenza vaccination were given orally and thus may not be accurate, and mismatching between circulating viruses and antigens used in the study may have occurred. Also, recent seasonal influenza infection may interfere with the determination of subclinical infection with influenza A(H5N2) virus because the NS1 protein is remarkably conserved in type A influenza viruses.

Cross-reactive antibodies in humans elicited from heterologous influenza viruses can complicate serologic, HA-based identification of influenza subtype. The NS1-ELISA method we describe may help determine the type more readily and improve diagnosis of subclinical infection in humans. Further, our findings indicate that occupational exposure to infected poultry may pose a risk for infection in humans.

Acknowledgments

We thank Ruben Donis and Masato Tashiro for their valuable suggestions and critical appraisal of the manuscript and the many medical officers and colleagues in Regional Centers of the Taiwan CDC who participated in sample collection and onsite investigation.

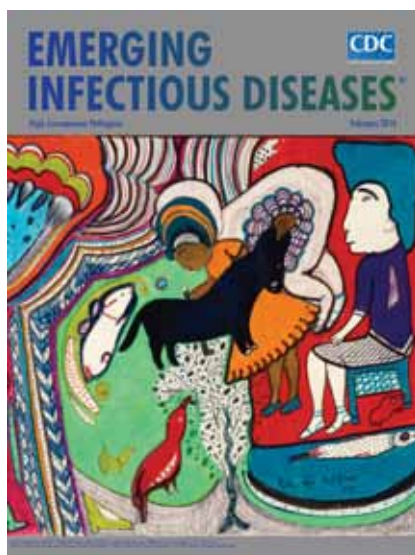
This study received partial financial support from Department of Health, Taiwan (DOH101-DC-2013).

Dr Wu is director of the Taiwan National Influenza Center and of the Center for Research, Diagnostics and Vaccine Development at Taiwan CDC. His research interests are influenza viruses and enteroviruses.

References

- Kalthoff D, Globig A, Beer M. (Highly pathogenic) avian influenza as a zoonotic agent. *Vet Microbiol.* 2010;140:237–45. <http://dx.doi.org/10.1016/j.vetmic.2009.08.022>
- Bean WJ, Kawaoka Y, Wood JM, Pearson JE, Webster RG. Characterization of virulent and avirulent A/chicken/Pennsylvania/83 influenza A viruses: potential role of defective interfering RNAs in nature. *J Virol.* 1985;54:151–60.
- Thompson PN, Sinclair M, Ganzevoort B. Risk factors for seropositivity to H5 avian influenza virus in ostrich farms in the Western Cape Province, South Africa. *Prev Vet Med.* 2008;86:139–52. <http://dx.doi.org/10.1016/j.prevetmed.2008.03.011>
- Soda K, Cheng MC, Yoshida H, Endo M, Lee SH, Okamoto M, et al. A low pathogenic H5N2 influenza virus isolated in Taiwan acquired high pathogenicity by consecutive passages in chickens. *J Vet Med Sci.* 2011;73:767–72. <http://dx.doi.org/10.1292/jvms.10-0532>
- Cheng MC, Soda K, Lee MS, Lee SH, Sakoda Y, Kida H, et al. Isolation and characterization of potentially pathogenic H5N2 influenza virus from a chicken in Taiwan in 2008. *Avian Dis.* 2010;54:885–93. <http://dx.doi.org/10.1637/9208-120609-Reg.1>
- Avian influenza (72): Taiwan LPAI H5N2. *ProMed.* 2011 Nov 18 [cited 2013 Sep 10]. <http://www.promedmail.org>, archive no. 20111122.3426.
- Huang SY, Chen CH, Hu YJ, Shih YL, Chang CC, Chao YN, et al. Overview on prevention and control measures for human infection with HPAI H5N2 virus in Taiwan in 2012. *Taiwan Epidemiol. Bull.* 2013;29:202–23.
- Ogata T, Yamazaki Y, Okabe N, Nakamura Y, Tashiro M, Nagata N, et al. Human H5N2 avian influenza infection in Japan and the factors associated with high H5N2-neutralizing antibody titer. *J Epidemiol.* 2008;18:160–6. <http://dx.doi.org/10.2188/jea.JE2007446>
- Yamazaki Y, Doy M, Okabe N, Yasui Y, Nakashima K, Fujieda T, et al. Serological survey of avian H5N2-subtype influenza virus infections in human populations. *Arch Virol.* 2009;154:421–7. <http://dx.doi.org/10.1007/s00705-009-0319-7>
- Kayali G, Ortiz EJ, Chorazy ML, Gray GC. Evidence of previous avian influenza infection among US turkey workers. *Zoonoses Public Health.* 2010;57:265–72. <http://dx.doi.org/10.1111/j.1863-2378.2009.01231.x>
- Shafir SC, Fuller T, Smith TB, Rimoin AW. A national study of individuals who handle migratory birds for evidence of avian and swine-origin influenza virus infections. *J Clin Virol.* 2012;54:364–7. <http://dx.doi.org/10.1016/j.jcv.2012.05.001>
- Ozaki H, Sugiura T, Sugita S, Imagawa H, Kida H. Detection of antibodies to the nonstructural protein (NS1) of influenza A virus allows distinction between vaccinated and infected horses. *Vet Microbiol.* 2001;82:111–9. [http://dx.doi.org/10.1016/S0378-1135\(01\)00366-2](http://dx.doi.org/10.1016/S0378-1135(01)00366-2)
- Tumpey TM, Alvarez R, Swayne DE, Suarez DL. Diagnostic approach for differentiating infected from vaccinated poultry on the basis of antibodies to NS1, the nonstructural protein of influenza A virus. *J Clin Microbiol.* 2005;43:676–83. <http://dx.doi.org/10.1128/JCM.43.2.676-683.2005>
- Avellaneda G, Mundt E, Lee CW, Jadhao S, Suarez DL. Differentiation of infected and vaccinated animals (DIVA) using the NS1 protein of avian influenza virus. *Avian Dis.* 2010;54(Suppl):278–86. <http://dx.doi.org/10.1637/8644-020409-Reg.1>
- Takeyama N, Minari K, Kajihara M, Isoda N, Sakamoto R, Sasaki T, et al. Detection of highly pathogenic avian influenza virus infection in vaccinated chicken flocks by monitoring antibodies against non-structural protein 1 (NS1). *Vet Microbiol.* 2011;147:283–91. <http://dx.doi.org/10.1016/j.vetmic.2010.07.002>

Address for correspondence: Feng-Yee Chang, Centers for Disease Control, No. 6, Linsen S. Rd, Taipei, 10050, Taiwan; email: fychang@cdc.gov.tw



February 2014: High-Consequence Pathogens

Including:

- Poxvirus Viability and Signatures in Historical Relics
- Novel Paramyxovirus Associated with Severe Acute Febrile Disease, South Sudan and Uganda, 2012
- Subtyping *Cryptosporidium ubiquitum*, a Zoonotic Pathogen Emerging in Humans
- Genomic Variability of Monkeypox Virus among Humans, Democratic Republic of the Congo

<http://wwwnc.cdc.gov/eid/content/20/2/contents.htm>

Influenza A(H5N2) Virus Antibodies in Humans after Contact with Infected Poultry, Taiwan, 2012

Technical Appendix

Materials

Viruses

Influenza A virus strains of different subtypes, including A/Chicken/Taiwan/1209/2003 (H5N2), A/Vietnam/1194/2004 (RG14, H5N1), A/California/07/2009 (H1N1pdm09) and A/Taiwan/4055/2009 (H3N2) were used for HI tests as antigens to investigate the existence of specific antibodies and the 2003 H5N2 isolate was also used for MN tests.

Vaccine

An inactivated H5N1 influenza vaccine (Aflunov®, Novartis Vaccines and Diagnostics, Siena, Italy) was used for the study subjects on a voluntary basis. The vaccine contains purified HA and NA surface antigens from influenza A/Vietnam/1194/2004 (H5N1) vaccine strain. Current circulating Asian H5N1 viruses have evolved into various clades with antigenic differences and are antigenically different from the vaccine strain.

Methods

Real-time RT-PCR

RNA was extracted from throat swab specimens by MagNa Pure LC extraction system (Roche). The extracted RNA was tested for influenza viruses by real-time RT-PCR (1). Briefly, this assay includes primers and probe sets to detect the M genes from all influenza A viruses and the HA genes of H, H3, and H5 subtypes

Hemagglutination inhibition (HI) test

Human sera were treated with receptor destroying enzyme (Denka Seiken, Japan) to remove non-specific hemagglutinins. Final concentrations of 1% horse (for H5N1 and H5N2) and 0.75% guinea pig erythrocytes (for H1N1pdm09 and H3N2) were used for HI tests (2). HI titers were expressed as the reciprocal of the highest dilution of serum that inhibited virus-induced hemagglutination.

Microneutralization (MN) test

MDCK cells were seeded in 96-well plates at the concentration of 1.5×10^4 cells/well. Two-fold serially diluted serum specimens were mixed with an equal volume of virus inoculums (100 TCID₅₀), followed by 1 hour of incubation at 37°C. After incubation, the mixture was added to confluent MDCK monolayers. Cells were cultured for 72 hours before the examination of cytopathic effect (CPE). The absence of CPE in individual wells was defined as protection. The assay was performed in quadruplicate.

NS1-ELISA test

An ELISA test for nonstructural protein 1 (NS1) antibody detection was developed in the study. Peptides of NS1³⁶⁻⁴⁸ (LRRDQKSLRGRGS, designated as peptide A), NS1¹⁶¹⁻¹⁷⁵

(SPLPSLPGHTDEDVK, peptide B) were used in the test. The peptide A was synthesized according to the previous study (3) with modifications to match that of the 2003 H5N2 virus and the peptide B was newly designed in this study and predicted by using Antibody Epitope Prediction in IEDB Analysis Resource (http://tools.immuneepitope.org/tools/bcell/iedb_input). ELISA plates were coated overnight at 4°C with each peptide (1 µg/well) diluted in carbonate-bicarbonate buffer (pH 9.6), followed by blocking with PBSB buffer (PBS with 1% bovine serum albumin). Serum specimens were heat inactivated at 56 °C for 30 minutes and then diluted 1:100 with PBSTB buffer (PBST with 1% bovine serum albumin). Serum antibodies bound to the coated peptides were detected by using horseradish peroxidase-labeled goat anti-human IgG antibodies (KPL, USA). After adding the TMB/E substrate (Millipore, USA) for 15-30 minutes, sulfuric acid was added to stop the reaction. The absorbance at 450 nm was measured and analyzed. Wells that coated with 1% BSA were used for controls. The normalized absorbance measurement of each serum specimen was calculated by using OD_{450nm} value of the peptide-coated well minus that of BSA-coated well. For each paired serum, if the value of the second sample was 30% higher than that of its first sample, it was defined as a positive anti-NS1 antibody response.

References

1. Yang JR, Lo J, Liu JL, Lin CH, Ho YL, Chen CJ, et al. Rapid SYBR green I and modified probe real-time reverse transcription-PCR assays identify influenza H1N1 viruses and distinguish between pandemic and seasonal strains. *J Clin Microbiol.* 2009;47:3714–6. [PubMed](#)
<http://dx.doi.org/10.1128/JCM.01646-09>
2. World Health Organization. Manual for the laboratory diagnosis and virological surveillance of influenza [cited 2013 Feb 8]. http://whqlibdoc.who.int/publications/2011/9789241548090_eng.pdf

3. Tumpey TM, Alvarez R, Swayne DE, Suarez DL. Diagnostic approach for differentiating infected from vaccinated poultry on the basis of antibodies to NS1, the nonstructural protein of influenza A virus. *J Clin Microbiol.* 2005;43:676–83. [PubMed http://dx.doi.org/10.1128/JCM.43.2.676-683.2005](http://dx.doi.org/10.1128/JCM.43.2.676-683.2005)