

Central Nervous System Involvement by Novel Clade 2.3.2.1e H5N1 Avian Influenza Virus in a Pediatric Patient

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Novel clade 2.3.2.1e A(H5N1) virus was detected in cerebrospinal fluid but not in respiratory, rectal swab, or blood samples of an 8-year-old boy presenting with meningoencephalitis without respiratory symptoms. Cerebrospinal fluid A(H5N1) hemagglutinin-specific antibody levels were higher than those of sera. Clinicians should be aware of emerging clade 2.3.2.1e A(H5N1)-associated meningoencephalitis.

Keywords. H5N1; highly pathogenic avian influenza; meningoencephalitis; Vietnam.

Highly pathogenic avian influenza (HPAI) A(H5N1) virus is a public health threat with pandemic potential. H5N1 infection in humans can result in severe respiratory disease but is rarely accompanied by central nervous system (CNS) involvement. Herein, we report on HPAI A(H5N1)-associated meningoencephalitis in the absence of respiratory symptoms in an 8-year-old boy.

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PATIENT PRESENTATION AND INVESTIGATIONS

In mid-April 2025, a previously healthy 8-year-old boy from Tay Ninh province in Vietnam, bordering Cambodia, was admitted to a local hospital with a 24-hour history of fever, severe headache, and vomiting, without any respiratory symptoms. Routine blood tests showed leukocytosis (23 700 cells/μL) and mildly elevated platelet level (456 000 platelets/μL) (Supplementary Table 1). Rapid NS1 test for dengue virus was negative. He was diagnosed with sepsis/meningoencephalitis. Two days later he was transferred to the Children's Hospital 1 (CH1), a tertiary referral hospital in Ho Chi Minh City. At CH1, he presented with fever (38.1°C) and neck stiffness with altered consciousness, but without apparent respiratory illness. His chest radiograph showed consolidation, corresponding to the interpretation of left lower lobe findings (Supplementary Figure 1). Chest computed tomography was not performed. However, he had no respiratory symptoms and had normal heart and respiratory rates (95 beats per minute and 22 breaths per minute, respectively), normal oxygen saturation on room air (95%), and normal auscultation of the lungs. The admission diagnosis was suspected meningoencephalitis. Empiric intravenous ceftriaxone, vancomycin, and acyclovir were started on the basis of clinical presentations.

Cerebrospinal fluid (CSF) collected on admission (day 3 of illness) showed pleocytosis (486 cells/μL) with neutrophil predominance, elevated lactate and protein concentrations, and hypoglycorrhachia (Table 1). Blood hematologic indices were within normal ranges (Supplementary Table 1). Brain magnetic resonance imaging (MRI) revealed dilated lateral ventricles (Figure 1A). Routine bacterial culture of admission CSF, urine, blood, and endotracheal aspirate samples was negative, but analysis of the admission CSF, using a multiplex real-time reverse-transcription polymerase chain reaction (PCR) platform targeting >60 pathogens [1], revealed influenza A virus (IAV) with a cycle threshold (Ct) value of 19. Subsequent confirmatory IAV and subtyping PCR testing of a second CSF sample collected on day 6 of illness onset using US Centers for Disease Control and Prevention assays returned IAV (Ct = 26) and A/H5 (Ct = 34, Supplementary Tables 2 and 3). PCR testing for IAV in urine, throat swabs, rectal swabs, endotracheal aspirate, and blood samples was all negative, while serial CSF samples collected until day 10 of hospitalization were all positive (Supplementary Table 3). Details about the commercially available diagnostic assays and the targeted pathogens are presented in the Supplementary Appendix and the Supplementary Table 3 footnote.

After A/H5 PCR results became available, epidemiological investigations were initiated and revealed that his family owned

Table 1. Laboratory Findings of Admission and Follow-up Cerebrospinal Fluid Samples

Laboratory Examination	Normal Range	CSF1 (Admission), Illness Day 3	CSF2, Illness Day 6	CSF3, Illness Day 8	CSF4, Illness Day 12	CSF5, Illness Day 18
Leukocyte count, cells/ μ L	≤ 5	486	418	550	118	160
Neutrophils, %	0	77	85	78	83	57
CSF glucose concentration, mmol/L	2.8–4.4	1.1	2.1	1.8	2.9	1.9
CSF/plasma glucose ratio, %	≥ 0.6	0.16	0.23	0.3	0.57	0.3
Lactate, mmol/L	1–2	10.4	8.7	4.9	2.73	3.29
Protein, g/L	< 0.4	3.6	11.7	6.2	2.7	3.06

Abbreviation: CSF, cerebrospinal fluid.

many young fighting cocks, which the patient treated as pets, with frequent close contact. Around 2 weeks before his illness, some sporadically died of unknown causes. The patient, however, did not have any respiratory symptoms within the 2 weeks preceding his present illness. The neighbors also reported that groups of 40–50 chickens died of unknown reasons at around the same time. No poultry samples were available for testing.

Mechanical ventilation was initiated 15 hours after admission to CH1 because of worsening coma and suspected cerebral edema. Subsequently, the patient's condition deteriorated, progressing to a deep coma with no response to stimuli and unstable hemodynamic observations, requiring vasopressor and inotropic support. Intravenous mannitol and hypertonic saline were administered to reduce elevated intracranial pressure. Based on the PCR results, MRI findings, and clinical features, the patient was diagnosed with meningoencephalitis due to IAV A(H5N1) virus infection and isolated in accordance with local public health measures. Acyclovir treatment was discontinued, and oseltamivir was commenced on day 3 of hospitalization, followed by intravenous immunoglobulin. Vancomycin was changed to linezolid, and levofloxacin was added on the basis of recurrent fever and elevated procalcitonin level (Table 1). Durations of antiviral and antibiotic administration are detailed in Supplementary Table 3. The patient recovered and was extubated on day 8 of hospitalization. Follow-up CSF remained abnormal until day 18 of illness (Table 1), when IAV RNA in the CSF was undetectable (Supplementary Table 4). As part of our routine care, obtaining clinical samples (especially CSF) for routine diagnosis was verbally agreed by the parents, and was based on the basis of clinical progression and local public health measures requiring that the patient tested negative before discharge. He was discharged with full recovery after 19 days of hospitalization.

WHOLE GENOME SEQUENCING AND SEQUENCE ANALYSIS

Direct sequencing of the second CSF sample recovered all 8 gene segments of the IAV genome. Laboratory workflow is detailed in the Supplementary Appendix. Phylogenetic analysis assigned the hemagglutinin (HA) gene segment to clade

2.3.2.1e (Figure 1B, Supplementary Figures 2 and 3, Supplementary Table 5), previously known as clade 2.3.2.1c of HPAI A(H5N1) viruses that circulate endemically in Southeast Asia. More specifically, the obtained sequences belonged to a novel reassortant clade 2.3.2.1e of HPAI A(H5N1) with gene segments coming from both clade 2.3.2.1c and clade 2.3.4.4b viruses [2] that emerged in late 2023, causing outbreaks in poultry and zoonotic infections in mammals, including 14 confirmed human cases (6 deaths) in Cambodia and 1 human case (fatal) in Vietnam. Likewise, all remaining segments were closely related to the corresponding segments of clade 2.3.2.1e reassortant genotype viruses (Supplementary Figure 3). Further in-depth phylogenetic and phylogeographic analysis is beyond the scope of this study and was hindered by a lack of contemporary sequences, especially from poultry in Vietnam. However, the placement of gene segments on the corresponding phylogenetic trees suggested that the viral strain is closely related to sequences recovered from recent human cases and poultry samples in Vietnam and Cambodia [2]. Several amino acid substitutions associated with the host specificity shift and mammalian adaptation were observed (Supplementary Table 6). Of these, 2 substitutions in the HA sequences, S123P and R167K, associated with increased binding of the virus to $\alpha 2,6$ receptors, were unique to the virus of the present study (Supplementary Table 6). Otherwise, the remaining mutations have previously been reported.

Using microsphere immunoassay, immunoglobulin G antibodies against H5 A/Cambodia/i0125001G/2024 HA1 subunit were detectable in plasma and CSF samples collected at day 11–12 of illness, with CSF antibody levels higher than those of plasma (Figure 1C). Specifically, the mean fluorescence intensity value increased from a borderline level to 2542 and 29 286 in the CSF as compared to from an undetectable level to 1363 and 8723 in the plasma. Anti-H5 hemagglutinin antibody was not detectable in negative control CSF and serum samples (Figure 1C).

For both parents, their throat swabs and plasma samples collected at enrollment were negative for IAV by PCR analysis. Likewise, their plasma samples were also negative for antibodies against the HA1 subunit (data not shown).

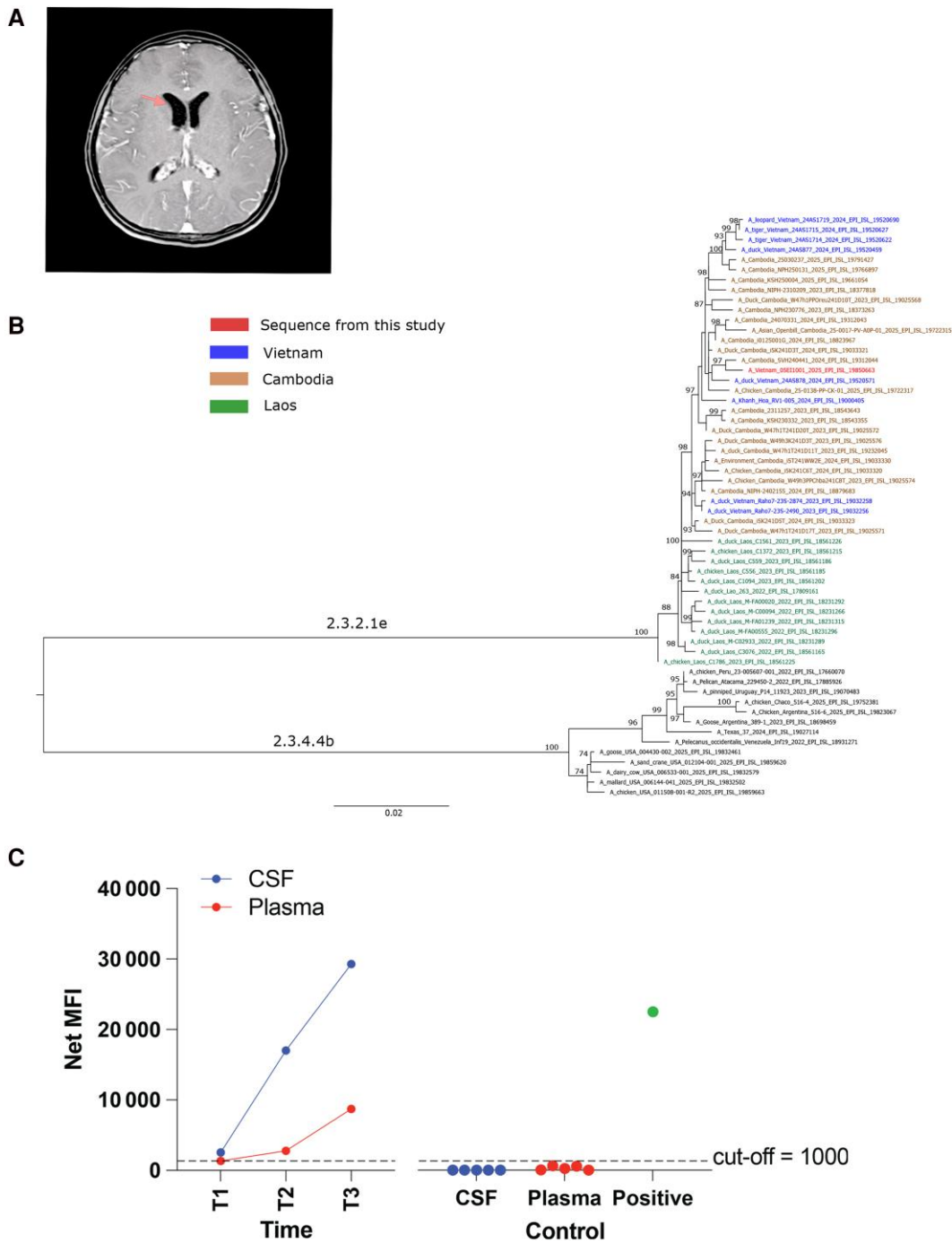


Figure 1. Results of brain image and laboratory investigations. *A*, T2 fluid-attenuated inversion recovery sequence after contrast administration of brain magnetic resonance imaging obtained on illness day 6 showing dilated lateral ventricles with a dimension of 14 mm, marked by red arrow (normal range: ≤ 10 mm). *B*, Reconstructed maximum likelihood (ML) trees of hemagglutinin-segment sequences obtained from this study and representatives of the corresponding gene segment sequences of clades 2.3.2.1e and 2.3.4.4b. *C*, Antibodies against HA1 subunit in serial CSF and plasma samples collected over the course of illness. In (*C*), the dashed line indicates assay cut-off. Negative control samples include 5 CSF samples from patients with central nervous system infection (bacterial meningitis due to *Enterococcus faecium* [$n = 1$], tuberculous meningitis [$n = 2$], cryptococcal meningitis [$n = 1$], and cerebral tumor [$n = 1$]), and 5 plasma samples from healthcare workers participating in a COVID-19 vaccine evaluation study (Supplementary Appendix). Positive control was derived from plasma sample of a patient with polymerase chain reaction–confirmed H5N1 infection. T1, T2, and T3 for CSF are samples collected on illness day 6, 12, and 18, respectively. T1, T2, and T3 for plasma are samples collected on illness day 6, 11, and 21, respectively. Abbreviations: CSF, cerebrospinal fluid; MFI, mean fluorescence intensity.

DISCUSSION

Influenza A(H5N1)-associated CNS infection in humans has rarely been reported but typically presents as a complication, following respiratory symptoms [3–5]. Notably, our patient presented with meningoencephalitis in the absence of respiratory symptoms. Additionally, unlike the previously reported patients, who had viral RNA detected in both CSF and non-CSF samples [3–5], our patient only had viral RNA detected in serial CSF samples in the absence of viral RNA detected in urine, blood, rectal swab, and respiratory samples. Low respiratory tract viral loads, transient viral replication in the respiratory tract, and/or delayed sample collection (illness day 6 onward) might explain the negative PCR findings in non-CSF samples, including the endotracheal aspirate sample. Notably, HPAI A(H5N1) viruses can infect human respiratory tissues by binding to receptors bearing sialic acids linked to galactose by α 2,3-linkages, which are found in the lungs and lower respiratory tract, supported by the chest radiograph findings suggestive of lower left lung pneumonia.

Intrathecal antibody production following seasonal influenza virus infection has been reported previously [6, 7]. Likewise, we showed that the titers of antibodies against A(H5N1) HA were higher in the CSF than in the plasma. This suggested that CSF antibodies were likely intrathecally produced as a consequence of viral invasion of the CNS, which can occur via hematogenous pathway by passing the blood–brain barrier, or via cranial nerves, especially the olfactory nerve route without viremia [3, 8, 9]. Based on our collective findings, it is likely that the A(H5N1) virus from poultry entered the CNS without establishing a significant infection phase in epithelial cells of the nasal cavity [10]. Therefore, future study should assess the mucosal immune response to A(H5N1) to further shed light on the disease pathogenesis.

The CSF findings of our patient showed neutrophil predominance and hypoglycorrhachia, which were more compatible with bacterial meningitis but inconsistent with findings from previous reports about human cases of A(H5N1) virus-associated CNS infection [4, 5, 11–15]. Head and sinus computed tomography, however, was not performed. Therefore, although routine culture and PCR were negative for common bacterial causes, we cannot exclude other possibilities (eg, parameningeal infections).

Mammalian-adapted mutations have been documented in some clade 2.3.4.4b viruses causing outbreaks in cows in the United States [16]. Likewise, we documented 2 substitutions (S123P and R167K) in the HA sequences associated with increased virus binding to α 2,6 receptors that are unique to the virus of the present study. These data emphasize the increasing risk of A(H5N1) virus adapting to mammals and becoming more neurologically virulent and more transmissible. However, molecular and serological testing of 180 households

of human cases in the United States was negative [16], demonstrating the absence of human-to-human transmission, supporting our findings.

Our study has some limitations. First, the microsphere immunoassay used in this study is an investigational assay. Therefore, the detection of CSF antibodies should be further validated using gold standard tests such as hemagglutination inhibition or live-virus neutralization assay. Second, epidemiological investigations were only carried out after HPAI A(H5N1) PCR results became available, which resulted in a delayed oseltamivir administration. Clinicians in endemic regions should be aware of meningoencephalitis associated with A(H5N1) infection.

In summary, we report on an HPAI A(H5N1) infection in a child presenting with meningoencephalitis in the absence of respiratory symptoms. Viral RNA was detected in CSF but not in respiratory, rectal swab, or blood samples. Testing for IAV and A(H5N1) virus should be considered in patients presenting with CNS infection with a history of exposure (eg, dead poultry). Clinicians should be aware of meningoencephalitis associated with A(H5N1) infection in the absence of respiratory symptoms.

Supplementary Data

Supplementary materials are available at [Open Forum Infectious Diseases](#) online. Consisting of data provided by the authors to benefit the reader, the posted materials are not copyedited and are the sole responsibility of the authors, so questions or comments should be addressed to the corresponding author.

Notes

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Patient consent. The patient and both his parents consented to participate in an ongoing investigation ([Supplementary Appendix](#)).

Data availability. The obtained viral sequence was submitted to GISAID under the ID number EPI_ISL_19850663. Clinical and laboratory data are detailed in the Supplementary Tables (where appropriate).

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Potential conflicts of interest. The authors: No reported conflicts of interest.

APPENDIX

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References

1. Phung NTN, Pham HT, Tran TT, et al. *Naegleria fowleri*: portrait of a cerebral killer. *Diagnostics (Basel)* **2025**; 15:89.
2. Siegers JY, Xie R, Edwards KM, et al. Resurgence of zoonotic highly pathogenic avian influenza A(H5N1) virus in Cambodia. *N Engl J Med* **2025**; 393:1650–2.
3. Bauer L, Benavides FFW, Veldhuis Kroeze EJB, de Wit E, van Riel D. The neuro-pathogenesis of highly pathogenic avian influenza H5Nx viruses in mammalian species including humans. *Trends Neurosci* **2023**; 46:953–70.
4. de Jong MD, Cam BV, Qui PT, et al. Fatal avian influenza A (H5N1) in a child presenting with diarrhea followed by coma. *N Engl J Med* **2005**; 352:686–91.
5. Mak GCK, Kwan MY-W, Mok CKP, Lo JYC, Peiris M, Leung CW. Influenza A(H5N1) virus infection in a child with encephalitis complicated by obstructive hydrocephalus. *Clin Infect Dis* **2018**; 66:136–9.
6. Salonen O, Koskineniemi M, Saari A, et al. Myelitis associated with influenza A virus infection. *J Neurovirol* **1997**; 3:83–5.
7. Fujimoto Y, Shibata M, Tsuyuki M, et al. Influenza A virus encephalopathy with symmetrical thalamic lesions. *Eur J Pediatr* **2000**; 159:319–21.
8. Xerra F, Cafarella G, Ferrante F, et al. Neurological manifestations of influenza virus and RSV infections in children. *Curr Respir Med Rev* **2025**; 21:8–19.
9. Bin N-R, Prescott SL, Horio N, Wang Y, Chiu IM, Liberles SD. An airway-to-brain sensory pathway mediates influenza-induced sickness. *Nature* **2023**; 615:660–7.
10. Siegers JY, van de Bildt MWG, Lin Z, et al. Viral factors important for efficient replication of influenza A viruses in cells of the central nervous system. *J Virol* **2019**; 93:e02273–18.
11. Zhang L, Liu K, Su Q, et al. Clinical features of the first critical case of acute encephalitis caused by the avian influenza A (H5N6) virus. *Emerg Microbes Infect* **2022**; 11:2437–46.
12. Chokephaibulkit K, Uiprasertkul M, Puthavathana P, et al. A child with avian influenza A (H5N1) infection. *Pediatr Infect Dis J* **2005**; 24:162–6.
13. Gao R, Dong L, Dong J, et al. A systematic molecular pathology study of a laboratory confirmed H5N1 human case. *PLoS One* **2010**; 5:e13315.
14. Gu J, Xie Z, Gao Z, et al. H5n1 infection of the respiratory tract and beyond: a molecular pathology study. *Lancet* **2007**; 370:1137–45.
15. Rajabali N, Lim T, Sokolowski C, Prevost JD, Lee EZ. Avian influenza A (H5N1) infection with respiratory failure and meningoencephalitis in a Canadian traveller. *Can J Infect Dis Med Microbiol* **2015**; 26:221–3.
16. Rolfes MA, Kniss K, Kirby MK, et al. Human infections with highly pathogenic avian influenza A(H5N1) viruses in the United States from March 2024 to May 2025. *Nat Med* **2025**; 31:3889–98.