Follow-up report No.1 (Final report)


Report Summary

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Epidemiology

The sample, from a wild mallard duck, was collected on 27 December 2016 as part of the hunter – harvested wild bird avian influenza surveillance program. Genome sequencing results showed that the Montana isolate is an Eurasian/American (EA/AM) H5N2 HPAI strain. The partial genome fragments were >99% similar to the virus isolated from a northern pintail duck in Washington State in December 2014. This detection of HPAI (EA/AM) H5N2 virus in a wild bird was NOT associated with any commercial poultry in the United States.

This event is considered closed.

Source of the outbreak(s) or origin of infection

• Contact with wild species

Measures applied

No Control Measures

Animals treated

Vaccination Prohibited

No

Future Reporting

The report and all its outbreaks have been resolved.