

NVSL AVIAN INFLUENZA UPDATE









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NATIONAL VETERINARY SERVICES LABORATORIES

NATIONAL POULTRY IMPROVEMENT PLAN GENERAL CONFERENCE COMMITTEE MEETING, PORTLAND, MAINE MAY 18, 2017



Al Surveillance in the United States: National System



NPIP breeding flocks

- Egg and Meat-Type chickens
- Turkeys, Exhibition Poultry, Upland Game birds and Waterfowl



NPIP production flocks

- Meat-type chickens & turkeys
- Table-Egg Layers
- Upland Game birds and Waterfowl



Live Bird Marketing System

- Producers, distributors, and retail markets
- Backyard flocks, auctions, swap meets, etc.



Other Diagnostics

- Passive surveillance, export testing, foreign animal disease investigations
- Wild bird surveillance

>2 million tests/year

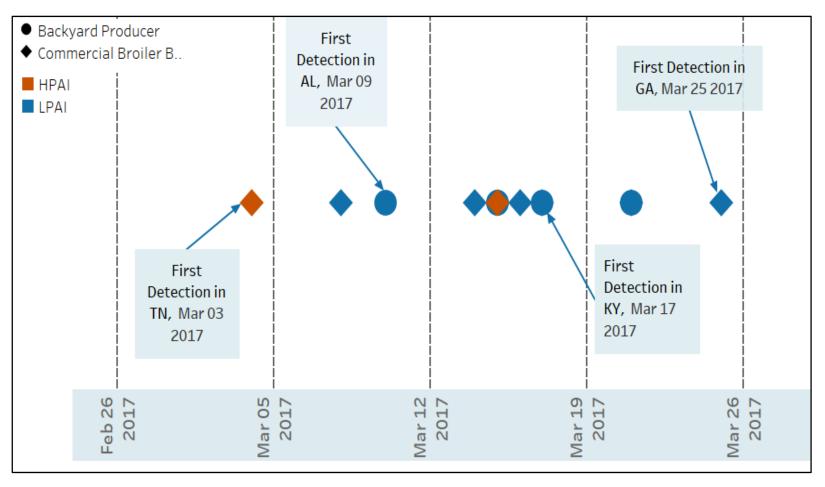


AM H7N9 2017

- H7N9 HPAI and LPAI started in early March 2017, identified initially due to clinical signs
- The AM H7N9 is of North American wild bird lineage and not related to the Asian lineage H7N9
- HPAI and LPAI viruses are highly similar across the entire genome except for the insertion in the HA gene of the HPAI virus:
 - likely derived from chicken host 28S ribosomal RNA
- A closely related H7N9 LPAI virus was recovered from a Wildlife Services Wild Bird Surveillance sample collected from a blue-winged teal in Wyoming as part of a live bird banding effort during September 2016.



Timeline of detections



USDA-APHIS (2017). Epidemiologic and other analyses of avian influenza: April 27, 2017 Report. USDA:APHIS:VS:STAS:Center for Epidemiology and Animal Health. Fort Collins, CO. April 2017. Doc #391.0317.



Diagnostic challenges

- Two important elements:
 - a) the HA-subtype, which is 'presumptive' based upon the NAHLN lab result and 'confirmed' based upon NVSL result, and
 - b) the virus pathotype, which is 'presumptive' based upon the clinical presentation of the flock compared to the USDA <u>HPAI case definition</u>, and 'confirmed' based upon the HA cleavage site sequence at NVSL.

	IAV	HA subtype	NA subtype	pathotype
NPIP AI AGID/ELISA	NP-Antibody	\Diamond	\Diamond	\Diamond
NVSL AI HI/NI	HA-Antibody	✓	✓	0
NAHLN IAV-M PCR	M-RNA	\Diamond	\Diamond	\Diamond
NAHLN H5/H7 PCR	HA-RNA	✓	\Diamond	\Diamond
NVSL Sequencing	✓	✓	✓	✓
NVSL chicken pathogenicty test	\Diamond	\Diamond	\Diamond	✓

^{✓ =} test can determine

^{○ =} test cannot determine



Overall findings: AM H7N9

- Molecular, epidemiologic, and serologic (antibody) data suggest the AM H7N9 virus circulated in the area undetected in poultry prior to the initial HPAI detection; and
- Molecular and epidemiologic support exists for
 - a) secondary spread from the first HPAI site to the second, and
 - b) more than one independent introduction of AM H7N9 LPAI, for example the available data show that LPAI/HPAI viruses from TN appear to have a common source and cluster separately from the AL viruses.

Recall H7N8 HPAI/LPAI Indiana January 2016

Missouri

🗂 Indiana

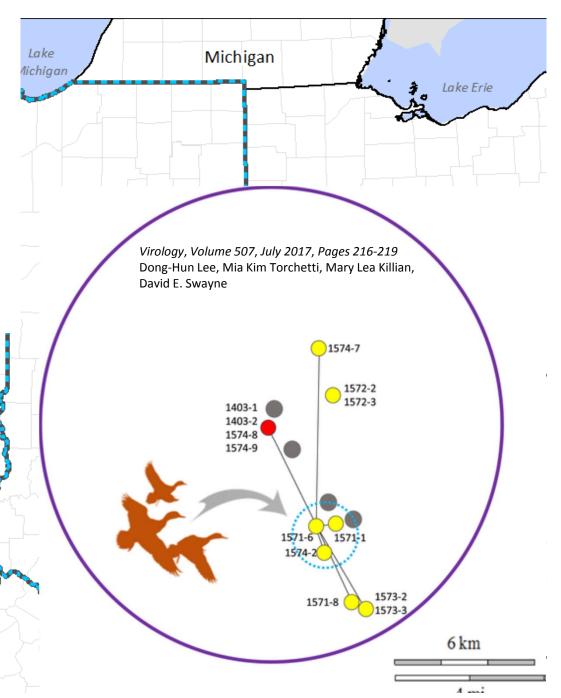
Bird Type:
Commercial

County Boundaries

State Boundaries

- 10 Jan 16: meat turkeys w/↓ water consumption
- 14 Jan 16: preliminary †H7 rRT-PCR and depopulation initiated the NEXT DAY
- Wild bird virus identified with 5/8 genes in common – high similarity
- Molecular analysis supports epidemiologic data for likely location of LPAI introduction

IIIInois



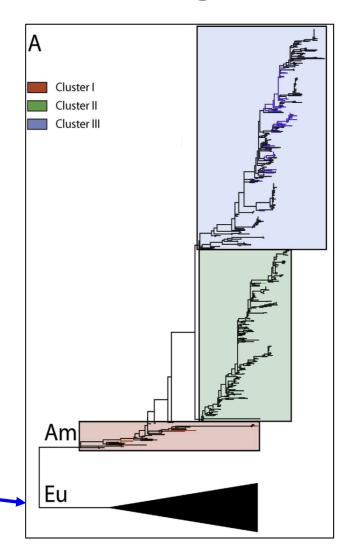
North American H7 Lineage

The recent H7 viruses are North American lineage from Cluster III (H7N8 and H7N9)

Schematic phylogenetic tree of the HA1 nucleotide sequences of H7 AIVs (maximum-likelihood method)
Boxes represent the three major genetic clusters; the Eurasian lineage (EU) is represented by the large black triangle. AM = North American lineage

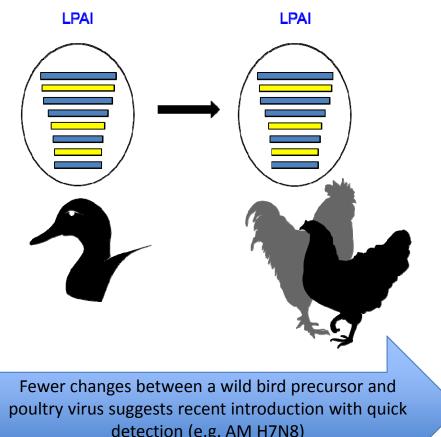
Courtesy of Xi-Feng Wan et al, Department of Basic Sciences, College of Veterinary Medicine, Mississippi State University

North American wild bird lineage viruses are <u>not</u> related to the Asian lineage H7N9



What can wild bird precursors tell us?

- Type and frequency of virus detection
- Distance to nearest wild bird relative helps with understanding timing of introduction



detection (e.g. AM H7N8)



H7N2 in Cats at NYC Shelter

Rare bird flu strain infects 45 cats in single Manhattan shelter and may have spread to recently adopted felines







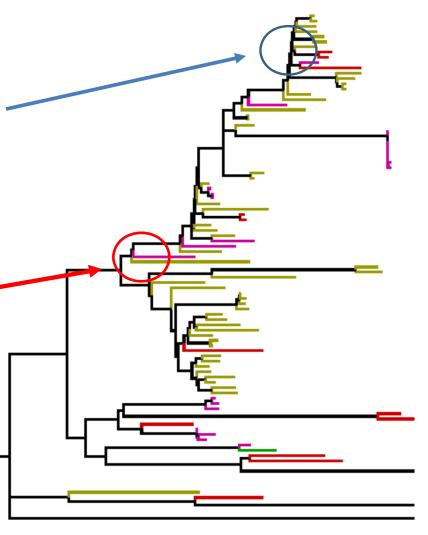






H5 events in US -2017

- March 3, 2017: H5N2 LPAI of wild bird lineage positive result from a commercial turkey flock in Wisconsin
- April 10, 2017: H5N2 LPAI virus in a primarily duck mixed species backyard flock in Idaho
- Both the HA and NA are distinct from the EA/AM H5N2 clade
 2.3.4.4 viruses from 2015



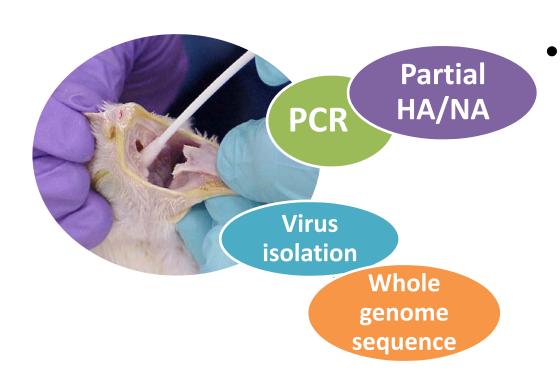
Rapid Response

- USDA APHIS Veterinary Services is committed to rapid depopulation where HPAI is suspected
- The National Animal Health Laboratory Network plays a crucial role;
 depopulation can be initiated at the state level with the following:
 - A non-negative H5 or H7 virus detection (and samples that have been forwarded to the NVSL for confirmation), and
 - A flock that meets the <u>case definition</u>, and
 - Agreement of State and Federal officials.
- NOTE: NVSL confirmation is required to determine subtype and pathotype for official disease declaration – forward non-negative samples ASAP



Fastest route for confirmation

Submit duplicate samples



NVSL leverages the Ct from NAHLN Lab PCR to target samples for rapid subtype/ pathotype by partial sequence where sufficient RNA is present





























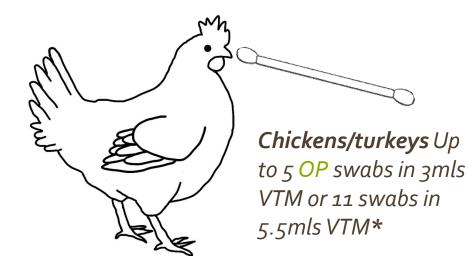
WI-AV-0020

Avian Samples for PCR and virus isolation

https://www.aphis.usda.go v/animal health/lab info services/downloads/WIAVO 020.pdf

Pool by:

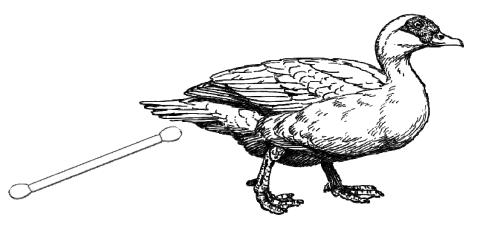
- the same premises
- the same species
- the same sampling route



Domestic waterfowl

Up to 5 CL swabs in 3mls VTM

If sampling free ranging waterfowl — 1 OP <u>and</u> 1 CL swab in SAMETUBE may be preferred (and is recommended for wild waterfowl)

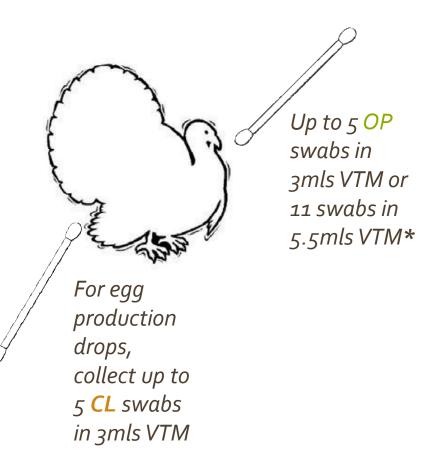




Alternate Sampling

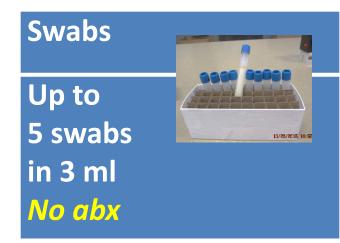


- OP swabs recommended for routine detection
- For egg production drops (potential for swine lineage viruses) consider additional sampling routes
 - Cloacal swabs (CL)
 - Oviduct swabs
 - Semen





How many swabs / how much media?



Purpose

FAD investigations where AI/ND are suspected and detection of bacterial diseases is needed (FADDs are supplied with this media)

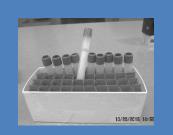
Option to collect 5 swabs/3ml and pool samples at the lab for AI testing allowing ongoing use of such samples for detection of bacterial diseases (refer to 3.1.2 and footnote #2 of the current version of NVSL WI-AV-0020)



How many swabs / how much media?

Swabs

Up to 5 swabs in 3 ml No abx



Either 5 swabs in 3ml (any for domestic species) or up to 11 swabs in 5.5 ml with abx for TR/OP swabs from gallinaceous poultry

Purpose

FAD investigations where AI/ND are suspected and detection of bacterial diseases is needed (FADDs are supplied with this media)

10K surveillance around an infected premise

Surveillance before a premise is positive

Surveillance outside a 10K ring

Surveillance after cleanup





Blood Collection for Antibody Detection

A minimum 0.5ml -1.0ml needed for HI/NI (not to exceed 1% bw of bird – contact the NVSL for testing options for small birds)

- Serum/plasma/yolk
 - 0.2 ml each needed for HI and NI
- Vein (wing, jugular, leg): up to 10 ml depending on bird size; 22–27 ga, 0.5–1" needle
- Heart (anesthesia required): large volume; 18–20 ga, 1.5–2" needle

Important tips -

- To increase the volume of sera
 obtained immediately after collecting
 blood, lay the tube on its side to
 increase surface area exposure to air
 at room temperature
- To obtain better overall sample quality and reduce hemolysis, remove the sera from the clot prior to shipping

Blood collection video

(stop at 1:50sec)

Helpful links:

http://onlinelibrary.wiley.com/enhanced/doi/10.1111/j.1557-9263.2011.00338.x/ http://www.fao.org/docrep/005/ac802e/ac802e0a.htm