

# A Brief Overview of Avian Influenza Virus







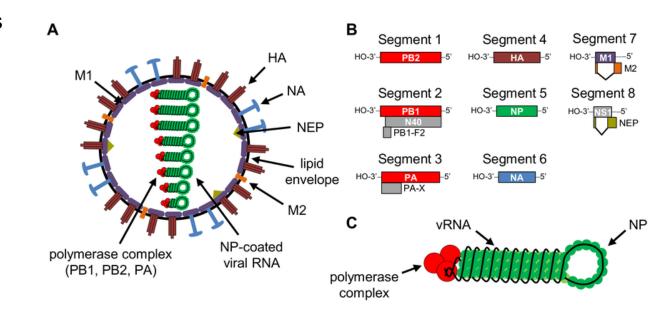


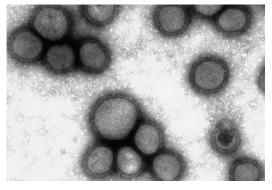
Darrell R. Kapczynski
U.S. National Poultry Research Center, ARS, USDA, Athens, Georgia, U.S.A
NASEM Heritable Genetic Modification of Food Animals- Feb 28<sup>th</sup>, 2024



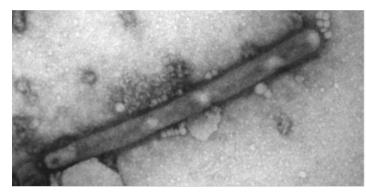
### Avian influenza virus

- □AIV is classified in the family Orthomyxoviridae, genus Influenza A (Type A).
- □ Contains a segmented, negative-sense RNA genome wrapped in enveloped, pleomorphic particles.
- ■8 RNA Segments coding for 10+ proteins
- □ Virus subtype defined by surface proteins
- -Hemagglutinin (HA) H1-H16- attachment
- -Neuraminidase (NA) N1-N9- release
- □ Receptor binding to host cells:
  - Avian influenza HA binds alpha 2,3 linked sialic acid gal
  - □ Human influenza HA binds alpha 2,6 linked sialic acid gal





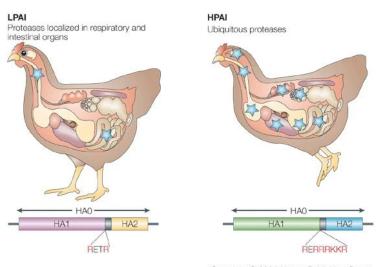




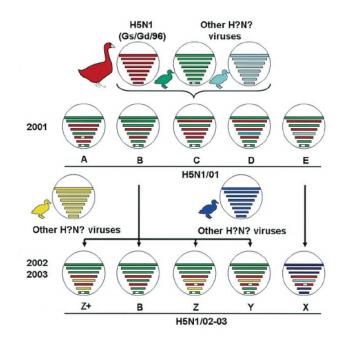
Wild bird origin

### Avian influenza virus

- □ The viruses are divided into low-pathogenic AI (LPAI) and highly pathogenic AI (HPAI) strains. HPAI strains result in high mortality, while LPAI causes milder symptoms, but have demonstrated the ability to mutate and evolve into HPAI.
- ■In molecular terms, HPAI viruses are defined as possessing an H5 or H7 subtype HA with a polybasic sequence at the cleavage site that confers HA cleavage in deep tissue with furin-like enzymes.
- □LPAIV->HPAIV Increase in basic amino acids at HA cleavage site. Genetic insertion of chicken 16S RNA (H7N3 Mexico, 2013).
- □Genetic drift-Substitution rate ~ 5 x 10<sup>-3</sup> per site per year (10-70 nucleotides per year across genome).
- □ Genetic shift-Reassortments between all segments. Segment swapping-infection by two or more viruses in a host.
  - Generates reassortants within wild birds.

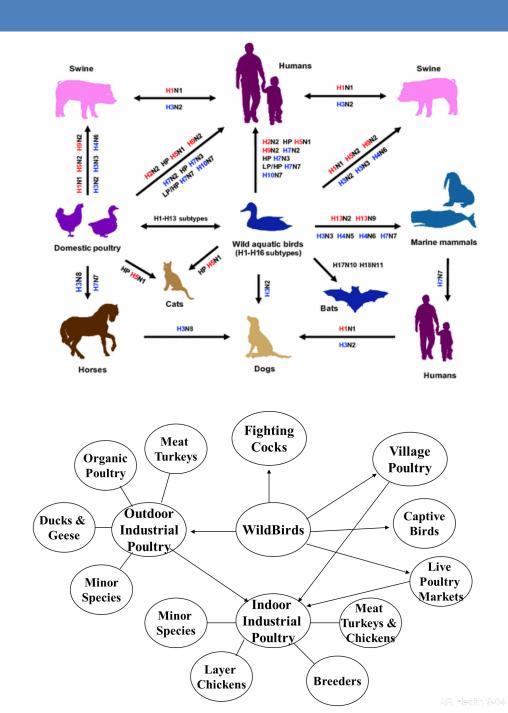




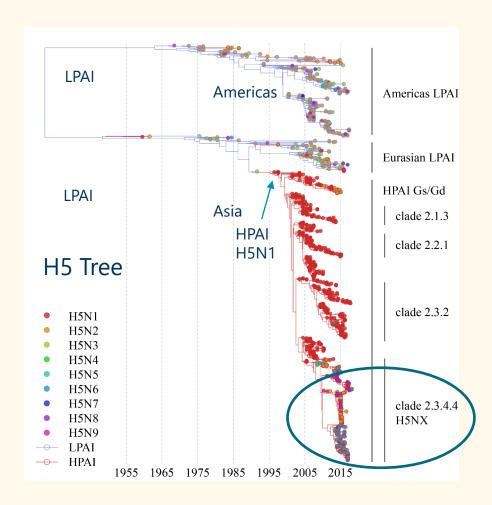


### Ecology of Avian Influenza Virus

- Avian influenza virus (AIV) is endemic in wild birds
- Transmission of AIV from wild birds to poultry species (ducks, chickens, turkeys) occurs commonly
- AIV on rare occasions may become adapted to and become endemic in poultry species (chickens and turkeys)
- AIV once adapted to chickens and turkeys can be difficult to eradicate



### A Brief History of High Path Bird Flu (H5)



Lycett, Duchatel, Digard (2019)
Phil Trans B 2019 Jun 24;374(1775):20180257
doi: 10.1098/rstb.2018.0257.

- HPAI H5N1 circulating in Asia since 1996 (Goose/Guangdong/1996)
- Major outbreak in wild birds (Qinghai lake) 2005
- Ongoing epidemics in Asia and Europe
- Spread between domestic birds, but also to/from wild birds
- Unusual HPAI H5 clade multiple NA subtypes



# Influenza A Viruses Separate Into Genetic Lineages



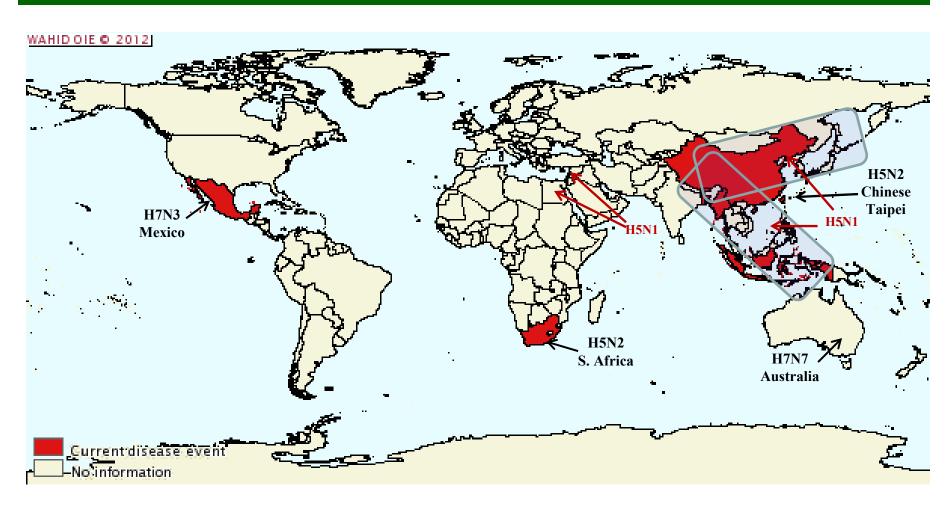


H15



Eurasian

# HPAI (2011-2013): 20 countries



#### H5N1 HPAI

16 countries – poultry, wild birds, humans

#### **H5N2 HPAI**

S. Africa – ostriches Chinese Taipei - poultry

#### H7N3 HPAI

Mexico - layers

#### H7N7 HPAI

Australia - layers

### First long range transmission by wild birds 2014-2015

Species adaption of virus to host:
Sick birds do fly

- if they are wild anseriformes
- if H5N8/X

First detection of Asian lineage H5 HPAIV in U.S.

North Pole Projection











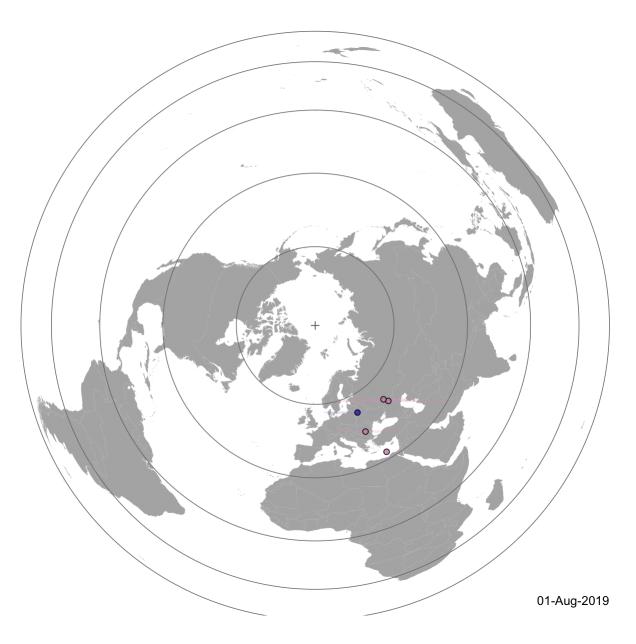


### H5NX 2020-2023 Reassortants From North Pole

- H5N1/2020/R2 Europe
- H5N1/2021/R2\_Europe
- H5N1/2021/R4 Asia
- H5N1/2021/R5\_Europe
- H5N1/2021/R7\_Europe
- H5N1/2022/R1\_NAmerica
- H5N1/2022/R10 NAmerica
- H5N1/2022/R12 Europe
- H5N1/2022/R13 NAmerica
- H5N1/2022/R2\_NAmerica
- H5N1/2022/R3\_NAmerica
- H5N1/2022/R4\_NAmerica
- H5N1/2022/R8 NAmerica
- H5N5/2020/R1 Europe
- H5N8/2020/R1\_Europe
- H5N8/2020/R2 Asia
- Other







#### Record avian flu outbreak sees 48m birds culled in UK and EU

By Malcolm Prior BBC News Rural Affairs Produ

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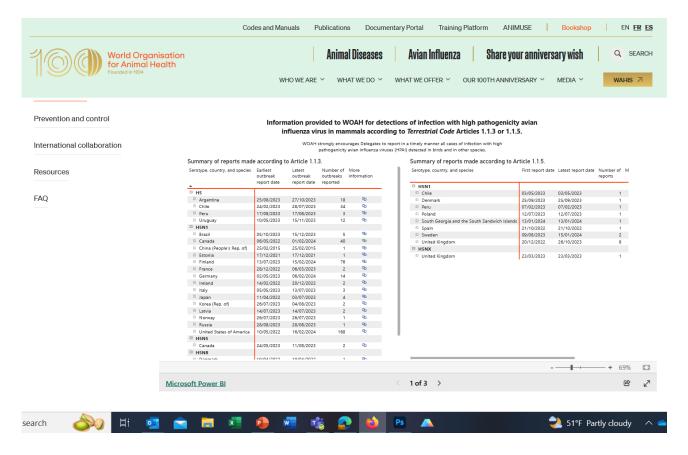




Dr. Samantha Lycett at Univ. of Edinburgh.

### Increased detections of H5N1 clade 2.3.4.4 HPAIV in mammals

Species jump into terrestrial and marine mammals. No interspecies transmission reported to date.



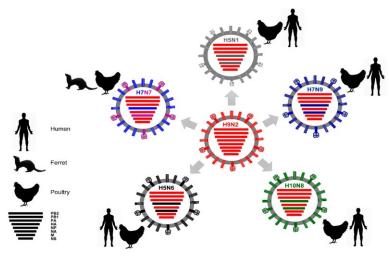
#### Mass sea lion deaths in Peru raise further fears bird flu is jumping between mammals

Scientists are monitoring the outbreak to see if the H5N1 virus was transmitted between the sea mammals or caught it directly from birds in the South American country

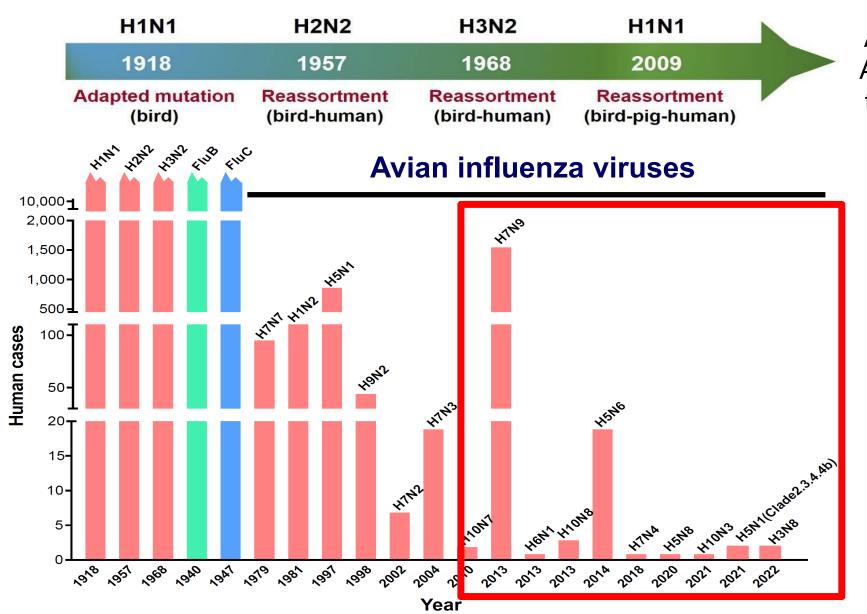


iNews 10/02/23

Scientists at the Paracas National Reserve inspecting a dead sea lion (Photo: Sernang/Getty



### Avian influenza virus and human infections



At least **15 subtypes** of AIV have been identified that could directly infect human.

### Strategies for control of HPAI

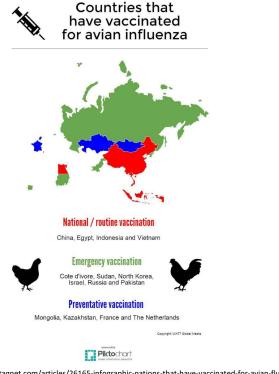
#### **Traditional**

#### **Biosecurity**





#### **Vaccination**

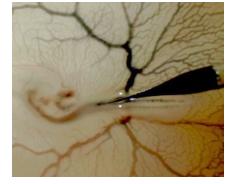


https://www.wattagnet.com/articles/26165-infographic-nations-that-have-vaccinated-for-avian-flu

**Depopulation policy** "stamping out"

### **Genetic engineering**

#### Disease resistance

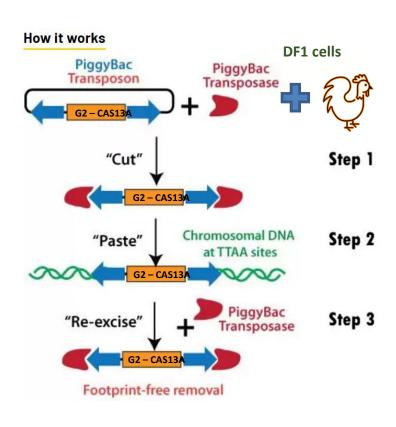


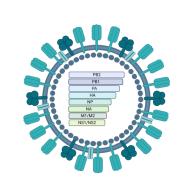


#### Targets to research

- Inhibit virus attachment
  - Viral receptor
  - HA binding
- Inhibit virus replication
  - Polymerase complex
  - ❖ ANP32 family
- Enhance innate immunity
  - Adaptive immunity too slow to control mortality
  - MX, TLR7, RIG-I
  - CRISPR/Cas 13 family

### Targeting AIV genome with CRISPR-Cas13A in vitro



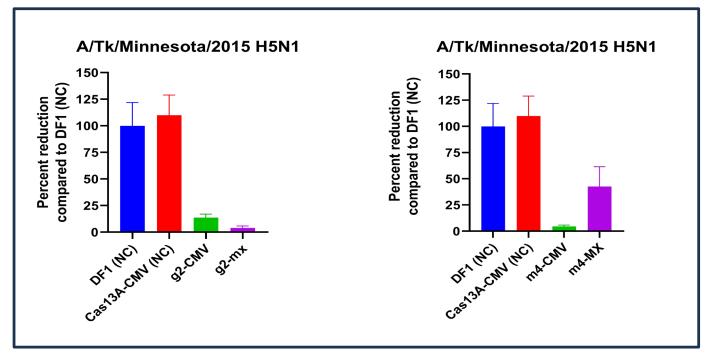


#### PB2

G2 = is the genome sequence (binds to mRNA) M4 = is complimentary sequence (binds to vgRNA)

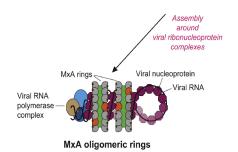


#### **HPAIV**

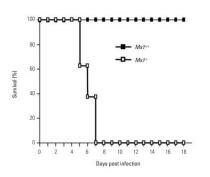


### Host-derived targets using Mouse Mx in vitro

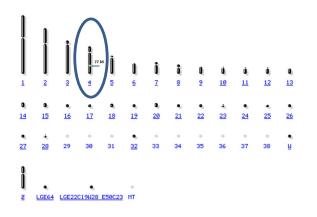
- Mx proteins are interferon-induced GTPases that directly interact with viral proteins to inhibit viral replication
- ❖ Mouse Mx1 is a critical determinant of influenza resistance
- Chicken Mx has conflicting reports on its effectiveness against AIV
- ❖ Inserted mouse Mx1 under control of the chicken Mx promoter



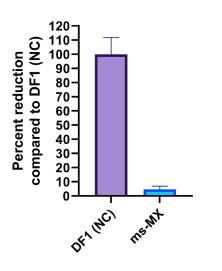
https://doi.org/10.1016/j.immuni.2011.07.012



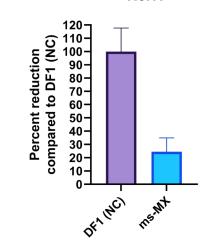
Chicken genome – insert at chromosome 4



#### A/Tk/Wisc/68 H5N9



#### A/AmWidgeon/South Carolina/21 H5N1



Virus name	strain	Pathogenicity	Percent Reduction compared to DF1 control
A/Tk/Wisc./68	H5N9	LPAI	90%
A/Ck/CA/03	H6N2	LPAI	80%
A/Ck/NJ/97	H9N2	LPAI	90%
A/AmWidgeon/SC/2021	H5N1	HPAI	75%
A/Ck/Jalisco/2012	H7N3	HPAI	90%
A/Tk/Minn./2015	H5N1	HPAI	95%
A/Tk/poland/2021 HPAI	H5N8	HPAI	70%

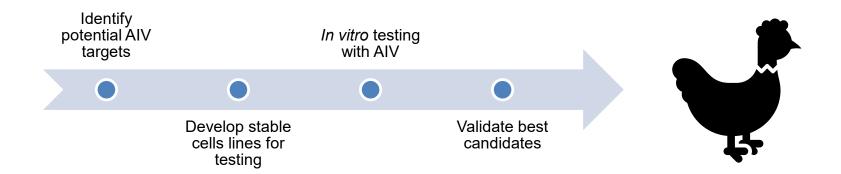
#### Conclusions & future directions

The current global transmission and infection landscape of AIV in birds and mammals supports evidence that these viruses are entrenched in migratory waterfowl.

Gene-editing approaches to genetically modify poultry offer a means to control highly pathogenic avian influenza virus and limit or restrict infections in birds, and limit transmission to other birds and mammals.

In the absence of vaccine usage, gene-editing offers an alternative avenue to breeding animals for disease resistance and farm biosecurity.

Do gene-edited animals with disease resistance create a novel pool with unique characteristics ?







A pig with a CRISPR-edited gene (ear tag #43) that makes it resistant to a deadly virus looks indistinguishable from the unedited pigs surrounding it. PIC



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Dr. Klaudia Chrzastek

Dr. Karen Segovia

Dr. David L. Suarez

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Dr. Kelsey Briggs

Ryan Sweeney





