WHO Global Update:

Public health risks from influenza viruses circulating in swine

Elizabeth Mumford WHO/HSE Department of Food Safety and Zoonoses





• ???????????

- "Increasing" number of human cases detected/reported
- National/International notification requirements



IHR (2005) Case definition: Human influenza caused by a new subtype

- State Parties to the IHR (2005) are required to immediately notify WHO of any
 <u>laboratory confirmed case</u> of a <u>recent human infection</u> caused by an influenza
 A virus with the <u>potential to cause a pandemic</u>. Evidence of illness is not required for this report.
- An influenza A virus is considered to have the potential to cause a pandemic if the virus has demonstrated the <u>capacity to infect a human</u> and if the heamagglutinin gene (or protein) is <u>not a variant or mutated form</u> of those, i.e. A/H1 or A/H3, circulating widely in the human population.
- An infection is considered recent if it has been confirmed by positive results from polymerase chain reaction (PCR), virus isolation, or paired acute and convalescent serologic tests. An antibody titre in a single serum is often not enough to confirm a recent infection, and should be assessed by reference to valid WHO case definitions for human infections with specific influenza A subtypes.



Human infections: USA (Jan 2011- Mar 2012)

- H3N2 TrR H3N2 (USA strain SIV) with A(H1N1)pdm09 M gene
 - 7 sporadic in 3 states
 - 6 exposed to pigs or contaminated environments
 - 1 no known exposure (potentially H2H, swine worker, untested)
 - 3 in epi-linked cluster (plus 2 ILI epi-linked no confirmation)

• H1N2, H1N1



Human cases

- Viruses appear to reflect those circulating in swine in region at the time of infection
- Variable, generally mild, clinical presentation (subclinical ←→ ILI)
- No deaths
- Most have exposure to swine or other infected humans
 - Some exposures unclear

Public health risk assessment for flu at the HAI (same as last year)

- Before human cases occur assessment based on antigenic and genetic characteristics of viruses identified/circulating in animals
- Mostly UNABLE to identify what constitutes an increased PH risk, due to:
 - PH risks associated with sequence mutations and markers??
 - lack of linked virological and epidemiological information
 - lack of "contextual" information (ecology, current pattern of virus circulation, management system, human innate/acquired immunity, etc)
 - Limited surveillance scope (humans and animals)
 - No global baseline

Overarching question (still)

What are the public health risks from influenza viruses in swine?



World Health

anization

WHO perspective: Two aspects (still)

- WHO's role re SIV in humans and at the humanswine interface
- What WHO would like to learn from AH through SIV surveillance or research



What we need to know: PH & interface

- Epidemiology and clinical
 - Baseline frequency and severity of human cases
 - Changes in frequency and/or severity of human infections
 - Activities putting humans at risk for infection/disease
 - Types of animal workers at risk for infection/disease
 - Differences between exposure risks and disease risks
 - Host factors (genetic, acquired immunity)

Virology

- Characteristics of viruses infecting humans (including antiviral sensitivity)
- Whether viruses infecting humans reflect the spectrum of circulating animal viruses or if they are a subset of the viruses circulating in animals
 - and if a subset, identification of any common differentiating characteristics.



What we would like to know: AH

• Epidemiology

- Transmission dynamics of SIVs, including seasonality
- Changes in epidemiology

Virology

- Distribution of subtypes and strains infecting swine (and other species!)
- Whether viruses circulating in swine are sensitive to antiviral drugs
 - Oseltamivir, adamantines, zanamivir
 - genetic screen follow up with phenotypic testing
- Viral factors associated with cross-species transmission to/from swine
 - especially from birds to swine
- Distribution and characteristics of H2 viruses in swine
- Changes: in distribution, circulation of "new" subtypes, when there are crossspecies events



This requires...

- On animal health side:
 - Epidemiological and virological surveillance and monitoring is vine = baseline and changes
- On public health side:
 - Epidemiological surveillance and monitor of high-contact humans for exposure (serological testing required) no disease (ILI/SARI surveillance required) = baseline and change (O)
 - A better understanding of a for orbital reactions (e.g. cross reactions, persistence of antibody, conditions for seroconversion)
 - Virological surveillar and monitoring in humans
- At the interval
 - Tests/regents to specifically distinguish strains serologically
 - Mechanism for sharing information
 - Mechanism for early joint assessment of emerging events



Naming

• Yet to find system that meets all needs and concerns

Request for tripartite to continue conversation



Summary

- Lack of understanding of global incidence of human infections with influenza viruses circulating in swine
- Little understanding of PH risks at the human-swine interface
 - Epidemiology
 - Impact

 Significant gaps in swine influenza virus (SIV) surveillance - AH and PH - worldwide



Thank you!

