

# The public health risk of influenza in pigs – recent insights, key knowledge gaps

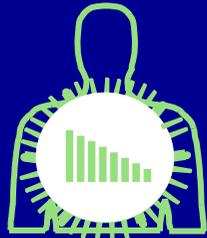
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# H1N1, H3N2 and H1N2 viruses established in swine populations worldwide

- SIVs are of human or avian origin, most viruses are reassortants with a mix of human/avian and swine-adapted genes
- Concurrent circulation of multiple subtypes and lineages, especially in swine-dense regions
- Distinct virus lineages in Northern America vs Europe (and Asia)

# H3N2 SIV in Europe and North America

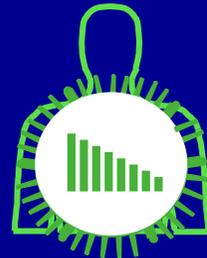
Seasonal human H3N2



Hong Kong/68



Philippines/82



Sydney/97



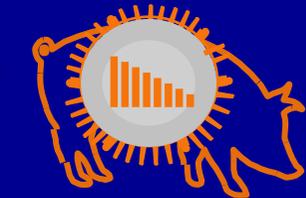
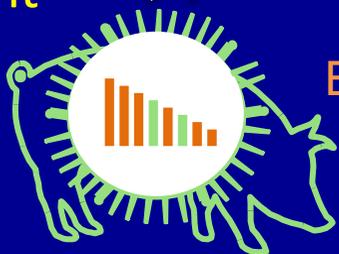
Perth/09



1970

mid 1980's

Reassortant H3N2

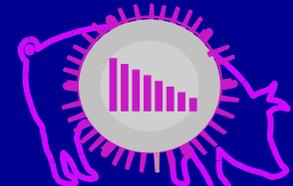
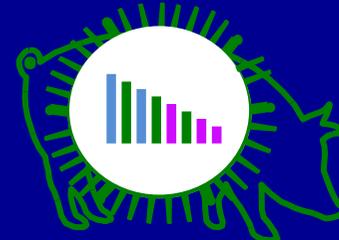


Eu swine H1N1 (avian-like)

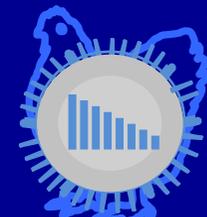


1998

"Triple" reassortant H3N2



Classical swine H1N1

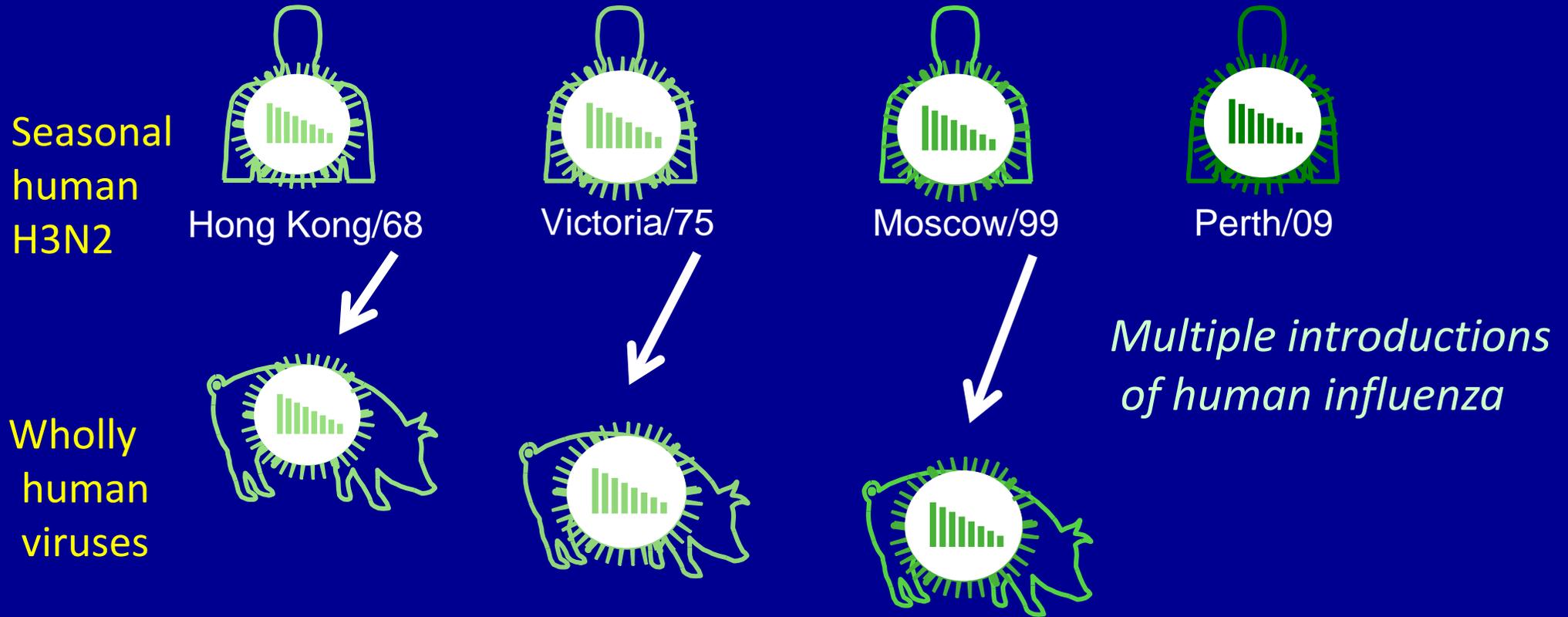


Avian

H3 related to older (1970-80) human viruses, not to recent ones (Kyriakis et al., ZPH 2011)

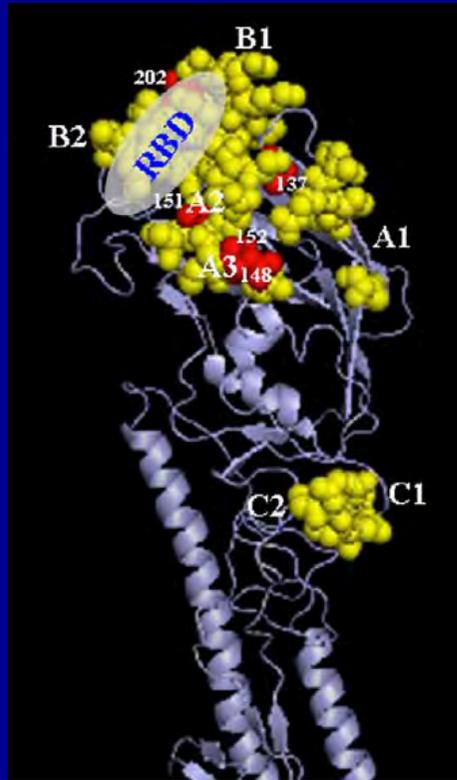


# H3N2 SIV in Asia (China)



- ❑ Various double and triple H3N2 reassortants (Yu et al. J. Clin. Microbiol. 2008, Sun et al. J. Clin. Virol. 2009, Shu et al. Virol. 1994)
- ❑ H3N2 reassortants with internal genes of avian viruses (H9N2 or H5N1) or avian N2 (Bi et al. EID 2010, Cong et al. Pone 2010)

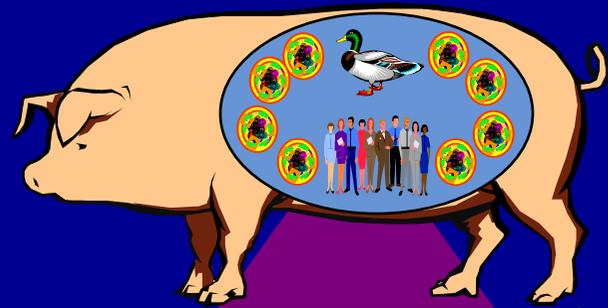
# Antigenic drift in SIV HA



Conserved amino acids and changes in HA1 of sw/Jilin/19/07 (H3N2) vs Moscow/10/99 (H3N2) (Cong et al. POne 2010)

- European swine H3N2: genetic drift at similar rate as human H3N2, but slower drift because changes outside antigenic domains (de Jong et al. Vaccine 2007)
- Substantial genetic/antigenic diversity between H1 lineages:
  - N. America: classical H1 vs more recent H1 clusters ( $\beta$ ,  $\gamma$ ,  $\delta 1$ ,  $\delta 2$ , 2009p)
  - Eu: avian-like H1N1 vs human-like H1N2 and 2009 pH1N1
- Need for exhaustive antigenic analyses, study of immunodominant epitopes

# Central dogma: pigs as unique intermediate hosts for pandemic influenza viruses

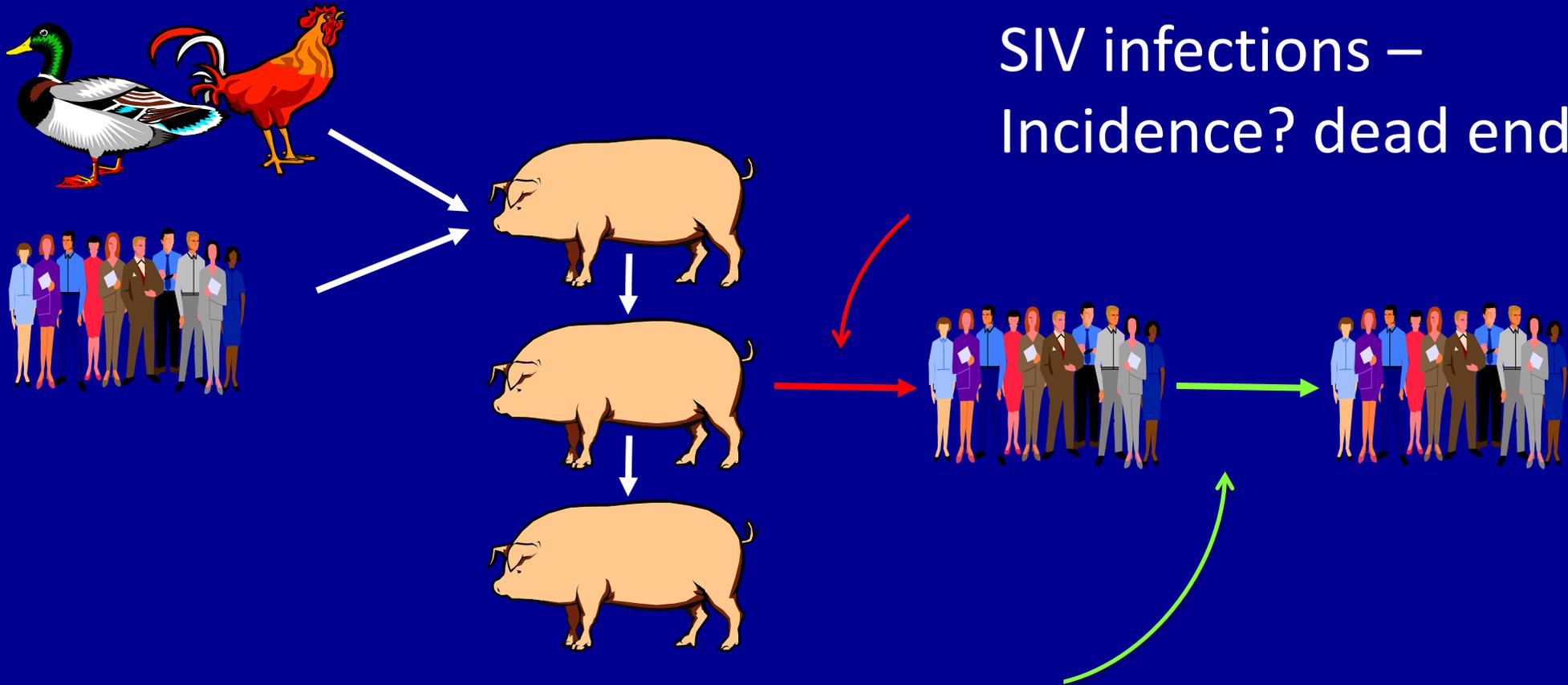


- ✓ Pigs susceptible to avian and human viruses
- ✓ Pigs have both avian-type ( $\alpha 2,3$ ) and human-type ( $\alpha 2,6$ ) sialic acid receptors

**Conclusion:** Avian viruses adapt to humans in the pig by mutation or reassortment



# Facts



✓ Occasional zoonotic SIV infections – Incidence? dead end!

✓ 2009 pH1N1 is first pandemic virus (that is almost certainly) of swine origin

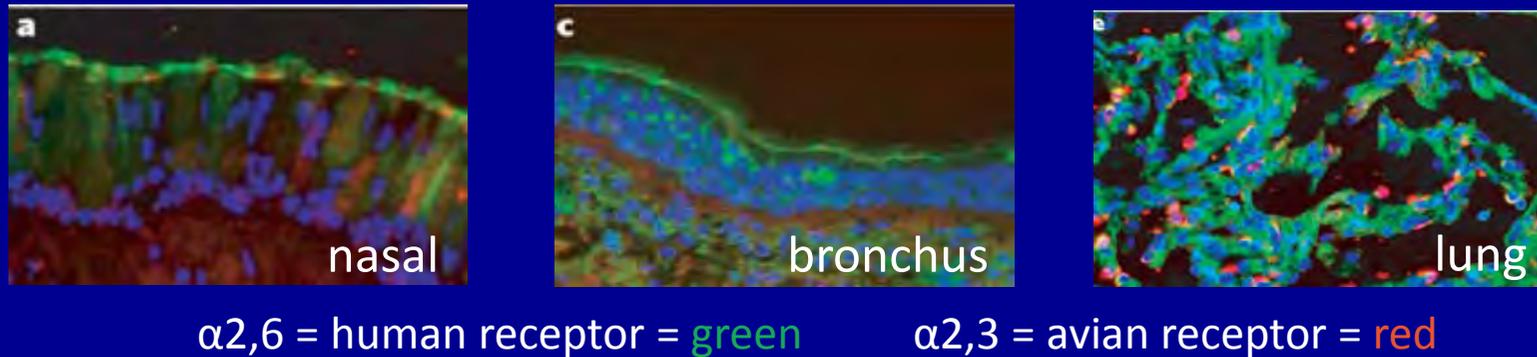
✓ H5N1 has infected humans directly, not through pigs

# Research Issue #1 – Factors that make influenza viruses adapted for replication in pigs, transmission between pigs and from pigs to humans



- Can we identify genetic markers of adaptation (e.g. mutations, sets of genes)?
- Can non-H1 or -H3 viruses (e.g. H5, H9) become adapted to swine or human populations?
- Are pigs preferred “mixing vessels” as compared to humans or birds?
- Is the barrier between pigs and humans weaker than that between birds and humans?

# Research Issue #1 cont– Significance of $\alpha 2,3$ (avian-type) and $\alpha 2,6$ (human-type) sialic acid receptors in porcine respiratory tract? (Ito et al 1998)



- Humans and some land-based poultry species also express both types of sialic acids (Shinya et al. 2006; Wan and Perez 2006; Kuchipudi et al. 2009)  
Expression pattern similar to pigs (Nelli et al. 2010, Van Poucke et al. 2010)
- Virus binding to a given sialic acid may not lead to productive infection of cells: co-receptors? other barriers of infection?

# Research Issue #2 – Cross-protective immunity between viruses from swine and humans (and birds)?: Significance of heterovariant/heterosubtypic immunity, mechanisms, viral targets



- Differences between mice and swine/men!
  - ✓ Novel generation vaccines (DNA, rec M2e, adenovector) not protective in pigs
  - ✓ Cross-protection between H1N1 and H3N2 limited to mice
- Post-infection immunity in swine is broad , not entirely dependent on HA-Abs

# Experimental cross-protection studies pigs

## Cross-protection between antigenically distinct H1N1 swine influenza viruses from Europe and North America

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Prior infection of pigs with swine influenza viruses is a barrier to infection with avian influenza viruses

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## Determinants of pathogenicity may also differ between mice and other species

- Reconstructed 1918 virus was lethal in mice, ferrets and macaques, but not in pigs  
*Weingartl et al. JVi 2009*
- Pathogenicity of SIVs with different PB2 genes and mutation of codon 627 in mice does not correlate with pathogenicity of SIVs in the pig  
*Ma et al. Virology 2000*

# Research Issue #3 – Determinants of pathogenicity



- Can H1 or H3 influenza viruses become highly pathogenic for swine or humans?
- Can H1 or H3 viruses induce a cytokine “storm”?
- NS1 is an IFN antagonist in continuous cell lines, but does it interfere with the IFN system in pigs and people???



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*Collaborative project*

Pathogenesis and transmission  
of influenza virus in pigs

*2010-14*



**FLUPIG**

- ✓ Research priorities for SI
- ✓ Harmonization of global surveillance for influenza in swine