


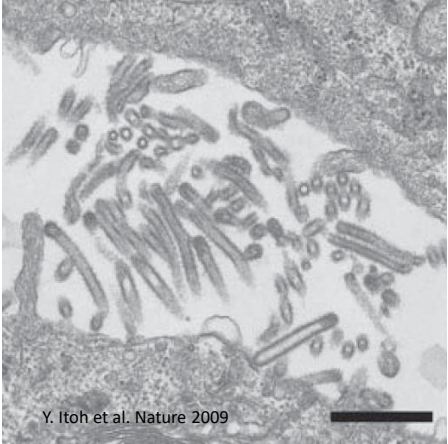
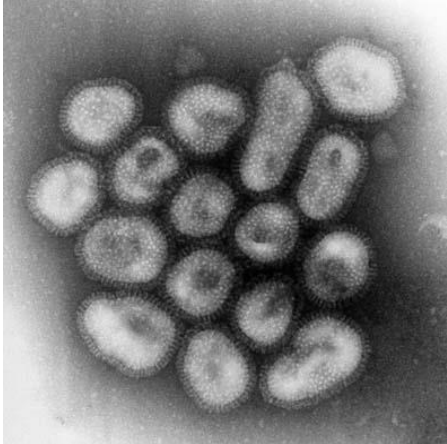
Avian Influenza Viruses with Pandemic Potential

Takato Odagiri
Director

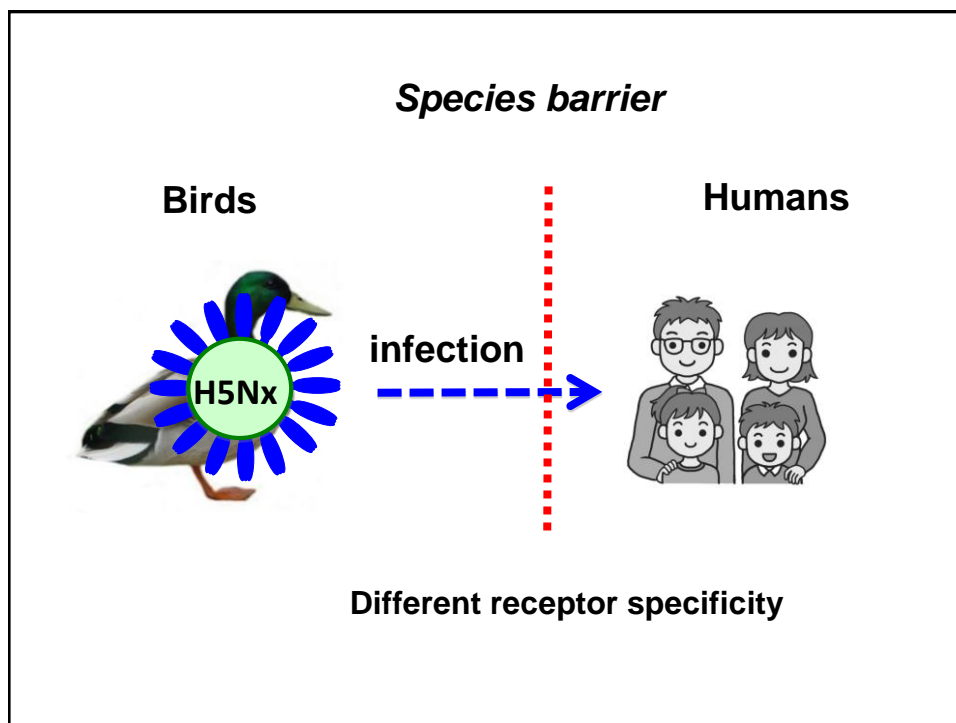
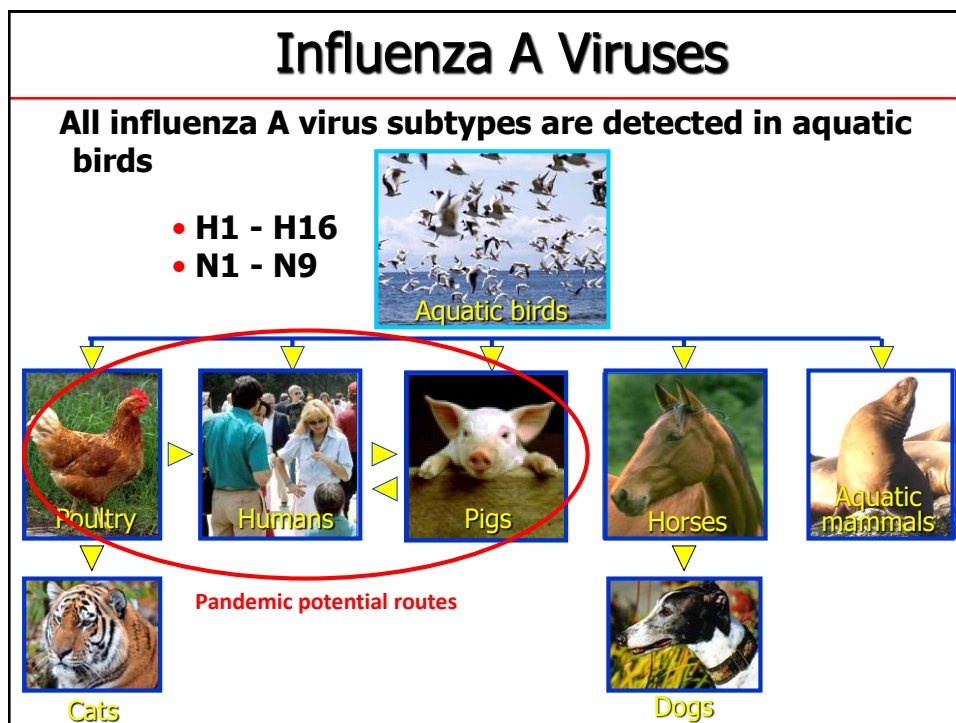
**WHO Collaborating Center on Influenza , Tokyo
and
Influenza Virus Research Center,
National Institute of Infectious Diseases,
Japan**

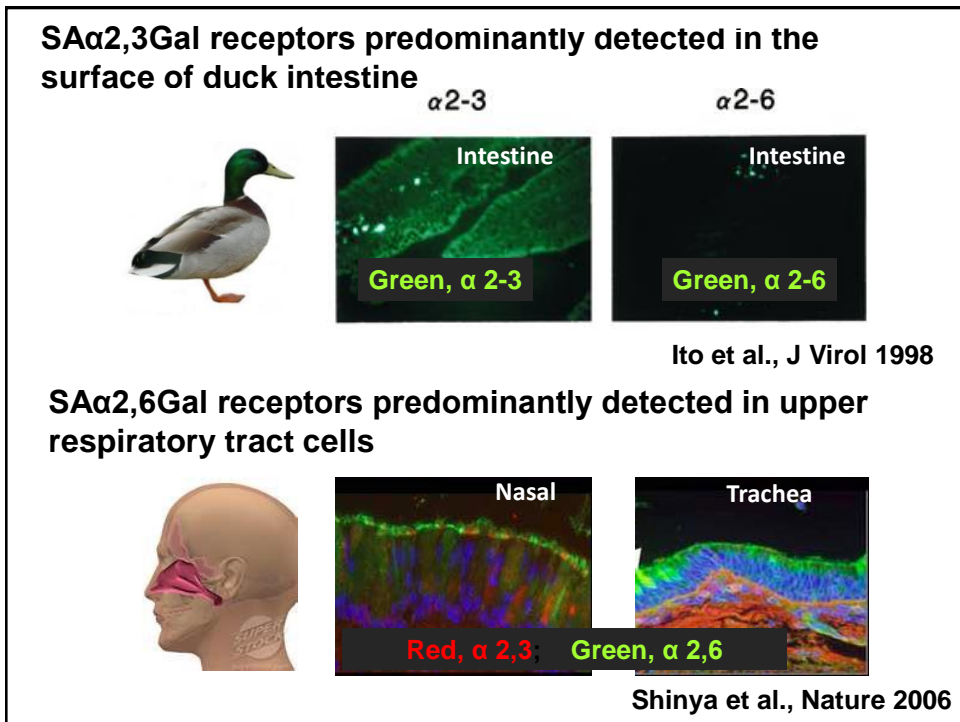
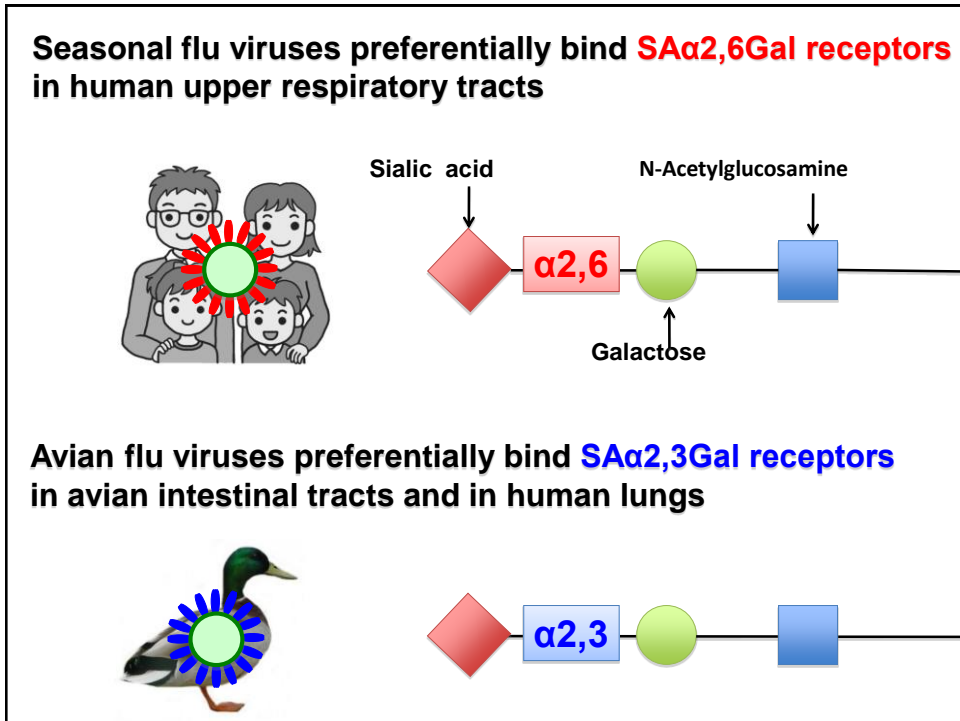


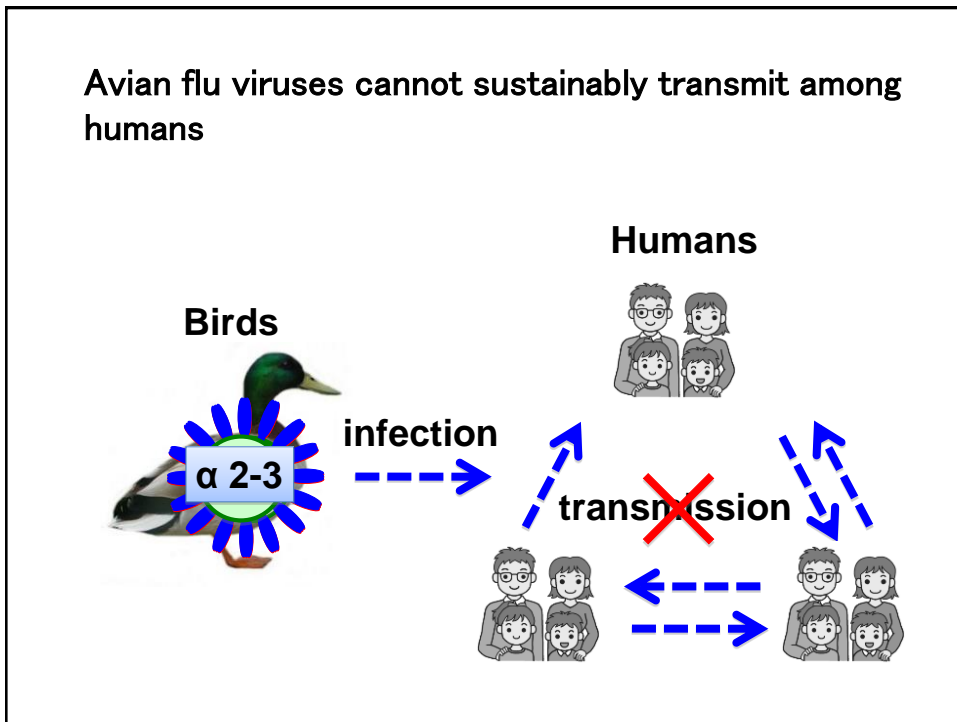
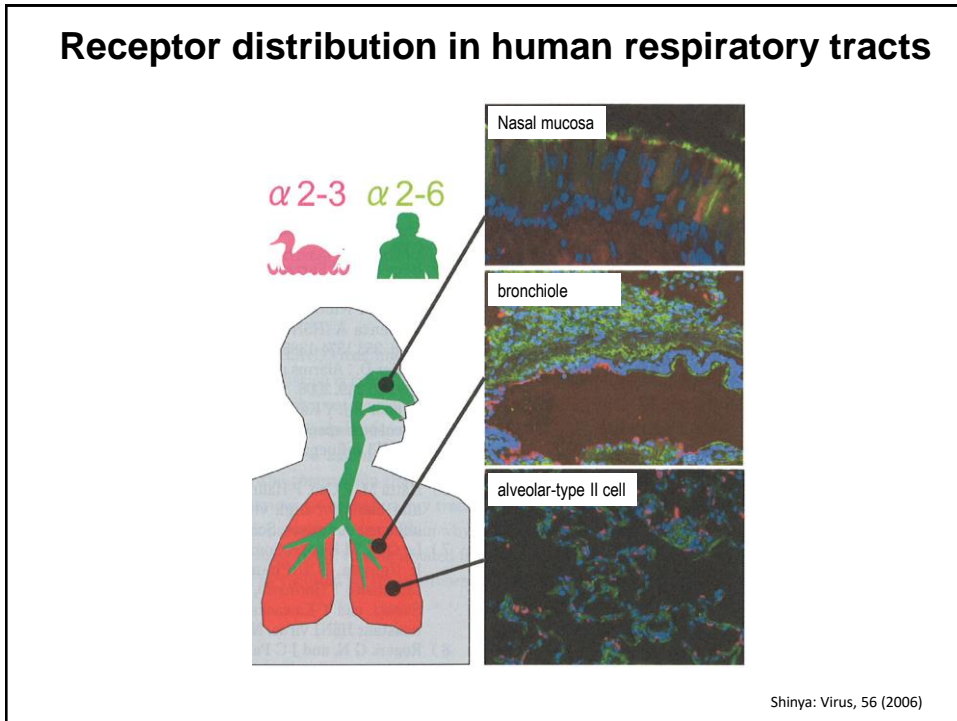
Influenza Viruses

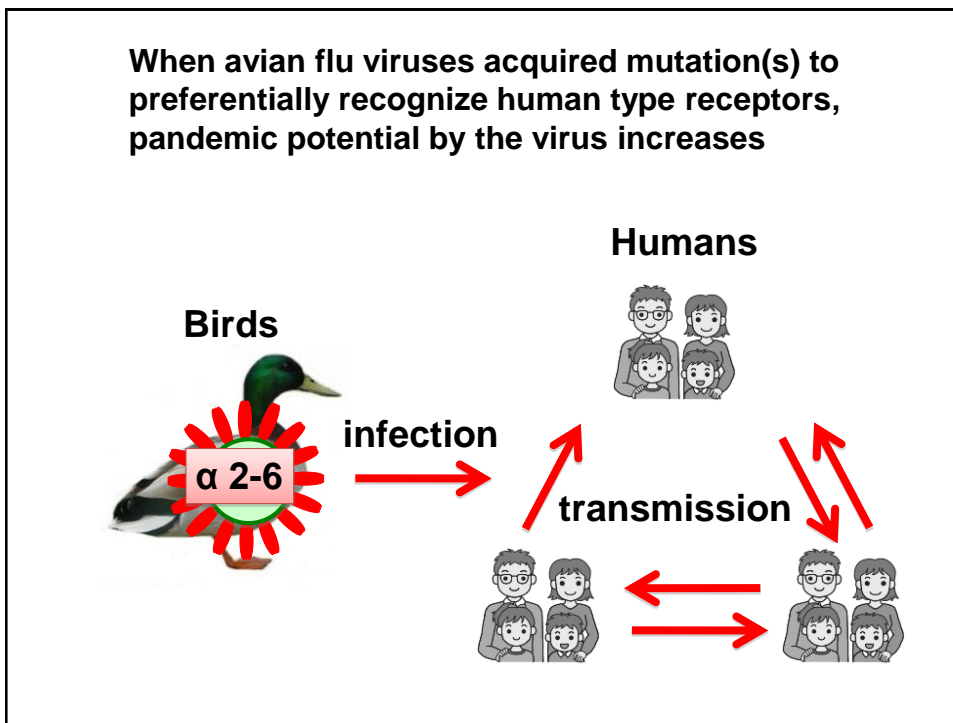


Y. Itoh et al. Nature 2009









Zoonotic Influenza A viruses currently detected in Animals and Humans in the World (since Sept 2016)

- Avian viruses
 - H5Nx :
 - Since 2003,
 - H5N1 (poultry, wild birds, **human (6)**) **(860)**
 - H5N2 (poultry, wild birds)
 - H5N5 (wild birds)
 - H5N6 (poultry, wild birds, **human (16)**)
 - H5N8 (poultry, wild birds)
 - Since 2013,
 - H7N9 : (poultry, **human (758)**, environment) **(1564)**
 - H7N2 : (poultry)
 - H9N2 : (environment, **human(4)**)
- Swine viruses
 - H1N1v (pig)
 - H1N2v (**human(2)**, pig)
 - H3N2v (**human(31)**, pig)

http://www.who.int/influenza/vaccines/virus/201709_zoonotic_vaccinevirusupdate.pdf?ua=1

Zoonotic Influenza A viruses currently detected in Animals and Humans in the World (since Sept 2016)

● Avian viruses



H5Nx :

H5N1 (poultry, wild birds, **human (6)**)(860)

H5N2 (poultry , wild birds)

H5N5 (wild birds)

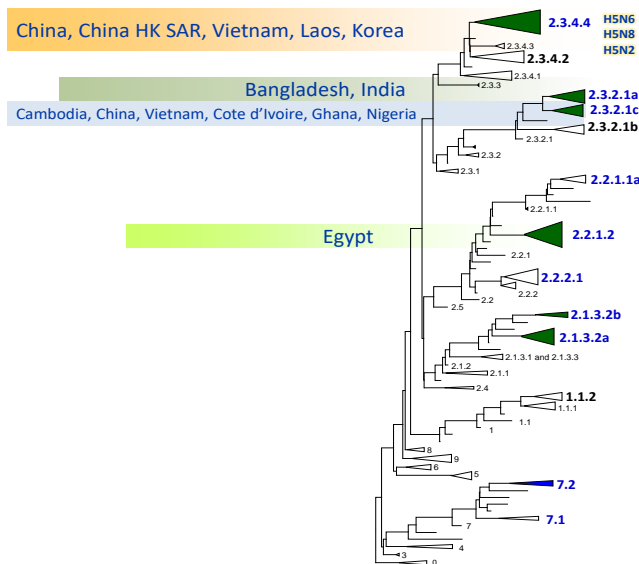
H5N6 (poultry, wild birds, **human (16)**)

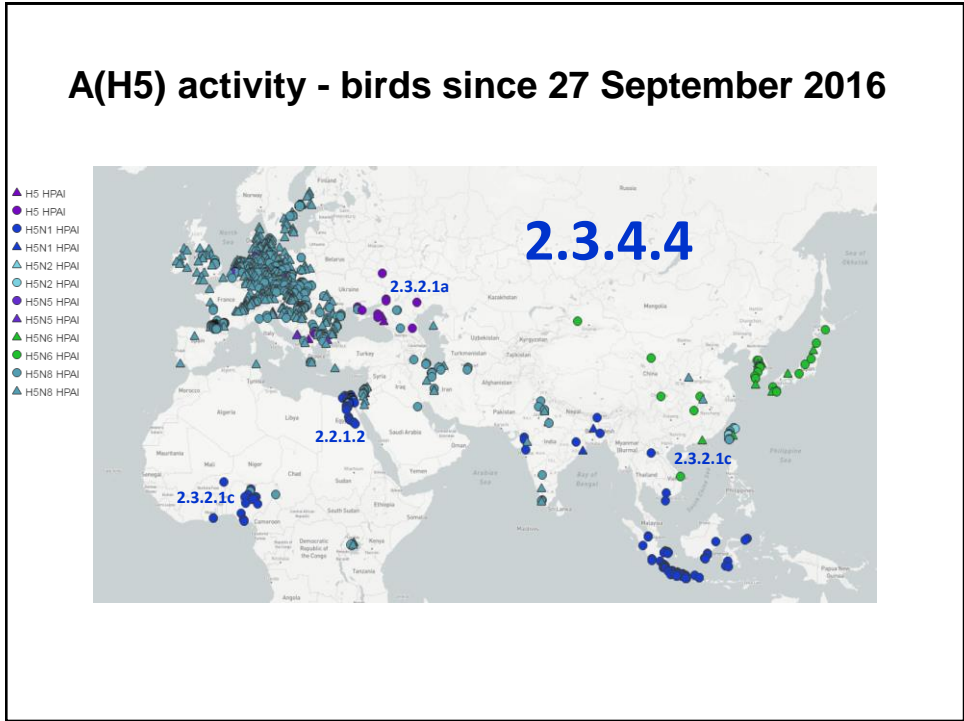
H5N8 (poultry, wild birds)

Since 2003,

http://www.who.int/influenza/vaccines/virus/201709_zoonotic_vaccinevirusupdate.pdf?ua=1

Evolution of A(H5) HA genes





Cumulative number of confirmed human cases for avian influenza A(H5N1) reported to WHO, 2003-2017

Country	2003-2009*		2010-2014**		2015		2016		2017		Total	
	cases	deaths	cases	deaths	cases	deaths	cases	deaths	cases	deaths	cases	deaths
Azerbaijan	8	5	0	0	0	0	0	0	0	0	8	5
Bangladesh	1	0	6	1	1	0	0	0	0	0	8	1
Cambodia	9	7	47	30	0	0	0	0	0	0	56	37
Canada	0	0	1	1	0	0	0	0	0	0	1	1
China	38	25	9	5	6	1	0	0	0	0	53	31
Djibouti	1	0	0	0	0	0	0	0	0	0	1	0
Egypt	90	27	120	50	136	39	10	3	3	1	359	120
Indonesia	162	134	35	31	2	2	0	0	0	0	199	167
Iraq	3	2	0	0	0	0	0	0	0	0	3	2
Lao People's Democratic Republic	2	2	0	0	0	0	0	0	0	0	2	2
Myanmar	1	0	0	0	0	0	0	0	0	0	1	0
Nigeria	1	1	0	0	0	0	0	0	0	0	1	1
Pakistan	3	1	0	0	0	0	0	0	0	0	3	1
Thailand	25	17	0	0	0	0	0	0	0	0	25	17
Turkey	12	4	0	0	0	0	0	0	0	0	12	4
Viet Nam	112	57	15	7	0	0	0	0	0	0	127	64
Total	468	282	233	125	145	42	10	3	3	1	859	453




http://www.who.int/influenza/human_animal_interface/2017_07_25_tableH5N1.pdf?ua=1

Available H5Nx CVVs provided by WHO CCs/ERLs

Table 2. Status of influenza A(H5) candidate vaccine virus development


Candidate vaccine viruses	Clade	Institution*	Available
A/Viet Nam/1203/2004 (CDC-RG; SJRG-161052)	1	CDC and SJCRH	Yes
A/Viet Nam/1194/2004 (NIBRG-14)	1	NIBSC	Yes
A/Cambodia/R0405050/2007 (NIBRG-88)	1.1	NIBSC	Yes
A/Cambodia/X0810301/2013 (IDCDC-RG34B)	1.1.2	CDC	Yes
A/duck/Hunan/795/2002 (SJRG-166614)	2.1.1	SJCRH/HKU	Yes
A/Indonesia/5/2005 (CDC-RG2)	2.1.3.2	CDC	Yes
A/Indonesia/NHHD11771/2011 (NHDRG-9)	2.1.3.2a	NIID	Yes
A/bar-headed goose/Qinghai/1A/2005 (SJRG-163222)	2.2	SJCRH/HKU	Yes
A/chicken/India/NIV33487/2006 (IBCDC-RG7)	2.2	CDC/NIV	Yes
A/whooper swan/Mongolia/244/2005 (SJRG-163243)	2.2	SJCRH	Yes
A/Egypt/2321-NAMRU3/2007 (IDCDC-RG11)	2.2.1	CDC	Yes
A/turkey/Turkey/1/2005 (NIBRG-23)	2.2.1	NIBSC	Yes
A/Egypt/N03072/2010 (IDCDC-RG29)	2.2.1	CDC	Yes
A/Egypt/3300-NAMRU3/2008 (IDCDC-RG13)	2.2.1.1	CDC	Yes
A/Egypt/N04915/2014 (NIBRG-306)	2.2.1.2	NIBSC	Yes
A/common magpie/Hong Kong/5052/2007 (SJRG-166615)	2.3.2.1	SJCRH/HKU	Yes
A/Hubei/1/2010 (IDCDC-RG30)	2.3.2.1a	CDC	Yes
A/duck/Bangladesh/19097/2013 (SJ007)	2.3.2.1a	SJCRH	Yes
A/barn swallow/Hong Kong/D10-1161/2010 (SJ003)	2.3.2.1b	SJCRH/HKU	Yes
A/duck/Viet Nam/NCVD-1584/2012 (NIBRG-301)	2.3.2.1c	NIBSC	Yes
A/chicken/Hong Kong/API156/2008 (SJ002)	2.3.4	SJCRH/HKU	Yes
A/Anhui/1/2005 (IBCDC-RG6)	2.3.4	CDC	Yes
A/duck/Laos/3295/2006 (CBER-RG1)	2.3.4	FDA	Yes
A/Japanese white eye/Hong Kong/1038/2006 (SJRG-164281)	2.3.4	SJCRH/HKU	Yes
A/chicken/Bangladesh/11rs1984-30/2011 (IDCDC-RG36)	2.3.4.2	CDC	Yes
A/Guizhou/1/2013 (IDCDC-RG35)	2.3.4.2	CDC/CCDC	Yes
A/Sichuan/26221/2014 (IDCDC-RG42A) (H5N6)	2.3.4.4	CDC/CCDC	Yes
A/gyrfalcon/Washington/41088-6/2014 (IDCDC-RG43A) (H5N8)	2.3.4.4	CDC	Yes
A/goose/Guizhou/337/2006 (SJRG-165396)	4	SJCRH/HKU	Yes
A/chicken/Viet Nam/NCVD-016/2008 (IDCDC-RG12)	7.1	CDC	Yes
A/chicken/Viet Nam/NCVY-03/2008 (IDCDC-RG25A)	7.1	CDC	Yes
A/environment/Hubei/950/2013	7.2	CDC/CCDC	Yes
Candidate vaccine viruses in preparation	Clade	Institution	Availability
A/chicken/Guizhou/1153/2016-like	2.3.2.1c	SJCRH/HKU	Pending
A/chicken/Ghana/20/2015-like	2.3.2.1c	CDC	Pending
A/chicken/Viet Nam/NCVD-15A59/2015-like (H5N6)	2.3.4.4	SJCRH	Pending
A/Hubei/29578/2016-like (H5N6)	2.3.4.4	CCDC	Pending
A/duck/Hyogo/1/2016-like (H5N6)	2.3.4.4	NIID	Pending

http://www.who.int/flu/vaccines/virus/201703_zoonotic_vaccinevirusupdate.pdf?ua=1




World Health Organization
WHO GISRS network

感染研GMP facilityでワクチン製造株の作製



動物への感染等で安全性試験
品質試験確認試験





ワクチン候補野生株の分与



国内外のワクチン製造所
研究機関

現在新型インフルエンザワクチン製造株
を作製・提供している海外施設

英国・国立生物製剤品質管理研究所
(NIBSC) BSL4実験施設

RG法で弱毒化したワクチン
高増殖株の開発

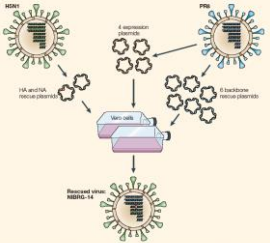


Figure 2 | Schematic diagram of the derivation of the H5N1 reference vaccine strain, NIBRG-04, by reverse genetics. The haemagglutinin (HA) and neuraminidase (NA) gene segments of the wild-

米国・セントジュード小児研究
病院 BSL3適合GMP施設

米国・疾病予防対策センター
(CDC) BSL3実験施設

感染研インフルエンザ
ウイルス研究センター

Zoonotic Influenza A viruses currently detected in Animals and Humans in the World (since Sept 2016)

● Avian viruses

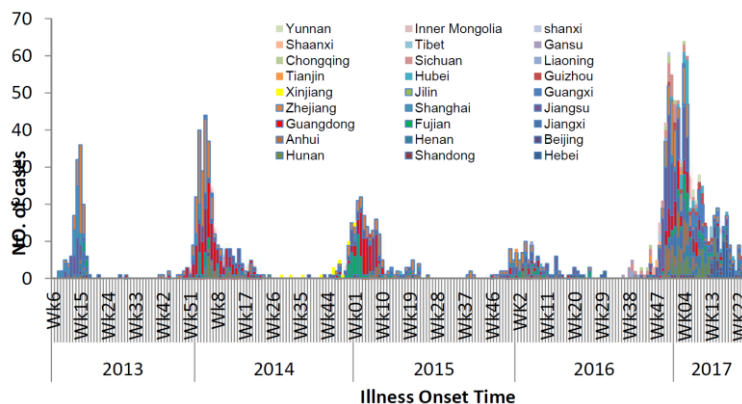


Since 2013,
H7N9 : (poultry, **human (758)**, environment) **(1564)**

http://www.who.int/influenza/vaccines/virus/201709_zoonotic_vaccinevirusupdate.pdf?ua=1

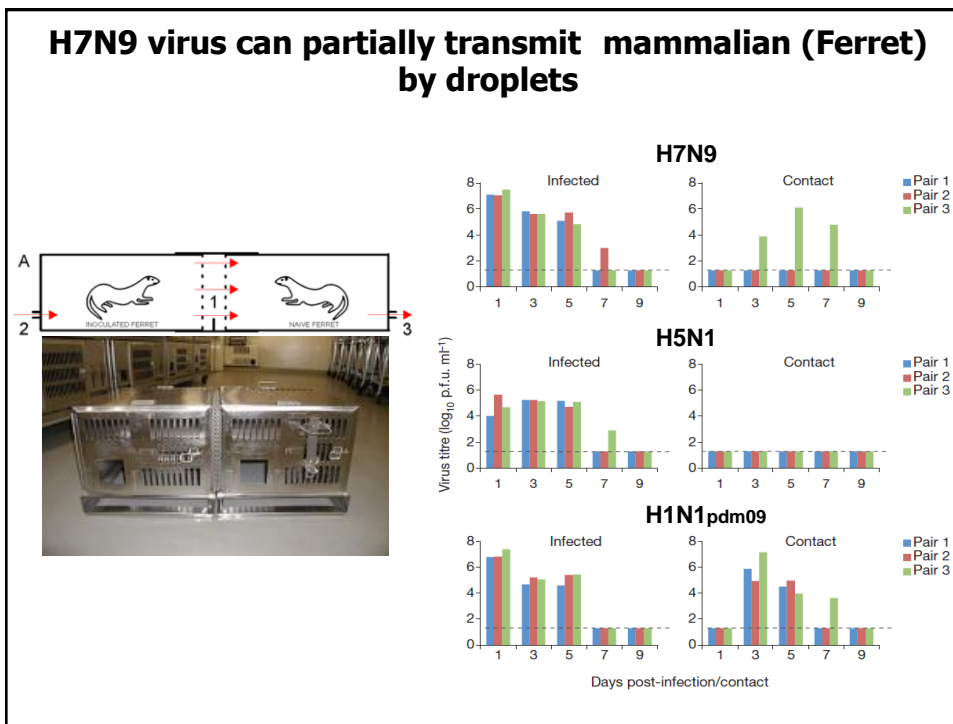
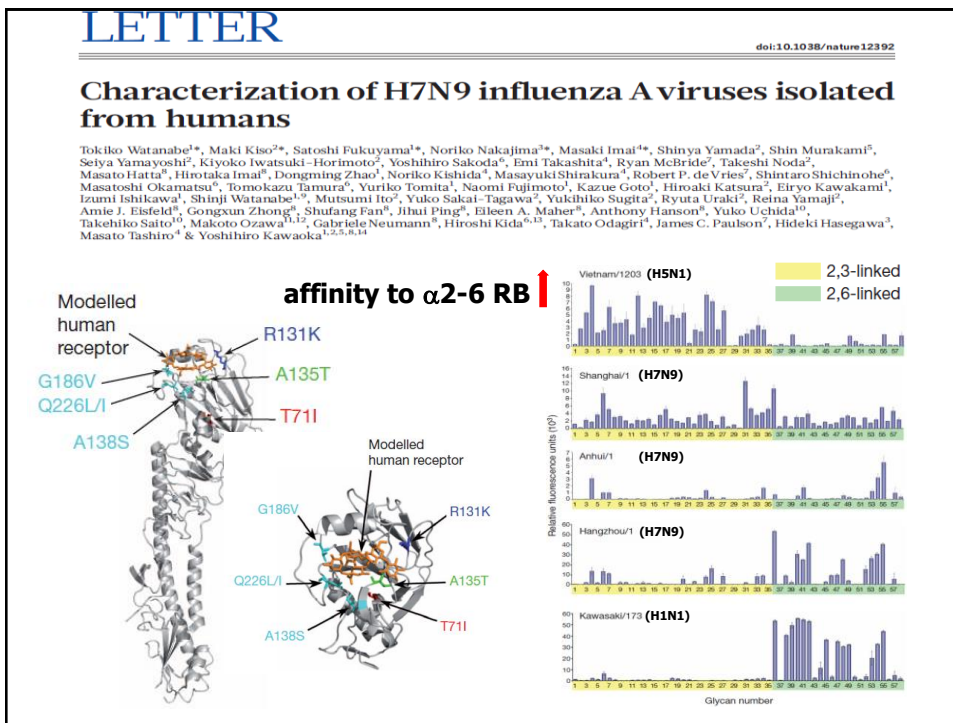
Epi-curve of human H7N9 cases by date of illness onset

19 Feb, 2013 to 31 Aug, 2017 (N=1531)



Province	First wave (2013.2-)		Second wave (2013.10-)		Third wave (2014.10-)		Forth wave (2015.10-)		Fifth wave (2016.10-)		Cumulative counts		
	Number of cases	Deaths	Number of cases	Deaths	Number of cases	Deaths	Number of cases	Deaths	Number of cases	Deaths	Number of cases	Deaths	Fatality rate(%)
Total	134	44	306	128	219	100	116	47	756	285	1531	604	39.5

Data provided by China CDC



Family clusters of human H7N9 cases

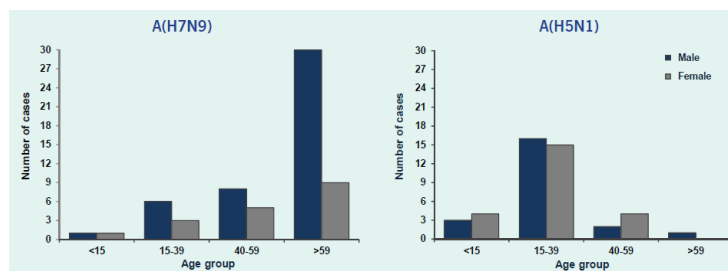
Route of transmission	NO.
Possibility of Human-to-human transmission	19+1
Co-exposure	3+4
Possibility of Human-to-human transmission or Co-exposure	9+4
Total	40

*New family clusters since Sep,2016 in red.

Data provided by China CDC

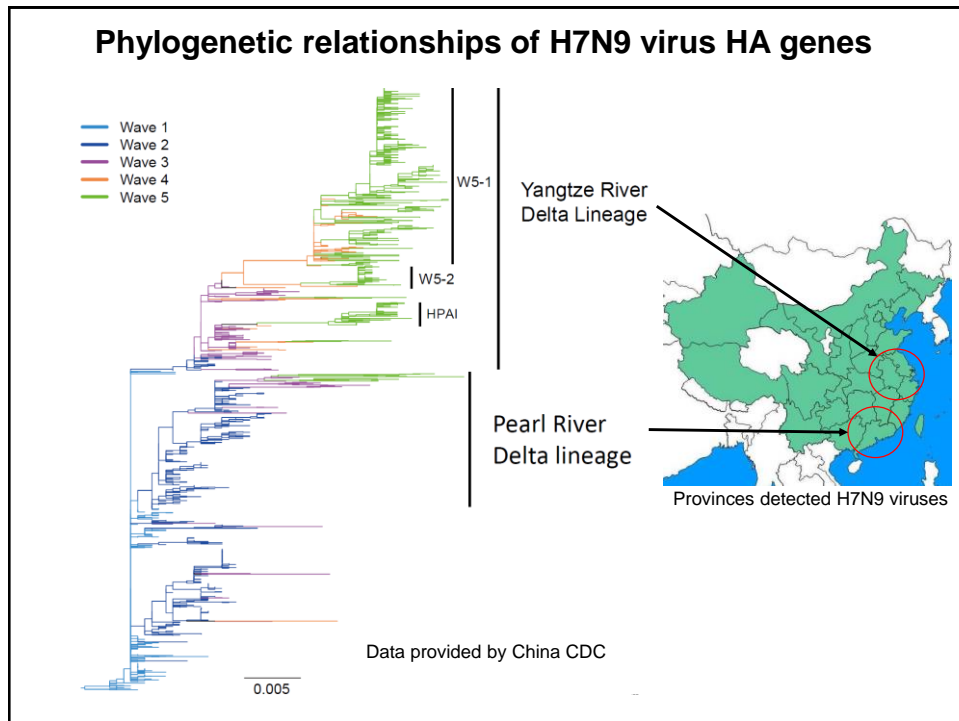
- Pandemic potential with H7N9 viruses is not low
- Sustained H-to-H transmission is limited so far

Age distribution of infected cases with A(H7N9) and A(H5N1) viruses



>45 years old: 75% of cases
<20 years old: 7% of cases

<20 years old: 50% of cases



Summary of A(H7N9) human cases reported to WHO since 27 September 2016

- 758 human cases of A(H7N9) virus infection (288 fatal) in 5th W
 - Total cases since 2013 = 1564 with 610 deaths
 - Chinese Taipei, Hong Kong Special Administrative Region and Macau Special Administrative Region reported travel-associated cases
 - CFR in reported cases = 39%
 - 68% exposed to live poultry or contaminated environments
 - 40 clusters confirmed, but not sustained human-to-human transmission
- The majority of A(H7N9) viruses detected in poultry and in humans are low pathogenic
- **Since 2016 high pathogenic A(H7N9) viruses were detected in poultry and in human (28 cases)**
- Most of H7N9 viruses recovered from patients were susceptible to antivirals of NA inhibitors

Available H7N9 CVVs provided by WHO CCs/ERLs

Table 5. Status of influenza A(H7N9) candidate vaccine virus development

Candidate vaccine virus	Type	Institution*	Available
A/Anhui/1/2013 (IDCDC-RG33A)	Reverse genetics	CDC	Yes
A/Anhui/1/2013 (NIBRG-268)	Reverse genetics	NIBSC	Yes
A/Anhui/1/2013 (NIIDRG-10.1)	Reverse genetics	NIID	Yes
A/Anhui/1/2013 (SJ005)	Reverse genetics	SJCRH	Yes
A/Shanghai/2/2013 (NIBRG-267)	Reverse genetics	NIBSC	Yes
A/Shanghai/2/2013 (CBER-RG4A)	Reverse genetics	FDA	Yes
A/Shanghai/2/2013 (IDCDC-RG32A)	Reverse genetics	CDC	Yes
A/Shanghai/2/2013 (IDCDC-RG32A.3)	Reverse genetics	CDC	Yes
Candidate vaccine viruses in preparation	Type	Institution	Availability
★ ● A/Guangdong/17SF003/2016-like	Reverse genetics	CCDC and NIBSC	Pending
★ ● A/Hunan/2650/2016-like	Reverse genetics	CCDC	Pending
★ ● A/Hong Kong/125/2017 (A/Hunan/2650/2016-like)	Reverse genetics	CDC and FDA	Pending

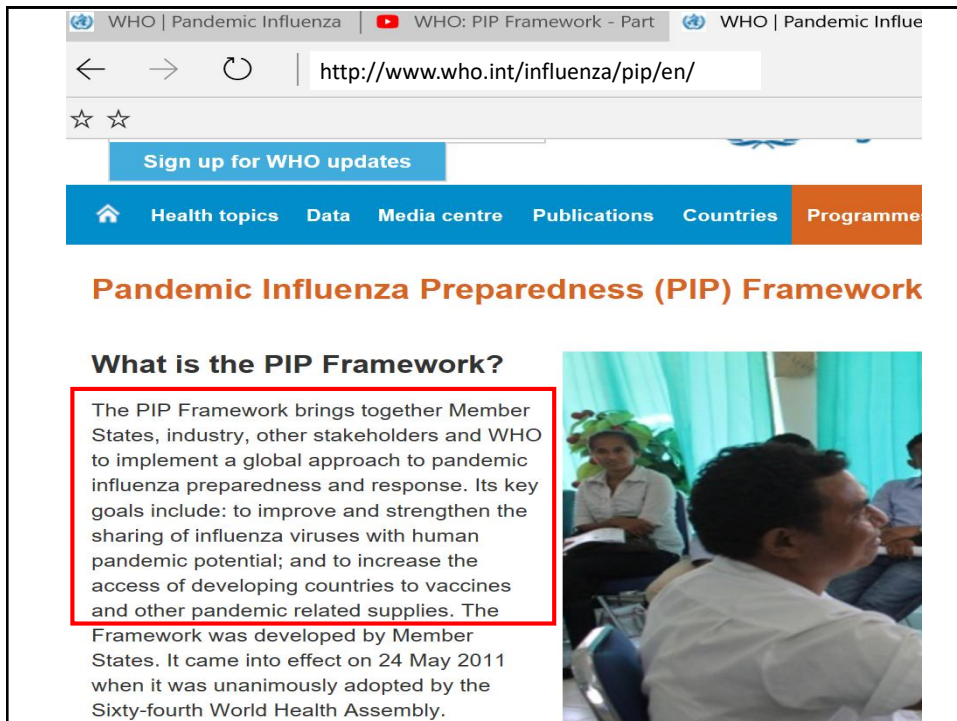
★ Yangtze river delta-lineage viruses (predominant-lineage)

● High-path virus

● Low-path virus

http://www.who.int/influenza/vaccines/virus/201703_zoonotic_vaccinevirusupdate.pdf?ua=1

To obtain CVVs and to use CVV for research and vaccine/diagnostic production, member states, industries, and stakeholders are required **Compliance of pandemic influenza preparedness (PIP)-FW**



WHO | Pandemic Influenza | WHO: PIP Framework - Part | WHO | Pandemic Influe

← → ↻ | <http://www.who.int/influenza/pip/en/>

☆ ☆


[Sign up for WHO updates](#)

🏠 [Health topics](#) [Data](#) [Media centre](#) [Publications](#) [Countries](#) [Programme](#)

Pandemic Influenza Preparedness (PIP) Framework

What is the PIP Framework?

The PIP Framework brings together Member States, industry, other stakeholders and WHO to implement a global approach to pandemic influenza preparedness and response. Its key goals include: to improve and strengthen the sharing of influenza viruses with human pandemic potential; and to increase the access of developing countries to vaccines and other pandemic related supplies. The Framework was developed by Member States. It came into effect on 24 May 2011 when it was unanimously adopted by the Sixty-fourth World Health Assembly.



WHO PIP-Framework is for;

- {
 - Virus sharing**
 - Equally access to vaccines and other benefits**

- **To get PIP-CVVs and reference viruses, members and stakeholders need to contract with WHO by S·MTA2**
 (<https://youtu.be/RMsR9ZbYN7I>)
- **Under the S·MTA2, members and stakeholders are required to response partnership contribution**
- **Partnership contribution is used for equal access of pandemic vaccine and for strengthen of pandemic preparedness**
 (<https://youtu.be/7M031gg1AnQ>)

