

OFFLU Technical Meeting

Coordinating world-wide surveillance for influenza in swine

University of Minnesota, Minneapolis, USA
March 19-20, 2014



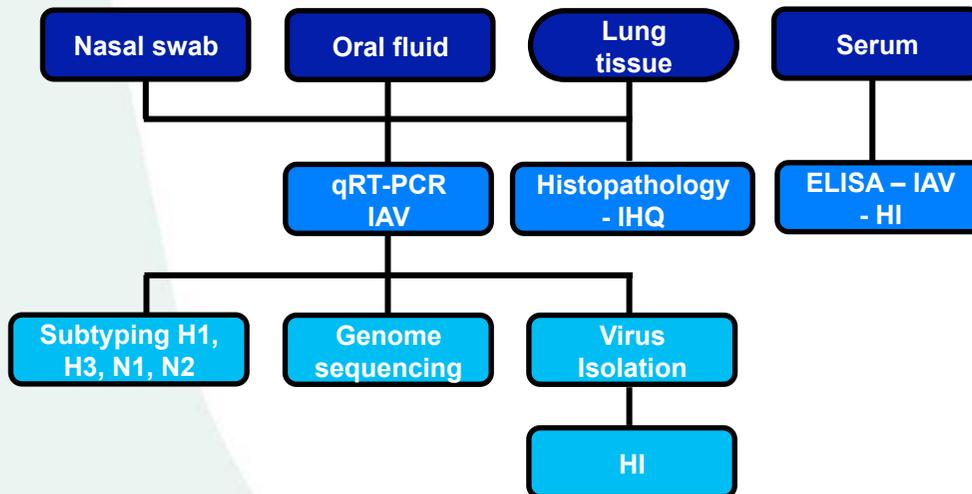
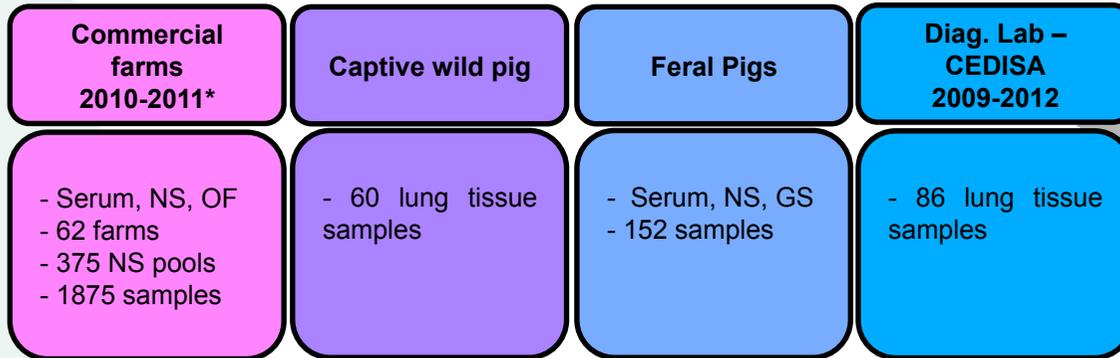
Brazil update

Janice Reis Ciacci Zanella

Brazilian Agricultural Research Corporation – EMBRAPA
Embrapa Swine and Poultry Research Center



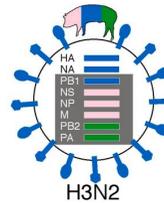
INFLUENZA A VIRUS INFECTION IN SWINE HERDS IN BRAZIL IN 2009 – 2012



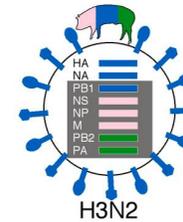
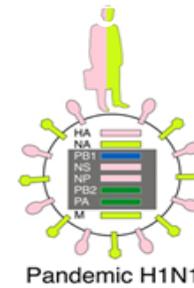
*87% of swine production

❖ 13 Farrow-to-finish farms (HI results):

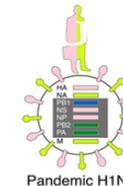
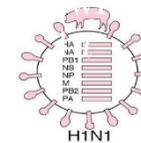
❖ Antibodies against **H3N2**: 4 farms



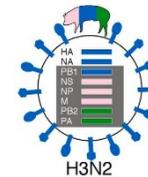
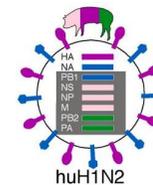
❖ Antibodies against **H1N1pdm09** and **H3N2**: 5 farms



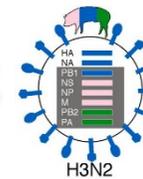
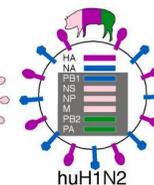
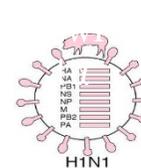
❖ Antibodies against **H1N1**, **H1N1pdm09** and **H3N2**: 2 farms



❖ Antibodies against **H1N2** and **H3N2**: 1 farm



❖ Antibodies against **H1N1**, **H1N2** and **H3N2**; 1 farm



Viral isolation samples

» **From lungs or nasal swabs: total of 68 IAV**

» Nasal swabs: 107/10, 83/10, 72/11.

Total: 03 samples

» Lungs: 104/09, 12a/10, 83/10, 89/10, 107/10, 131/10 (2 samples), 136/10, 170b/10 (2 samples), 170c/10, 170d/10, 170e/10 (3 samples), 170f/10, 170h/10, 31/11 (2 samples), 66/11, 70/11, 71/11, 85/11, 95/11, 138/11, 146/11, 146b/11, 152/11, 173/11, 18/12, 37/12 (6 samples), 42/12, 198/12 (10 samples), 263/12, 93/13, 119c/13, 137c/13, 183/13 (5 samples). **Total: 65 samples**

** In green = genome sequencing data*

Genome Sequencing

- » Sequencing data for HA, M and NA (some genomes with complete genome, other partial sequencing).

37 samples sequenced:

- » **HA, NA and M segments: 18 samples**
- » **NA and M segments: 10 samples**
- » ***Eight gene segments: 9 samples.**

**Illumina platform (MiSeq)*

Genomic Sequencing of HA, NA and M - 2009-2011

IAV Sample	Subtype	HA	NA	M
A/swine/Brazil/107-3A/2010	H1N1	H1N1pdm09	H1N1pdm09	H1N1pdm09
A/swine/Brazil/72-11-507/2011	H1N1	H1N1pdm09	H1N1pdm09	H1N1pdm09
A/swine/Brazil/132-09/2009	H1N1	H1N1pdm09	H1N1pdm09	H1N1pdm09
A/swine/Brazil/12A10/2010	H1N1	H1N1pdm09	H1N1pdm09	H1N1pdm09
A/swine/Brazil/136-10/2010	H1N1	H1N1pdm09	H1N1pdm09	H1N1pdm09
A/swine/Brazil/173-11-4/2011	H1N1	H1N1pdm09	H1N1pdm09	H1N1pdm09
A/swine/Brazil/170h-10/2009	H1N1	H1N1pdm09	H1N1pdm09	H1N1pdm09
A/swine/Brazil/31-11-1/2011	H1N2	H1 δ	N2	H1N1pdm09
A/swine/Brazil/31-11-3/2011	H1N2	H1 δ	N2	H1N1pdm09
A/wild boar/Brazil/214-11-13D/2011	H1N2	H1 δ	N2	H1N1pdm09
A/swine/Brazil/185-11-7/2011	H1N2	H1 δ	N2	H1N1pdm09
A/swine/Brazil/232-11-13/2011	H1N2	H1 δ	N2	H1N1pdm09
A/swine/Brazil/232-11-14/2011	H1N2	H1 δ	N2	H1N1pdm09
A/swine/Brazil/365-11-6/2011	H3N2	H3	N2	H1N1pdm09
A/sw/Brazil/355-11-6/2011	H3N2	H3	N2	H1N1pdm09
A/sw/Brazil/365-11-7/2011	H3N2	H3	N2	H1N1pdm09
A/swine/Brazil/231-11-1/2011**	H3N2	H3	N2	H1N1pdm09

**In the same farm it was identified H3N2, H1N1pdm09 and H1N2

Feral pigs

and

Captive wild boar



Captive wild boars



Contents lists available at ScienceDirect

Veterinary Microbiology

journal homepage: www.elsevier.com/locate/vetmic



- HA and NA δ cluster
- Internal genes derived from H1N1pdm09

Genomic analysis of influenza A virus from captive wild boars in Brazil reveals a human-like H1N2 influenza virus

Natalha Biondo^a, Rejane Schaefer^{b,*}, Danielle Gava^b,
Mauricio E. Cantão^b, Simone Silveira^b, Marcos A.Z. Mores^b,
Janice R. Ciacci-Zanella^b, David E.S.N. Barcellos^a

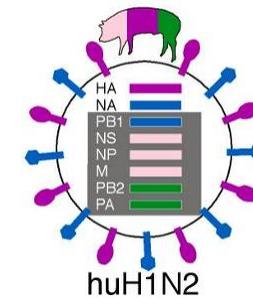
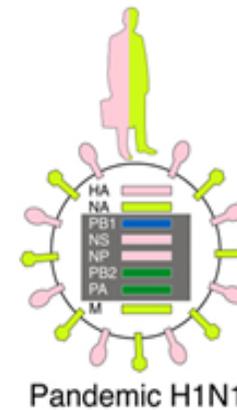
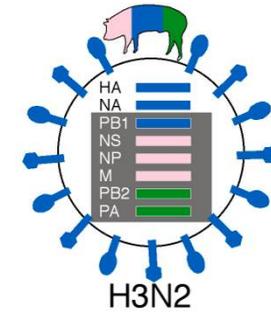
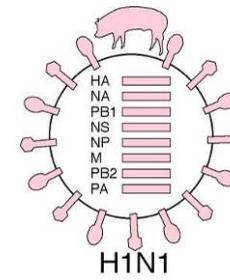
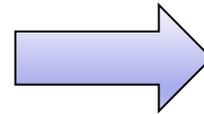
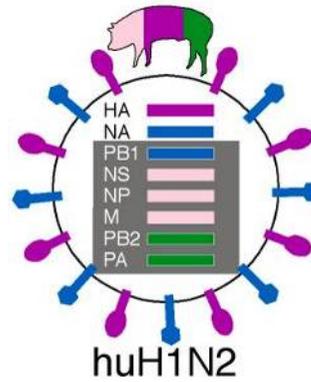
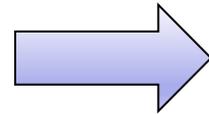
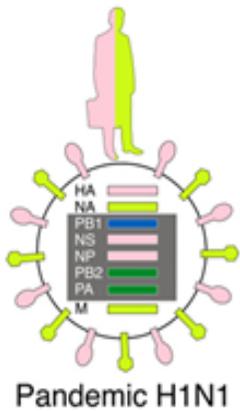
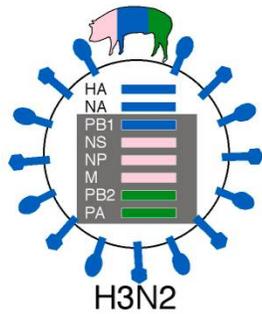
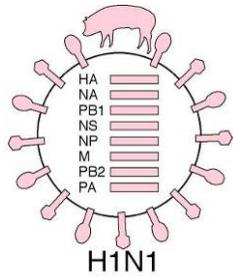
Table 1

Most closely relatives to A/wild boar/Brazil/214-11-13D/2011 determined by BLAST search at NCBI (<http://www.ncbi.nlm.nih.gov/blast/Blast.cgi>).

Gene	GenBank accession number	Isolate	Subtype	Identity (%)
PB2	CY122668.1	A/Singapore/GP1132/2009	H1N1	99
	CY122700.1	A/Singapore/GP1146/2009	H1N1	99
PB1	CY055303.1	A/Singapore/GN285/2009	H1N1	99
	CY053625.1	A/Russia/165/2009	H1N1	99
PA	KC833448.1	A/swine/Thailand/UD400/2009	H1N1	99
	CY045233.2	A/Taiwan/126/2009	H1N1	99
HA	CY125172.1	A/New York/26/2002	Mixed	97
	CY003696.1	A/New York/489/2003	H1N2	97
NP	CY045235.2	A/Taiwan/126/2009	H1N1	99
	HQ728111.1	A/swine/Taiwan/CH-1204/2009	H1N1	99
NA	AF533744.1	A/Neuquen/V690/98	H3N2	96
	AF533741.1	A/Cordoba/V391/98	H3N2	96
M	CY069644.1	A/Singapore/527/2009	H1N1	99
	CY069629.1	A/Singapore/471/2009	H1N1	99
NS	CY050371.1	A/Korea/S1/2009	H1N1	100
	GU108490.1	A/Zhejiang-Yiwu/11/2009	H1N1	100

Before 2010

After 2010





Thank you

janice.zanella@embrapa.br



Ministry of
Agriculture, Livestock
and Food Supply

