



Instituto Nacional de Tecnología Agropecuaria  
Centro de Investigación en Ciencias Veterinarias y Agronómicas  
Instituto de Virología "Dr. Schlein Rivenson"



UNIVERSITY OF  
MARYLAND



# Swine Influenza in South and Central America

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Institute of Virology - INTA

Swine Influenza  
Technical Meeting  
Minnesota, March 19<sup>th</sup> 2014



# Last Technical Meeting (Rome, 2013)



## **Colombia** (Universidad Nacional de Colombia)

Seroprevalence aprox. 50% Cauca Valley region and Antioquia

Isolates: pdm H1N1, probably cH1N1

## **Chile**

Overall seropositivity of 48% to H1N1 strains and 22% to H3N2 strains

Producers have began vaccinating sows (during the gestation and maternity period)

Isolates: cH1N1, pdm H1N1, swH3N2

## **Guatemala**

Overall seropositivity of 19% to NP Elisa

Isolates: pdm H1N1, huH3N2

## **Argentina** Viruses isolated till 2012

Differential seroprevalence between farms depending in the infection status

HIs: 90% pdmH1N1, 10% H3

Isolates: pdmH1N1, huH3N2 + pdm internal genes,  $\delta$ 2 H1N1 + pdm internal genes,  $\delta$ 1 H1N2 + pdm internal genes

# South and Central America SIV Surveillance



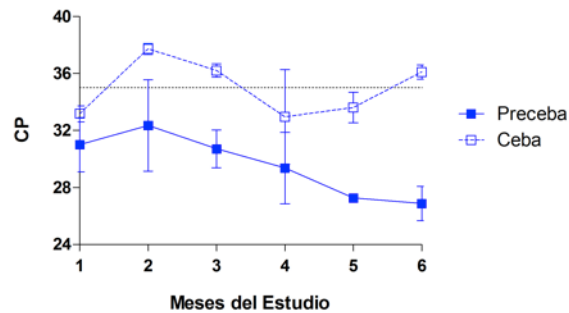
# Colombia

- Started to implement Oral Fluid analysis
- SIV, PRRS and PCV2 by Real Time PCR

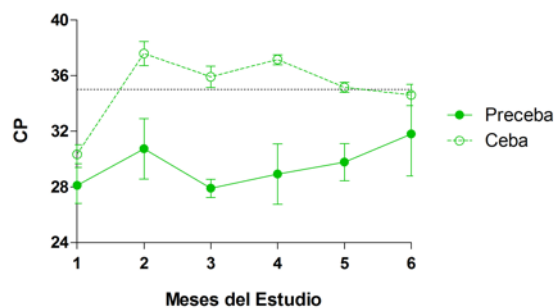


## Post weaning vs fattening

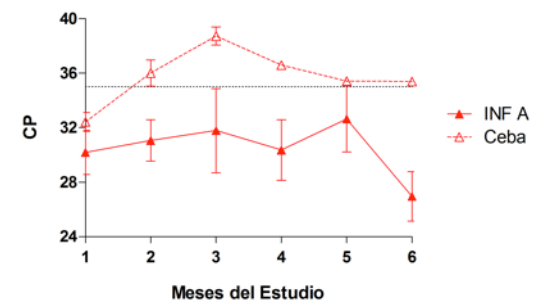
Variación del CP entre grupos en la Granja A



Variación del CP entre grupos en la Granja B



Variación del CP entre grupos en la Granja C





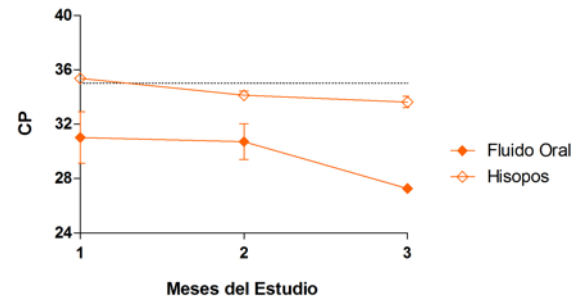
# Post weaning vs fattening + Swabs vs OF



Farm A

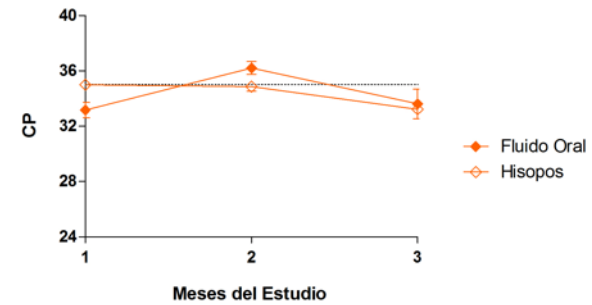
## Post weaning

Variación del CP entre pruebas en la granja A en el grupo Preceba



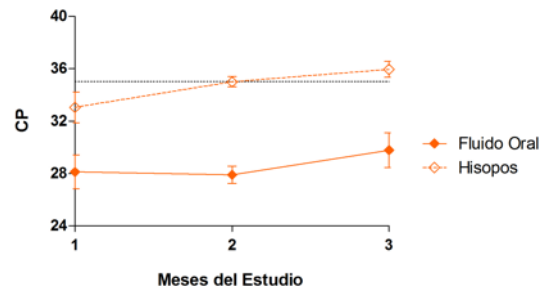
## Fattening

Variación del CP entre pruebas en la granja A en el grupo Ceba

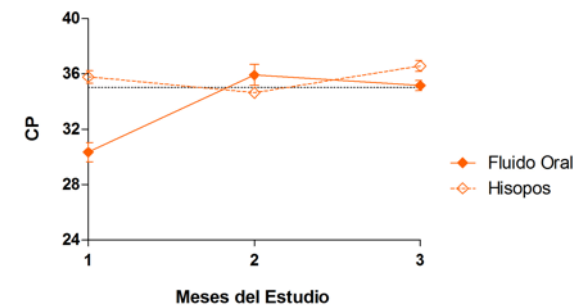


Farm B

Variación del CP entre pruebas en la granja B en el grupo Preceba

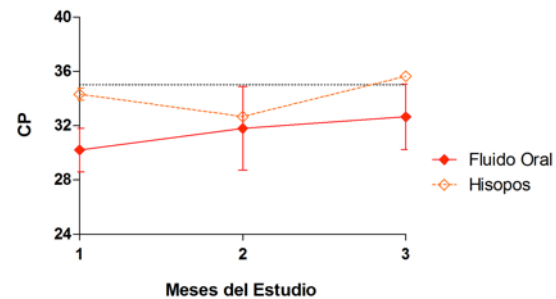


Variación del CP entre pruebas en la granja B en el grupo Ceba

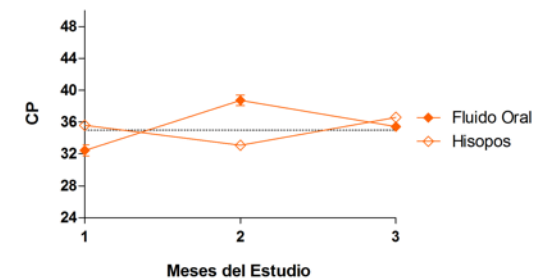


Farm C

Variación del CP entre pruebas en la granja C en el grupo Preceba



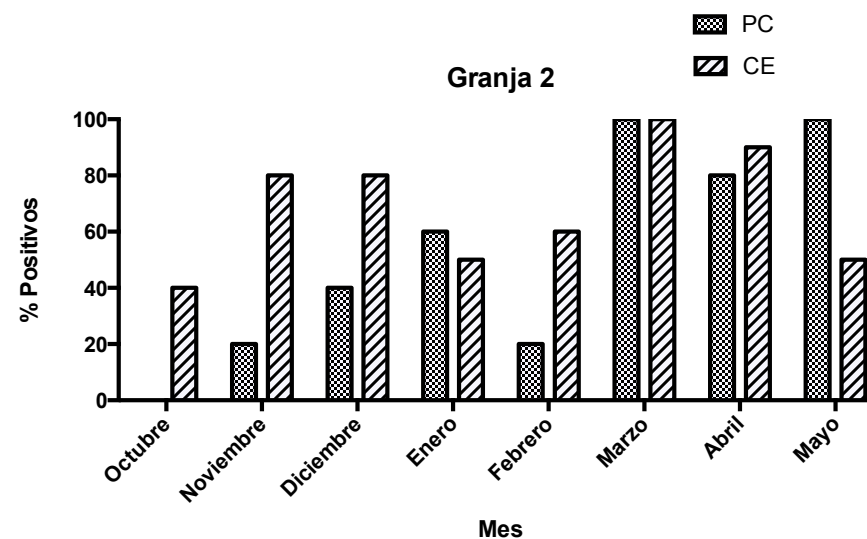
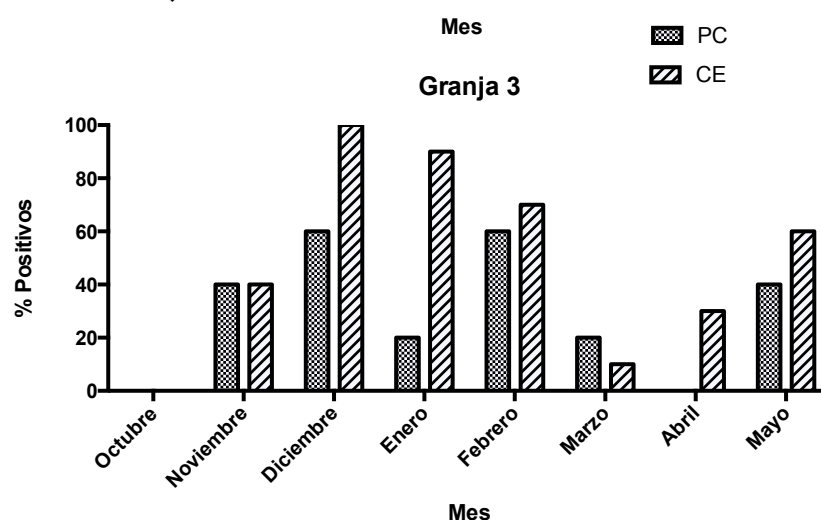
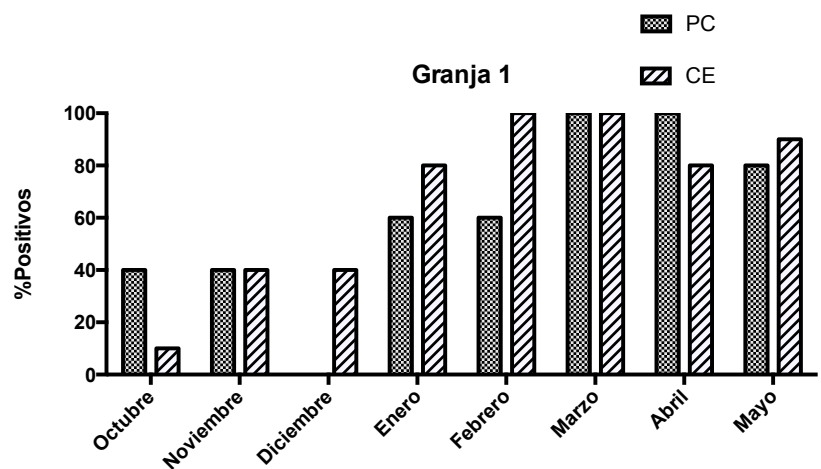
Variación del CP entre pruebas en la granja C en el grupo Ceba





# Serology

## Post weaning vs fattening + Seasonality



No evident seasonality



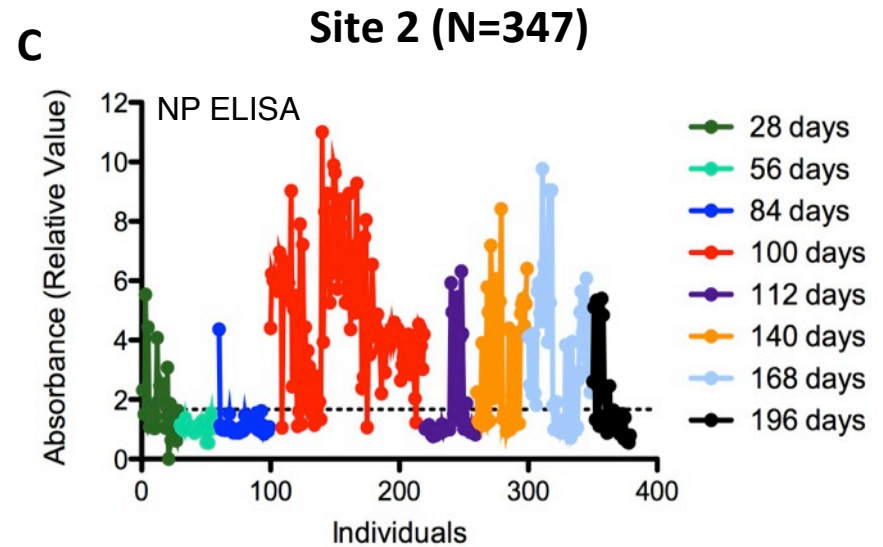
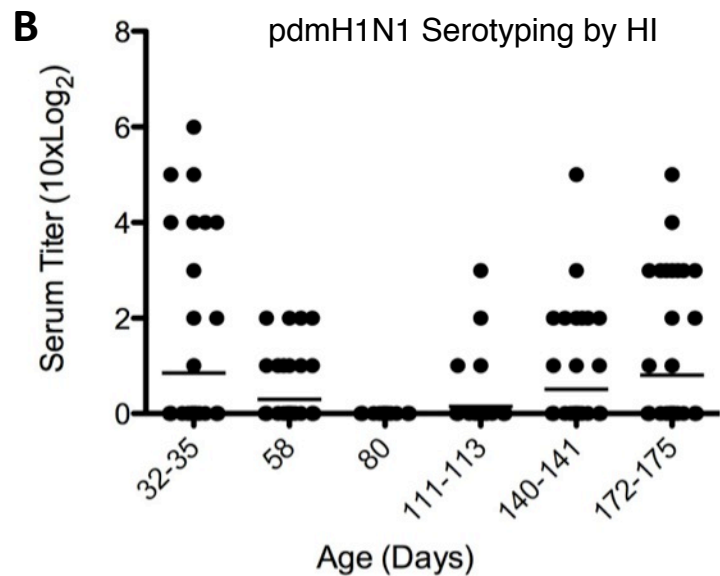
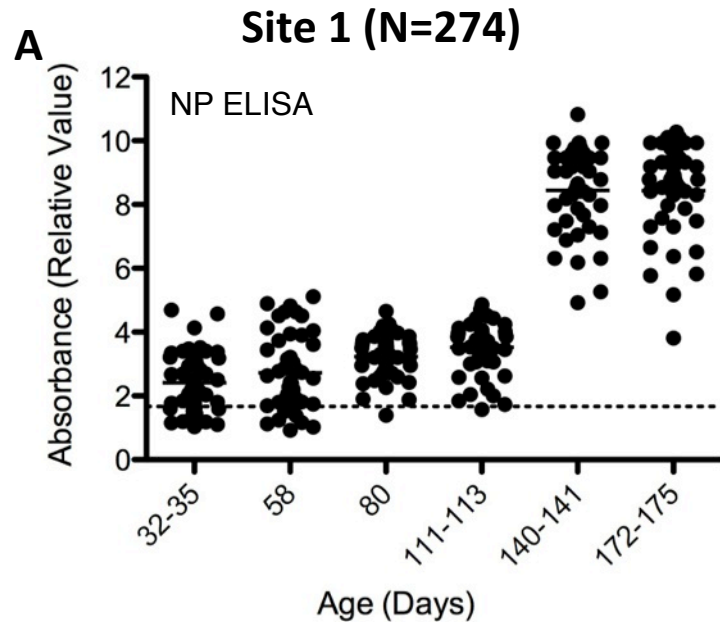
## Circulation of swine influenza viruses in Chile

- Since 2010 four different strains of swine influenza viruses have been identified to be circulating in Chile: pdmH1N1-like, H1N1, H1N2 and H3N2.
- A serological study done in 2009 of 13 production sites revealed an overall seropositivity of 48% to H1N1 strains and 22% to H3N2 strains.\*
- Recent studies by our group in collaboration with ASPROCER (Association of Swine Producers of Chile), have found swine farms with seropositivity to Influenza A virus ranging from 53% - 90.5%.
- High seroprevalence to influenza was detected in animals 100 days old and older, with the main susceptible/seronegative population identified in animals of 56-84 days.

\* Unpublished data, **Dr. Alvaro Ruiz**, University of Concepcion



# Age distribution of seropositive animals in two farms in central Chile



- Seroprevalence to pdmH1N1 ranges from 18% - 51.7%.
- Little cross-reactivity to classical swine strains.
- Serotyping to other strains currently ongoing.





## Circulation of swine influenza viruses in Chile

- During 2011 and 2012 part of the swine industry in Chile began to use a commercial adjuvanted vaccine made in the United States, containing five strains.
- Current serological data indicate the circulation of at least 2 influenza virus strains in most production sites, with pdmH1N1 being highly prevalent.
- HI data suggest potential drifted pdmH1N1 strains (currently under study).
- The antigenic diversity and lack of cross-reactivity of the commercial vaccine strains and the current strains circulating in the country, motivated an effort to design and make homologous vaccines.
- Current mitigation strategies in most industrialized production farms grouped by ASPROCER include the use of a homologous US made vaccine, which contains 4 components.



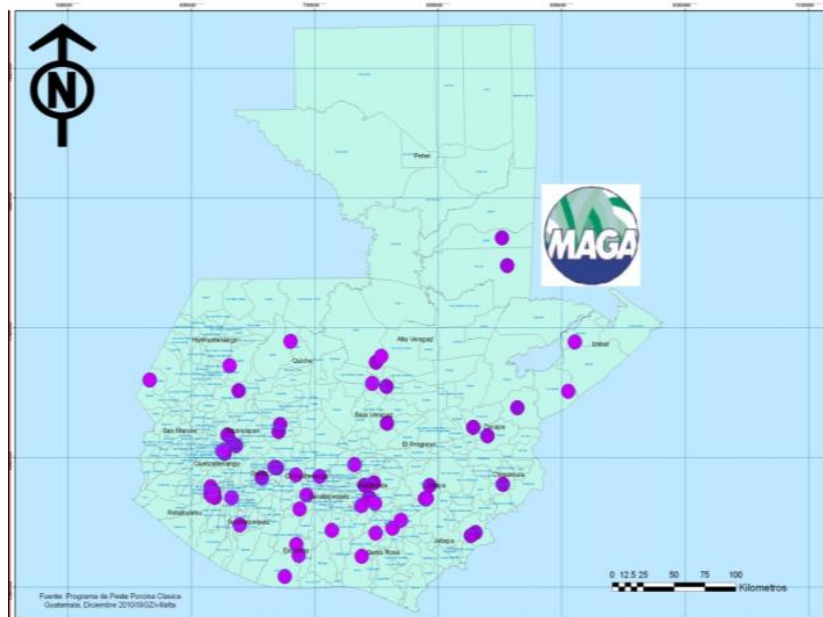
# Guatemala



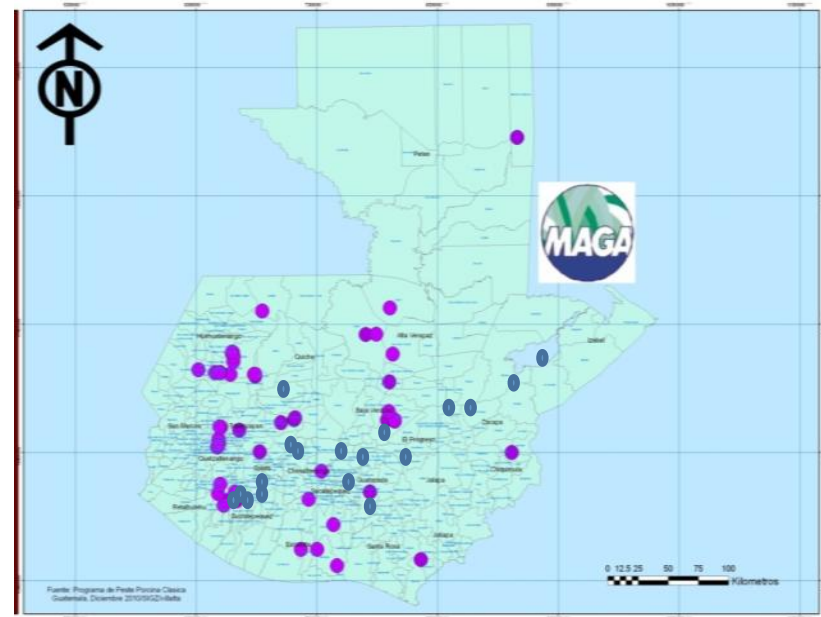
Nation-wide cross-sectional survey  
multi-stage random sampling:  
2010 – October (n=500) (MAGA, FAO)  
2011 – June to August (n=499) (MAGA)

- Case definition: Farms or backyard pigs (pig productions units) where  $\geq 10\%$  of the exposed population have respiratory clinical signs
  - Nasal swabs (Virus detection by rRT-PCR)
  - Serum samples (Antibody detection by ELISA and HI)

## Commercial farms



## Backyard pigs



**Spatial analysis.** Global methods were used to detect the presence of purely spatial clusters of positive farms (Ripleys K function)

2) Vigilancia de influenza porcina: en 2013 no se contó con fondos asignados para continuar el muestreo en cerdos. Sin embargo, en el laboratorio en Guatemala se implementó el cultivo de células MDCK y los protocolos para aislamiento viral a partir de muestras de hisopados nasales. Así mismo se implementó el ensayo de inhibición de hemaglutinación para la identificación de anticuerpos hace distintos grupos antigénicos de influenza H1 y H3.

Las muestras colectadas en 2011 y 2012, positivas para Influenza tipo A por la prueba de rRT-PCR en tiempo real, fueron probadas para aislamiento viral. No se obtuvieron aislados virales. Los resultados fueron presentados a las autoridades de Salud Animal del Ministerio de Agricultura, Ganadería y Alimentación (MAGA) y discutidos para la mejora de futuros estudios en colaboración.



# Estimated virus prevalence was comparable between years

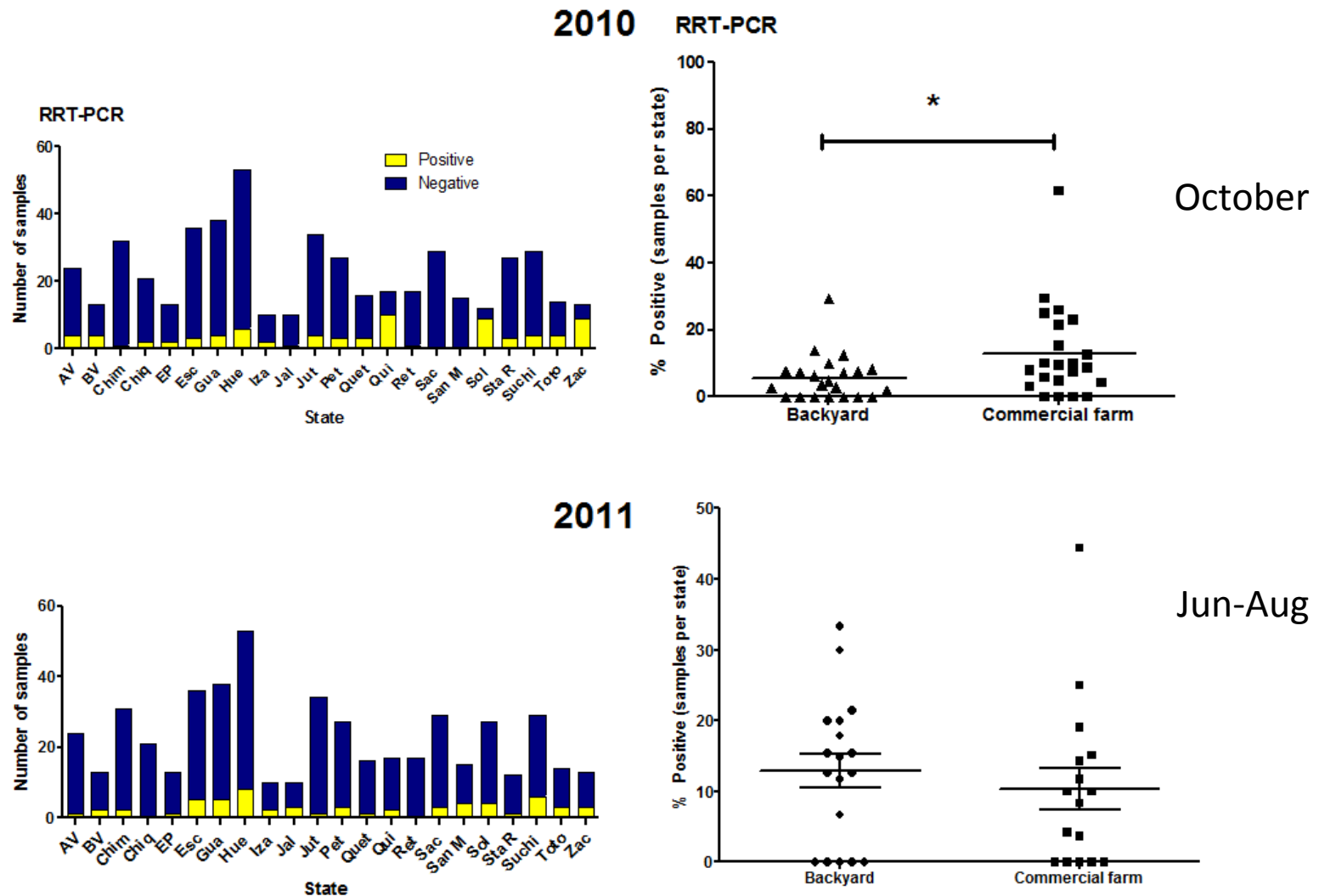
RRT-PCR estimated Influenza A prevalence in sampled farms from Guatemala

	Prevalence		Total Average
	2010 (n=500)	2011 (n=499)	
<b>RRT-PCR</b>			
Influenza A positive	16%	12%	14%
Sick positive animals	17%	9%	13%
Healthy positive animals	10%	12%	11%
<b>IDEXX ELISA</b>			
Positive for influenza A antibodies	10%	1%	6%
Sick positive animals	9%	0%	4%
Healthy positive animals	36%	2%	19%

(A. González-Reiche, Unpublished)



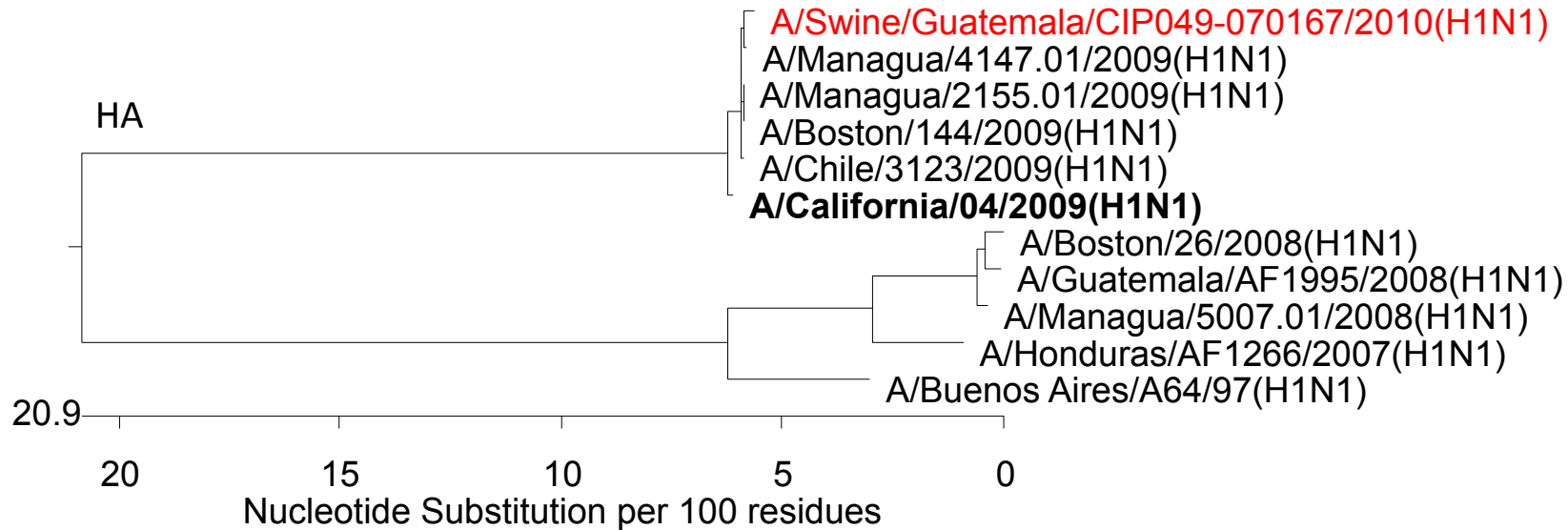
# Influenza A RRT-PCR prevalence in pigs varies between states and type of production unit





## The viruses...

fully pandemic H1N1 viruses were isolated from pigs



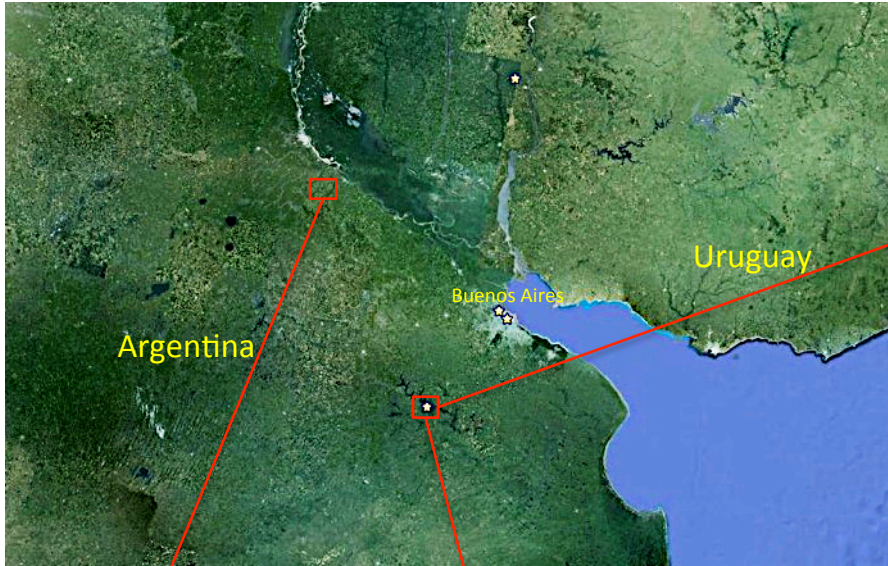
H3N2 isolates are related to seasonal human viruses

### A/Swine/Guatemala/CIP049-040078/2010(H3N2)

Segment	Most Similar Sequence in BLAST	%ID
PB2	A/Mexico City/WRAIR3580T/2010(H3N2)	99%
PB1	A/Mexico City/WRAIR3577T/2010(H3N2)	98%
PA	A/Thailand/CU-B657/2009(H3N2)	99%
HA	A/California/NHRC0004/2011(H3N2)	99%
NP	A/Mexico City/WRAIR4139N/2010(H3N2)	97%
NA	A/Mexico City/WRAIR4139N/2010(H3N2)	97%
M	A/Uganda/MUWRP-070/2009(H3N2)	100%
NS	A/Mexico/UASLP-013/2008(H3N2)	99%

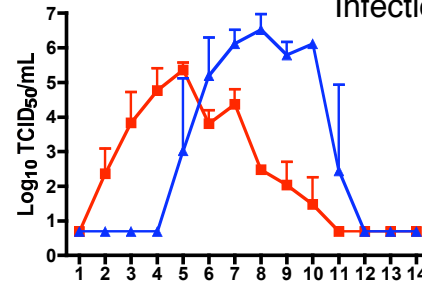
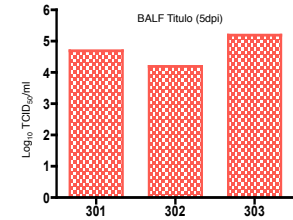
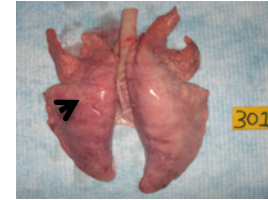


# ARGENTINA



Evidence of a non-contemporary human H3N2 adapted to be transmitted among pigs

Farm A  
Buenos Aires  
6000 sows  
Nov-2008



Infection and Transmission of A2/08 (H3N2)

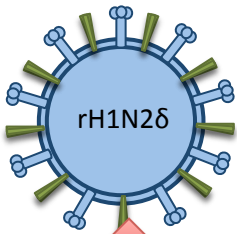
—■— Infected  
—▲— Direct Contact

Journal of General Virology (2011), 92, 2871–2878 DOI: 10.1099/vir.0.08590-0

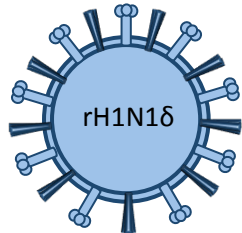
Outbreak of swine influenza in Argentina reveals a non-contemporary human H3N2 virus highly transmissible among pigs

Javier A. Cappuccino,<sup>1†</sup> Lindomar Pena,<sup>2†</sup> Marina Dibarbora,<sup>3</sup> Agustina Rimondi,<sup>3</sup> Pablo Pflieger,<sup>1</sup> Lucas Insarralde,<sup>1</sup> María A. Quiroga,<sup>1</sup> Mariana Machuca,<sup>1</sup> María I. Craig,<sup>3</sup> Valeria Olivera,<sup>3</sup> Ashok Chockalingam,<sup>3</sup> Carlos J. Perfumo,<sup>1</sup> Daniel R. Perez<sup>2</sup> and Ariel Pereda<sup>3</sup>

Farm B  
Santa Fe  
4000 sows  
May-2010



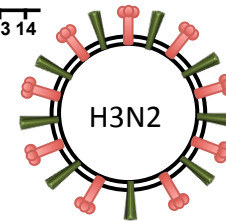
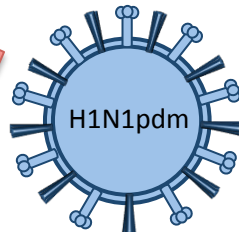
Farm C  
Buenos Aires  
9000 sows  
Oct-2009



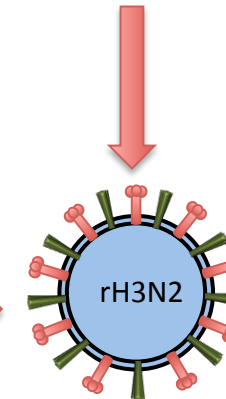
DISPATCHES

**Pandemic (H1N1) 2009 Outbreak on Pig Farm, Argentina**

Ariel Pereda,<sup>1</sup> Javier Cappuccino,<sup>1</sup> María A. Quiroga, Elsa Baumeister, Lucas Insarralde, Mariela Ibar, Ramón Sanguinetti, María L. Cannilla, Débora Franzese, Oscar E. Escobar Cabrera, María I. Craig, Agustina Rimondi, Mariana Machuca, Rosa T. Debenedetti, Carlos Zenobi, Leonardo Barral, Rodrigo Balzano, Santiago Capalbo, Adriana Rizzo, and Carlos J. Perfumo



Farm D  
Buenos Aires  
9000 sows  
Jan-2012



Recombinant H3N2 + pdm

Influenza Other Respi. Viruses, 2011, Nov;5(6):408-12. doi: 10.1111/1750-2659.2011.00246.x. Epub 2011 Apr 7.

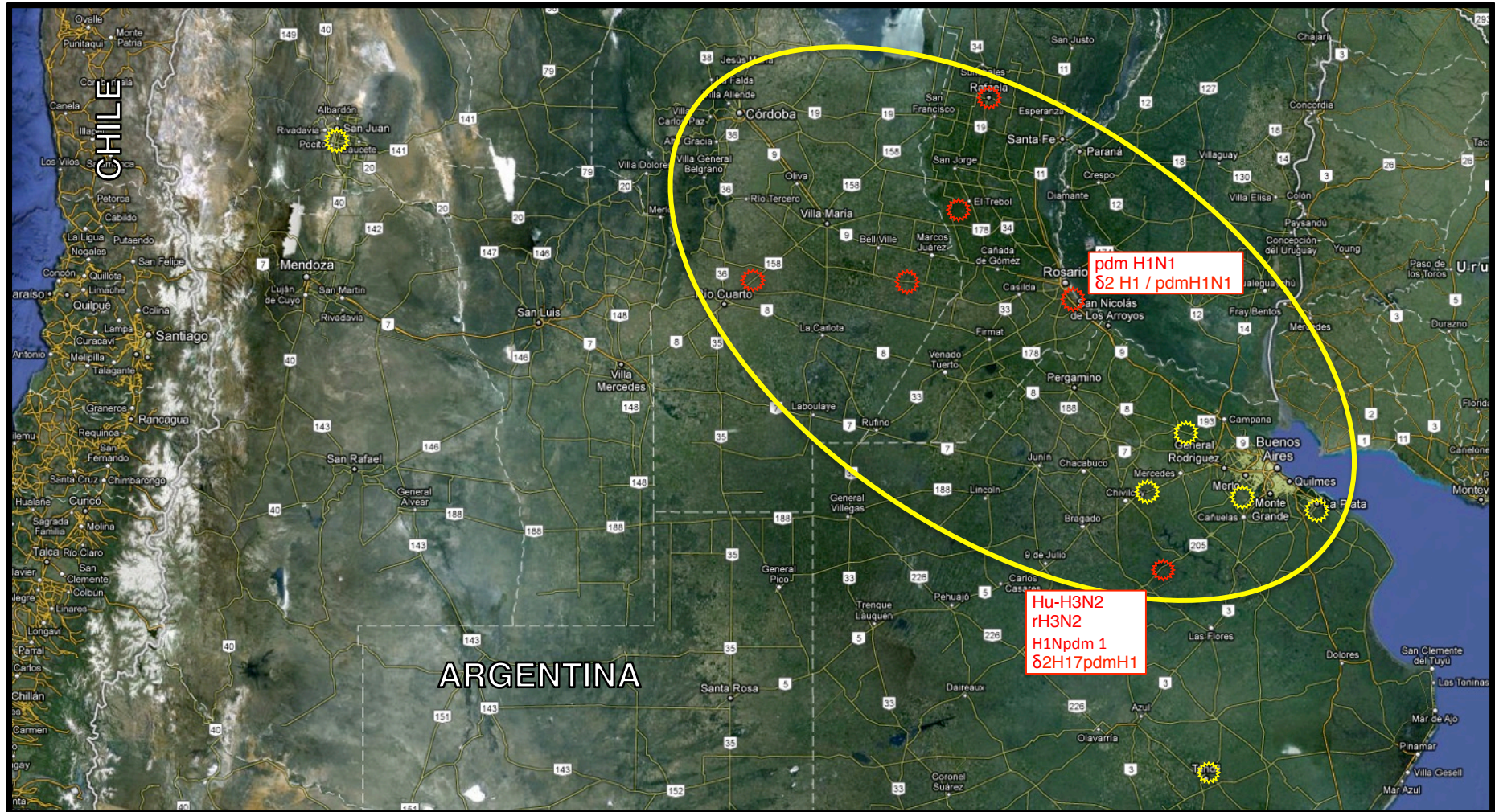
Evidence of reassortment of pandemic H1N1 influenza virus in swine in Argentina: are we facing the expansion of potential epicenters of influenza emergence?



Pereda A, Rimondi A, Cappuccino J, Sanguinetti B, Angel M, Ye J, Sutton T, Dibarbora M, Olivera V, Craig M, Quiroga M, Machuca M, Ferrero A, Perfumo C, Perez DR.

Laboratorio Aves y Porcinos, Instituto de Virología CICVyA-Instituto Nacional de Tecnología Agropecuaria, Buenos Aires, Argentina. apereda@cnia.inta.gov.ar



# GITEP study 2012 Sampling sites



-  rRT-PCR/isolate = Positive - Serology = Positive
-  rRT-PCR/isolate = Negative - Serology = Positive

## Analyzed samples

- 1199 nasal and/or bronchial swabs
  - 59 lungs
- 10% of sows stock in Argentina





# 2012 : Estudio del comportamiento de la infección por el virus de influenza en cerdos previa implementación de vacunas en Argentina.

10% del stock de hembras del país

9 granjas

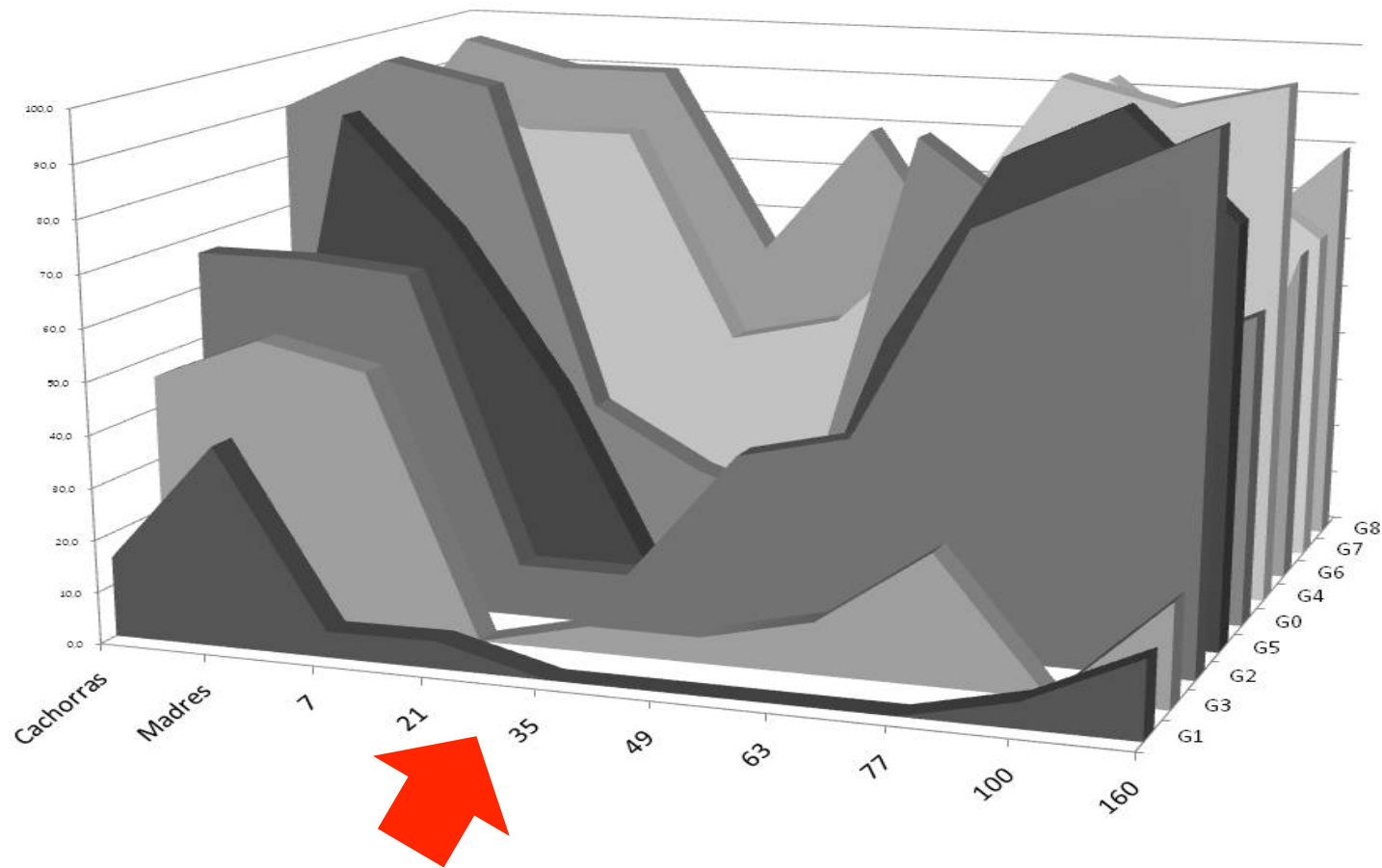
(Santa Fe, Córdoba, Buenos Aires y San Luis)



Granja	Signos clínicos	Edad (días)	Aislamiento
G0	SI	63	SI (rH3N2)
G1	NO		NO
G2	SI	49 y 63	SI (pH1N1)
G3	SI	35 y 49	SI (pH1N1)
G4	SI	21 y 140	SI (pH1N1)
G5	SI	50 y 120	NO
G6	SI	25	NO
G7	SI	35	SI (pH1N1)
G8	SI	35	NO

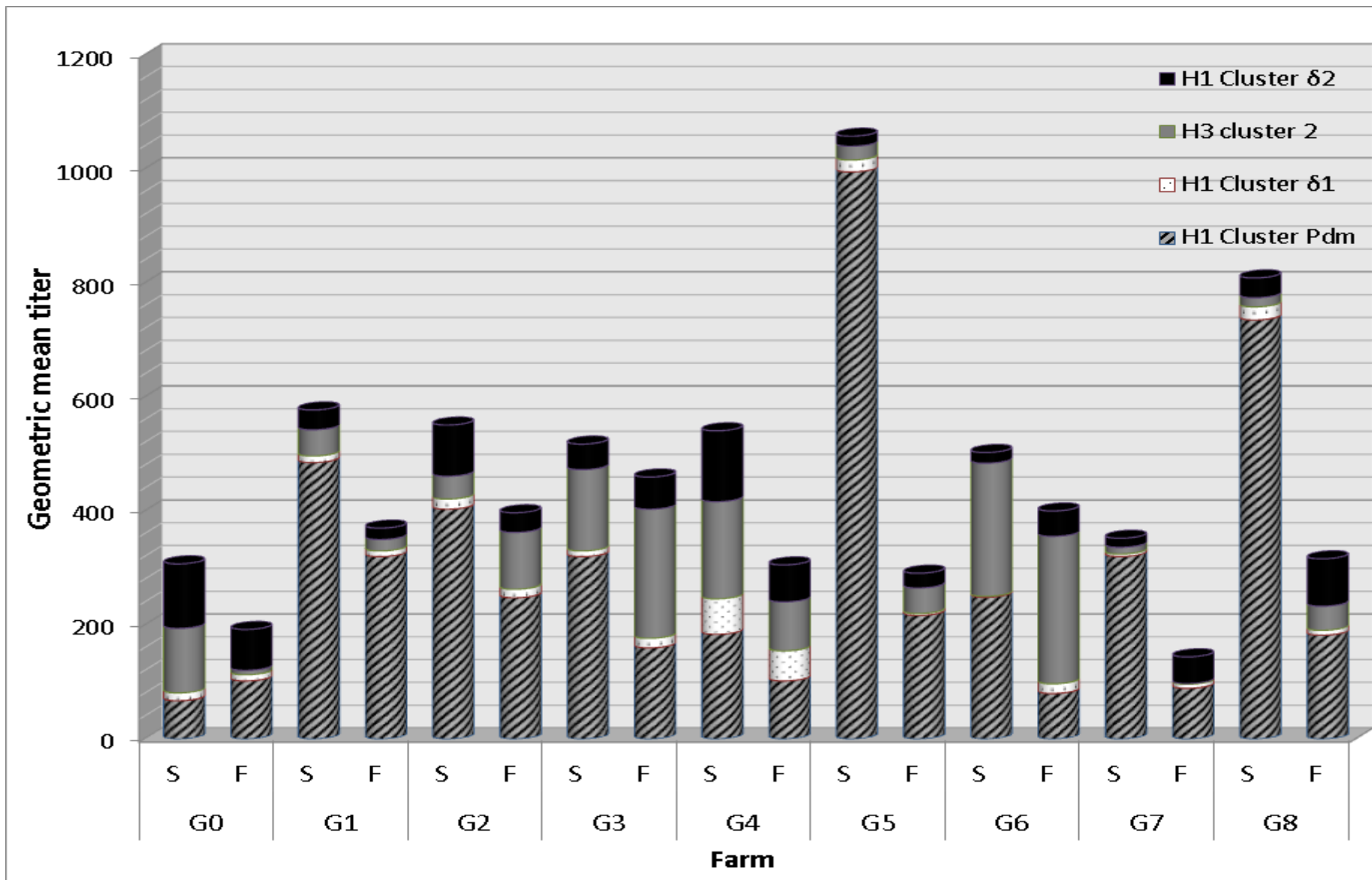
8/9 granjas signos clínicos en destete  
2/9 granjas signos clínicos en engorde

Granja	Lesiones Macroscópicas	Lesiones Microscópicas	rt PCR +	Aislamiento
G2	2/9	3/9	1/2	1
G4	4/7	3/7	1/3	1
G7	1/1	1/1	1/1	1
G8	5/17	1/5	1/3	0



- 48% del total de los sueros positivos
- Se observó una caída de anticuerpos calostrales hacia los 21 a 35 días y un aumento entre los 49 y los 100 días.
- El porcentaje de cerdas positivas varió entre 25-100%, mientras que en el engorde se observaron porcentajes de entre 15 a 100% de animales positivos.

# Subtipos circulantes (HI)

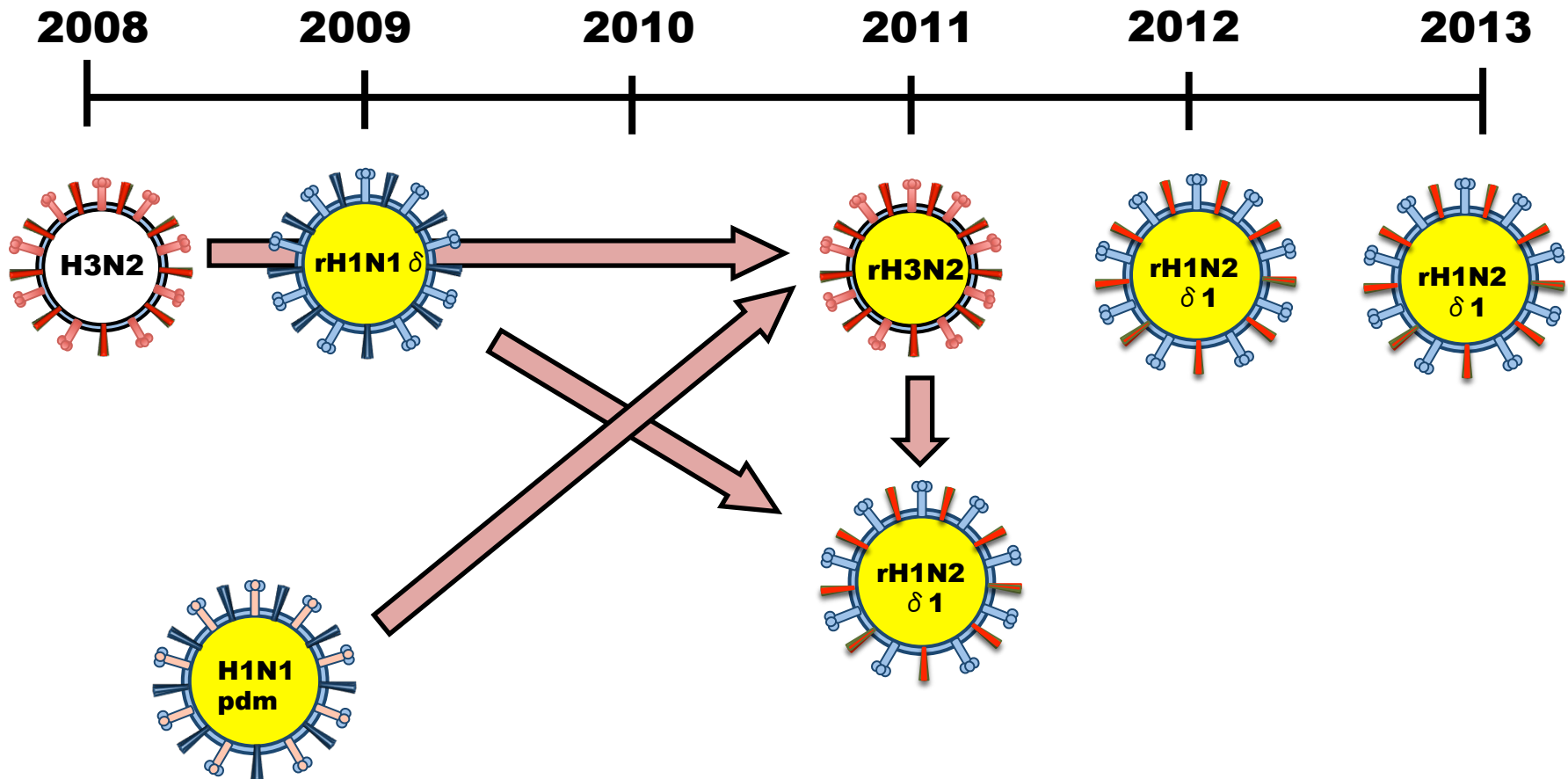


## Conclusiones trabajo 2012

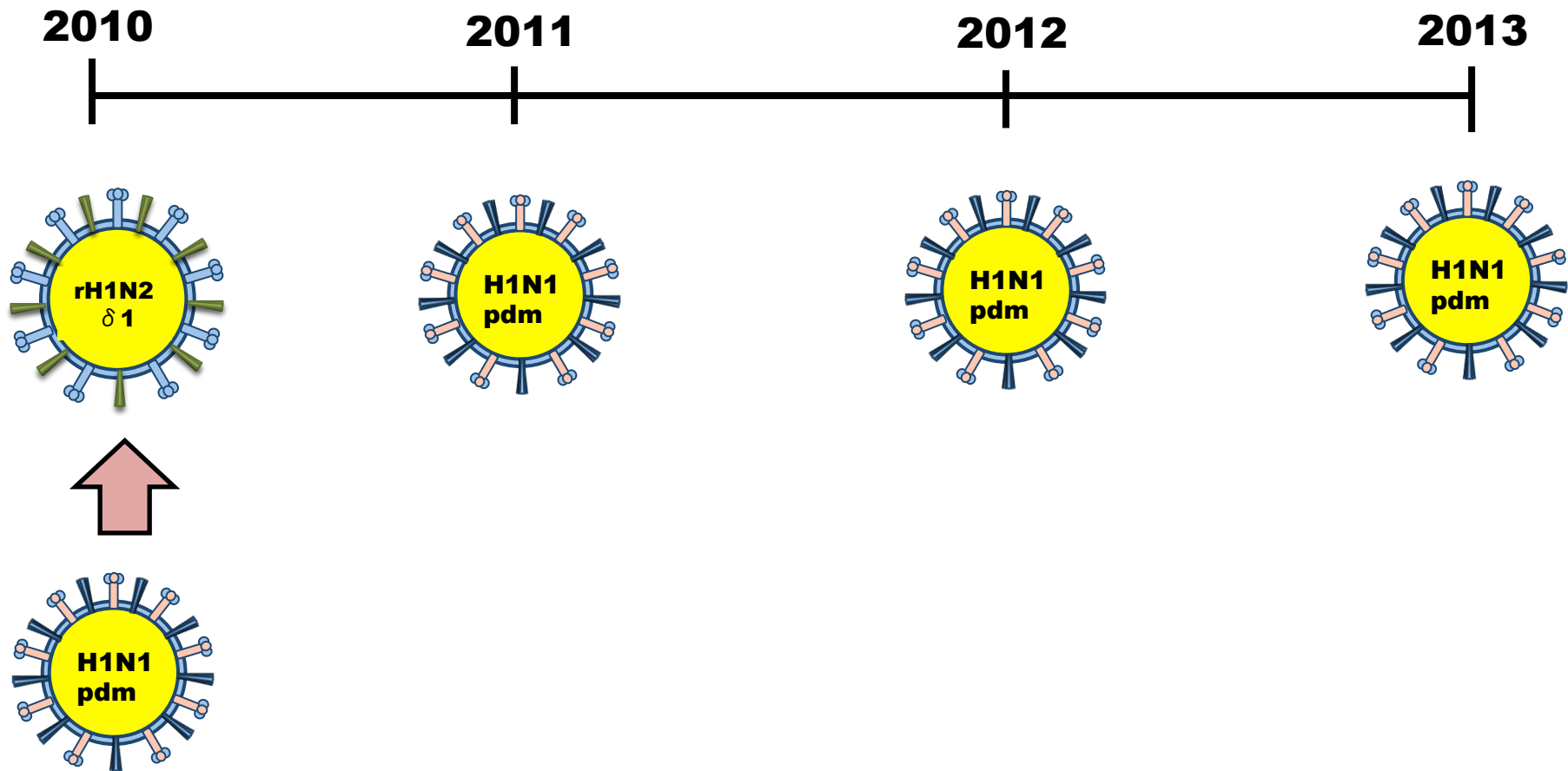
- Existe una infección endémica de virus de influenza que afecta particularmente en la etapa de posdestete a lo largo del año
- Los estudios virológicos demuestran, que si bien existe una predominancia de H1N1 pdm (4 aislamientos), existe circulación de otros subtipos y recombinaciones entre ellos (rH3N2)
- En relación a la serología se observó una caída de anticuerpos hacia los 21 a 35 días y un aumento entre los 49 y los 100 días
- Los resultados del estudio de HI indican una co-circulación de más de un subtipo viral.

# Relevamiento continuo de virus de Influenza en granjas porcinas

## Granja 1



**Granja 2**





## Vigilancia Influenza porcina 2013

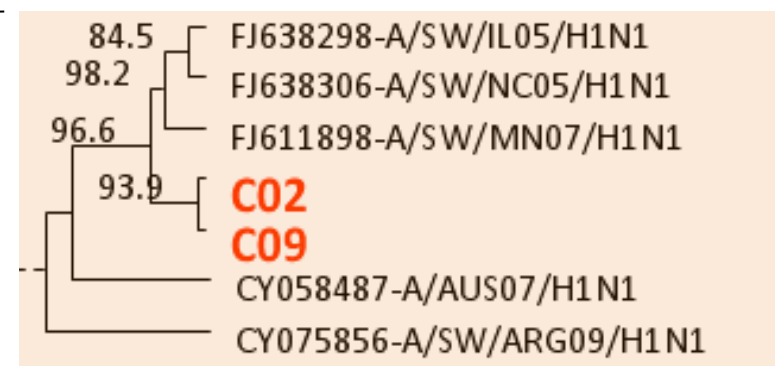
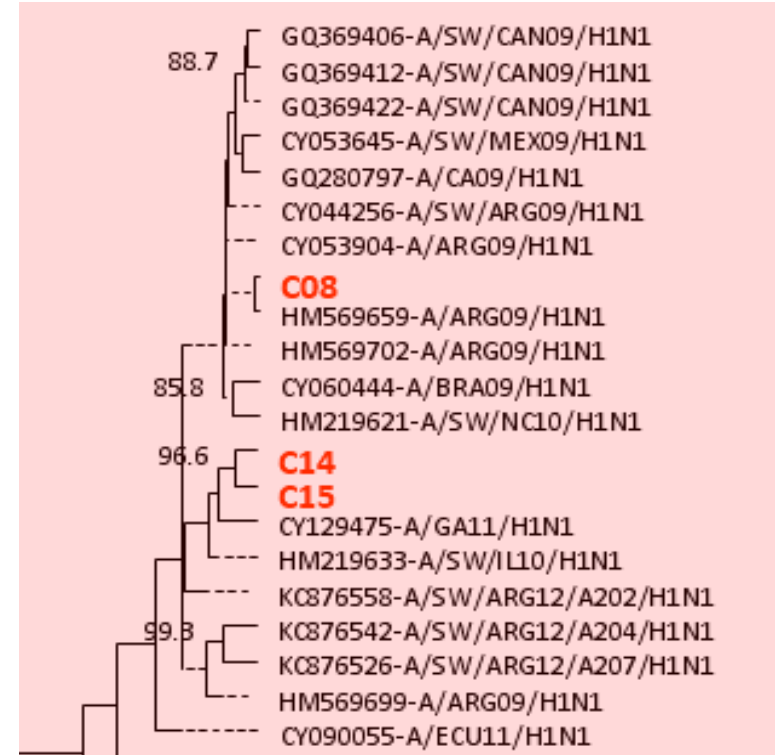
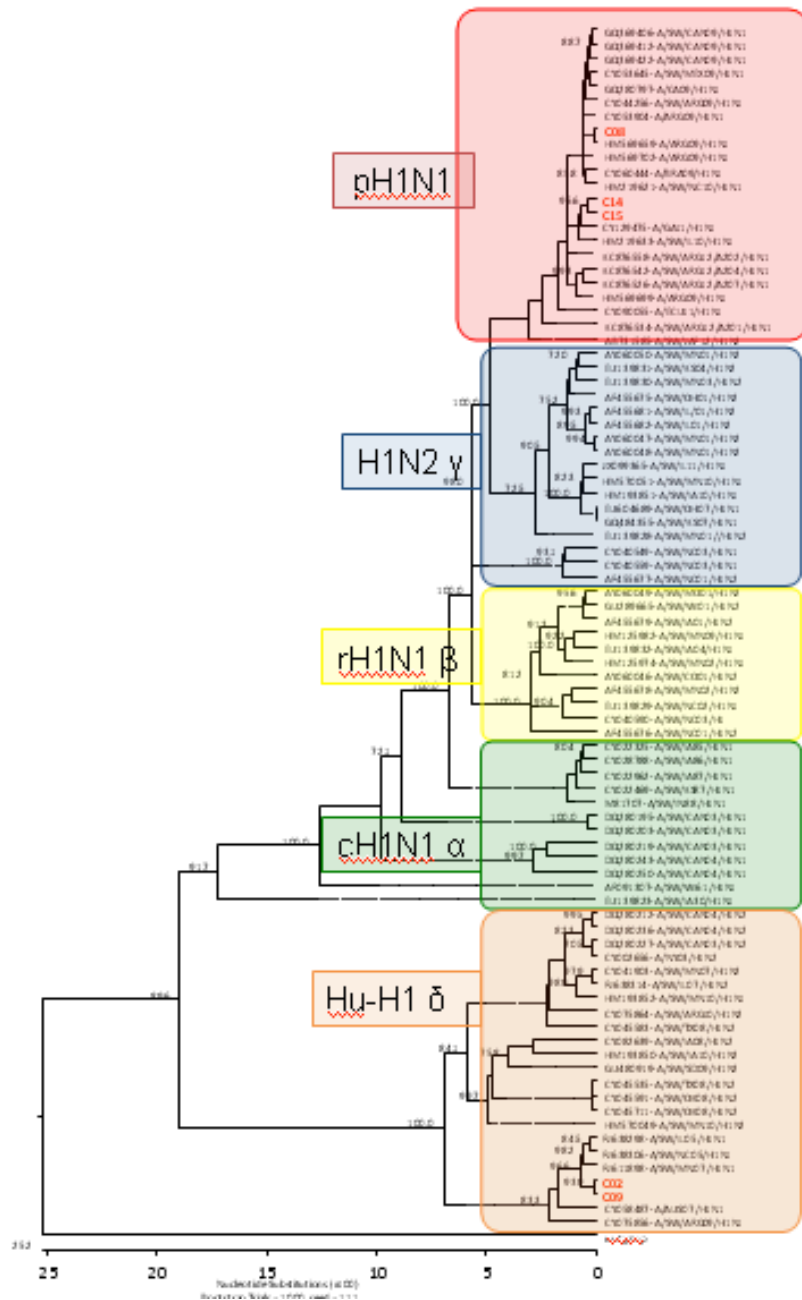
Nº de muestras procesadas: 320      Nº de muestras positivas: 120/320

Nº Total de aislamientos: 33

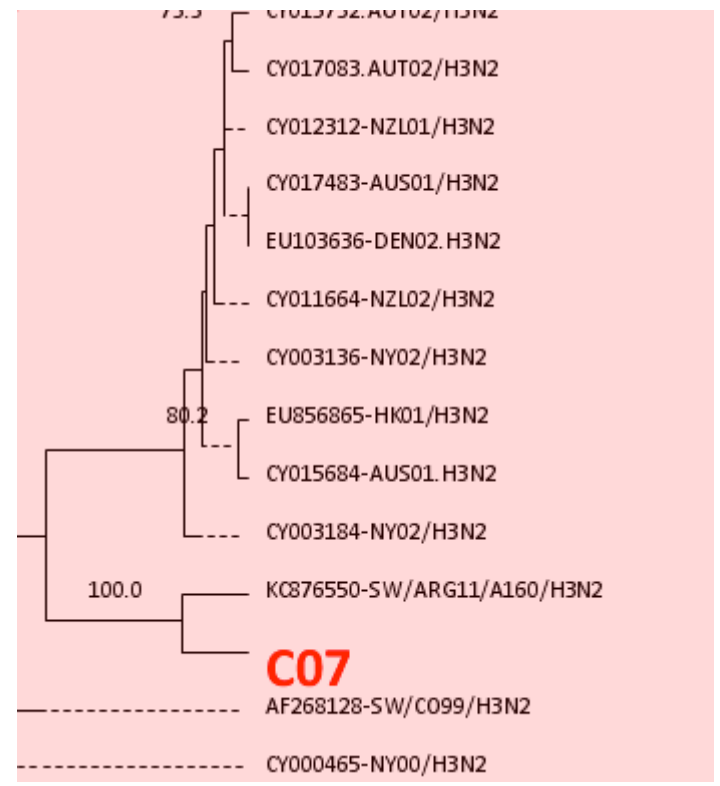
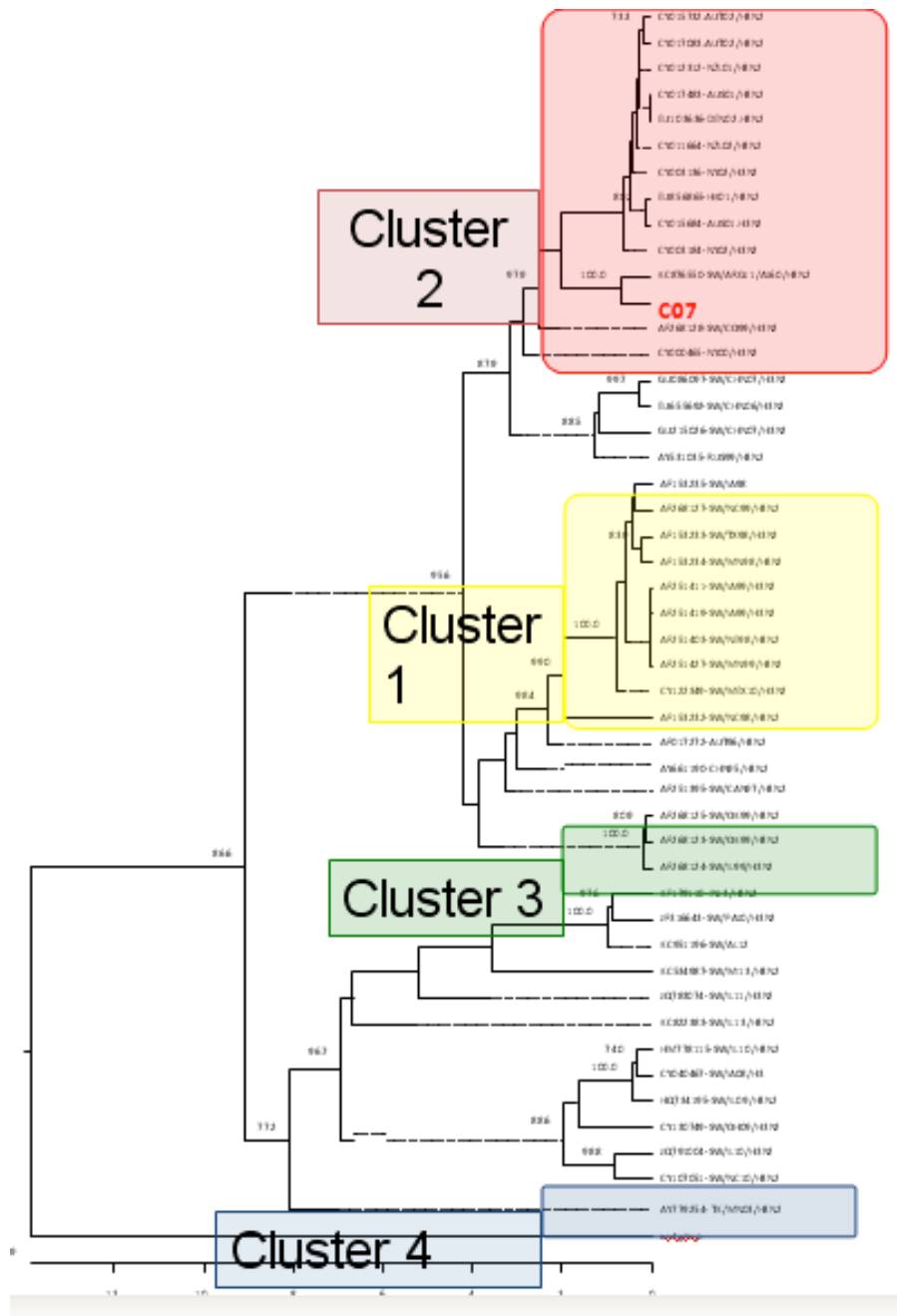
Protocolo	Fecha	Procedencia	Subtipo
<b>C02</b>	Feb-13	BsAs	<b>δH1N2</b>
<b>C07</b>	Abr-13	BsAs	<b>rH3N2</b>
<b>C08</b>	Jun-13	Sta.Fe	<b>pH1N1</b>
<b>C09</b>	Jun-13	BsAs	<b>δH1N2</b>
<b>C14</b>	Jul-13	Sta.Fe	<b>pH1N1</b>
<b>C15</b>	Ago-13	Sta.Fe	<b>pH1N1</b>



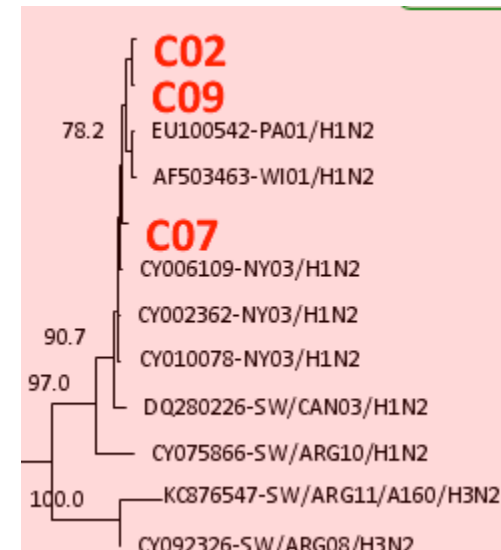
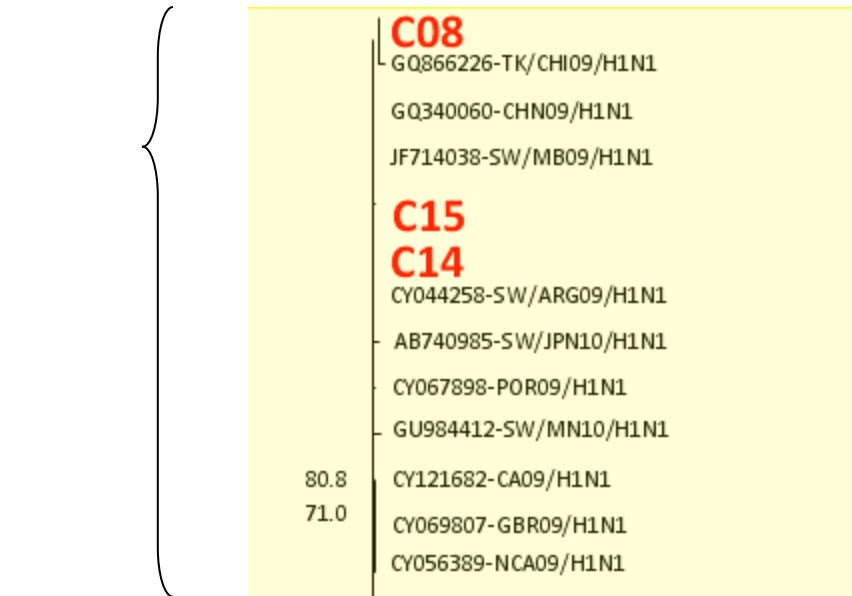
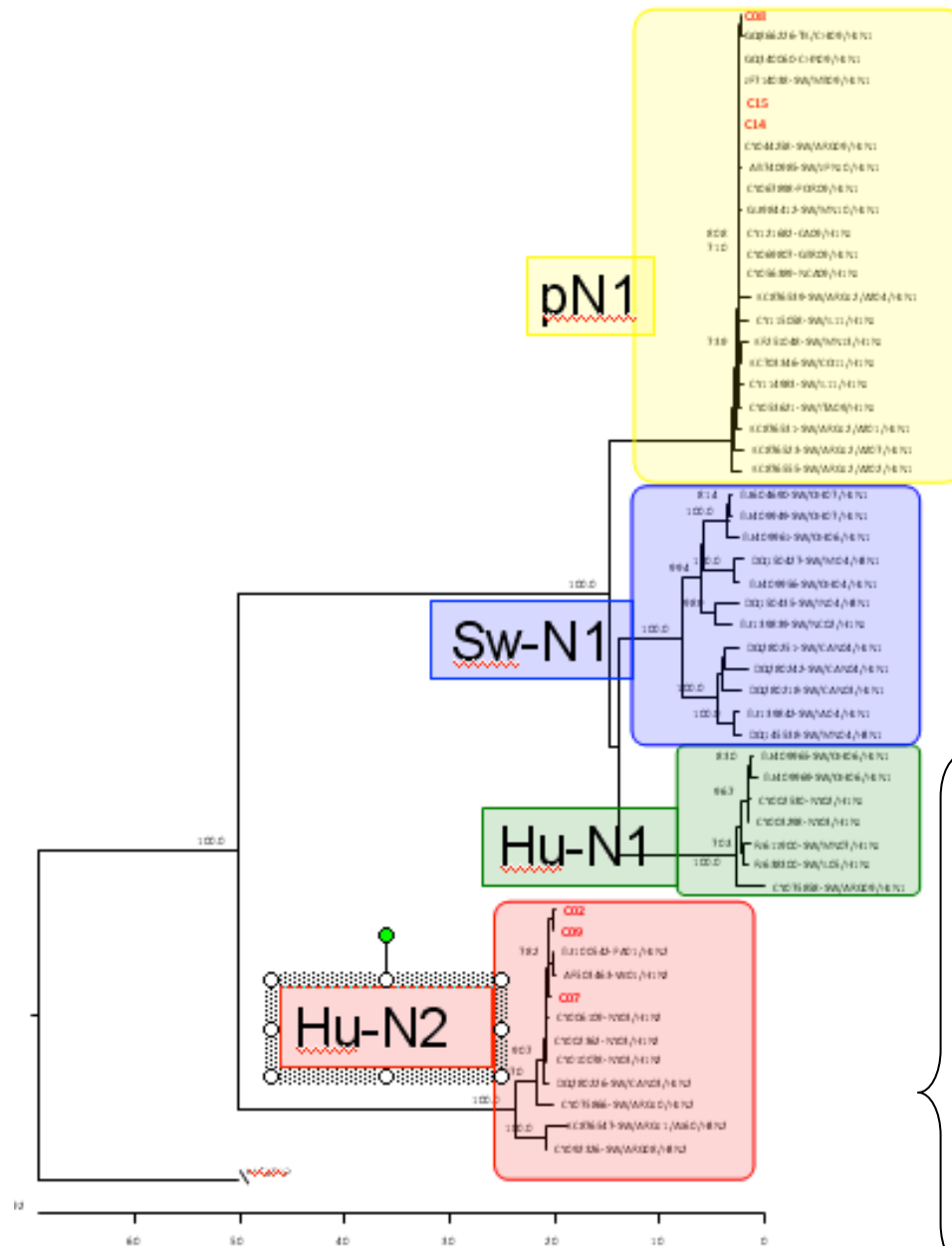
# Árbol filogenético para H1



# Árbol filogenético para H3

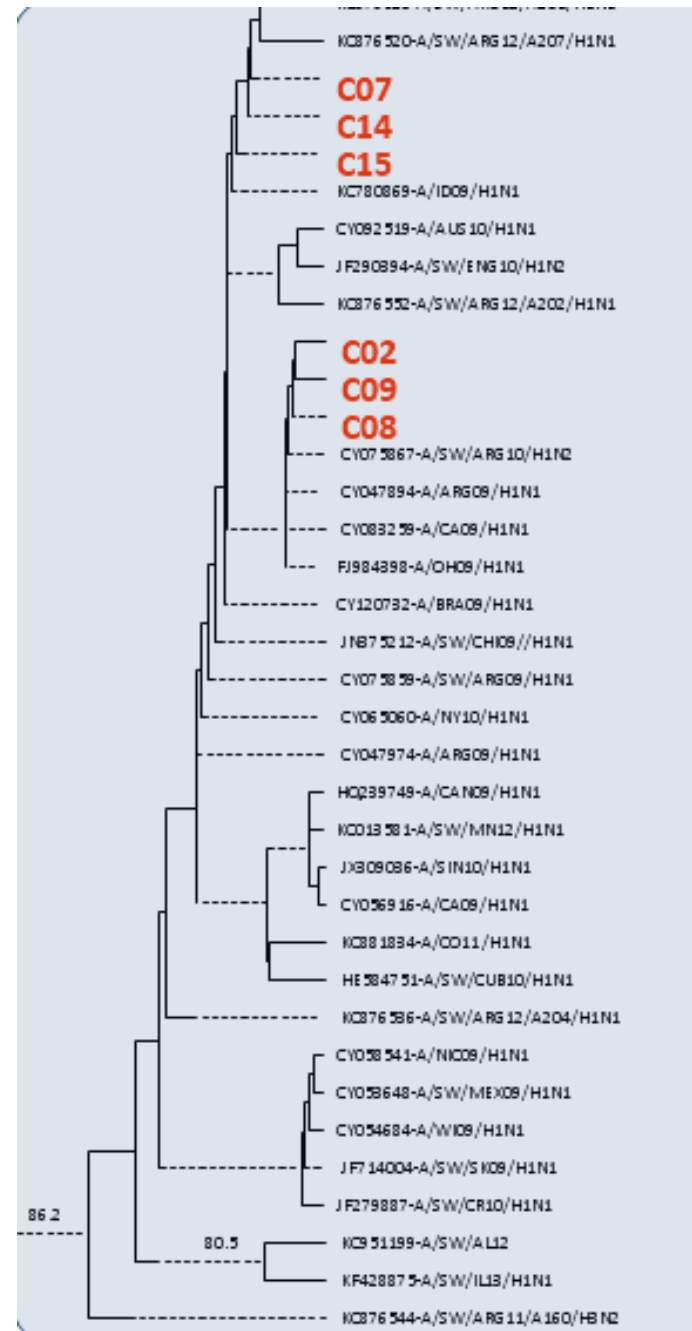
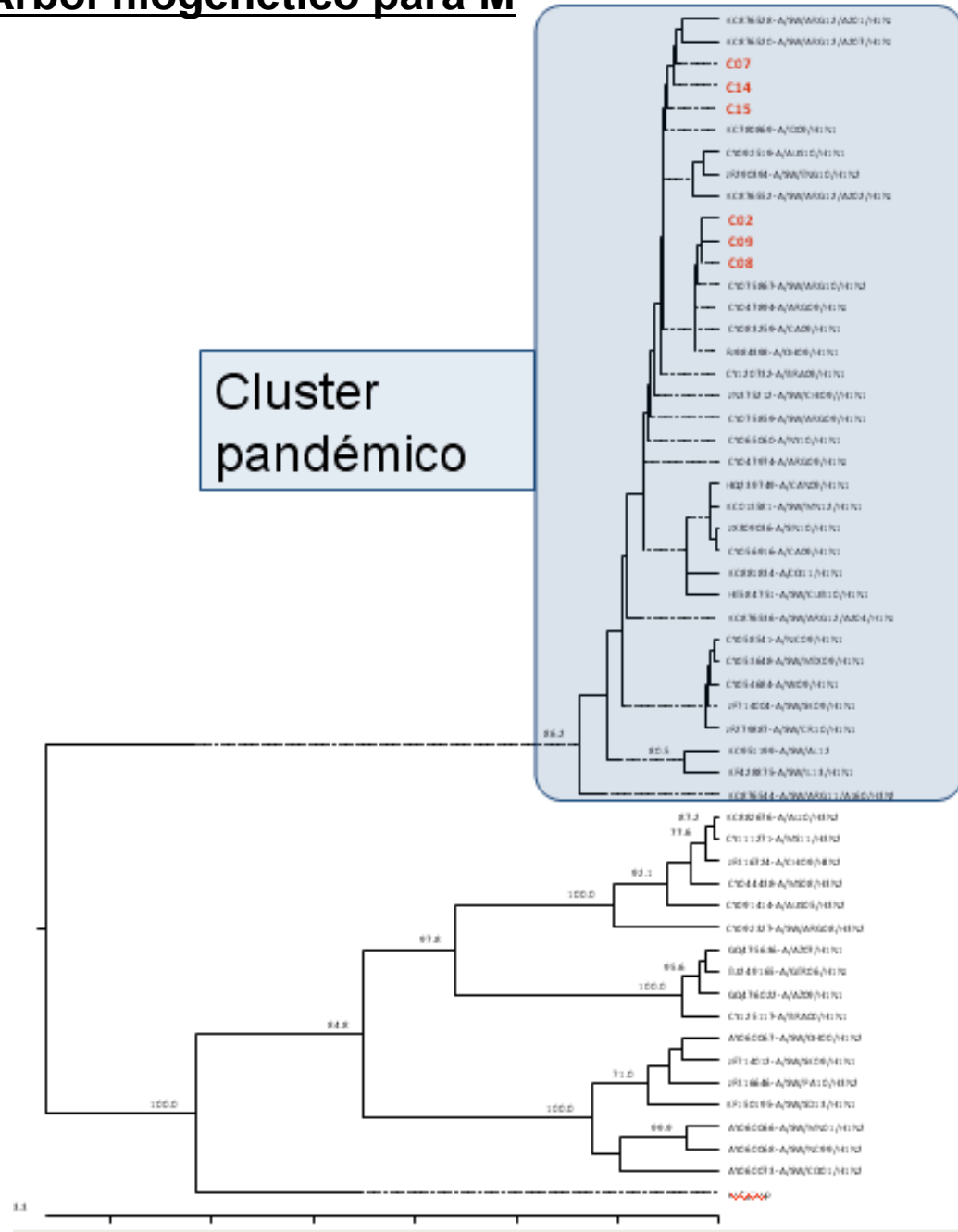


# Árbol filogenético para NA

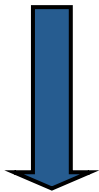
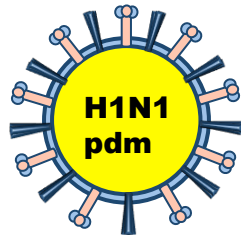


# Árbol filogenético para M

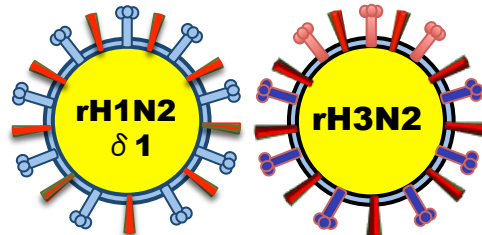
Cluster  
pandémico



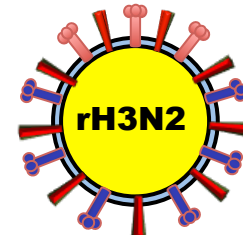
## Conclusiones 2013



**Subtipo  
predominante en  
Argentina**



**Nuevas evidencias  
de reasociación**

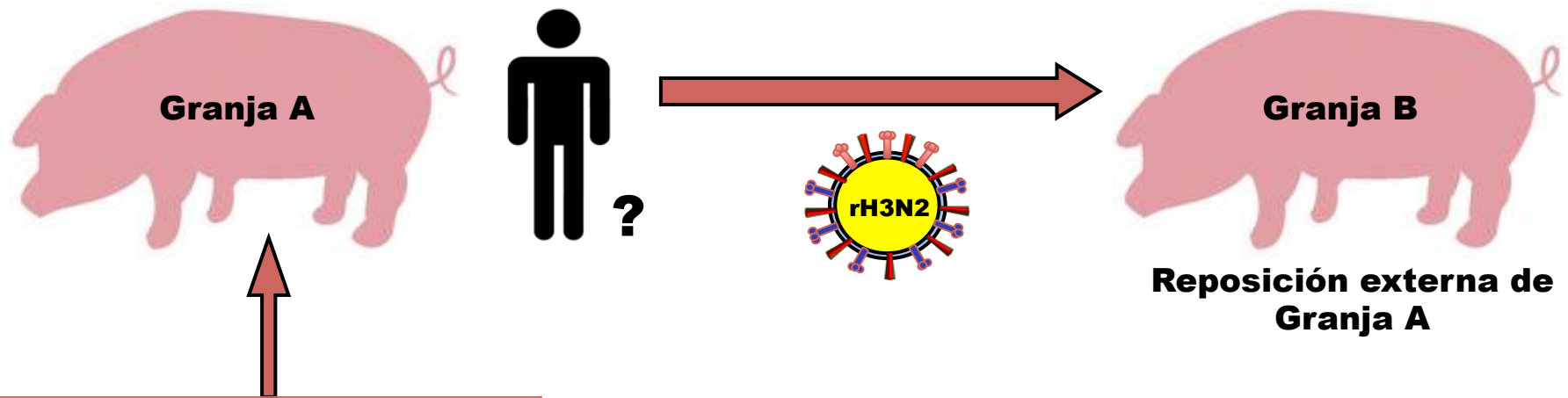


**Alta capacidad  
de transmisión**

**¿Eficacia de las vacunas?**

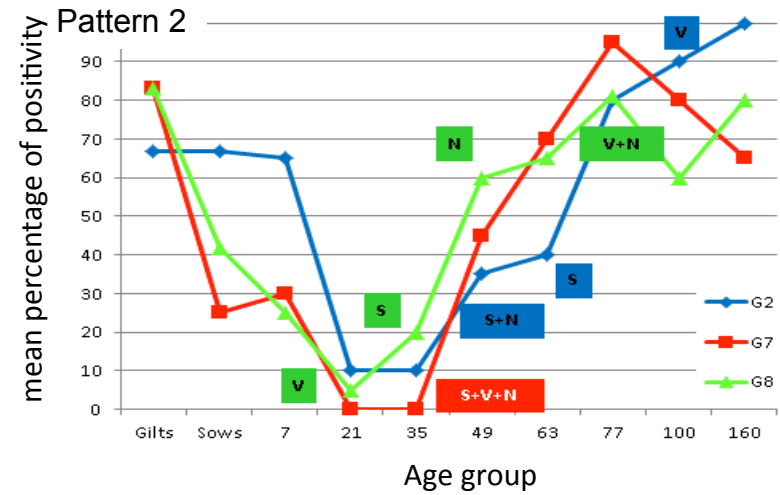
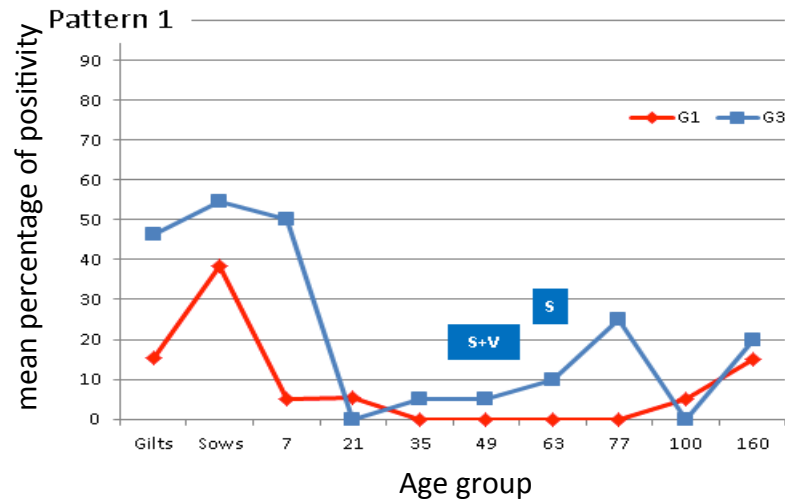
## Evidencia de infección con el mismo virus en dos granjas NO relacionadas

- Abril 2013
- **Granja B** Pcia. BsAs 300 madres, reposición externa
- Brote agudo sintomatología respiratoria (tos -T° 40-42)
- Todas las categorías afectadas
- Rápida transmisión entre galpones y rápida recuperación de la granja (15-20ds)
- Aislamiento virus de Influenza en todas las categorías: **rH3N2**



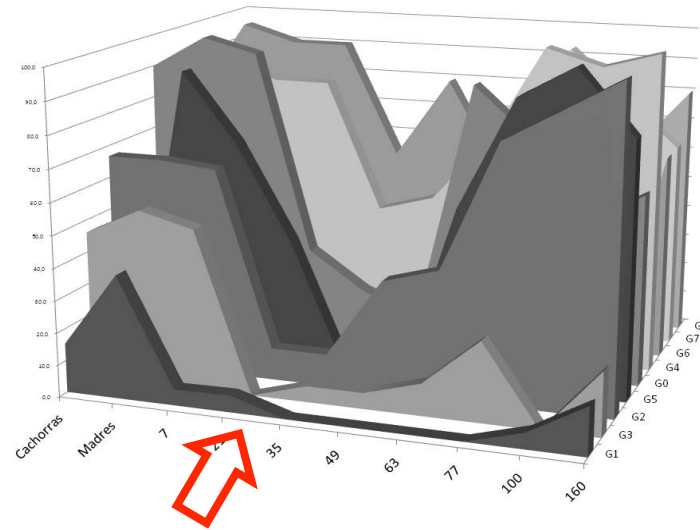
**Subtipos aislados en Granja A :**

	Description	Max score	Total score	Query cover	E value	Ident	Accession
<input type="checkbox"/>	<a href="#">Influenza A virus (A/swine/Argentina/CIP051-A160/2011(H3N2)) segment 4 hemagglutinin (HA) ge</a>	3024	3024	99%	0.0	99%	<a href="#">KC876550.1</a>
<input type="checkbox"/>	<a href="#">Influenza A virus (A/swine/Argentina/CIP051-A2/2008(H3N2)) hemagglutinin (HA) gene, complete</a>	2952	2952	99%	0.0	98%	<a href="#">CY092324.1</a>



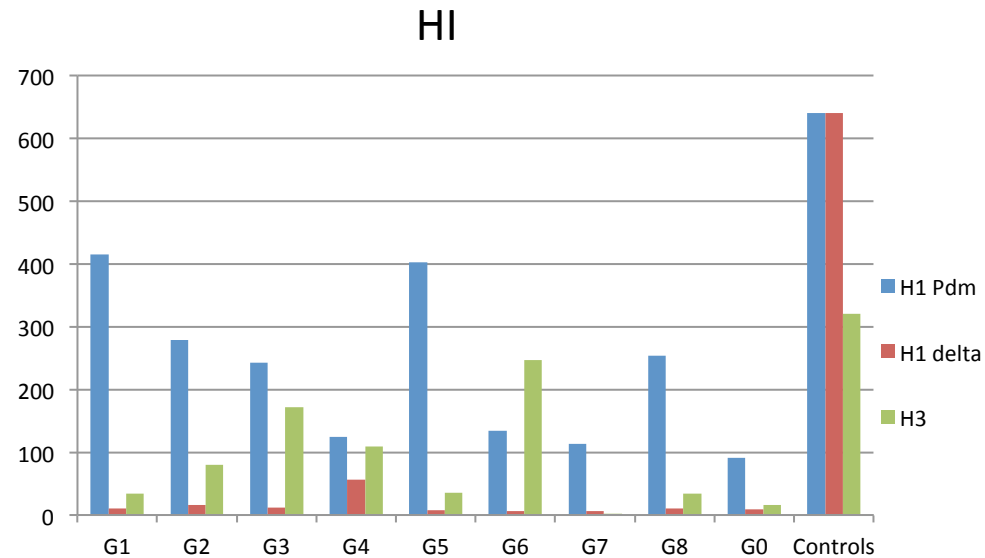
S: clinical signs. V: virus isolation. N: virus isolation from pneumonic lung lesions at necropsy

### ELISA





10% rH3N2  
90% H1N1pdm



## Isolated Viruses in Argentina

- **Non contemporaneous human origin H3N2**  
A/swine/Argentina/CIP051-A2/08 (H3N2)
- **pdm H1N1**  
A/swine/Argentina/SAGiles-31215/2009 (H1N1)
- Reassortant Viruses :
  - **Human Like  $\delta$ 2 H1 SIV (HA+NA) – pdm H1N1 (internal genes)**  
A/Swine/Argentina/CIP051-BsAs76/2009 (H1N1)  
A/Swine/Argentina/CIP051-SantaFe/2010 (H1N2)
  - **Pig adapted Hu H3N2 (HA+NA) – pdm H1N1 (internal genes)**

Antropozoonoses

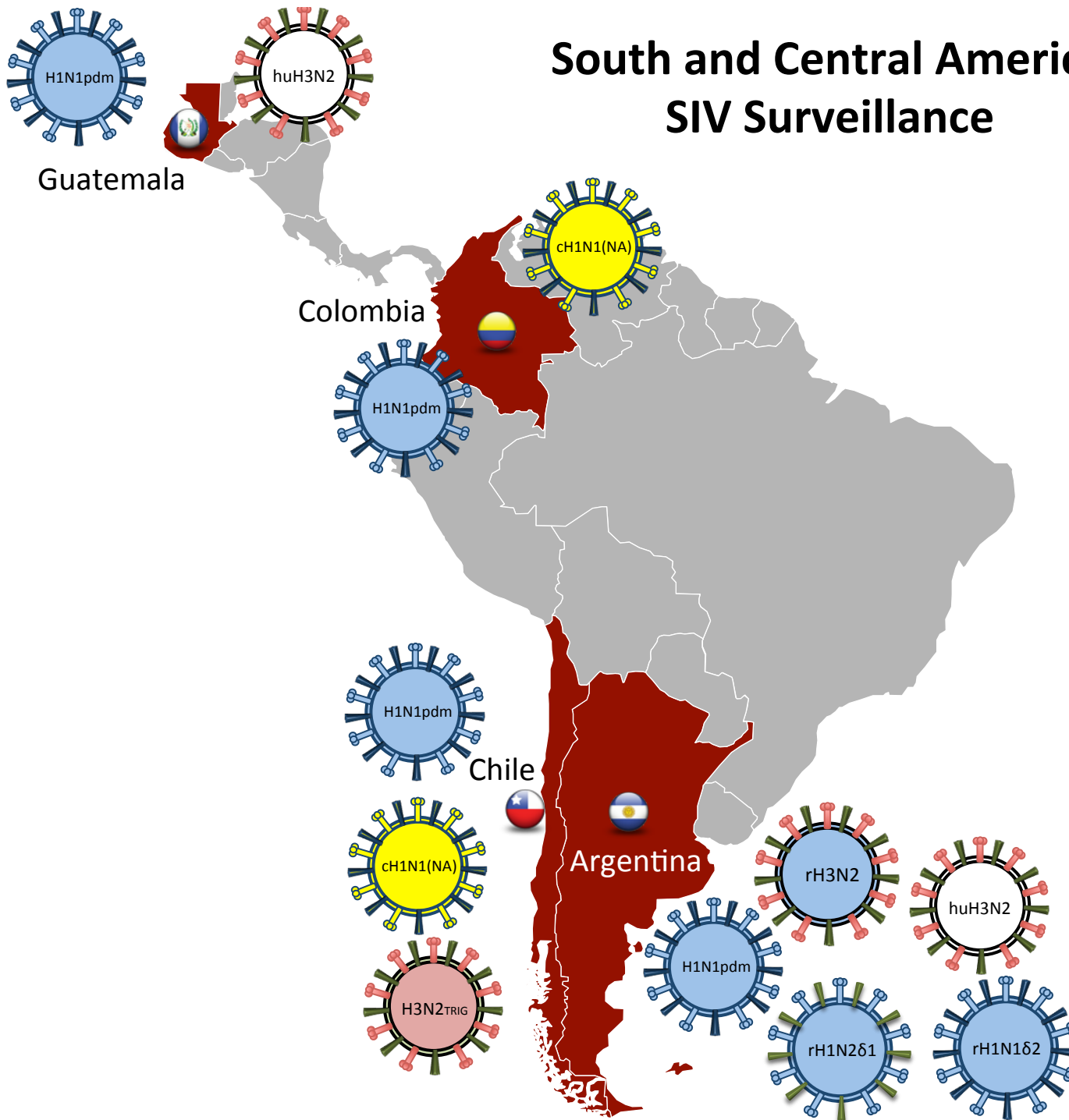


# Summary and perspectives:



- **From collected data we know that:**
  - Influenza circulates in pigs in Central and South America: year to year variation suggests that levels of influenza transmission may vary along the year.
  - Mostly of the viruses isolated have a human origin and the principal subtype is the pandemic H1N1 virus.
  - These viruses started to reassort between them, at least in Argentina (and Brazil).
  - In some countries, the sera collected before the 2009 pandemic outbreak have very low HI reactivity against the pandemic virus, which explains the spread of this virus, but also demonstrate the lack of activity of this virus in the region before this pandemic virus appears in 2009.
- **But we still don't know:**
  - The seasonality of influenza transmission in pigs.
  - The ecological factors important for transmission.
  - Relatedness of human cases of influenza to circulation of the viruses in pigs
  - Antigenic relatedness.
- **What we need for the region:**
  - More training in epidemiological/risk assessment and lab testing.
  - More awareness of the relevance of SIV by institutional presence (an agenda between FAO and/or OIE local vet officers with OFFLU).
  - More financial support to ensure the continuity of these efforts.
  - Local reference laboratory for South and Central America.

# South and Central America SIV Surveillance



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