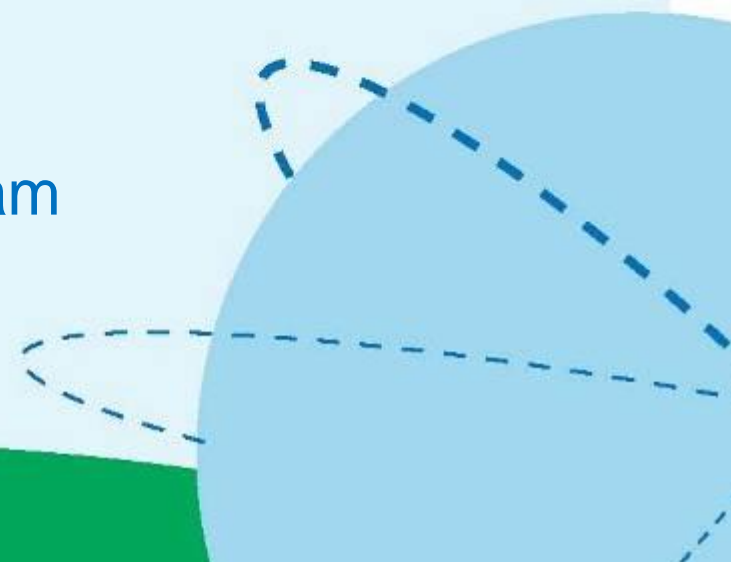




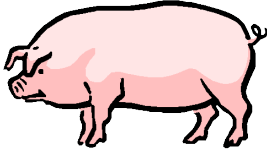



*OFFLU swine influenza virus meeting
27 – 28 March 2017
FAO Headquarters, Rome, Italy*

Tung Nguyen
Department of Animal Health, Vietnam



Influenza Surveillance in Viet Nam 2017

					
Human			Poultry	Pig	Wildlife
Hospital SARI/ILI	Hospital EBS	PREDICT Human Illness Monitoring	DAH/FAO Influenza Surveillance	DAH/FAO Influenza Surveillance	PREDICT Wildlife Surveillance



**Coordinated Surveillance "LISN" will add
More capacity, more sharing, more knowlege, more analysis**


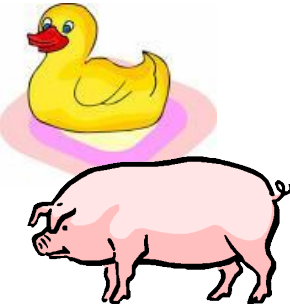

- More influenza virus characterization
- Test existing specimens for other viruses
- Strengthen capacity for surveillance and laboratory diagnostics
- Share outputs across sectors for situation analysis and risk assessment

Coordinated Surveillance for Influenza and Potential Pandemic Pathogens (LISN = Longitudinal influenza surveillance network)

Objectives:

1. Alignment of the surveillance localities, sampling time and laboratory testing algorithm.
2. Monitor and characterize influenza viruses and other potential pandemic pathogens.
3. Regular information sharing and risk assessment using the aligned surveillance results.

Stakeholders

	Central	Regional	Provincial
	GDPM	NIHE (NIC, Epi) Pasteur I (NIC, Epi)	PDH PPPMC Hospital
	DAH NCVD	RAHO2 (Epi, Lab) RAHO6 (Epi, Lab) RAHO7 (Epi, Lab)	DARD Sub-DAH
		NIHE (Predict) Pasteur I (Predict) RAHO2 (Predict) RAHO6 (Predict)	

Influenza Surveillance in Humans

Objectives	Surveillance Strategy in Viet Nam
Identify unusual events	<ul style="list-style-type: none"> - Event-based surveillance <ul style="list-style-type: none"> - Reporting from healthcare facilities, labs, communities - Media monitoring - Indicator-based surveillance <ul style="list-style-type: none"> - Nationally notifiable diseases - Sentinel surveillance (ILI and SARI*) - Characterization of viruses
Monitor national influenza activity/trends & inform national influenza control policy	
Contribute to global influenza control (pandemic monitoring & vaccine determination)	

*ILI= influenza-like illness; SARI = Severe acute respiratory infections

Sentinel Surveillance

ILI Surveillance

- From 2006 - 2015, US-CDC supported establishment of the system
- Since 2016, sentinel surveillance is maintained by government funds

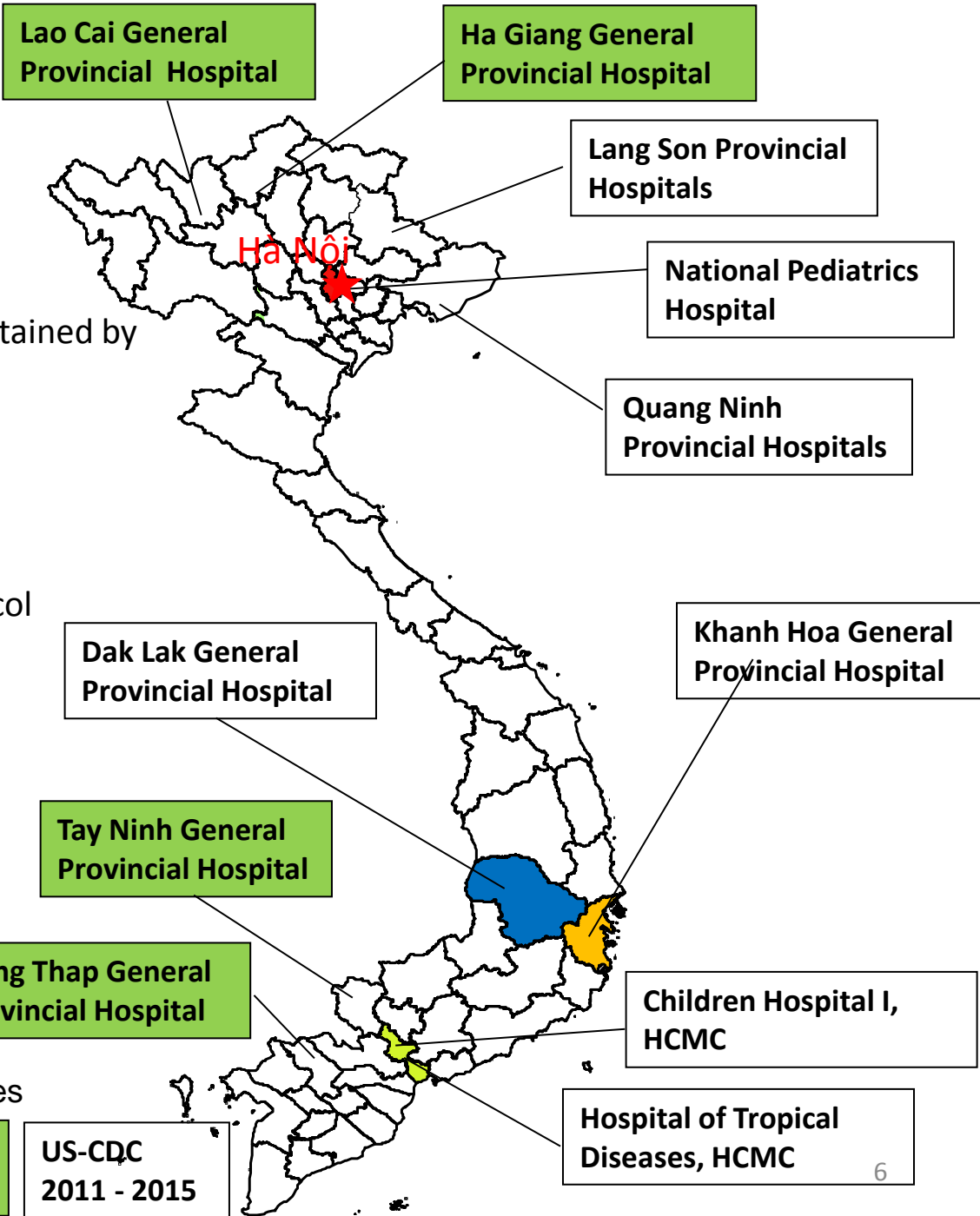
SARI surveillance

- Began in 2011
- January 2016: new “harmonized” protocol
- At least 10 sites
- Coordinated by MOH,
- Supported by CDC and WHO
- Test for
 - Influenza A (subtypes of H1pdm09, H3, H5 and H7) and influenza B
 - Respiratory syncytial virus (RSV);
 - Human metapneumovirus (HMPV);
 - Parainfluenza virus (PIV) 1, 2, 3;
 - Adenovirus (AdV);
 - Rhinovirus (RV)

Supported sites

WHO/USAID 2014 - 2015

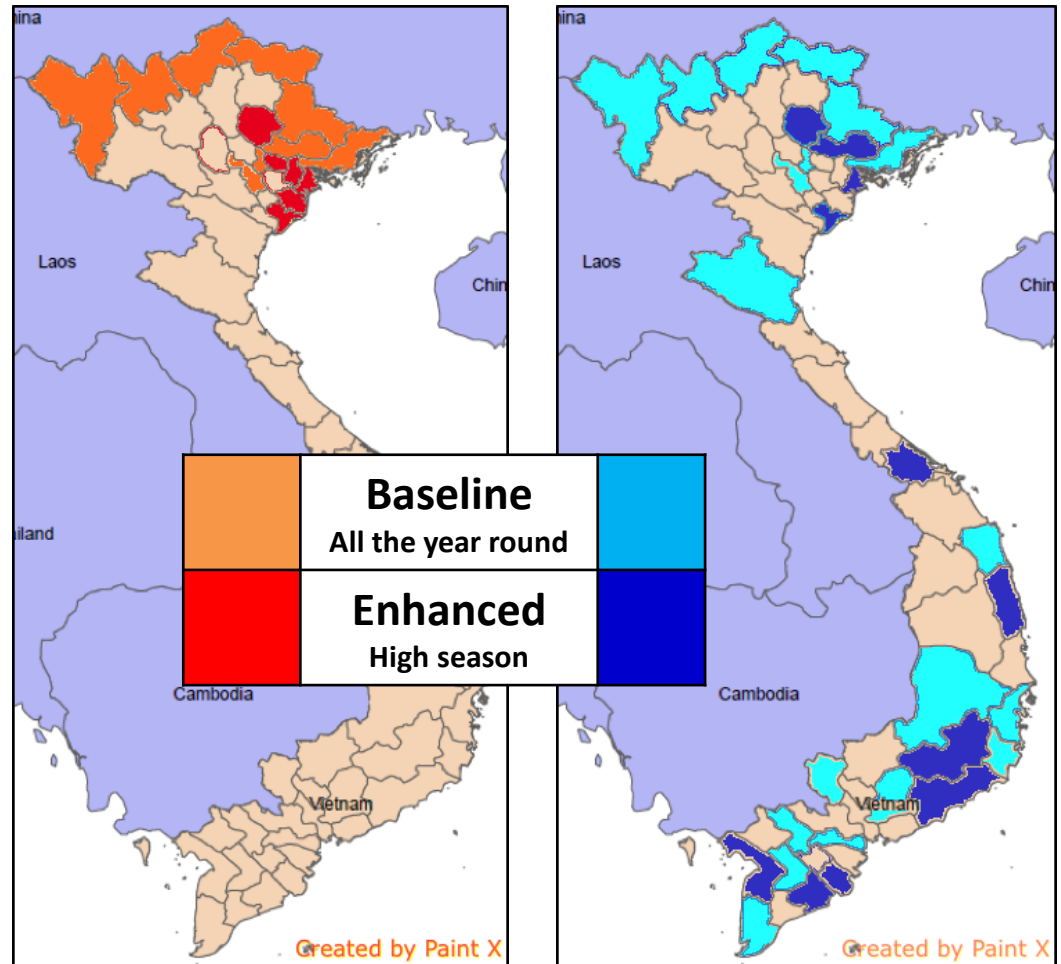
US-CDC 2011 - 2015



Avian Influenza Surveillance 2017

Objectives

- 1. Early detection of H7N9** viruses in poultry and LBMs in high-risk areas for rapid response;
- 2. To understand the epidemiology and the evolution of HPAI H5** viruses in poultry to inform prevention and control strategies, including vaccine selection; and to evaluate the performance of H5Nx control programme
- 3. To understand the gene pool and the evolution of influenza A** viruses in poultry and pigs for early warning of emergence of pandemic influenza A



Detection (H7N9)

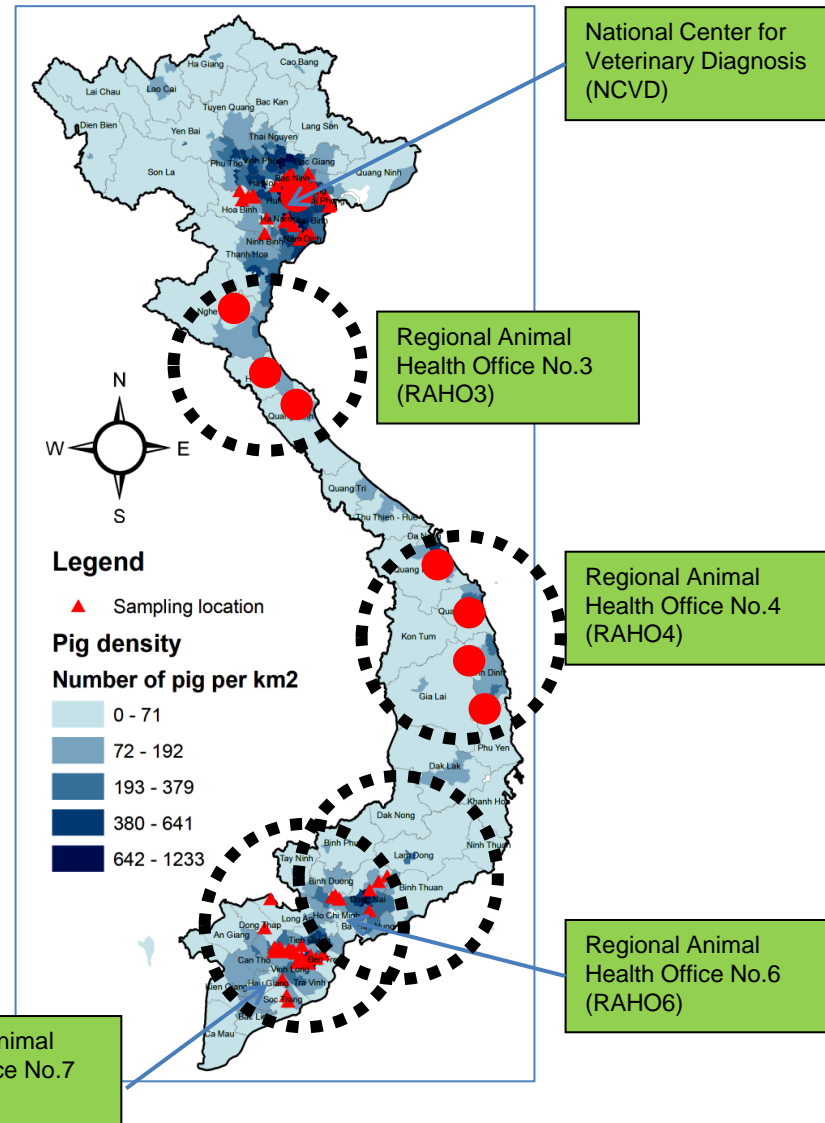
Monitoring (H5)

Swine influenza surveillance

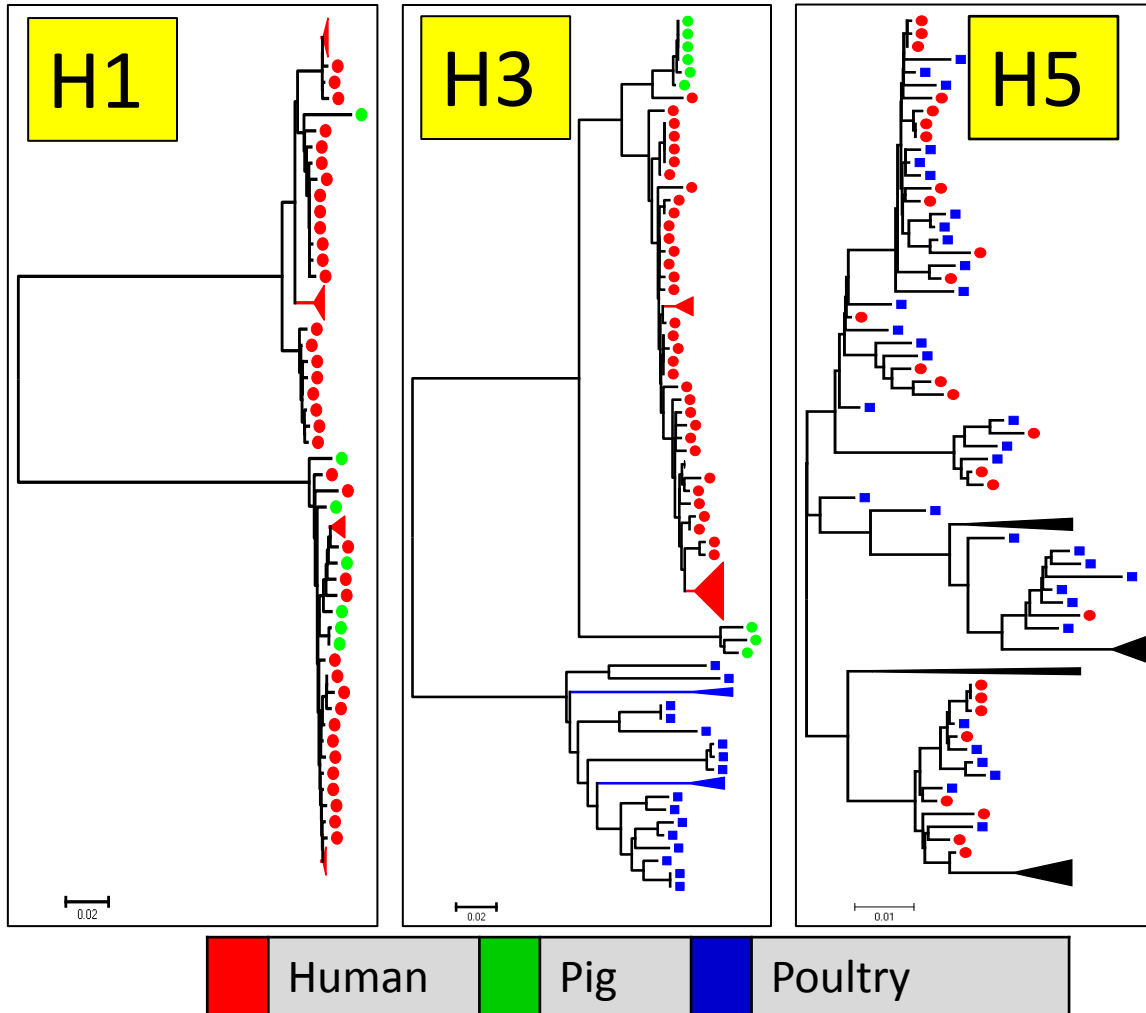
- Surveillance of influenza A virus was initiated in 2013 in order to understand better on;
 - **gene pool** of influenza A virus in pig population
 - **risk factors for virus transmission, circulation, and evolution** across sectors based on production systems and value chains
- Risk-based surveillance targeted at high risk population i.e. large commercial breeding/fattening farms with low farm biosecurity

Sampling Design

- Province:
 - High density of pig population
 - 2012 – 2013; 12 provinces
 - 2014 – 2015; 19 provinces
- Farm:
 - Large scale and low biosecurity
 - Breeding: 100-200 sows
 - Fattening: >1000 pigs
- Animals and samples:
 - 5-8 & 9-12 weeks old:
 - 30 nasal swabs/farm from each age group
 - Swabs from 60 animals/farm
 - Sera from 10 piglets/farm (+5 sows in breeding farms)

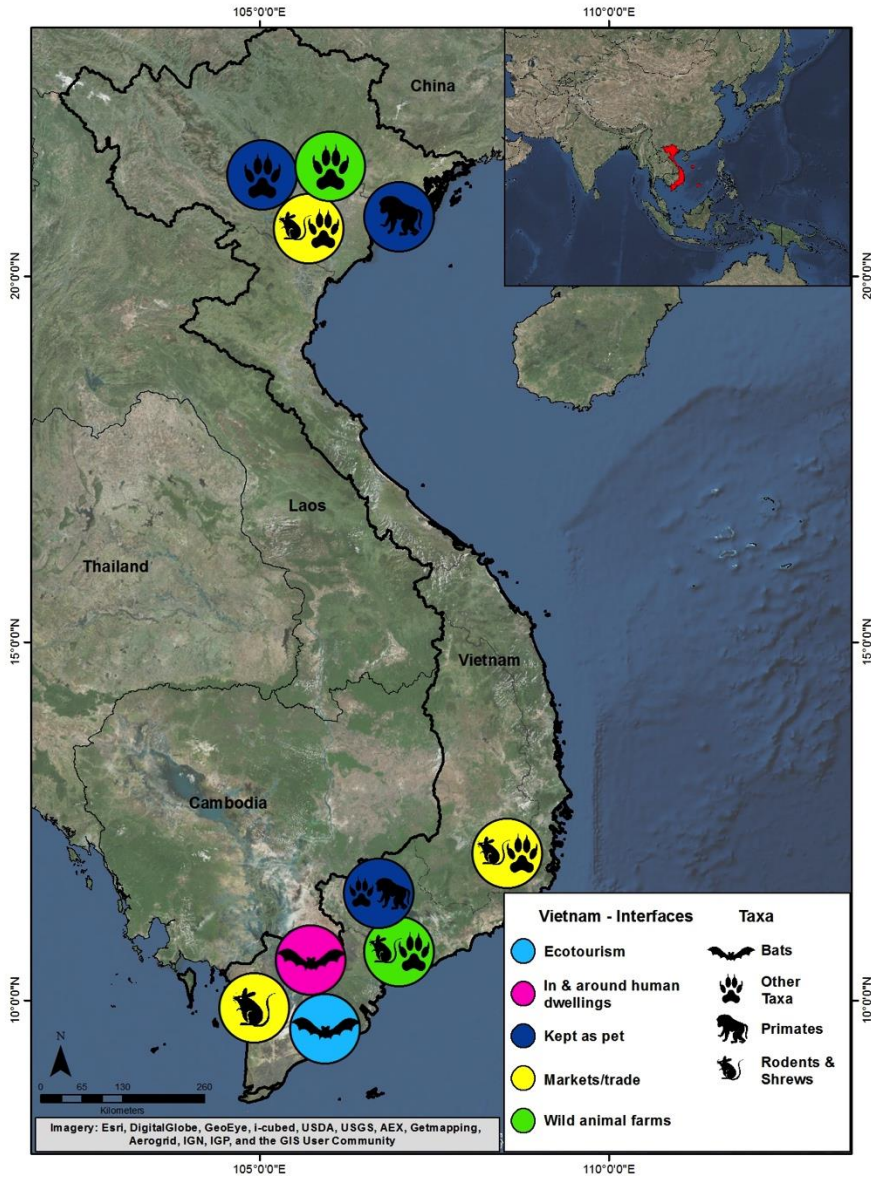


Flu A Viruses in Human, Pig, and Poultry in Viet Nam (Publicly available sequences in July 2016)



No of HA gene sequence available In public database (Genebank+GISAID)			
Sub-type	Poultry	Pig	Human
H1	0	5+	250+
H3	20+	5+	400+
H5	500+	0	50+
H7	3	0	0
H9	0	0	0

Wildlife Surveillance in Viet Nam

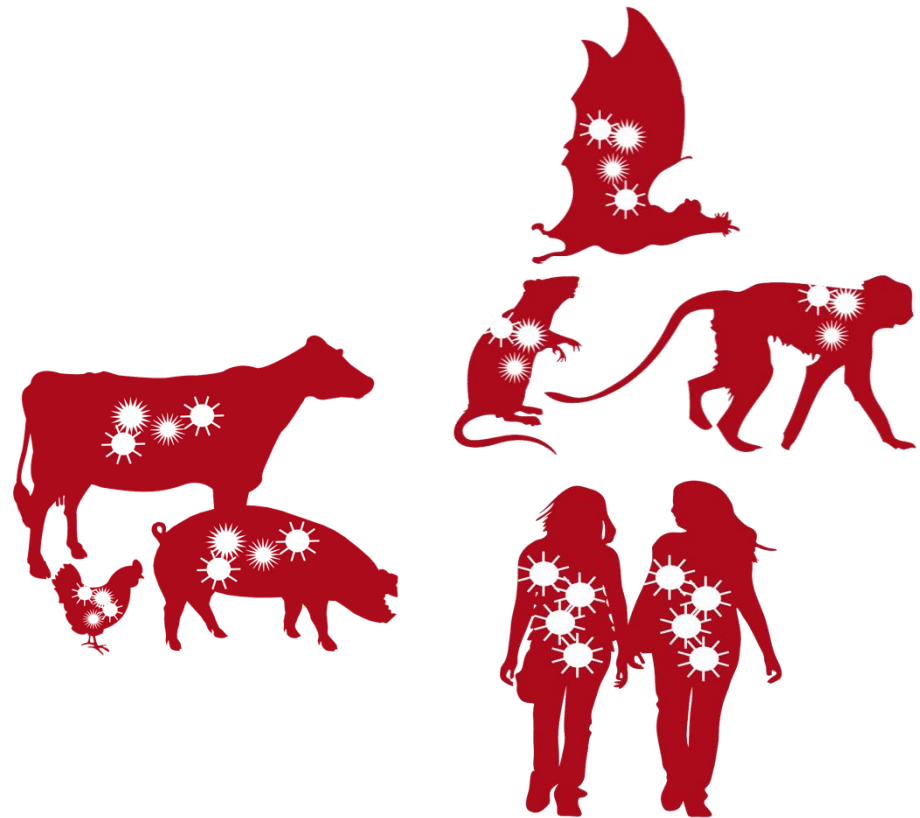


- Surveillance targeting “high-risk” interfaces for disease transmission between wildlife and humans
- Screening for viral pathogens of pandemic potential
- Building local capacity for surveillance & laboratory diagnostics

PREDICT2 - Focus

Specimens and Viral Testing

- Viral Families (n=12; 4 in all countries)
 - **Corona**, Hanta, Arena, Alpha, **Paramyxo**, **Flavi**, **Rhabdo**, **Influenza**, Seadorna, Picorna, **Filo**, Reo (Retro) viruses
- Specimen type
 - Oropharyngeal swab/saliva
 - Rectal swab/feces
 - Urine, blood



- * All taxa:
 - * Same sample type
 - * Same suit of viral families



PREDICT

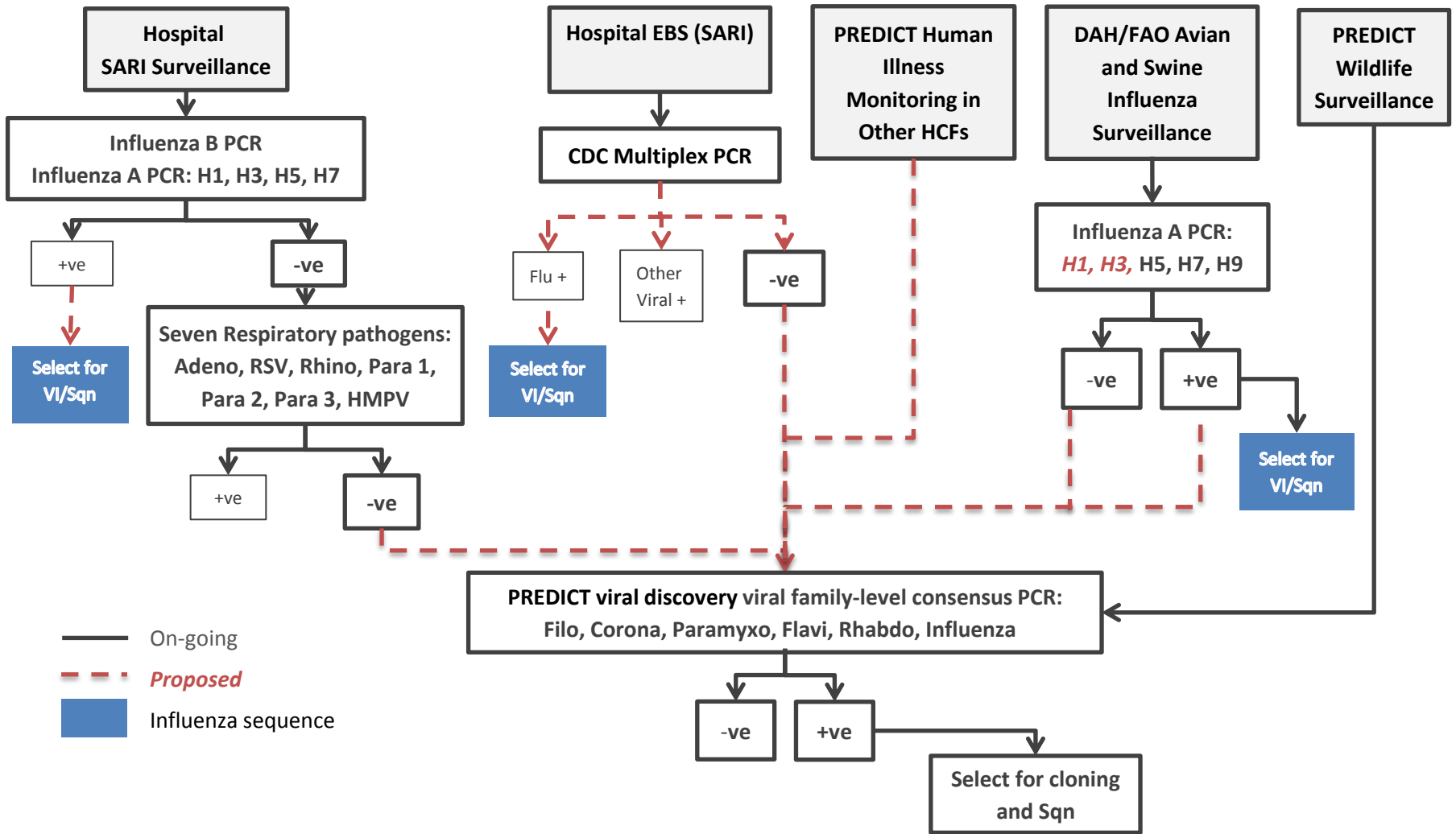


Wildlife Trade in Viet Nam

- Source Country
- Consumer Country
- Transit Country



Coordinated Surveillance for Influenza and Potential Pandemic Pathogens



Provincial Level (PPMC, Sub-DAH)	Regional Level (NIHE, PI-HCM, RAHO2,6,7)	National Level (GDPM, DAH, CITES)
<ul style="list-style-type: none"> Quarterly coordination meeting between PPMC and S-DAH to review progress and trouble shoot 	<ul style="list-style-type: none"> Six monthly meeting to review virological and epidemiological data 	<ul style="list-style-type: none"> Annual meeting to review risk assessment results and identify potential intervention

“I don’t know where to go, can’t do it alone...”

- Pandemic detection and prevention
 - Longitudinal influenza evolution
 - Pathogen discovery
- Strengthened capacity
 - Laboratories e.g. sequencing
 - Risk assessment
 - Routine information sharing (4-way linking, circular 16)
- Data / Information
 - Epidemiological linked samples / data
 - Viruses characteristics
 - Situational analysis and joint risk assessment

