

6th Annual CEIRS meeting

July 29 – August 1, 2012

New York, NY

New York Marriott Marquis, Times Square New York, NY

Sponsored by: NIH/NIAID CEIRS Program

Hosted by:
Mount Sinai School of Medicine

Center for Research on Influenza Pathogenesis (CRIP)

Welcome

6th Annual National Institute of Allergy and Infectious Diseases (NIAID) Centers of Excellence for Influenza Research and Surveillance (CEIRS) Network Meeting

> July 29-August 1, 2012 Mount Sinai School of Medicine Marriott Marquis, Times Square, New York, NY

Dear Colleague,

On behalf of the Influenza Program at the National Institutes of Health/National Institute of Allergy and Infectious Diseases/Division of Microbiology and Infectious Diseases (NIH/NIAID/DMID), and the Center for Research on Influenza Pathogenesis (CRIP), we welcome you to the 6th Annual Meeting of the Centers of Excellence for Influenza Research and Surveillance (CEIRS) at "The Crossroads of the World" in New York City's Times Square.

This meeting brings together representatives of the CEIRS network, including the foremost research scientists, public health experts, government representatives, wildlife biologists and veterinarians from all over the world to exchange and discuss critical scientific information on influenza virus infection in animals and in humans. The meeting will focus on progress made on the basic biology of influenza viruses, emerging scientific questions and future directions of the multiyear collaborative CEIRS contracts at: Mount Sinai School of Medicine, St. Jude Children's Research Hospital, University of Rochester, University of Minnesota, and Emory University.

Since the establishment of the CEIRS network, we have witnessed the 2009 H1N1 human pandemic, the establishment of H5N1 viruses as endemic viruses in poultry in different regions of the world, and the intense debate on the benefits and risks of experiments involving modified H5N1 viruses that are transmissible in ferrets. During this time, the CEIRS network has made significant scientific advances, identifying pathways and mechanisms involved in virus pathogenesis that could be used in antiviral development and expanding our understanding of the ecology and diversity of influenza viruses in animal reservoirs. In addition, the network has participated in the discovery of the conserved stem region of the HA as a potential Achilles' heel of the virus that might be a valuable contribution toward developing improved therapeutic and prophylactic measures against this continuously evolving pathogen. At the meeting, we will hear about progress and new discoveries in these and other areas of influenza research. The CEIRS network continues to highlight the value of an intercollaborative research program that contributes to understanding influenza virus biology and developing rational approaches to prevent infection, disease, and transmission in humans and animals.

We anticipate an exciting meeting covering broad areas of the influenza field, such as molecular virology, pathogenesis, immunology, transmission, and human and animal surveillance. There will be 56 speakers, dozens of posters, and plenty of opportunities to form and/or strengthen collaborations, develop interactions and exchange ideas and information, in both formal sessions and informal get-togethers. We hope that you will also enjoy this opportunity to explore New York and enjoy this unique city.

Sincerely,

Adolfo García-Sastre, PhD PI/Director CRIP Diane Post, PhD Program Officer NIAID/DMID



6th Annual CEIRS Network Meeting

July 29 – August 1, 2012 New York Marriott Marquis, Times Square 1535 Broadway New York, NY 10036

Arrival

Sunday July 29, 2012

5:00 – 7:00 pm Registration – Astor Ballroom Foyer, 7th Floor

7:00 – 9:00 pm Welcome Reception – Manhattan Ballroom, 8th Floor

Monday July 30, 2012

7:30-8:30	Registration – 7 th Floor Foyer Breakfast – 7th Floor Foyer	
8:30-8:35	Welcome - Adolfo Garcia-Sastre, CRIP Director	
8:35 – 8:50	CEIRS Update - Diane Post (NIAID)	

Session #1 - Dual Use Research of Concern

Session Chair: Carole Heilman (NIAID)

8:50 – 9:30 The USG Policy for Oversight of Life Sciences Dual Use Research of

Concern-Implementation Issues- Dennis Dixon (NIAID)

9:30 – 10:30 Panel discussion - "USG policy implementation at the institution level"

 Discuss what the institutions are currently doing, mitigation strategies and responsible communication

Convener: Carole Heilman (NIAID)

Panelists:

Adolfo Garcia – Sastre, Ron Fouchier, Yoshi Kawaoka, Richard Webby, Mark

Tompkins, Nancy Cox

General Discussion

10:30 – 11:00 **Break – 7**th Floor Foyer

Session #2 – Transmission

Session Chair: Peter Palese (CRIP)

11:00 – 11:30 Transmission of influenza A/H5N1 virus via aerosol or respiratory droplets –

Sander Herfst (CRIP)



11:30 – 12:00	The Potential for Respiratory Droplet–Transmissible A/H5N1 Influenza Virus to Evolve in a Mammalian Host – Derek Smith (CRIP)			
12:00 – 1:00	Lunch – Cantor/Jolson, 9th Floor			
1:00 – 1:15	Alternative reassortment events leading to a transmissible H9N2 influenza virus in the ferret model – Brian Kimble (CRIP)			
1:15 – 1:30	Comparable fitness and transmissibility between oseltamivir-resistant 2009 pandemic and seasonal H1N1 viruses with the H275Y neuraminidase mutation – Hui Ling Yen (SJCEIRS)			
1:30 – 1:45	Molecular factors affecting the transmission of the pH1N1 influenza virus – John Steel (IPIRC)			
1:45 – 2:00	Transmission of influenza A H1N1/2009 virus in China - Dhanasekaran Vijaykrishna (SJCEIRS)			
Session #3 – 2:00 – 3:00	Panel Discussion: Risk Assessment and CEIRS Role "Risk Assessment of Influenza viruses and CEIRS role" Convener: Stacey Schultz-Cherry Panelists: (10 min each) Nancy Cox: "Development of an Influenza Risk Assessment Tool" Carol Cardona: "Phenotyping Influenza Viruses" Hassan Zaraket: "Risk assessment studies on Avian H9N2 influenza viruses from different lineages" Scott Krauss: "Risk assessment studies on H2 viruses"			
	General Discussion			
3:00 – 3:30	Break – 7 th Floor Foyer			
Session #4 - Influenza Pathogenesis Session Chair: Malik Peiris (SJCEIRS)				
3:30 – 3:55	Secondary bacterial infection and the use of novel therapeutics in influenza virus infected-obese mice – Erik Karlsson (SJCEIRS)			
3:55 – 4:20	Responses and outcomes to influenza virus infection in context – S. Mark Tompkins (IPIRC)			
4:20 – 4:35	Effect of human mesenchymal stem cells on enhancing alveolar fluid clearance in human alveolar epithelium injured by influenza H5N1 virus infection – Michael Chan (SJCEIRS)			
4:35 – 4:50	Pathogenesis of influenza virus infection is regulated by the disease progression stage of allergic asthma – Amali Samarasinghe (SJCEIRS)			
4:50 – 5:05	Effect of Route of Inoculation and Type of Duck on the Pathogenicity of H5N1 Highly Pathogenic Influenza Viruses in Domestic Ducks – Mary Pantin Jackwood (MCEIRS)			



5:05 – 6:30	Poster Viewing – Empire Complex
6:30 - 9:30	Dinner – Cantor/Jolson, 9th Floor

Tuesday July 31, 2012

7:30-8:00	Breakfast –7 th Floor Foyer
8:00-8:05	Welcome and housekeeping

Session #5 - Molecular Virology

Session Chair: Dick Compans (IPIRC)

8:05 – 8:30	New insights on NS1 function – Adolfo Garcia-Sastre (CRIP)
8:30 – 8:55	Cleavage Activation of the HA of Human Influenza Virus Subtypes – Gary Whittaker & Karen Conrad (NYICE)
8:55 – 9:20	Role of PB1-F2 in the context of swine influenza strains in the pig model – Daniel Perez (CRIP)
9:20 – 9:35	Influenza HA subtypes demonstrate divergent phenotypes for cleavage activation and pH of fusion: implications for host range and adaptation to humans – Summer Galloway (IPIRC)
9:35 – 9:50	Influenza virus vs. host cell: a matter of life and death – Jerry Aldridge (SJCEIRS)
9:50 – 10:05	Functional consequences of influenza virus morphology – Anice Lowen (IPIRC)
10:05 – 10:30	Special Topic : Impact of <u>non-influenza viruses</u> in respiratory disease in families – John Treanor (NYICE)
10:30 – 11:00	Break – 7 th Floor Foyer

Session #6 - Open Discussion

11:00 – 12:00	"The Way Forward in Influenza Research: A Dialogue with the NIAID Director" Convener: Dr. Anthony Fauci
12:00 – 1:00	Lunch - Cantor/Jolson, 9th Floor

Session #7 - Clinical Surveillance and Transmission

Session Chair: John Treanor (NYICE)



1:00 – 1:20	Modes of influenza transmission in households – Ben Cowling (SJCEIRS)
1:20 – 1:35	Faceoff: Using RFID in the ED to Fingerprint Influenza Transmission for Surveillance in Humans – Vicki Hertzberg (IPIRC)
1:35 – 1:50	Local and systemic immune responses predict clinical outcomes in influenza infection independently of age and viral titer – Paul Thomas (SJCEIRS)
1:50 – 2:05	Measuring H5N1 influenza prevalence in humans in Egypt - Ghazi Kayali (SJCEIRS)
	- Influenza Immunology :: Paul Thomas (SJCEIRS) & Dave Topham (NYICE)
2:05 – 2:25	Automated analysis of influenza-specific human responses to vaccination – Tim Mosmann (NYICE)
2:25 – 2:45	Serological memory and bone marrow plasma cells after influenza vaccination – Carl Davis (IPIRC)
2:45 – 3:05	Vaccination and re-vaccination of human subjects with drifted variants of experimental inactivated subunit H5 influenza vaccine: Enhanced antibody secreting memory B cells and serum antibody responses in previously vaccinated subjects – Dave Topham (NYICE)
3:05 – 3:25	Break – 7 th Floor Foyer
3:05 – 3:25 3:25 – 3:45	Break – 7 th Floor Foyer Linked B and T cell epitopes in influenza virus hemagglutinin – Joshy Jacob (IPIRC)
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3:25 – 3:45	Linked B and T cell epitopes in influenza virus hemagglutinin – Joshy Jacob (IPIRC) The Specificity and Function of CD4 T cells in Response to Influenza Virus
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3:25 - 3:45 3:45 - 4:05 4:05 - 4:20 4:20 - 4:35 4:35 - 4:50	Linked B and T cell epitopes in influenza virus hemagglutinin – Joshy Jacob (IPIRC) The Specificity and Function of CD4 T cells in Response to Influenza Virus Infection and vaccination – Andrea Sant (NYICE) CD4 T cells respond to inactivated and live attenuated influenza vaccine with different kinetics – Xi Li (NYICE) IFNy-IL-2+ producing bias in human CD4 T cell cytokine profiles in response to recent influenza infection – Jason Weaver (NYICE) Multiple Distinct Forms of CD8+ T cell Cross-reactivity and Specificities revealed after 2009 H1N1 Influenza A Virus Infection in Mice – Hailong Guo (NYICE) Mapping drift of human H2N2 influenza viruses between 1957 and 1968 –



Wednesday August 1, 2012

7:30-8:00	Breakfast – 7 th Floor Foyer
9:00-12:15	Concurrent data manager breakout session (Soho Complex)
8:00-8:05	Welcome and housekeeping

Session #9 Panel Discussion: CEIRS Training Program

8:05 – 9:00 "CEIRS Training Program"

Convener: Adolfo Garcia-Sastre

Panelists: (10 min each)

Randy Albrecht "Highlights of the CEIRS training program"

Felix Santiago "The use of recombinant hemagglutinin proteins in the study of

influenza vaccination"

Martin Linster "Difficulties rescuing mutant viruses: improving reverse

genetics protocols"

Dan Dlugolenski "Phenotypic comparison of viral reassortments derived from

co-infection of swine epithelial cells"

Karl Ciuoderis "Surveillance and characterization of circulating Influenza A

viruses in the Llanos region of Colombia"

General Discussion

Session #10 CEIRS Surveillance Related Research - Avian

Session chairs: Robert Webster (SJCEIRS) & TBD (MCEIRS)

9:00 – 9:25	Influenza Surveillance: What a Novel Idea? - Mike Osterholm (MCEIRS)
9:25 – 9:50	Wild Bird Surveillance for Influenza A Viruses: Moving Forward – Dave Stallknecht (MCEIRS)
9:50 – 10:05	Surveillance for Avian Influenza Viruses in Live Poultry Markets in Tanzania: A Descriptive Epidemiologic Study - Peter Msoffe (MCEIRS)
10:05 – 10:25	Break – 7 th Floor Foyer
10:25 – 10:40	Understanding Temporal and Spatial Patterns in Influenza A Virus Infections in Northwest Ohio Using Migratory Patterns of Mallards (Anas platyrhynchos) -
	Anthony Fries (MCEIRS)
10:40 – 10:55	Anthony Fries (MCEIRS) Understanding Factors that Influence Low-Pathogenic Al Virus Infection in Ducks and Gulls – Justin Brown (MCEIRS)



11:10 – 11:25	A point mutation in H5N1 HPAIV HA affects its processing and cell fusion activity - Chinglai Yang (IPIRC)
11:25 – 11:50	Long-term evolutionary consequences of viral migration between wild bird populations – Justin Bahl (SJCEIRS)
11:50 – 12:15	Avian influenza A virus ecology and surveillance in wild birds in Georgia: 2009-2011 - Nicola Lewis (CRIP)

Session #11 CEIRS Surveillance Related Research - Swine and Other species Session chairs: Dick Slemons (MCEIRS)

1:15 – 1:40	Genetic Diversity and Transmission Dynamics in Swine Populations - Montse Torremorell (MCEIRS)
1:40 – 1:55	Genesis, establishment and risk assessment of pandemic 2009 reassortant viruses emerging from pigs in China – Huachen Maria Zhu (SJCEIRS)
1:55 – 2:10	Swine influenza surveillance in Guatemala: A nation-wide survey, 2010-2011 – Ana Silvia Gonzalez Reiche (CRIP)
2:10 – 2:25	Detection of Influenza Viruses among Pigs in a Kenyan Slaughterhouse - Kariuki Njenga (MCEIRS)
2:25 – 2:40	Potential Public Health Risk From Subclinical Influenza A Virus Infections in Swine at Agricultural Exhibitions in Ohio – Andrew Bowman (MCEIRS)
2:40 – 2:55	Characterization of pandemic H1N1 influenza viruses isolated from Northern Elephant Seals - Ignacio Mena (CRIP)
2:55 – 3:10	Canine influenza (H3N2) Outbreak in a Small-Animal Hospital in Thailand - Napawan Bunpapong (MCEIRS)
3:10 – 3:30	Program Highlights and Concluding Remarks
3:30	Adjourn

6TH ANNUAL CEIRS NETWORK MEETING POSTERS

 $\begin{tabular}{ll} Monday, July 30-5:05-6:30PM & Transmission, Immunology/Pathogenesis \& Vaccines \\ Tuesday, July 31-5:15-6:30PM & Surveillance \\ \end{tabular}$

TRANSMISSION

01	Barman, S.	SJCEIRS	Pathogenicity and Transmissibility of North American Triple Reassortant Swine Influenza A Viruses in Ferrets	
02	Campbell, P.J.	IPIRC	Role of the M Segment in 2009 Pandemic H1N1 Replication and Transmission	
03	Allerson, M.	MCEIRS	Impact of Immunity on Influenza A Virus Transmission in Pig Populations	
04	Allerson, M.	MCEIRS	Indirect Transmission of Influenza A Virus in Pig Populations under Two Different Biosecurity Settings	
05	Liu, Q.	SJCEIRS	Increased replication and transmissibility of reassortant H9 influenza viruses with genes from pandemic H1N1 virus in pigs	
06	Steel, J.	IPIRC	Molecular factors affecting the transmission of the pH1N1 influenza virus	
07	Yoon, SW.	SJCEIRS	Role of hemagglutinin residues in the emergence of the 2009 H1N1 pandemic virus	
VIROLOGY				
08	Whittaker, G.R.	NYICE	Acquisition of a novel eleven amino acid insertion directly N-terminal to a tetrabasic cleavage site confers intracellular cleavage of an H7N7 influenza virus hemagglutinin	
09	Dlugolenski, D.	IPIRC	Influenza virus coinfection of human and swine epithelial cell lines results in differential reassortment patterns and generation of novel recombinants	
10	Dlugolenski, D.	IPIRC	Pteropus alecto epithelial kidney cells are susceptible to infection with human and avian strains of influenza A virus and allow for novel reassortments	
11	Russell, C.J.	SJCEIRS	Energy barrier for HA activation helps determine influenza virus biology	

12	Takimoto, T.	NYICE	Identification of Influenza A PB2 Residues Involved in Enhanced Polymerase Activity in Mammalian Cells at Low Temperature
13	Yen, HL.	SJCEIRS	Comparative Mutational Analyses of Influenza viruses
14	Costello, D.A.	NYICE	Individual Virion Studies of Influenza Virus Fusion to Biomimetic Membranes
15	Lowen, A.C.	IPIRC	Functional consequences of Influenza virus morphology
16	Tse, L.V.	NYICE	Modification of the pH1N1 influenza hemagglutinin cleavage site modulates viral growth kinetics by increasing the protease spectrum for viral activation
17	Finch, C.L.	CRIP	Reverse genetics of Influenza B virus, Victoria lineage
18	Aggarwal, S.	NYICE	The E627K mutation in the PB2 subunit of influenza A polymerase affects promoter binding at higher temperatures
19	Duan, S.	SJCEIRS	How did they get so fit? – The molecular determinants of the naturally-occurring neuraminidase inhibitor-resistant H1N1 influenza A viruses
20	Gao, Q.	CRIP	The influenza A virus PB2, PA, NP and M segments play a pivotal role during genome packaging
21	Fabrizio, T.P.	SJCEIRS	The propensity of swine viruses to reassort and grow with human glycoproteins
22	Noble, E.	NYICE	Endonuclease Substrate Selectivity Characterized with Full- Length PA of Influenza A Virus Polymerase
23	Ye, L.	IPIRC	Mutation of a glycosylation site in the HA of SX H5N1 HPAIV eliminates the need of polybasic residues for trypsin-independent processing and cell fusion activities
24	Diaz, A.	MCEIRS	Whole Genome Genetic Diversity of Influenza A Virus within and between Pigs
25	Bialas, K.	NYICE	Influenza virion morphology correlates with efficiency of viral spread <i>in vitro</i>
IMM	UNOLOGY/PATHO	GENESIS	
26	Ramos, I.	CRIP	Contribution of dsRNA and CPSF Binding Domains of

Influenza Virus NS1 to the Inhibition of Type I IFN	
Production and Activation of human Dendritic Cells	

			Troduction and receivation of Haman Defiance Cells
27	Alam, S.	NYICE	The specificity of CD4 T cells and its influence to promote a specific antibody response after influenza infection
28	Sánchez, M.T.	CRIP	Bimolecular fluorescence complementation assay, a new tool to study viral and host cell protein interactions
29	Richard, M.	CRIP	Replication of avian influenza viruses in respiratory and intestinal tracts explants of chicken and ducks
30	Ayllon, J.	CRIP	Differential activation of heterotypic PI3K complexes by influenza virus NS1
31	Sreevatsan, S.	MCEIRS	Generation of Novel Single Chain Variable Fragments Against Hemagglutinin 3 for Influenza Surveillance in Swine
32	Bortz, E.	CRIP	DDX17 regulates highly pathogenic avian influenza virus replication in human cells according to PB2 genotype at position 627
33	Martínez-Romero, C.	CRIP	Residues 200 and 227 in the hemagglutinin of swine-origin H1N1 2009 influenza virus affect receptor binding and murine tropism
34	Dlugolenski, D.	IPIRC	Differential induction of antiviral genes in primary human and swine bronchial epithelial cells upon infection with Influenza A viruses
35	Xu, K.	CRIP	Isolation and characterization of an H9N2 influenza virus isolated in Argentina
36	Varga, Z.T.	CRIP	The Influenza Virus PB1-F2 Protein Inhibits the Induction of Type I Interferon by Binding to MAVS and Decreasing the Mitochondrial Membrane Potential
37	Minicucci, L.	MCEIRS	Avian Influenza at the Human-Animal Interface: Study Design and Virus Selection to Assess Exposure Risks in Wildlife and Conservation Professionals
38	Dash, P. & Sanders, C.J.	SJCEIRS	Early Replication and Host Response Characteristics Derived From the Parental Swine Strains Determine the 2009 Pandemic H1N1 Phenotype
39	Rajsbaum, R.	CRIP	Species-Specific Inhibition of RIG-I Ubiquitination and IFN Induction by the Influenza A Virus NS1 Protein

40	Wong, SS.	SJCEIRS	Flu09: A serological and virological profile of influenza infected children and their household contacts
41	Schmolke, M.	CRIP	Length of the NS1 linker region determines pathogenicity of a highly pathogenic H5N1 influenza A virus
42	Bui, V.N.	CRIP	H4N8 subtype influenza viruses isolated from shorebirds and causing severe respiratory disease in mice are PB1 reassortants: strategies to identify the viral genetics responsible for murine pathogenesis
43	Angel, M.	CRIP	In vivo characterization of influenza gene reassortment and constellation selection
44	Albrecht, R.A.	CRIP	Studies on the proteins encoded by the influenza A virus NS segment: potential contribution of the NEP to host tropism
45	McGargill, M.A.	SJCEIRS	Enhancing immunity to highly pathogenic H5N1 avian influenza A viruses
46	Ridenour, C.	IPIRC	Amino acid exchanges associated with continuous passage of a LPAIV chicken isolate in ducks suggests a strong selection process during adaptation to waterfowl
47	de Graaf, M.	CRIP	Characterization of influenza virus receptors in the respiratory tract of ferrets
48	Ye, J.	CRIP	Emergence of 2009 pandemic H1N1 with S186P in HA alters its phenotype?
SURVEILLANCE			
49	Oshansky, C.M.	SJCEIRS	Age-associated immune factors during natural acute influenza infection as predictors of clinical outcome
50	Jairak, W.	MCEIRS	Swine Influenza Surveillance during the 2011-2012 Season in Thailand
51	Jairak, W.	MCEIRS	Serologic Surveillance of Influenza A Virus Antibodies in Dogs and Cats in Thailand
52	Holden-Wiltse, J.	NYICE	<u>B</u> io- <u>L</u> ab <u>I</u> nformatics <u>Server</u> (<u>BLIS</u>) A Data Integration System for Clinical Influenza Surveillance & Vaccination Studies with
53	Nolting, J.	MCEIRS	Biological Assay Data Using the Open-Source LabKey Server H14 Influenza A Virus: Where Have You Been?
54	González-Reiche, A.S	S. CRIP	Phylogenetic analysis of Type A Influenza viruses isolated

from waterfowl in Guatemala

55	Spackman, E.	MCEIRS	Assuring the Quality of Laboratory Testing Across the CEIRS Surveillance Network	
56	Cowling, B.J.	SJCEIRS	Assessing the severity and impact of 2009-H1N1	
57	Leyva-Grado, V.	CRIP	Serological evidence of influenza virus infection in South American guinea pigs	
58	Smith, G.J.D.	SJCEIRS	Evolutionary genomics of pandemic H1N1/2009 influenza in Singapore	
59	Chiwanga, G.	MCEIRS	Surveillance for Avian Influenza Viruses in Live Poultry Market in Tanzania: Serological Evidence of Avian Influenza Viruses in Marketed Domestic Chickens	
60	Neumann, G.	CRIP	Influenza Virus Surveillance in Avian Species in Vietnam in 2010-2011	
61	King, CC.	IPIRC	Establishment of Chicken H6N1 Avian Influenza Viruses in Taiwan	
62	Shiilegdamba, E.	MCEIRS	High and Low Pathogenic Avian Influenza Virus Surveillance among Wild Birds in Mongolia from 2007-2011	
63	Perera, H.K.K.	SJCEIRS	Longitudinal study of swine influenza in Sri Lanka	
64	Ogawa, H.	CRIP	Surveillance of avian influenza virus in migratory waterfowls in eastern Hokkaido, Japan, during 2009 to 2010	
65	Slemons, R.	MCEIRS	Avian Influenza Virus in Sea Duck Populations	
66	Fourment, M.	SJCEIRS	Evolutionary dynamics and interspecies transmission of H2 influenza A viruses	
67	Sayfutdinova S.	CRIP	Influenza surveillance in North-Eastern Eurasia in 2008-2011	
68	Bunpapong, N.	MCEIRS	Survey of Influenza A Viruses in Free-Grazing Ducks in Thailand, 2011-2012	
69	Shao, H.	CRIP	A monoclonal antibody-based ELISA for serologic surveillance of H9 avian influenza virus	
VACCINES				

CD4 T cell expansion predicts pH1N1 vaccine neutralizing

70

Nayak, J.L.

NYICE

antibody response

71	Margine, I.	CRIP	An influenza B virus vaccine based on a conserved region of the hemagglutinin protein	
72	Sutton, T.C.	CRIP	Development of an Influenza-vectored Respiratory Syncytial Virus vaccine	
73	Richards, K.A.	NYICE	Trivalent inactivated influenza vaccines induce broad immunological reactivity to both internal virion components and influenza surface proteins	
74	Chen, H.	CRIP	Speeding up influenza vaccine development: partial and full plasmid-free reverse genetics systems	
75	Cox, A.	NYICE	The role of influenza virus polymerase stability in vaccine safety	
76	Miller, M.S.	CRIP	1976 and 2009 H1N1 Influenza Virus Vaccines Boost Anti- Hemagglutinin Stalk Antibodies in Humans	
77	Baker, S.F.	NYICE	Protection against lethal influenza challenge with a viral mimic	
78	Roumanes, D.	NYICE	CD4 T cells in elderly subjects respond in vivo and secrete effector cytokines in vitro in response to seasonal influenza vaccination	

Last Name	First Name	Affiliation	E-mail
Aggarwal	Shilpa	University of Rochester	shilpa_aggarwal@urmc.rochester.edu
Alam	Shabnam	University of Rochester	shabnam_alam@urmc.rochester.edu
Albrecht	Randy	Mount Sinai School of Medicine	randy.albrecht@mssm.edu
Aldridge	Jerry	St. Jude Children's Research Hospital	jerry.aldridge@stjude.org
Allerson	Matthew	University of Minnesota	alle0482@umn.edu
Amonsin	Alongkorn	Chulalongkorn University	alongkorn.a@chula.ac.th
Anderson	Janet	University of Minnesota	ander172@umn.edu
Angel	Matthew	University of Maryland	matthew.g.angel@gmail.com
Auchincloss	Hugh	National Institutes of Health	hugh.auchincloss@nih.gov
Ayllon	Juan	Mount Sinai School of Medicine	juan.ayllon@mssm.edu
Bahl	Justin	Duke-National University of Singapore	justin.bahl@duke-nus.edu.sg
Baker	Steven	University of Rochester	steven_baker@urmc.rochester.edu
Bao	Yiming	National Institutes of Health	bao@ncbi.nlm.nih.gov
Barman	Subrata	St. Jude Children's Research Hospital	subrata.barman@stjude.org
Baumgarth	Nicole	University of California, Davis	nbaumgarth@ucdavis.edu
Belicha	Alan	Mount Sinai School of Medicine	alan.belicha@mssm.edu
Bender	Jeffrey	University of Minnesota	bende002@umn.edu
Bernaciak	Teresa	National Institutes of Health	bernaciaktm@niaid.nih.gov
Bialas	Kristy	University of Rochester	kristy_bialas@urmc.rochester.edu
Boonyapisitsopa	Supanat	Chulalongkorn University	supanat.b@gmail.com
Bortz	Eric	Mount Sinai School of Medicine	eric.bortz@mssm.edu
Bouvier	Nicole	Mount Sinai School of Medicine	nicole.bouvier@mssm.edu
Bowman	Andrew	Ohio State University	bowman.214@osu.edu
Brown	Justin	University of Georgia	jubrown1@uga.edu
Bui	Vuong Nghia	Obihiro University	buinghiavuong@gmail.com
Bunpapong	Napawan Napawan	Chulalongkorn University	napawan.nb@gmail.com
Campbell	Patricia	Emory University	patricia.campbell2@emory.edu
Camping	Ryan	Mount Sinai School of Medicine	ryan.camping@mssm.edu
Cardona	Carol	University of Minnesota	ccardona@umn.edu
Casadevall	Arturo	Albert Einstein College of Medicine	arturo.casadevall@einstein.yu.edu
Cassingham	Tanya	Emory University	tcassin@emory.edu
Chan	Michael	The University of Hong Kong	mchan@hku.hk
Chaves	Francisco	·	
Chen		University of Rochester University of Maryland	francisco_chaves@urmc.rochester.edu vetchj@umd.edu
	Hongjun	· · · · · · · · · · · · · · · · · · ·	3
Chiwanga	Gaspar Karl	Sokoine University of Agriculture University of Wisconsin, Madison	chiwanga2000@yahoo.com
Ciuoderis	Richard		adolfmvz@gmail.com
Compans Conrad	Karen	Emory University	rcompan@emory.edu ksc64@cornell.edu
		Cornell University	dac325@cornell.edu
Costello	Deirdre	Cornell University	
Cowling	Ben	The University of Hong Kong	bcowling@hku.hk
Cox	Andrew	University of Rochester	andrew_cox@urmc.rochester.edu
Cox	Nancy	Centers for Disease Control & Prevention	nancy.cox@cdc.hhs.gov
Dash	Pradyot	St. Jude Children's Research Hospital	pradyot.dash@stjude.org
Davis	Carl	Emory University	cwdavi2@emory.edu
de Graaf	Miranda	Erasmus Medical Center	m.degraaf@erasmusmc.nl
Dhanasekaran	Vijaykrishna	Duke-National University of Singapore	vijay.dhanasekaran@duke-nus.edu.sg
Diaz	Andres	University of Minnesota	diazj002@umn.edu
Dixon	Dennis	National Institutes of Health	dd24a@nih.gov
Dlugolenski	Dan	University of Georgia	dandlugo@gmail.com
Duan	Susu	St. Jude Children's Research Hospital	susu.duan@stjude.org
Duchene	Alain	University of Minnesota	alain@ccbr.umn.edu
Dugan	Vivien	J. Craig Venter Institute	vdugan@jcvi.org
Eggink	Dirk	Mount Sinai School of Medicine	dirk.eggink@mssm.edu
Elia	Richard	St. Jude Children's Research Hospital	richard.elia@stjude.org
Fabrizio	Thomas	St. Jude Children's Research Hospital	thomas.fabrizio@stjude.org
Fauci	Anthony	National Institutes of Health	anthony.fauci@nih.gov
Fernandez-Sesma	Ana	Mount Sinai School of Medicine	ana.sesma@mssm.edu
Finn	Michael	National Institutes of Health	michael.finn@nih.gov
Fouchier	Ron	Erasmus Medical Center	r.fouchier@erasmusmc.nl
Fourment	Mathieu	Duke-National University of Singapore	mfourment@duke-nus.edu.sg
Francis	Doreen	University of Rochester	doreen_francis@urmc.rochester.edu

Last Name	First Name	Affiliation	E-mail
Fries	Anthony	Ohio State University	fries.41@osu.edu
Galloway	Summer	Emory University	sgallo2@emory.edu
Ganesan	Natarajan	National Institutes of Health	ganesann@niaid.nih.gov
Gao	Qinshan	Mount Sinai School of Medicine	qinshan.gao@mssm.edu
García-Sastre	Adolfo	Mount Sinai School of Medicine	adolfo.garcia-sastre@mssm.edu
Gellin	Bruce	U.S. Department of Health & Human Services	bruce.gellin@hhs.gov
Gilbert	Martin	Wildlife Conservation Society	mgilbert@wcs.org
Gilmandyar	Dzhamala	University of Rochester	dzhamala_gilmandyar@urmc.rochester.edu
Glowinski	Irene	National Institutes of Health	iglowinski@niaid.nih.gov
Goff	Peter	Mount Sinai School of Medicine	petegoff@gmail.com
Gonzalez Reiche	Ana	University of Maryland	areiche1@umd.edu
Govorkova	Elena	St. Jude Children's Research Hospital	elena.govorkova@stjude.org
Gu	Zhiping	Northrop Grumman	zhiping.gu@ngc.com
Guo	Hailong	University of Rochester	hailong_guo@urmc.rochester.edu
Hai	Rong	Mount Sinai School of Medicine	rong.hai@mssm.edu
Halpin	Rebecca	J. Craig Venter Institute	rhalpin@jcvi.org
Heaton	Nicholas	Mount Sinai School of Medicine	nicholas.heaton@mssm.edu
Hegngi	Fidelis	U.S. Department of Agriculture	fidelis.n.hegngi@aphis.usda.gov
Heilman	Carole	National Institutes of Health	cheilman@niaid.nih.gov
Hersft	Sander	Erasmus Medical Center	s.herfst@erasmusmc.nl
Hertzberg	Vickie	Emory University	vhertzb@emory.edu
Holden-Wiltse	Jeanne	University of Rochester	jeanne_wiltse@urmc.rochester.edu
Hussein	Islam	Massachusetts Institute of Technology	ihus@mit.edu
Jabbarzadeh	Shirin	Emory University	sjabba2@emory.edu
Jacob	Joshy	Emory University Emory University	jjacob3@emory.edu
Jairak	Waleemas	Chulalongkorn University	waleemas.wj@gmail.com
Karlsson	Erik	St. Jude Children's Research Hospital	erik.karlsson@stjude.org
Karisson	Yoshihiro	University of Wisconsin, Madison	kawaokay@svm.vetmed.wisc.edu
Kayali	Ghazi	St. Jude Children's Research Hospital	ghazi.kayali@stjude.org
Kiambi	Stella	Kenya Ministry of Livestock Development	gaichugi@yahoo.com
Kimble	Brian	University of Maryland	brian.kimble@gmail.com
King	Chwan-Chuen	National Taiwan University	chwanchuen@gmail.com
Klem	Ed Ed	Northrop Grumman	C
Knowlden	Zackery	University of Rochester	ed.klem@ngc.com
		Mount Sinai School of Medicine	zackery_knowlden@urmc.rochester.edu florian.krammer@mssm.edu
Krammer	Florian Scott		
Krauss		St. Jude Children's Research Hospital	scott.krauss@stjude.org
Kumar	Sanjeev	Northrop Grumman	skumar@virusbrc.org
LaPaglia Finch	Courtney	University of Maryland	clapagli@umd.edu
Larsen	Christopher	Vecna Technologies	clarsen@vecna.com
Lebarbenchon	Camille	University of La Réunion	camille.lebarbenchon@univ-reunion.fr
Lee	Eun Mi	National Institutes of Health	leee12@niaid.nih.gov
Levings	Randall	U.S. Department of Agriculture	randall.l.levings@aphis.usda.gov
Lewis	Nicola	University of Cambridge	nsl25@cam.ac.uk
Leyva-Grado	Victor	Mount Sinai School of Medicine	victor.leyva-grado@mssm.edu
Li	Xi	University of Rochester	xi_li@urmc.rochester.edu
Linster	Martin	Erasmus Medical Center	linster@erasmusmc.nl
Lipkin	Ian	Columbia University	wil2001@columbia.edu
Liu	Qinfang	Kansas State University	qfl@vet.k-state.edu
Lowen	Anice	Emory University	anice.lowen@emory.edu
Ma	Wenjun	Kansas State University	wma@vet.k-state.edu
Macken	Catherine	Los Alamos National Laboratory	cmacken@lanl.gov
Maher	Eileen	University of Wisconsin, Madison	eamaher@wisc.edu
Manicassamy	Balaji	Mount Sinai School of Medicine	balaji.manicassamy@mssm.edu
Margine	Irina	Mount Sinai School of Medicine	irina.margine@mssm.edu
Martinez	Carles	Mount Sinai School of Medicine	carles.martinez@mssm.edu
Martinez	Luis	University of Rochester	luis_martinez@urmc.rochester.edu
Mason	Robin	National Institutes of Health	rmason@niaid.nih.gov
Mathur	Punam	National Institutes of Health	mathurpu@niaid.nih.gov
Mazel-Sanchez	Béryl	Mount Sinai School of Medicine	beryl.mazel-sanchez@mssm.edu
McGargill	Maureen	St. Jude Children's Research Hospital	maureen.mcgargill@stjude.org

Last Name	First Name	Affiliation	E-mail
Medina Silva	Rafael	Pontifical Catholic University of Chile	rmedinas@med.puc.cl
Mena	Nacho	Mount Sinai School of Medicine	nacho.mena@mssm.edu
Mertens	Eva	University of California, Davis	evamertens@ucdavis.edu
Miller	Matthew	Mount Sinai School of Medicine	matthew.miller@mssm.edu
Milosovich	Karina	University of Minnesota	milo0014@umn.edu
Minicucci	Larissa	University of Minnesota	minic001@umn.edu
Moriarty	Megan	University of California, Davis	memoriarty@ucdavis.edu
Mosmann	Tim	University of Rochester	tim_mosmann@urmc.rochester.edu
Msoffe	Peter	Sokoine University of Agriculture	makengamsoffe@yahoo.co.uk
Munyua	Peninah	Kenya-Centers for Disease Control & Prevention	pmunyua@ke.cdc.gov
Nayak	Jennifer	University of Rochester	jennifer_nayak@urmc.rochester.edu
Nelson	Martha	National Institutes of Health	martha.nelson@nih.gov
Neu	Donna	University of Rochester	donna_neu@urmc.rochester.edu
Neumann	Gabriele	University of Wisconsin, Madison	neumanng@vetmed.wisc.edu
Njenga	Kariuki	Kenya-Centers for Disease Control & Prevention	knjenga@ke.cdc.gov
Noble	Erin	University of Rochester	erin_noble@urmc.rochester.edu
Nolting	Jacqueline	Ohio State University	nolting.4@osu.edu
Ogawa	Haruko	Obihiro University	hogawa@obihiro.ac.jp
Orenstein	Walter	Emory University	worenst@emory.edu
Oshansky	Christine	St. Jude Children's Research Hospital	christine.oshansky-weilnau@stjude.org
Osoro	Eric	Ministry of Public Health and Sanitation, Kenya	osoroe@yahoo.com
Osterholm	Michael	University of Minnesota	mto@umn.edu
Ostrowsky	Julie	University of Minnesota	jto@umn.edu
Otiang	Elkanah	Kenya Medical Research Institute-CDC	eotiang@kemricdc.org
Palese	Peter	Mount Sinai School of Medicine	peter.palese@mssm.edu
Pantin-Jackwood	Mary	U.S. Department of Agriculture	mary.pantin-jackwood@ars.usda.gov
Paquiot	David	Mount Sinai School of Medicine	david.paquiot@mssm.edu
Parker	Jerry	St. Jude Children's Research Hospital	jerry.parker@stjude.org
Patel	Jenish	Mount Sinai School of Medicine	jenish.patel@mssm.edu
Pavlisin	Amzie	Mount Sinai School of Medicine	amzie.pavlisin@mssm.edu
Peiris	Malik	The University of Hong Kong	malik@hku.hk
Perera	Harsha	University of Kelaniya, Sri Lanka	harshaperera@gmail.com
Perez	Daniel	University of Maryland	dperez1@umd.edu
Pickett	Brett	J. Craig Venter Institute	bpickett@jcvi.org
Plotsky	Andrea	Emory University	agplots@emory.edu
Poon	Leo	The University of Hong Kong	llmpoon@hku.hk
Post	Diane	National Institutes of Health	postd@niaid.nih.gov
Poulson	Rebecca	University of Georgia	rpoulson@uga.edu
Puryear	Wendy	Massachusetts Institute of Technology	wpuryear@mit.edu
Rajsbaum	Ricardo	Mount Sinai School of Medicine	ricardo.rajsbaum@mssm.edu
Ramos-Lopez	Irene	Mount Sinai School of Medicine	irene.ramos-lopez@mssm.edu
Richard	Mathilde	Erasmus Medical Center	m.richard@erasmusmc.nl
Richards	Katie	University of Rochester	katherine_skelly@urmc.rochester.edu
Richt	Juergen	Kansas State University	iricht@vet.k-state.edu
Ridenour	Callie	University of Georgia	callier@uga.edu
Roumanes	David	University of Rochester	david_roumanes@urmc.rochester.edu
Runstadler	Jonathan	Massachusetts Institute of Technology	jrun@mit.edu
Russell	Charles	St. Jude Children's Research Hospital	charles.russell@stjude.org
Samarasinghe	Amali	St. Jude Children's Research Hospital	amali.samarasinghe@stjude.org
Sanchez	Maria	Mount Sinai School of Medicine	maria.t.sanchez@mssm.edu
Sanders	Catherine	St. Jude Children's Research Hospital	catherine.sanders@stjude.org
Sant	Andrea	University of Rochester	andrea_sant@urmc.rochester.edu
Santiago	Felix	University of Rochester	felix_santiago@urmc.rochester.edu
Sayfutdinova	Sofya	Vector Labs	sayfutdinovas@gmail.com
Sayrutdinova Scheuermann	Richard		
Scheuermann Schmolke	Mirco	J. Craig Venter Institute Mount Sinai School of Medicine	rscheuermann@jcvi.org mirco.schmolke@mssm.edu
Schultz-Cherry Shortonelov	Stacey Alexander	St. Jude Children's Research Hospital	stacey.schultz-cherry@stjude.org
Shestopalov		Vector Labs Wildlife Conservation Society	shestopalov2@ngs.ru
Shiilegdamba	Enkhtuvshin	Wildlife Conservation Society	eshiilegdamba@wcs.org
Siedelman	Lori	University of Minnesota	sied0020@umn.edu
Slemons	Richard	Ohio State University	slemons.1@osu.edu

Last Name	First Name	Affiliation	E-mail
Smith	Derek	University of Cambridge	d.smith@zoo.cam.ac.uk
Smith	Gavin	Duke-National University of Singapore	gavin.smith@duke-nus.edu.sg
Solorzano	Alicia	University of Medicine and Dentistry of New Jersey	solorzal@umdnj.edu
Spackman	Erica	U.S. Department of Agriculture	erica.spackman@ars.usda.gov
Spiro	David	National Institutes of Health	david.spiro@nih.gov
Sreevatsan	Srinand	University of Minnesota	sreev001@umn.edu
Stallknecht	David	University of Georgia	dstall@uga.edu
Steel	John	Emory University	john.steel@emory.edu
Steinhauer	David	Emory University	dsteinh@emory.edu
Stewart	Lucy	J. Craig Venter Institute	lstewart@jcvi.org
Stockwell	Tim	J. Craig Venter Institute	tstockwell@jcvi.org
Suarez	David	U.S. Department of Agriculture	david.suarez@ars.usda.gov
Subbarao	Kanta	National Institutes of Health	kanta.subbarao@nih.gov
Sun	Guangyu	Vecna Technologies	gsun@vecna.com
Sutton	Troy	University of Maryland	tsutton@umd.edu
Takimoto	Toru	University of Rochester	toru_takimoto@urmc.rochester.edu
Tan	Gene	Mount Sinai School of Medicine	gene.tan@mssm.edu
Theamboonlers	Apiradee	Chulalongkorn University	apiradee.t@chula.ac.th
Thomas	Paul	St. Jude Children's Research Hospital	paul.thomas@stjude.org
Tompkins	Mark	University of Georgia	smt@uga.edu
Topham	David	University of Rochester	david_topham@urmc.rochester.edu
Torremorell	Montseurrat	University of Minnesota	torr0033@umn.edu
Treanor	John	University of Rochester	john_treanor@urmc.rochester.edu
Tse	Long Ping Victor	Cornell University	lt273@cornell.edu
Varga	Zsuzsanna	Mount Sinai School of Medicine	zsuzsanna.varga@mssm.edu
Versteeg	Gijs	Mount Sinai School of Medicine	gijs.versteeg@mssm.edu
Wan	Xiu-Feng	Mississippi State University	wan@cvm.msstate.edu
Wang	Taia	Mount Sinai School of Medicine	taia.wang@mssm.edu
Weaver	Jason	University of Rochester	jason_weaver@urmc.rochester.edu
Webby	Richard	St. Jude Children's Research Hospital	richard.webby@stjude.org
Webster	Robert	St. Jude Children's Research Hospital	robert.webster@stjude.org
Whittaker	Gary	Cornell University	gary.whittaker@cornell.edu
Williams	Jeffrey	University of Rochester	jeffrey_williams@urmc.rochester.edu
Wong	Diana	The University of Hong Kong	dywong2@hku.hk
Wong	Sook-San	St. Jude Children's Research Hospital	sook-san.wong@stjude.org
Xu	Kemin	University of Maryland	kmxu@umd.edu
Yang	Chinglai	Emory University	chyang@emory.edu
Yao	Qing Alison	National Institutes of Health	yaoal@niaid.nih.gov
Ye	Ling	Emory University	yling@emory.edu
Yen	Hui-Ling	The University of Hong Kong	hyen@hku.hk
Yondola	Mark	Mount Sinai School of Medicine	mark.yondola@mssm.edu
Yoon	Sun-Woo	St. Jude Children's Research Hospital	sun-woo.yoon@stjude.org
York	Ian	Centers for Disease Control & Prevention	ite1@cdc.gov
Zanin	Mark	St. Jude Children's Research Hospital	mark.zanin@stjude.org
Zaraket	Hassan	St. Jude Children's Research Hospital	hassan.zaraket@stjude.org
Zhou	Liwei	Northrop Grumman	liwei.zhou@ngc.com
Zhu	Maria	The University of Hong Kong	zhuhch@hku.hk
		1	***************************************