

Swine influenza activities in China

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Current situation

- 1.3 billion pigs: 43% of the total in the world
- Different breeding styles: backyard to large scale farms
- Closely contact with poultry
- Vaccines are not used
- Systemic surveillance is not in place

Virus isolated during 2002-2008

Year	Subtypes
2002	H1N1, H3N2, H9N2
2003	H1N1, H3N2, H9N2, H5N1 (also isolated one in 2001)
2004	H1N1, H1N2, H3N2, H9N2
2005	H1N1, H3N2, H9N2
2006	H1N1, H1N2, H3N2, H9N2
2007	H1N1, H1N2, H3N2, H9N2
2008	H1N1, H1N2, H3N2, H9N2

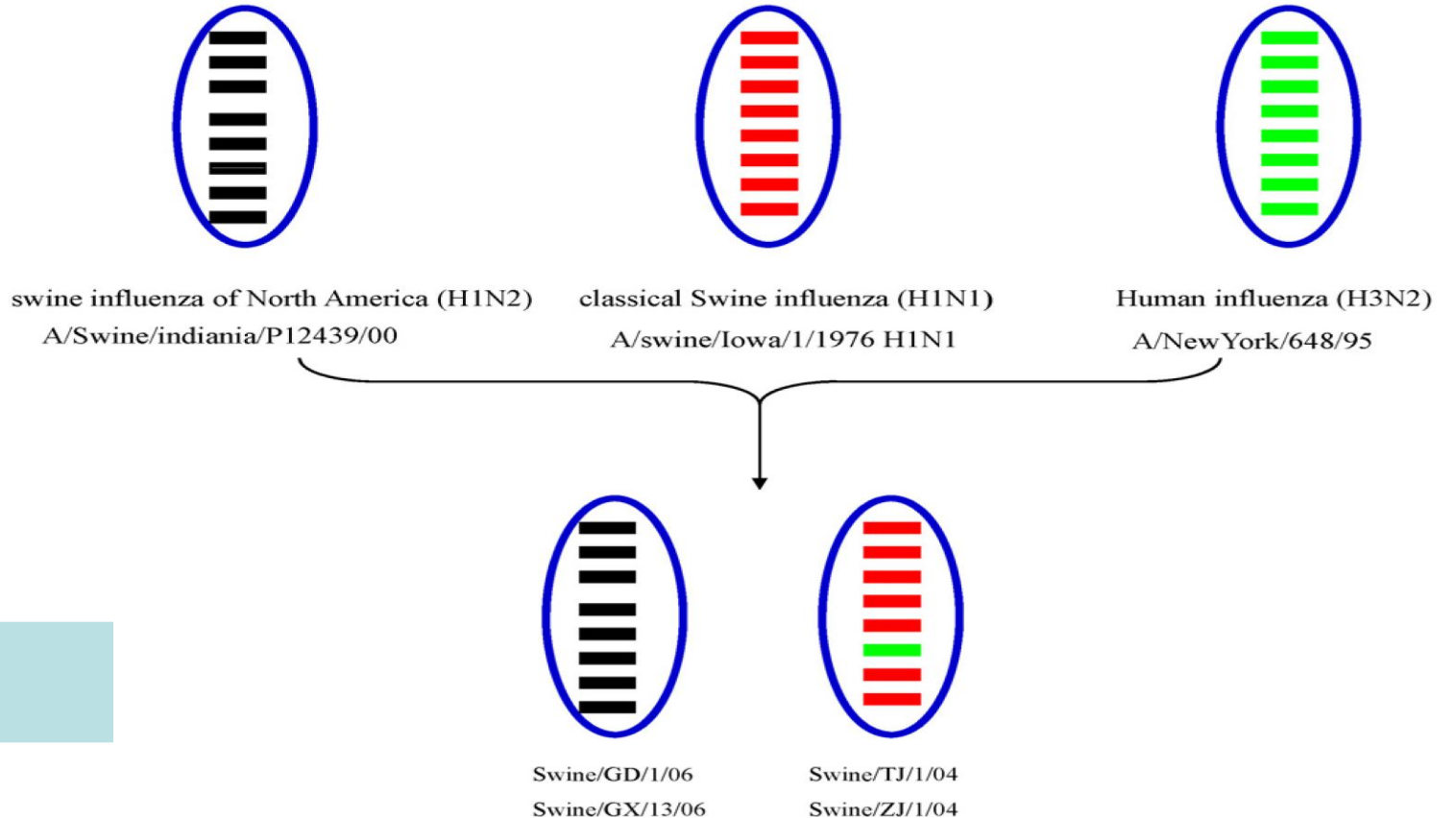
Serologic Surveillance in 2008

Place	Samples	Positive rate(%)			
		H1 subtype	H3 subtype	H5 subtype	H9 subtype
Henan	4150	23.7	50.8	0.1	0.2
Shandong	785	5.6	55.5	0.1	0.1
Jiangsu	590	19.3	58.8	0	0
Shanghai	394	34.4	54.7	0	0
Hubei	178	0	64.6	0	0
Hunan	105	2.86	77.1	0	0
Chongqing	176	15.8	76.3	0	0
Heilongjiang	175	22.7	62.7	0	0
Hebei	412	12.5	45.5	0	0
Guangdong	238	3.0	75.8	0	0
Total	7203	14.0	62.2	0.02	0.03

Short communication

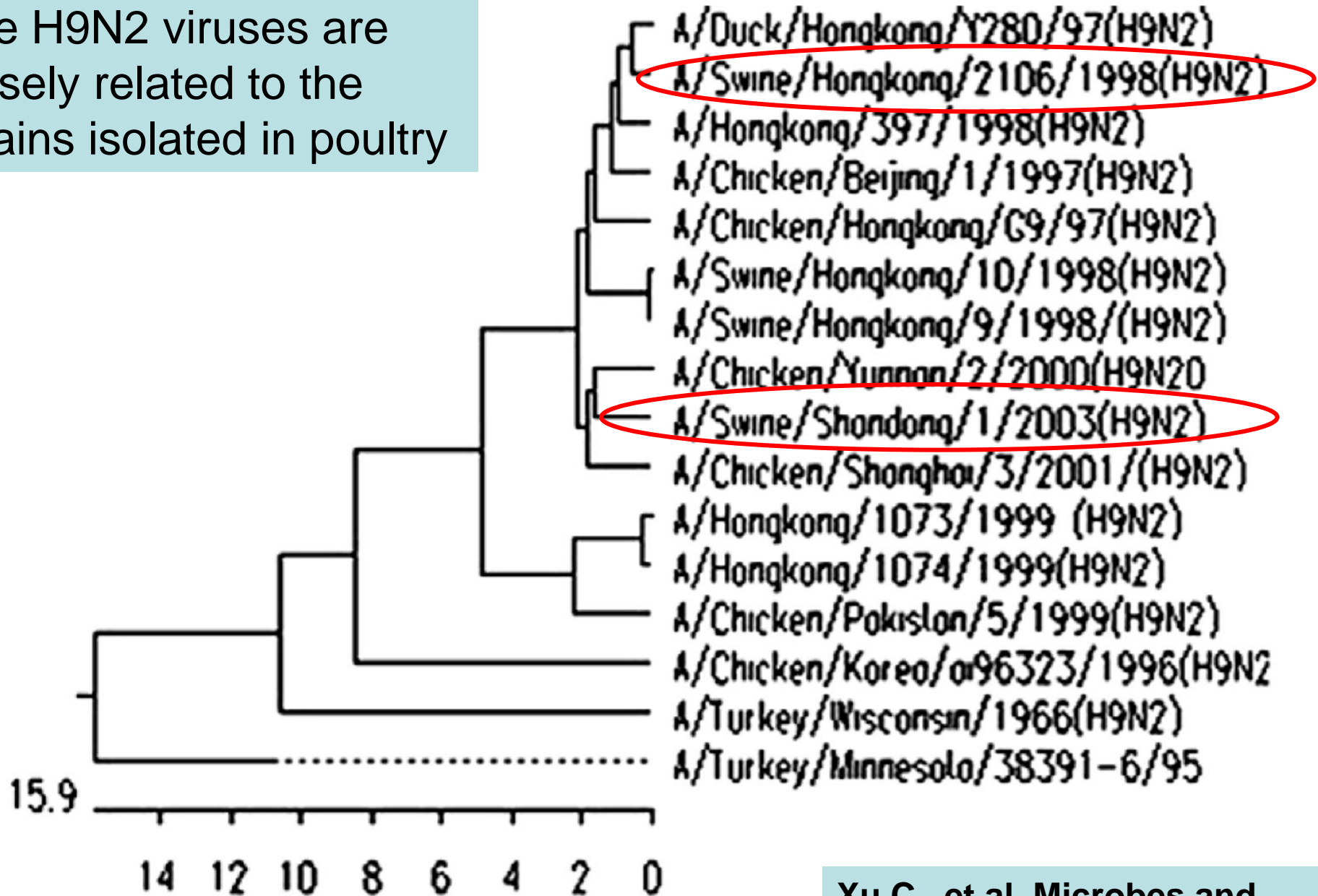
Two genotypes of H1N2 swine influenza viruses appeared among pigs in China

Chuantian Xu^{a,b}, Qiyun Zhu^a, Huanliang Yang^a, Xiumei Zhang^b, Chuanling Qiao^a, Yan Chen^a, Xiaoguang Xin^a, Hualan Chen^{a,*}



H1N2 viruses

The H9N2 viruses are closely related to the strains isolated in poultry



Xu C., et al. Microbes and infections, 2004

TABLE 2. Amino acid differences between two H5N1 swine influenza viruses and the duck virus DK/ZJ/00

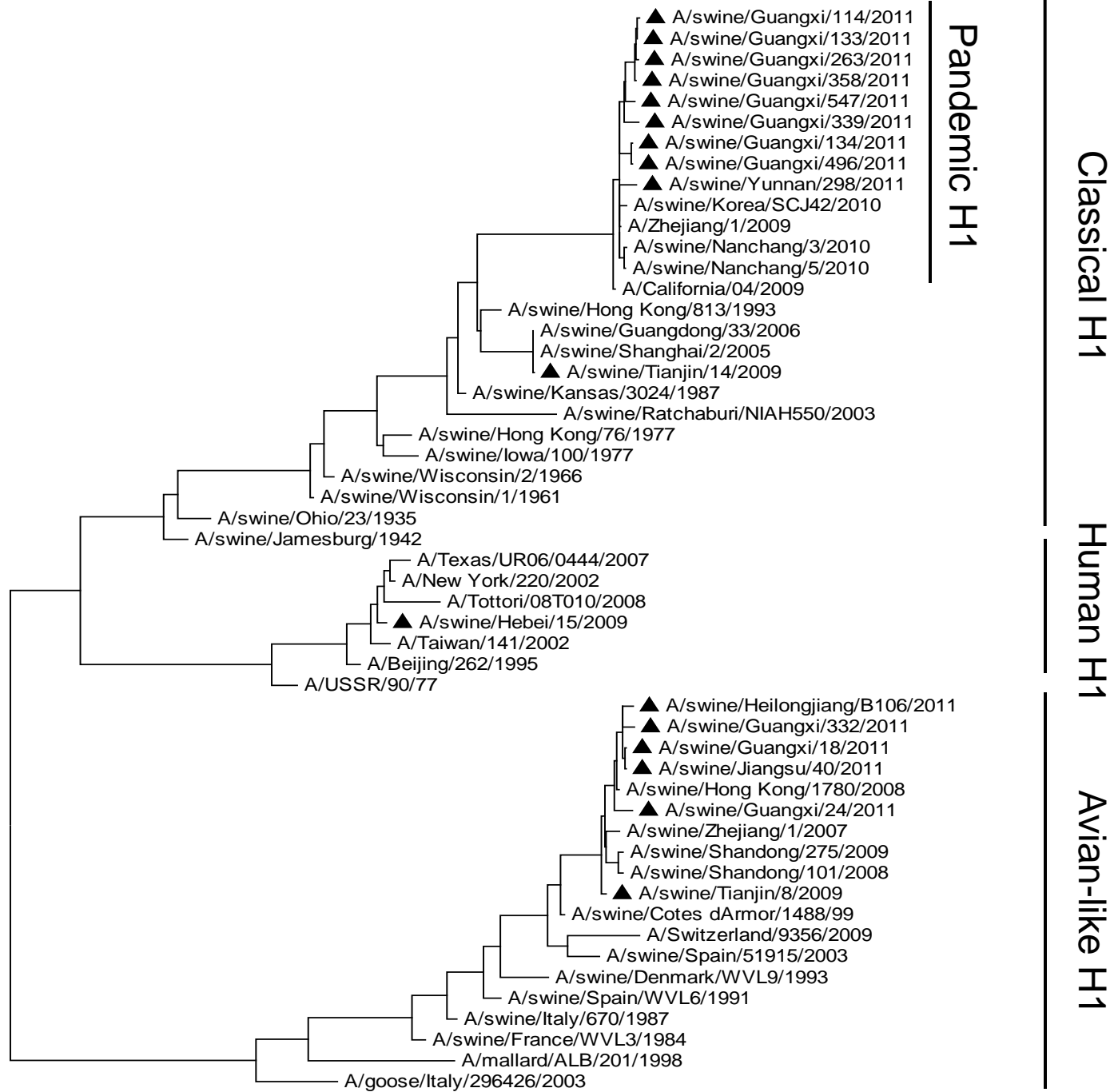
Protein	Amino acid position	Amino acid in virus ^a		
		DK/ZJ/00	SW/FJ/01	SW/FJ/03
PB2	51	V	M	M
	92	P	S	S
	164	L	M	M
	562	I	I	S
	679	L	P	P
	725	P	L	L
PB1	456	L	H	H
	486	L	R	R
PA	54	V	I	I
	330	I	V	V
	384	Y	C	C
	459	V	I	I
HA	16	S	G	G
	100	N	S	S
	168	K	K	E
	254	S	A	A
	444	L	I	L
NP	65	S	R	R
	257	T	I	I
	284	V	A	A
NA	34	V	I	I
M2	26	L	S	L
	65	T	T	A
	88	I	M	M
NS1	18	V	I	I
	69	P	L	L
	88	I	M	M
NS2	191–195	EALQR	<i>EALQR</i>	
	39–43	KLYRD	<i>KLYRD</i>	

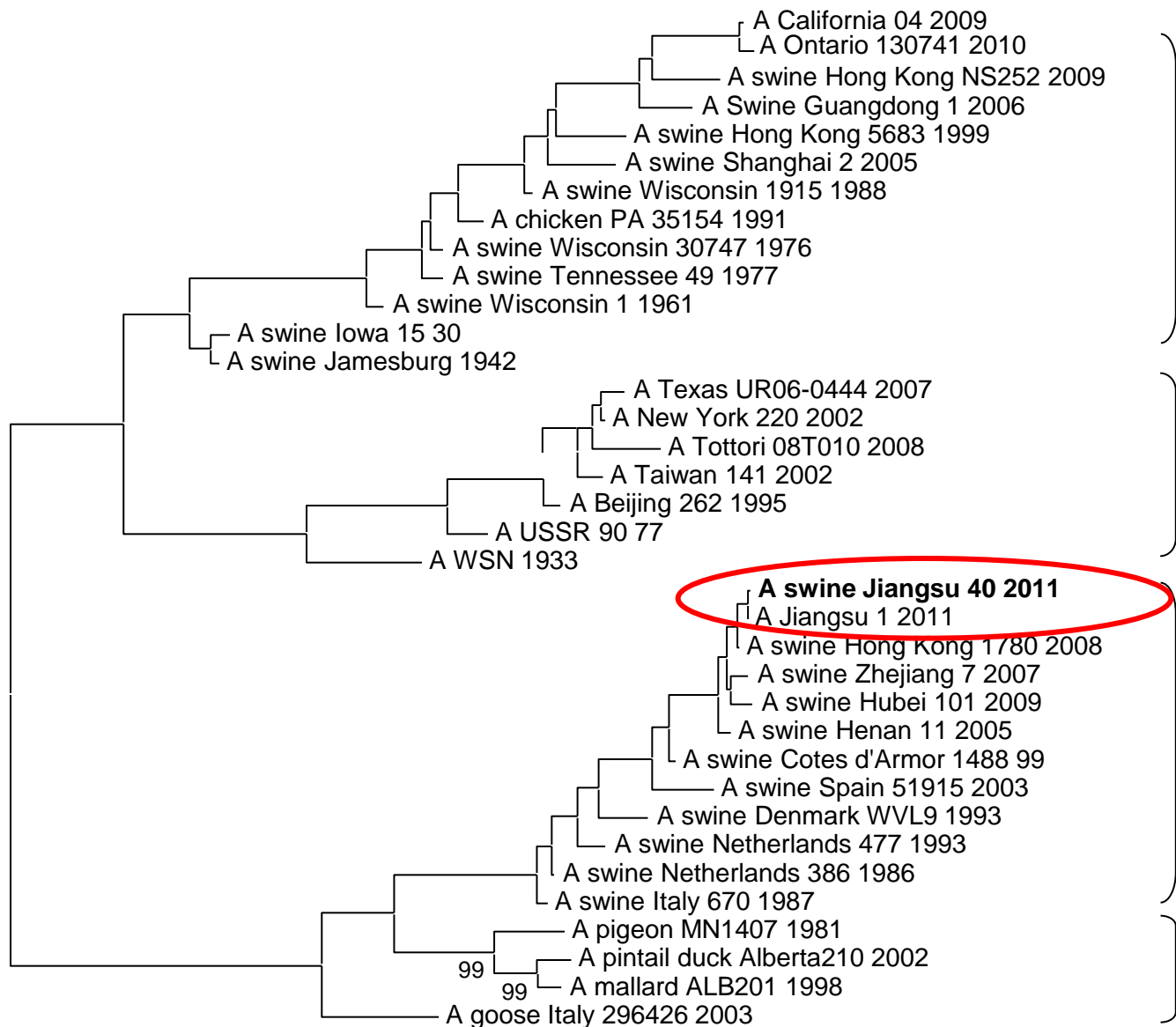
The H5N1 viruses isolated from pigs are genetically similar to the viruses isolated in ducks in China

Zhu Q et al.,
Journal of Virology,
2008

Total 55 SIVs were isolated from these samples, including 40 H1N1 SIVs, 11 H1N2 SIVs, and 4 H3N2 SIVs.

H1N1 HA genes





Classical swine H1N1

Human seasonal H1N1

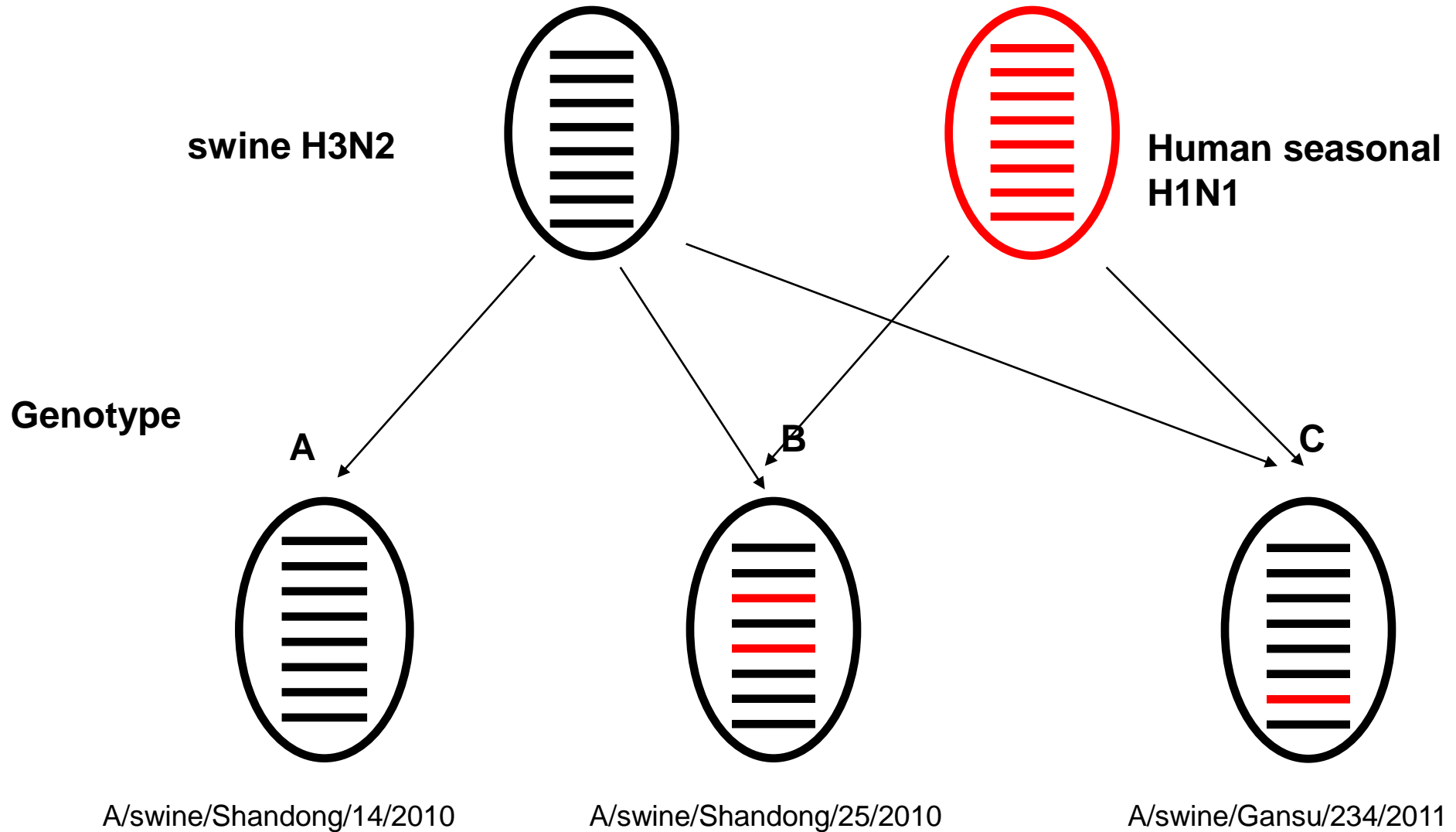
Caused human infection in Jiangsu Province in 2011

Avian-like H1N1

Avian H1N1

0.05

Different genotypes of H3N2 SIVs were detected in China



H1N1 swine influenza vaccine

Development of an inactivated vaccine (SC/PR8) against the infection of pandemic 2009 H1N1 viruses and endemic swine H1N1 or H1N2 influenza viruses in pigs.

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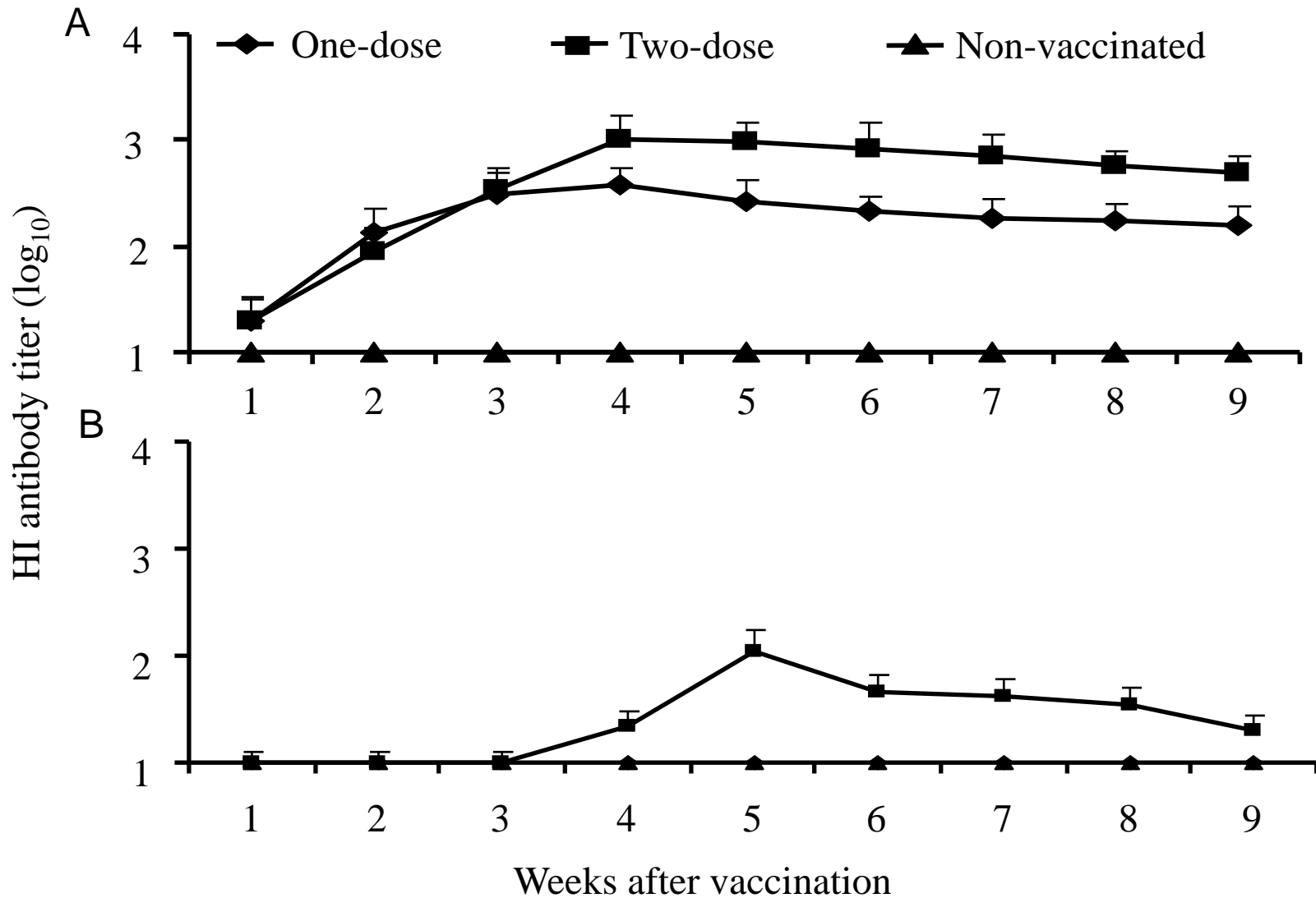
journal homepage: www.elsevier.com/locate/vetmic



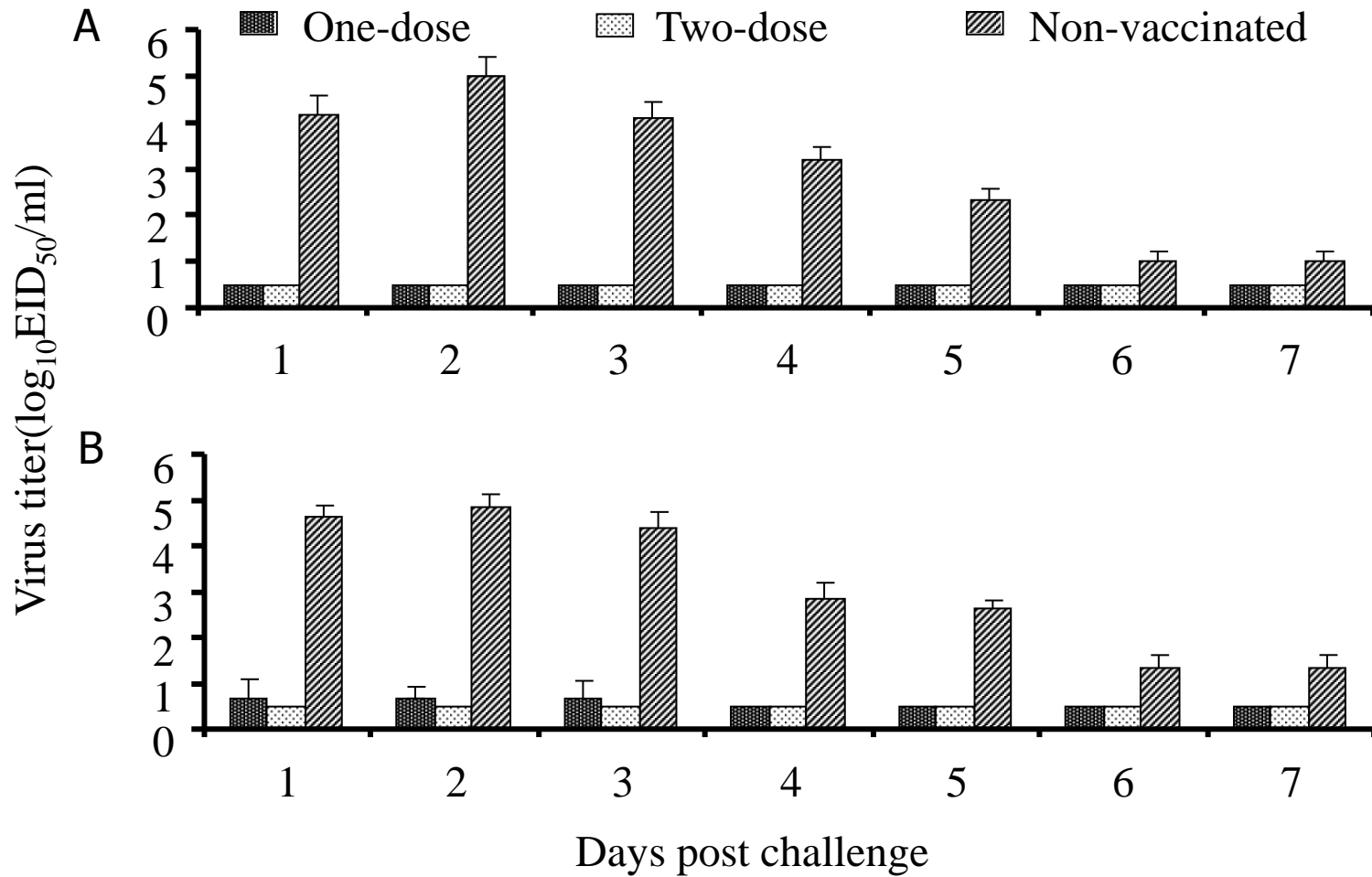
Reassortant H1N1 influenza virus vaccines protect pigs against pandemic H1N1 influenza virus and H1N2 swine influenza virus challenge

Huanliang Yang^{a,1}, Yan Chen^{a,1}, Jianzhong Shi^a, Jing Guo^a, Xiaoguang Xin^a, Jian Zhang^a, Dayan Wang^b, Yuelong Shu^b, Chuanling Qiao^{a,*}, Hualan Chen^{a,*}

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Antibody responses induced by SC/PR8 inactivated vaccines in the pigs. (A) Antibody titers against SC/09 virus. (B) Antibody titers against GD/06 virus.



Virus shedding in pigs after challenge with different swine influenza viruses.
 (A) SC/09 virus. (B) GD/06 virus.

Summary

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- The co-existing of 2009 pandemic H1N1 influenza virus and other subtypes viruses (e.g. avian like H1N1, H9N2 and H5N1) in the animals may produce new reassortants with pandemic potential.

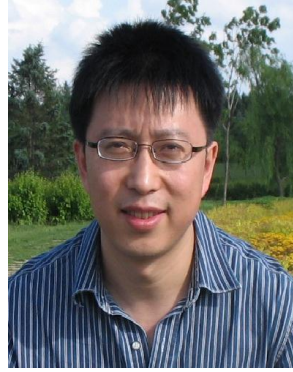
Summary

- Multiple subtypes of influenza viruses have been detected in pigs in China.
- The co-existing of 2009 pandemic H1N1 influenza virus and other subtypes viruses (e.g. avian like H1N1, H9N2 and H5N1) in the animals may produce new reassortants with pandemic potential.
- Systemic surveillance and analysis of SIVs are now high priorities in our research activities.

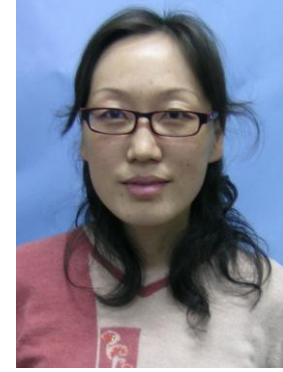
Acknowledgments



Dr. Chuanling Qiao



Dr. Huanliang Yang

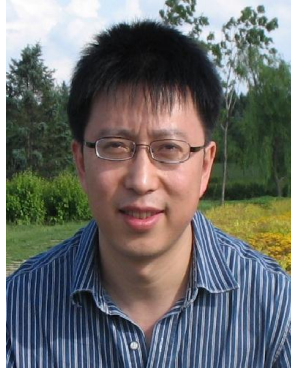


Dr. Yan Chen

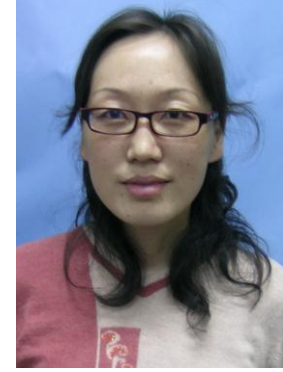
Acknowledgments



Dr. Chuanling Qiao



Dr. Huanliang Yang



Dr. Yan Chen

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- 973 programs (MOST)