



Building Capacity for Biodiversity and Human Health

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What do we concern for biodiversity and human health?

— —health sector's perspective

- Health benefits of biodiversity
- Health risks of biodiversity

Hypothesis for biodiversity and human health

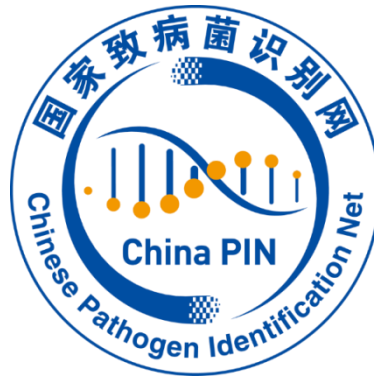
- Nonlinear effects of biodiversity on human health
 - Positive – health benefits
 - Negative – health risks
 - More positive than negative

Strategic plan for biodiversity and human health

- Maximize the positive effects (health benefits) of biodiversity
- Minimize the negative effects (health risks) of biodiversity

Road map for biodiversity and human health

- Surveillance and data integration
- Assessment on the effects of biodiversity
- Public health action for risk communication and risk management
- Public health action to maximize the benefits
- Minimize the impacts of public health action on biodiversity as much as possible

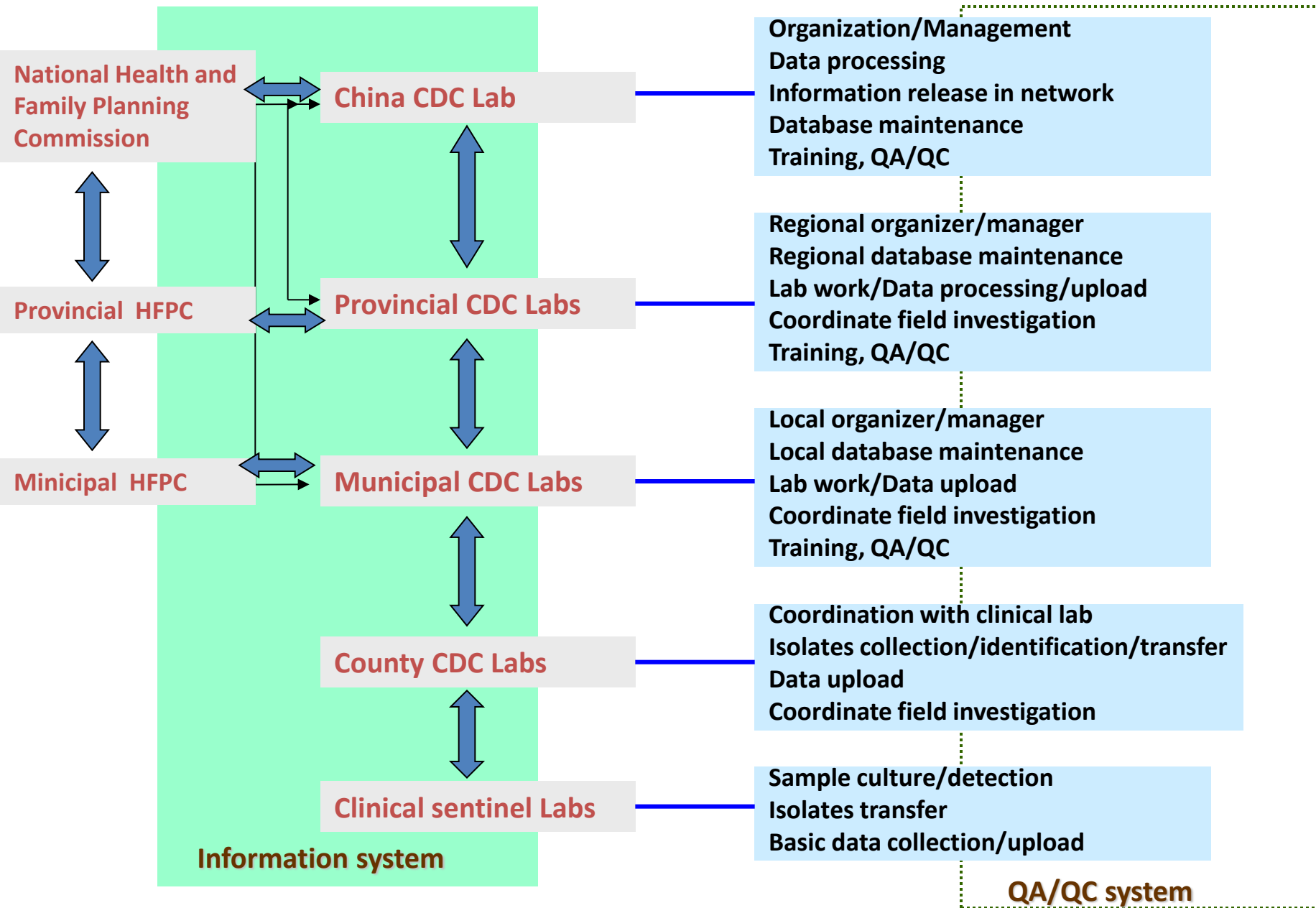


Chinese Pathogen Identification Net (China PIN)

Identify pathogens, Identify outbreaks, Identify outbreaks

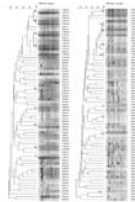
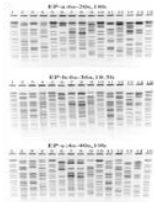
The national laboratory-based surveillance network for bacterial pathogens

China PIN

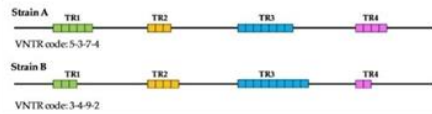


Technology

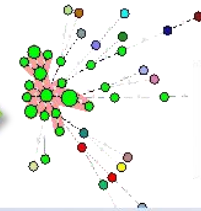
脉冲场凝胶电泳分析 (PFGE)



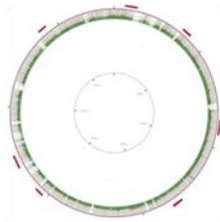
多位点可变数串联重复序列分析 (MLVA)



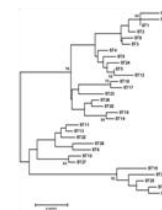
Strain	TR1	TR2	TR3	TR4
Strain A	5	3	7	4
Strain B	3	4	9	2



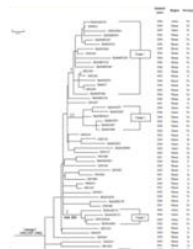
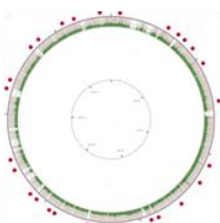
多基因位点测序列分型 (MLST或PLST)



```
rpoB_allele4 328 GAA TAGGCTCAAGGCAATTCCACT 389-  
rpoB_allele5 355 GAA TAGGCTCAAGGCAATTCCACT 416-  
rpoB_allele3 373 GAA TAGGCTCAAGGCAATTCCACT 434-  
rpoB_allele1 373 GAA TAGGCTCAAGGCAATTCCACT 434-  
rpoB_allele2 373 GAA TAGGCTCAAGGCAATTCCACT 434-
```



基于全基因组测序分型 (Genome sequencing)





Information System

Provincial users



VPN

Intranet

Analyse Server



Application Server

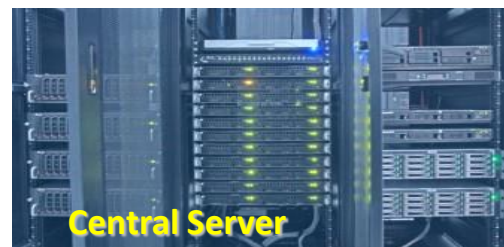
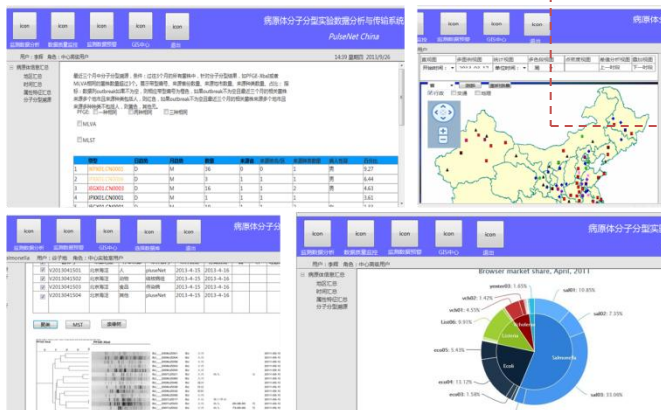
Central Data Management Server

Internet

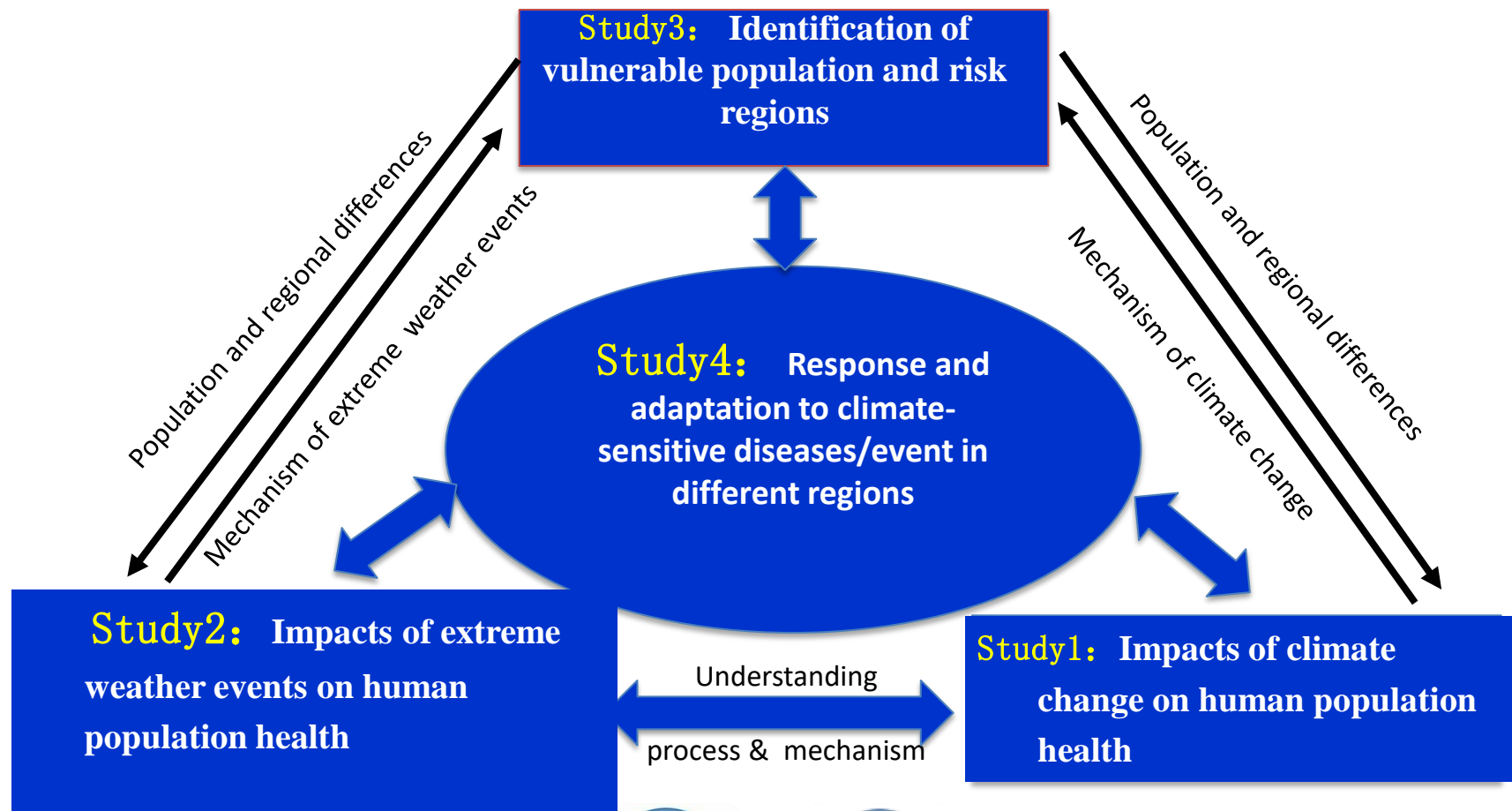
VPN



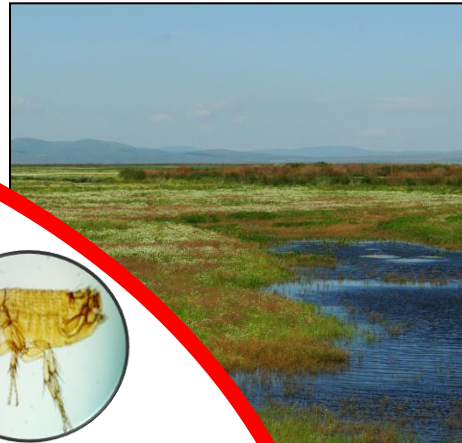
Intra users



Climate change impacts on health and adaptation mechanism



Plague ecology



Plague in China

PNAS

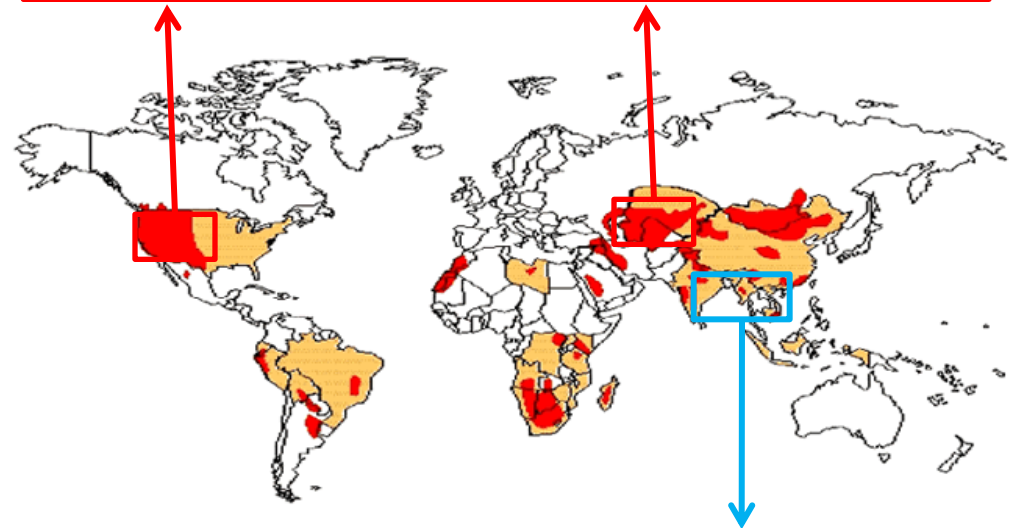
Nonlinear effect of climate on plague during the third pandemic in China

Lei Xu^{a,b,1}, Qiyong Liu^{c,1}, Leif Chr. Stige^d, Tamara Ben Ari^d, Xiye Fang^e, Kung-Sik Chan^f, Shuchun Wang^{a,2}, Nils Chr. Stenseth^{d,2}, and Zhibin Zhang^{a,2}

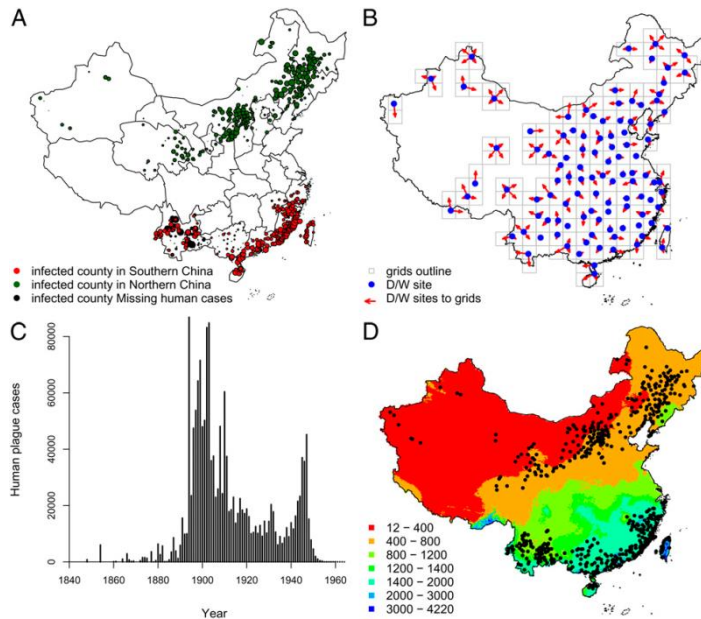
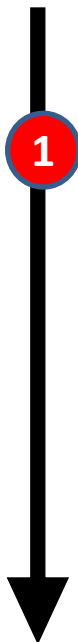
^aState Key Laboratory of Integrated Management on Pest Insects and Rodents, Institute of Zoology, Chinese Academy of Sciences, Beijing 100101, China; ^bState Key Laboratory for Infectious Diseases Prevention and Control, National Institute for Communicable Disease Control and Prevention, Chinese Centre for Disease Control and Prevention, Beijing 102206, China; ^cCentre for Ecological and Evolutionary Synthesis, Department of Biology, University of Oslo, Blindern, 0316 Oslo, Norway; ^dNational Institute for Communicable Disease Control and Prevention, Chinese Centre for Disease Control and Prevention, Beijing 102206, China; ^eDepartment of Statistics and Actuarial Sciences, University of Iowa, Iowa City, IA 52242; and ^fGraduate University of Chinese Academy of Sciences, Beijing 100049, China

- **Central Asia: positive**
- **Vietnam: negative**
- **WHY this difference?**

U. S. A : Parmenter et. al. , 1999; Enscoe et.al. , 2002)
Kazakhstan: (Stenseth, 2006; Kausrud, 2007; 2010)



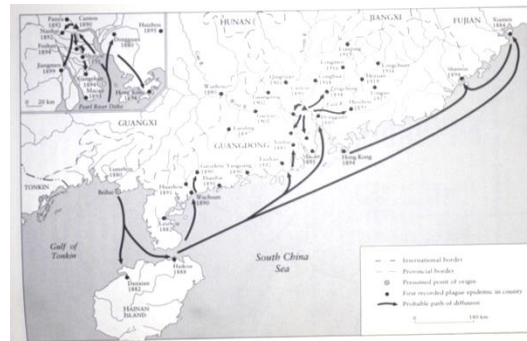
Viatan: (Cavanaugh, 1972; Pham, 2009)



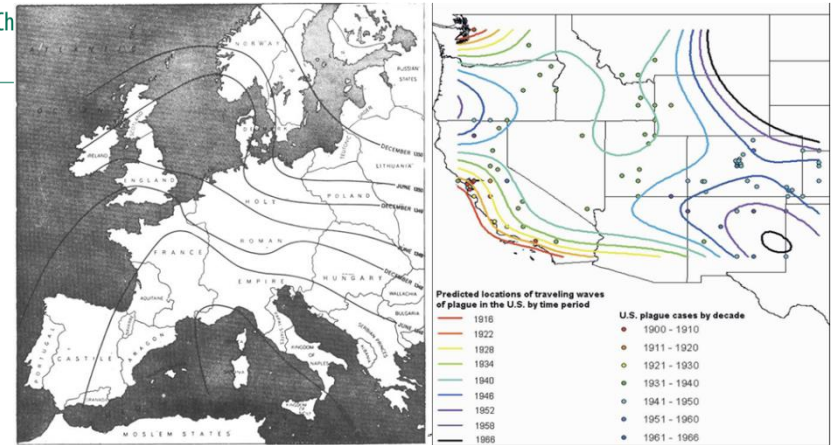
Plague in China

Wet climate and transportation routes accelerate spread of human plague

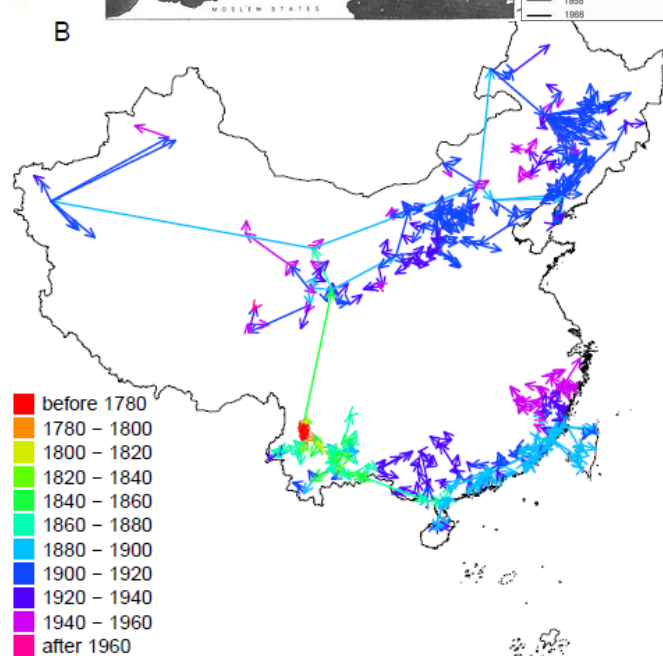
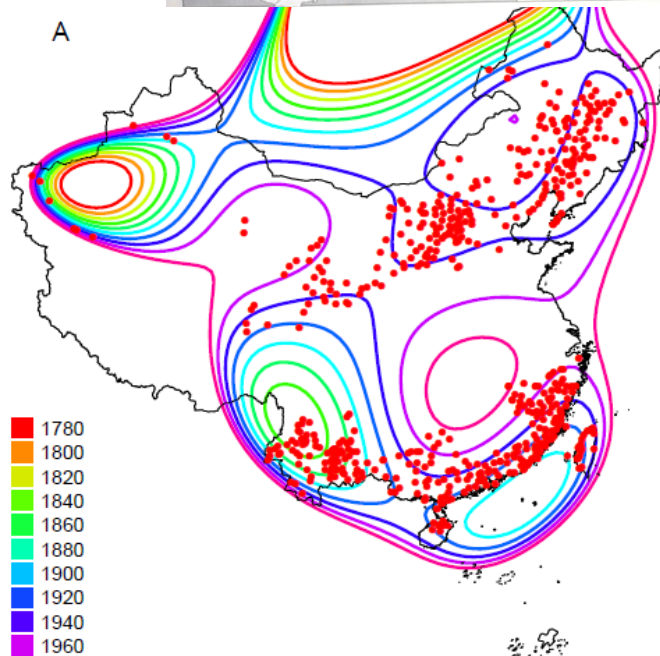
Lei Xu^{1,2}, Leif Chr. Stige², Kyrre Linné Kausrud², Tamara Ben Ari²,
Shuchun Wang³, Xiye Fang³, Boris V. Schmid², Qiyong Liu³, Nils Ch
and Zhibin Zhang¹



A



B



Plague in China

PROCEEDINGS B

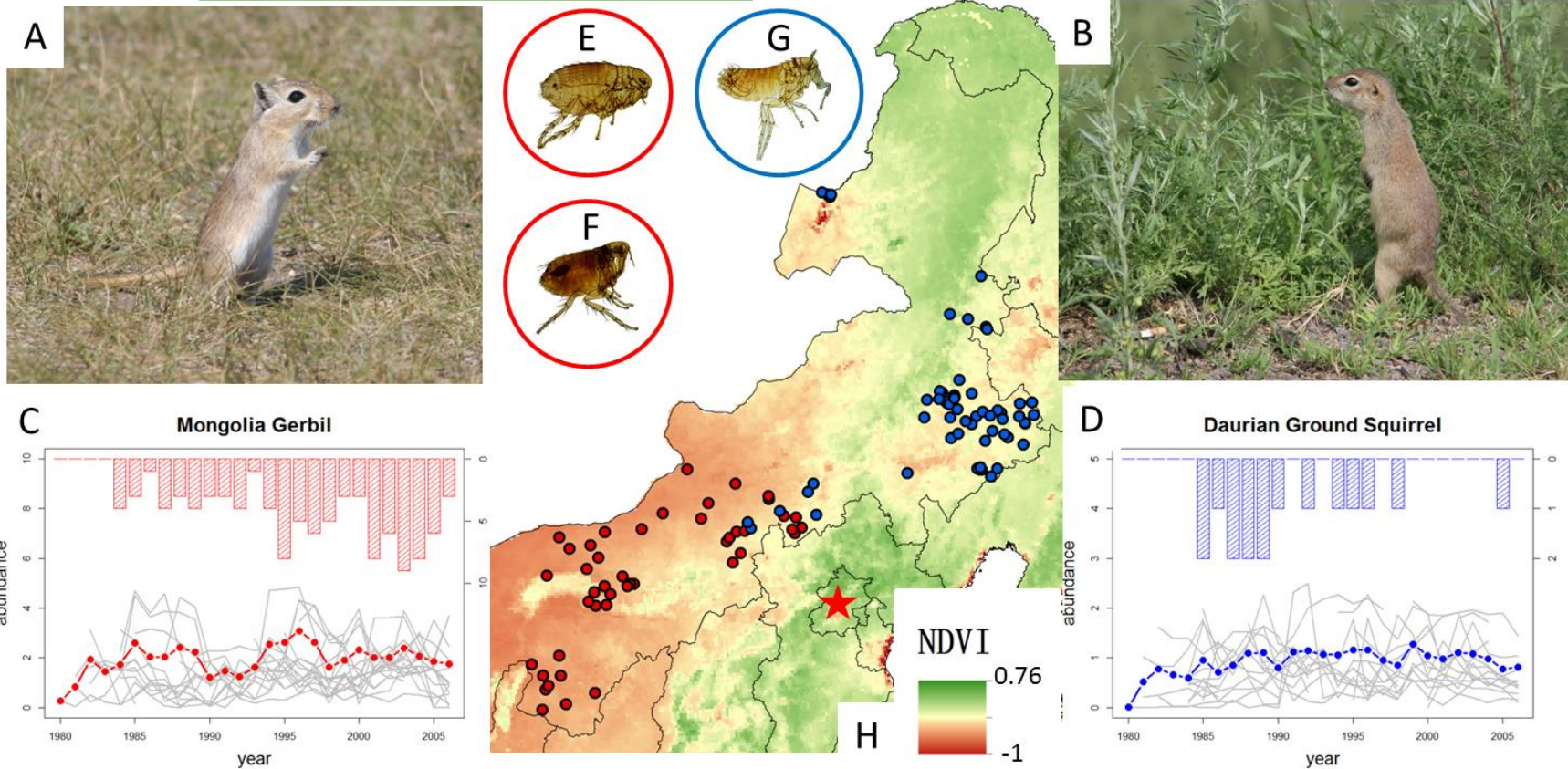
rspb.royalsocietypublishing.org

Research



The trophic responses of two different rodent–vector–plague systems to climate change

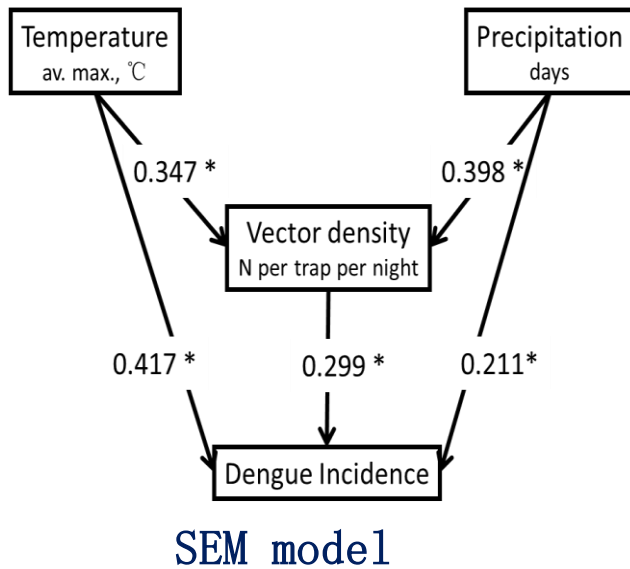
Lei Xu¹, Boris V. Schmid², Jun Liu³, Xiaoyan Si³, Nils Chr. Stenseth² and Zhibin Zhang¹



Plague surveillance locations for the **Mongolian gerbil natural plague foci** and the **Daurian ground squirrel natural plague foci** in Inner Mongolia, with the background color indicating the average vegetation that was measured by NDVI.



Climate variation drives dengue dynamics



Climate variation drives dengue dynamics

Lei Xu^{a,b,1}, Leif C. Stige^{b,1}, Kung-Sik Chan^c, Jie Zhou^d, Jun Yang^a, Shaowei Sang^a, Ming Wang^e, Zhicong Yang^e, Ziqiang Yan^e, Tong Jiang^f, Liang Lu^a, Yujuan Yue^a, Xiaobo Liu^a, Hualiang Lin^g, Jianguo Xu^{a,2}, Qiyong Liu^{a,h,1,2}, and Nils Chr. Stenseth^{b,2}

^aState Key Laboratory of Infectious Disease Prevention and Control, National Institute for Communicable Disease Control and Prevention, Chinese Center for Disease Control and Prevention, Beijing 102206, People's Republic of China; ^bCentre for Ecological and Evolutionary Synthesis (CEES), Department of Biosciences, University of Oslo, N-0316 Oslo, Norway; ^cDepartment of Statistics and Actuarial Science, University of Iowa, Iowa City, IA 52242; ^dInstitute of Psychology, Chinese Academy of Sciences, Beijing 100101, People's Republic of China; ^eTropical Diseases Research Base of State Key Laboratory of Infectious Disease Prevention and Control, Guangzhou Center for Disease Control and Prevention, Guangzhou 510440, People's Republic of China; ^fNational Climate Center, China Meteorological Administration, Beijing 100081, People's Republic of China; ^gGuangdong Provincial Institute of Public Health, Guangdong Provincial Center for Disease Control and Prevention, Guangzhou 511430, People's Republic of China; ^hShandong University Climate Change and Health Center, Jinan 250012, People's Republic of China; and ¹World Health Organization Collaborating Centre for Vector Surveillance and Management Beijing 102206, People's Republic of China

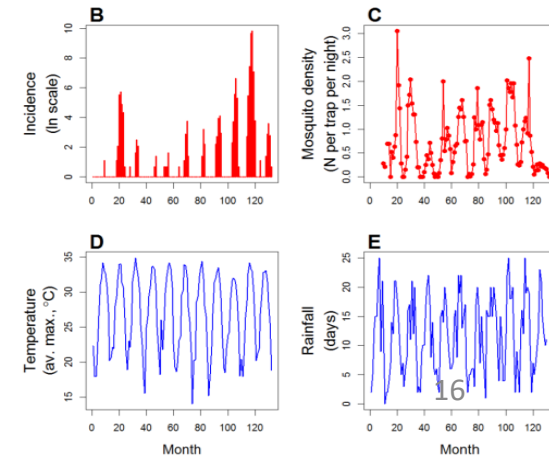
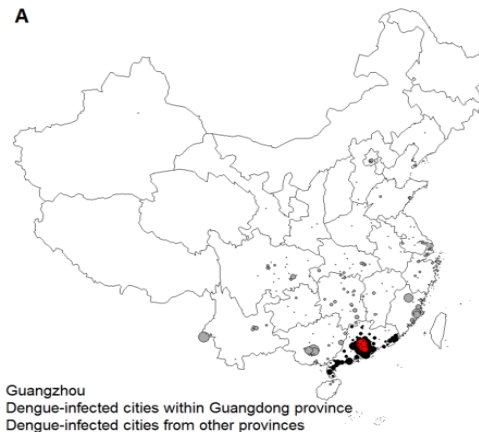
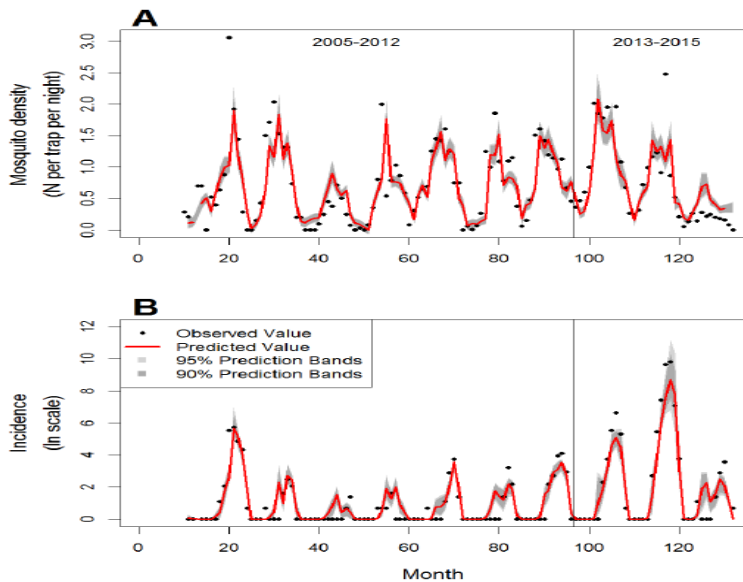
Contributed by Nils C. Stenseth, November 10, 2016 (sent for review September 10, 2016; reviewed by Robert A. Cheke and Kenneth L. Gage)

Climate conditions, through the effects of **rainfall** and **temperature** on mosquito abundance and dengue transmission rate, play key roles in explaining the temporal dynamics of dengue incidence in the human population.

Significance

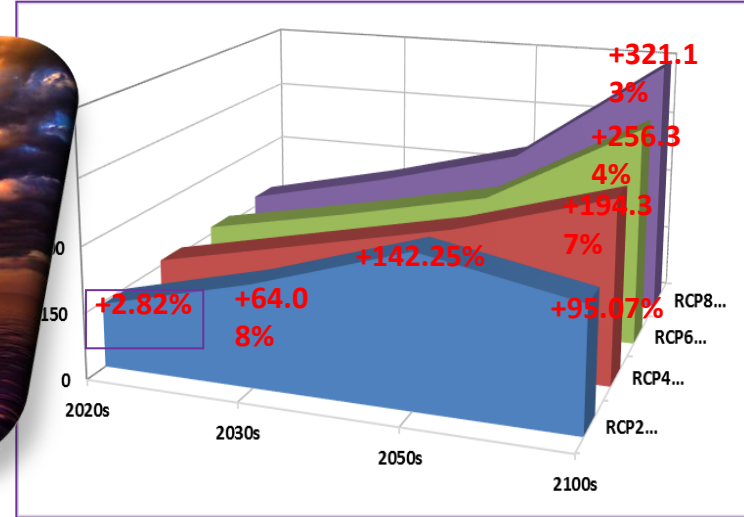
Dengue is a vector-borne infectious disease threatening human health on a global scale. Due to climate change, globalization and other factors, dengue has increasingly spread to new countries and over larger areas, from tropical to temperate zones. In this study, we found that climate has both direct effects on dengue incidence and indirect effects mediated by mosquito density, as mosquitoes are the vectors of dengue. The quantitative results derived from this study may be helpful towards advancing our understanding of how climate influences vector-borne diseases, and prove useful for the control and prevention of dengue fever.

Reserved for Publication Footnotes

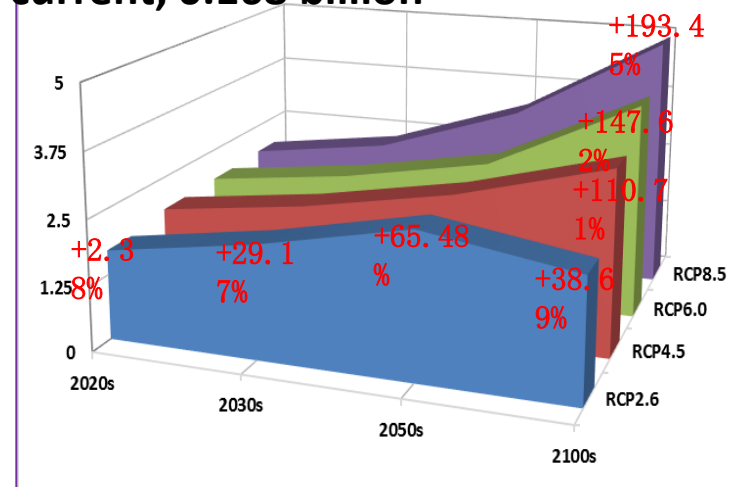


The impact of climate change on dengue fever

Number of high risk counties or districts : at present, 142

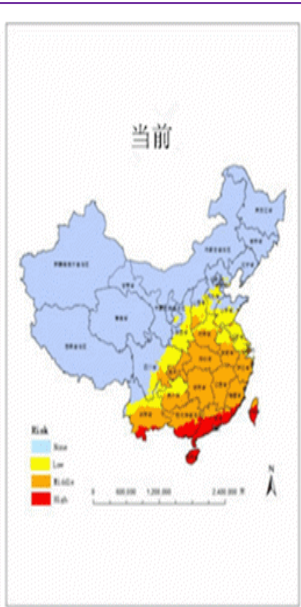


Number of high risk population : current, 0.168 billion

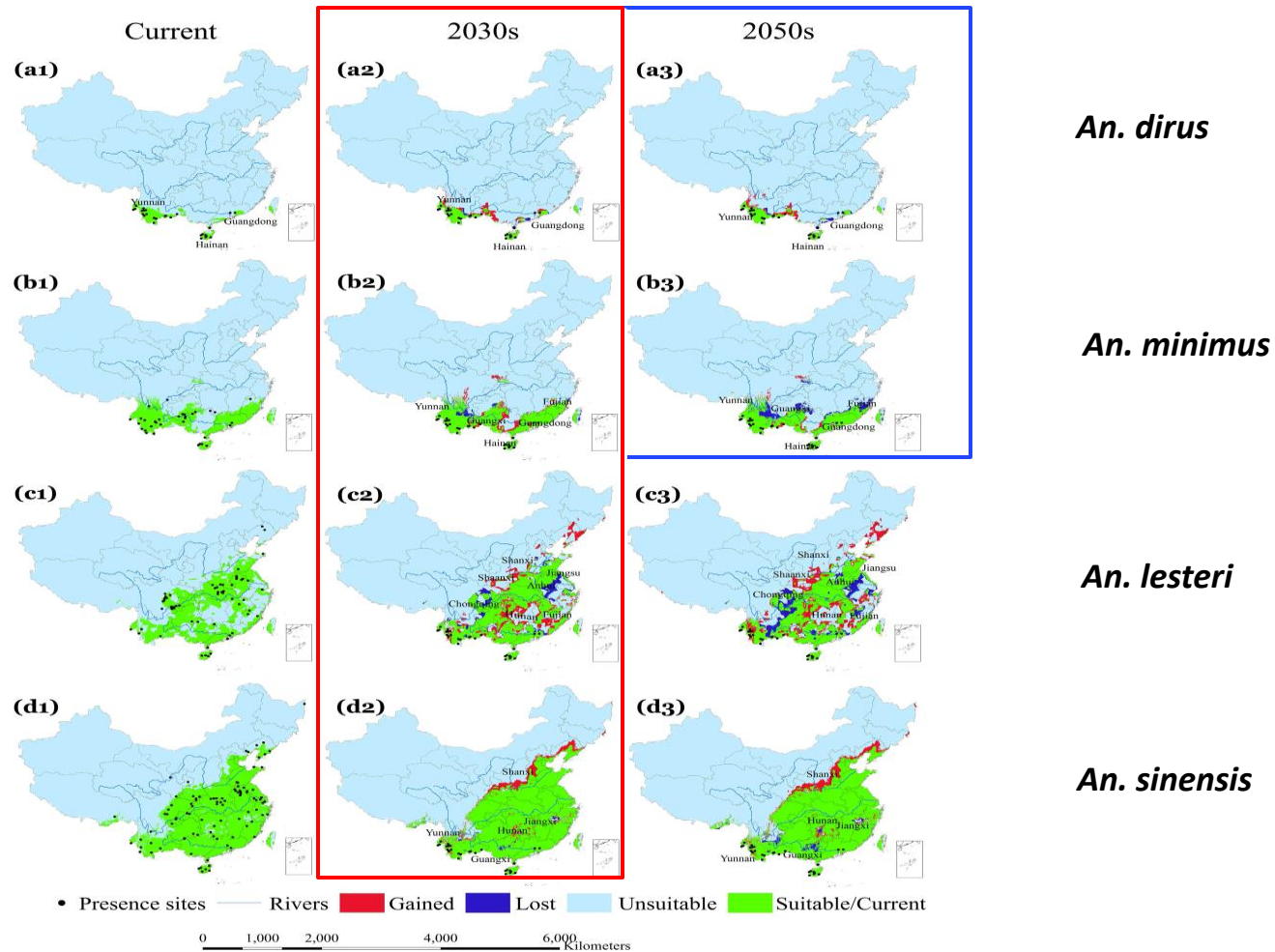


Different RCP scenarios:

- ✓ northward extension of risk area
- ✓ increase markedly of risk population

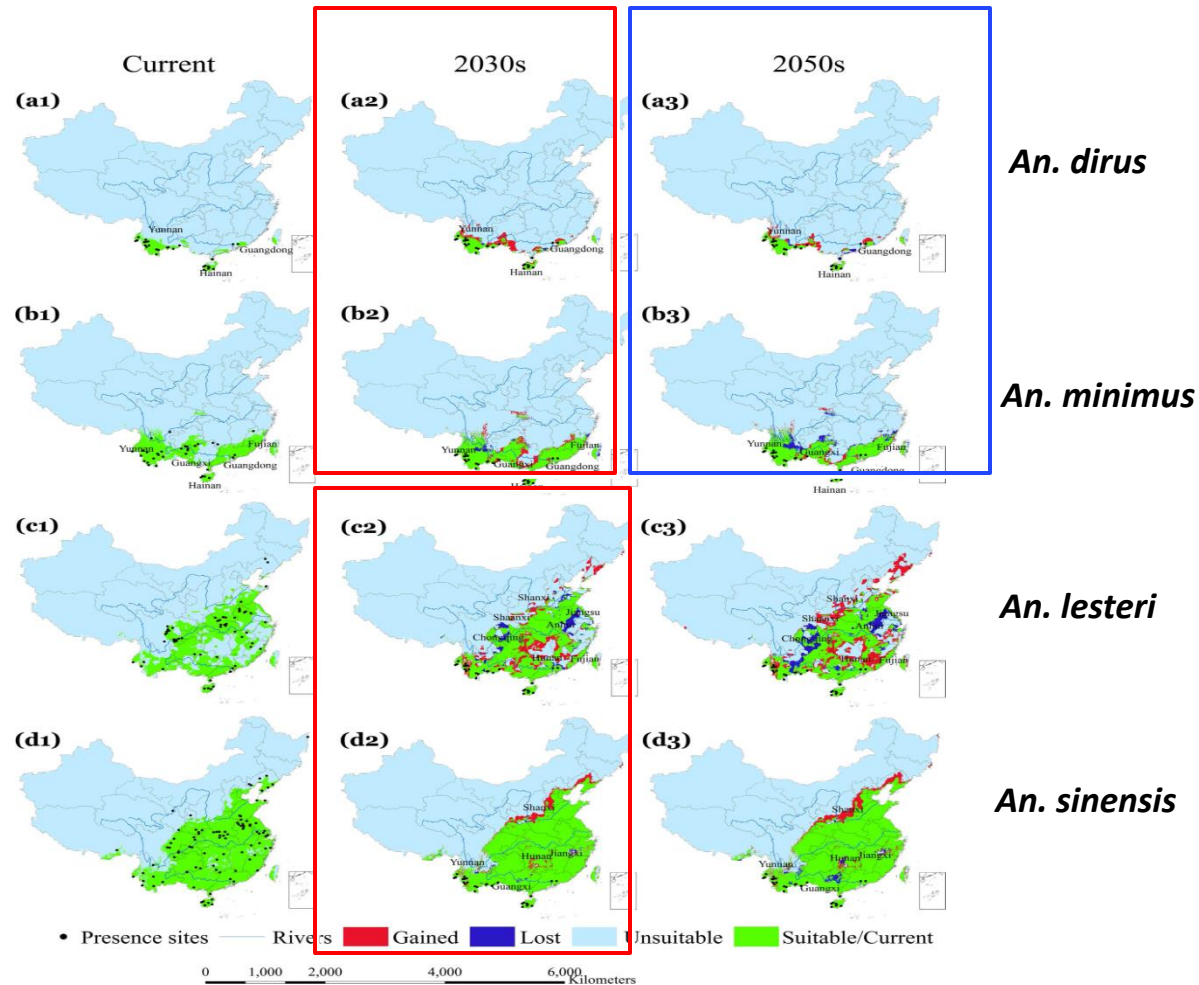


Project the malaria vector distribution under climate change scenarios in China (RCP 2.6)



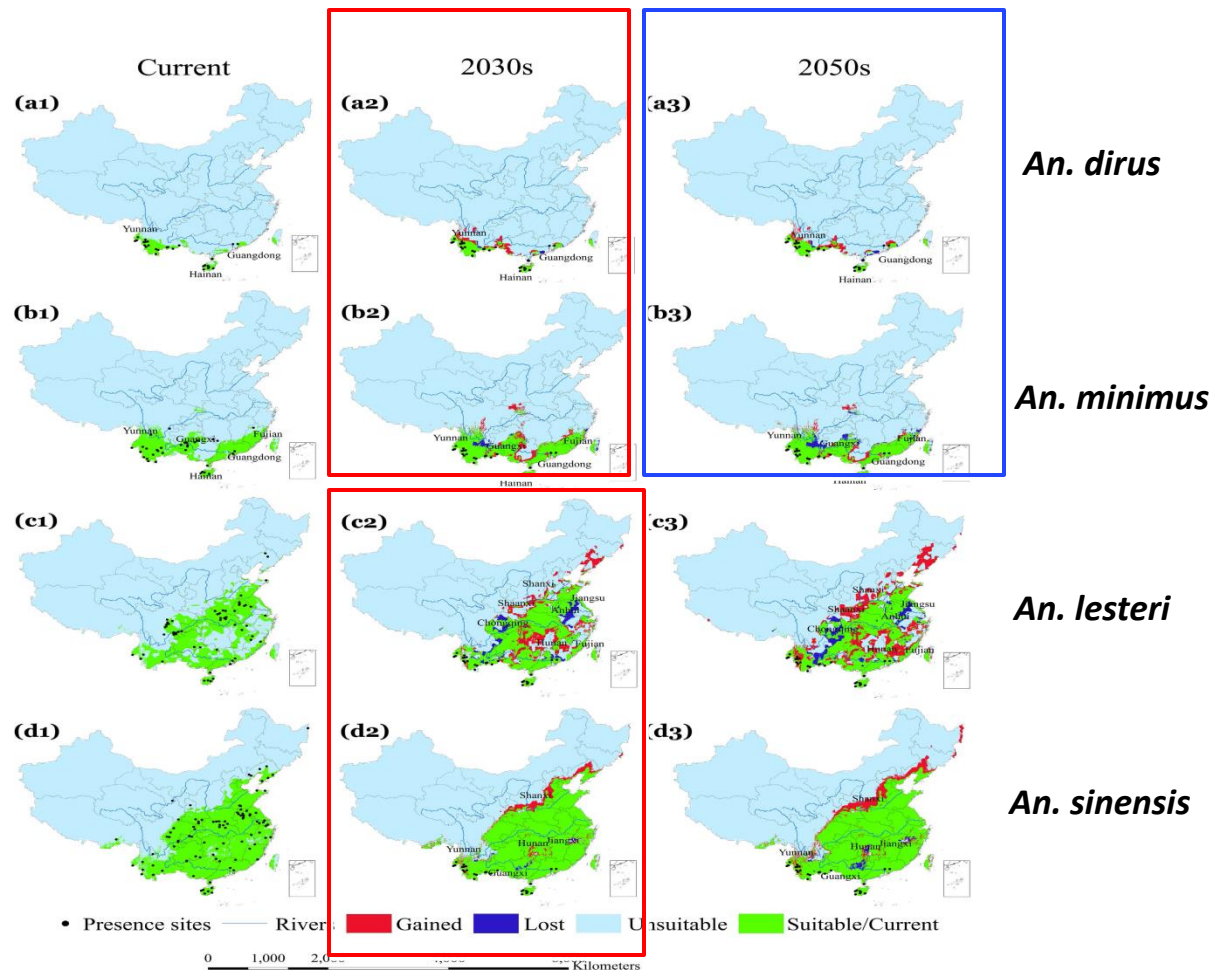
Potential current (suitable and unsuitable) and future (suitable/stable, lost, gained, and unsuitable) environmentally suitable area (ESA) for the four dominant vectors under **RCP2.6**, the lowest greenhouse gas emission scenario.

Project the malaria vector distribution under climate change scenarios in China (RCP 4.5)



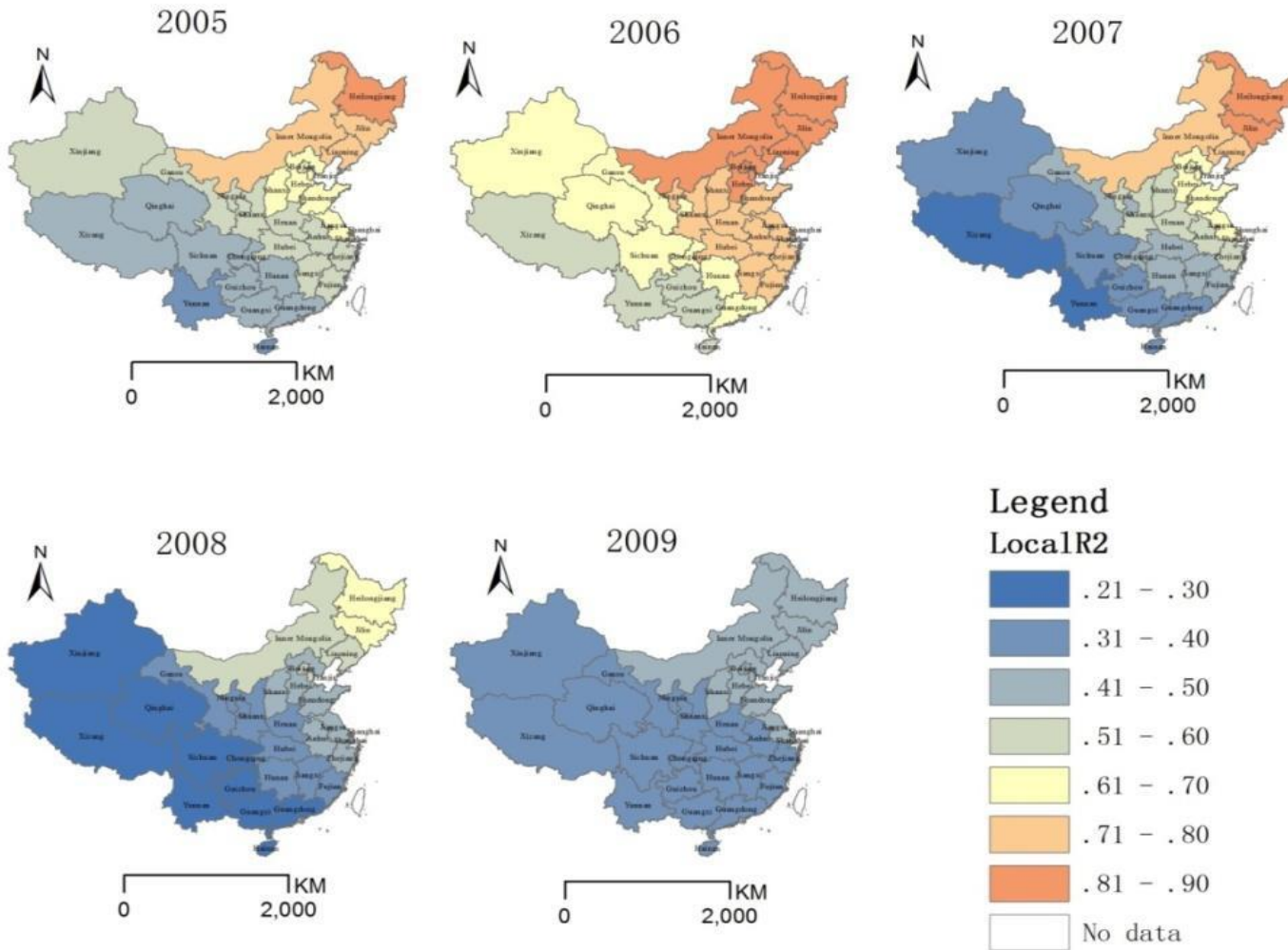
Potential current (suitable and unsuitable) and future (suitable/stable, lost, gained, and unsuitable) environmentally suitable area (ESA) for the four dominant vectors under **RCP4.5, the moderate greenhouse gas emission scenario.**

Project the malaria vector distribution under climate change scenarios in China (RCP 8.5)



Potential current (suitable and unsuitable) and future (suitable/stable, lost, gained, and unsuitable) environmentally suitable area (ESA) for the four dominant vectors under **RCP8.5**, the worst greenhouse gas emission scenario.

Heterogeneity analysis of HFRS in China by geographically weighted regression (GWR)



The incidence of HFRS from 2005 to 2009 is influenced by temperature, precipitation, relative humidity, January NDVI in the same year, August NDVI in the previous year, and elevation, et al, and heterogeneity in spatical distribution.

Addressing the public health impact of climate change-associated infectious diseases in China

- Title: Building capacity to curb the public health impact of emerging and re-emerging infectious diseases due to climate change in China
- Funding: Australian Government's AusAID Development Research Awards Scheme
 - The Australian Aid program is improving the lives of millions of people in the Asia Pacific region



British Embassy
Beijing

China Prosperity Strategic Programme Fund (SPF)



Perceptive assessment of health risks caused by climate change, air pollution and **health co-benefits** of low carbon transition in China

Principal investigator: Liu Qiyong



Next steps for biodiversity & human health

- Workshop
- Case study
- Policy
- Action plan
- Actions & activities
- Guidelines

Thank you for your attention!

