

# Gryphon: A Hybrid Agent-Based Modeling and Simulation Platform for Infectious Diseases

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Quantum Leap Innovations



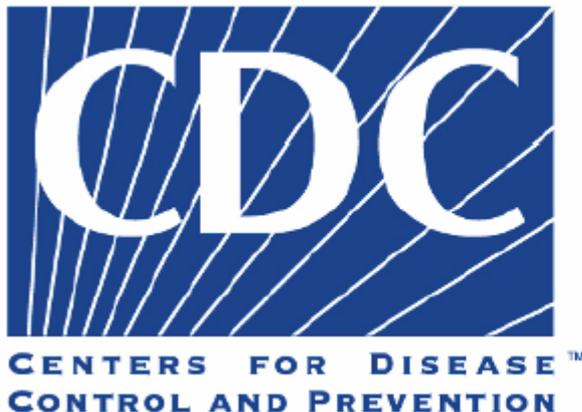
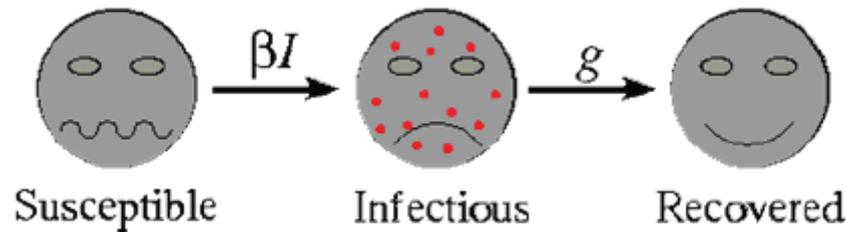
# Agenda

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- Overview of infectious disease modeling
- Gryphon modeling and simulation technologies
- Validation studies
- Conclusion

# Overview of Infectious Disease Modeling

- Compartmental models by sets of differential equations
  - Susceptible-infectious-recovered (SIR) or susceptible-exposed-infectious-recovered (SEIR) models



## Examples:

FluAid, FluSurge (CDC, Atlanta, GA)

DoDIM (QLHS)

# Overview of Infectious Disease Modeling

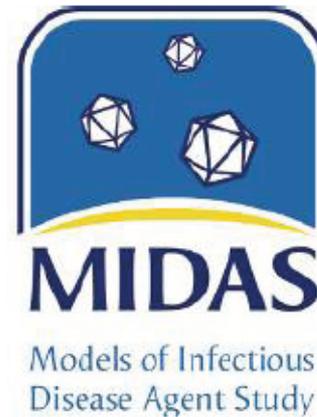
- Highly complex individual-based models
  - Represent daily activities and connections of individuals via transmission networks
  - Often resorts to supercomputers and makes it impractical for quick what-if analyses of interventions or treatments under different conditions.

## Examples:

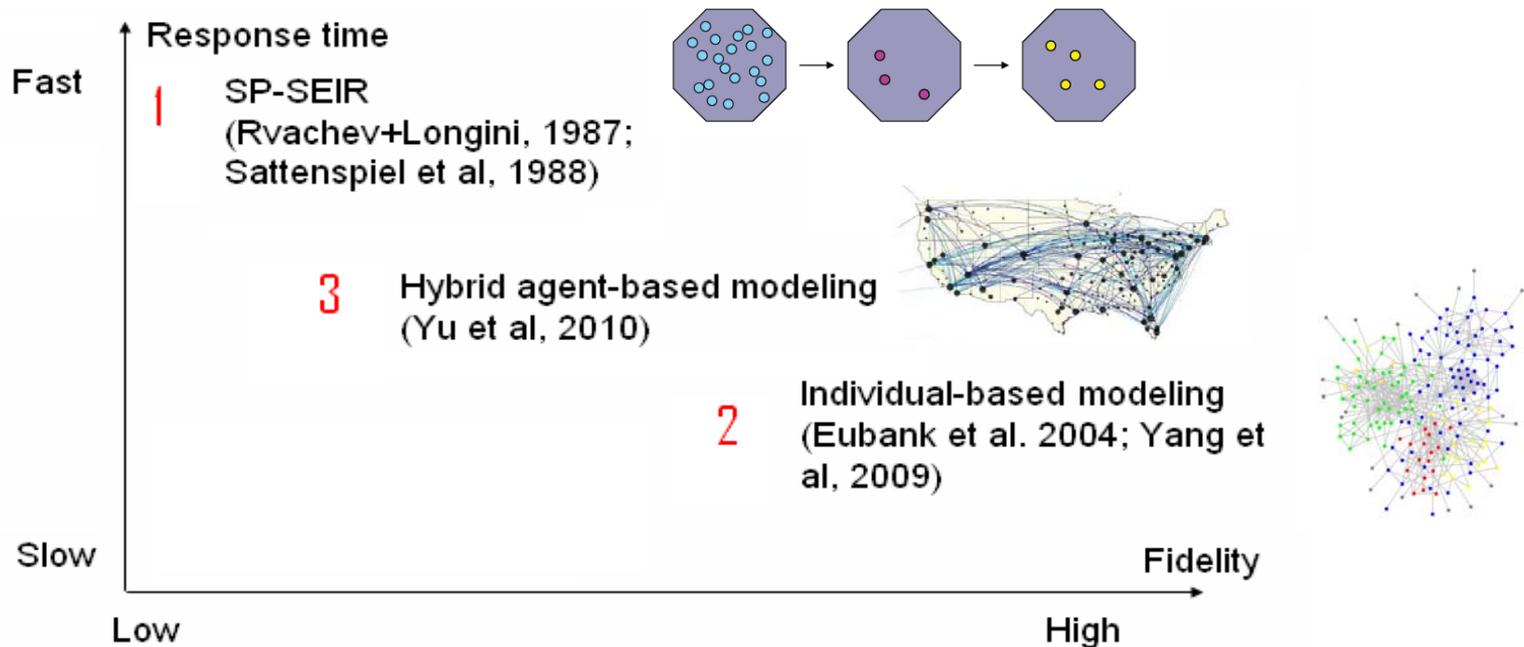
EpiSims (NIH/MIDAS, LANL, NM)

EpiCast (NIH/MIDAS, LANL, NM)

CIP-DSS (NIH/MIDAS, LANL, NM)



# Gryphon Technologies

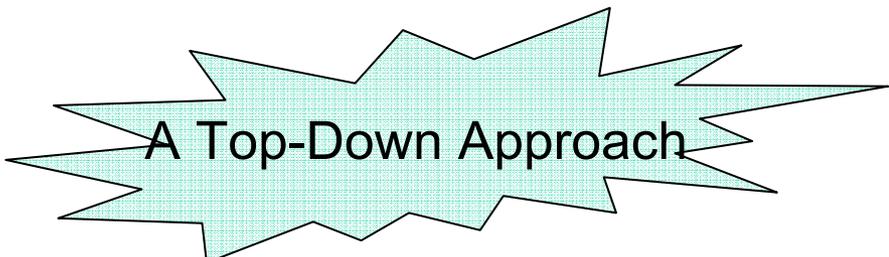


1. Compartmental models by sets of differential equations
2. Highly complex individual-based models
3. **Gryphon<sup>®</sup>**: Hybrid agent-based modeling and Simulation

# Hybrid Agent-Based Modeling

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- Based on groups, not individuals
  - Behaviors of groups are modeled by rules (**Behaviors**)
  - Populations are decomposed into primary, secondary groups (**Decomposition**)

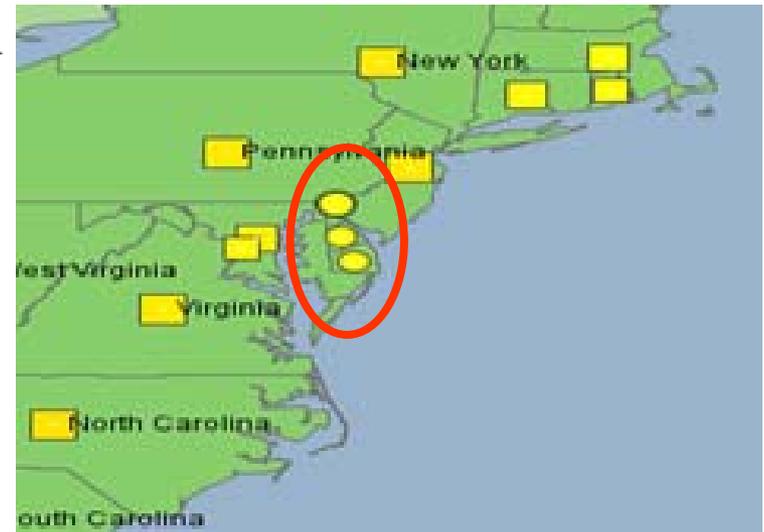
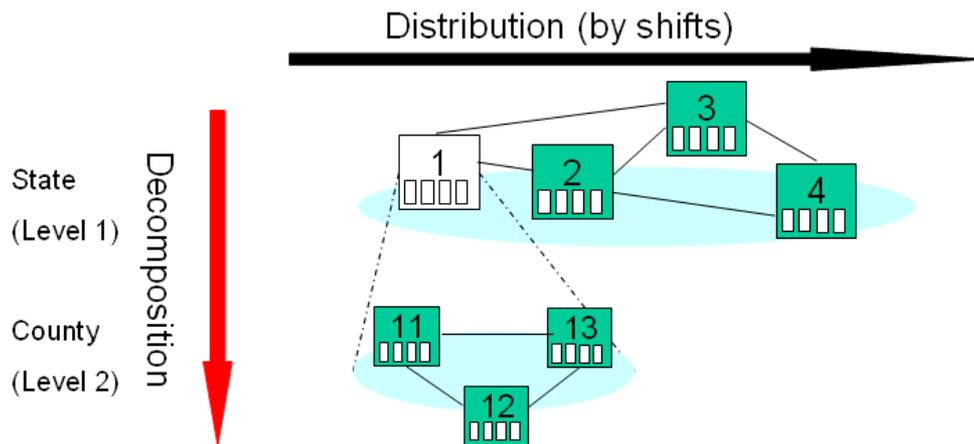


A Top-Down Approach

- Groups move among different locations and mix with each other at each location (**Distribution**)
- The groups for mixing at a location are determined by the behaviors of different groups, not a migration matrix in Structured Population-SEIR (**Mixing**)

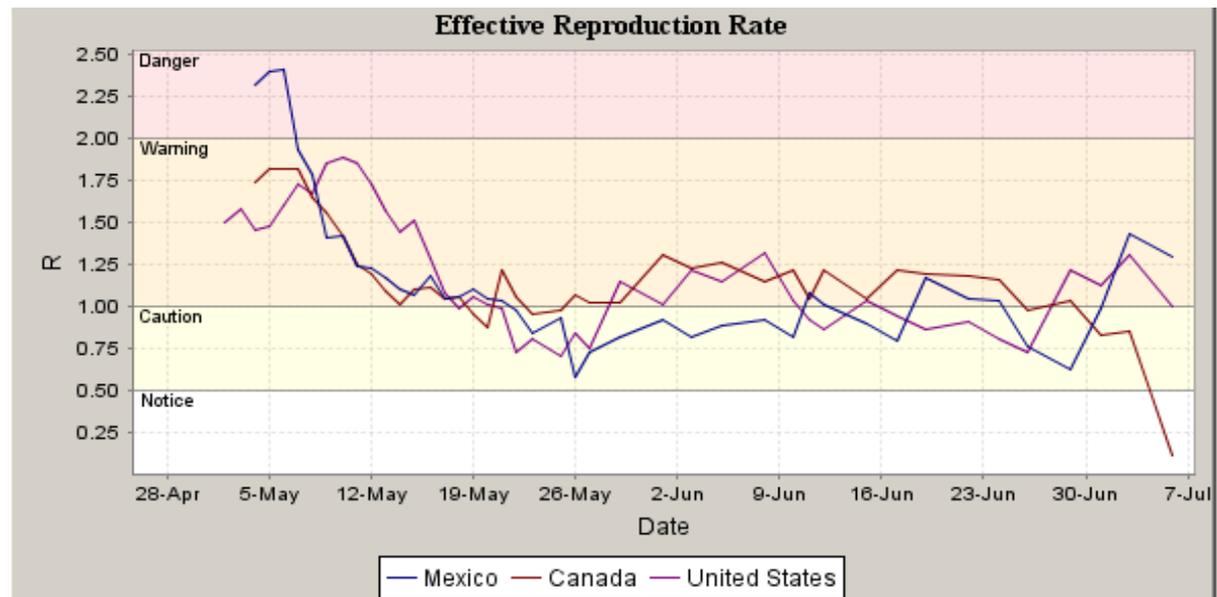
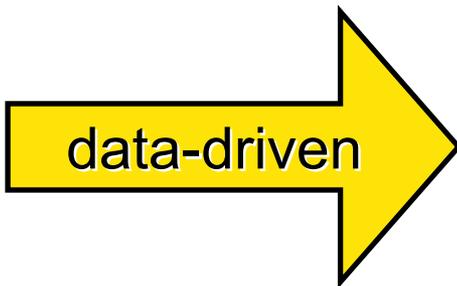
# Benefits of a Top-Down Approach

- Multi-scale modeling and simulation
  - State, county, age groups (top-down approach)
  - The level of abstraction for each group can be configured



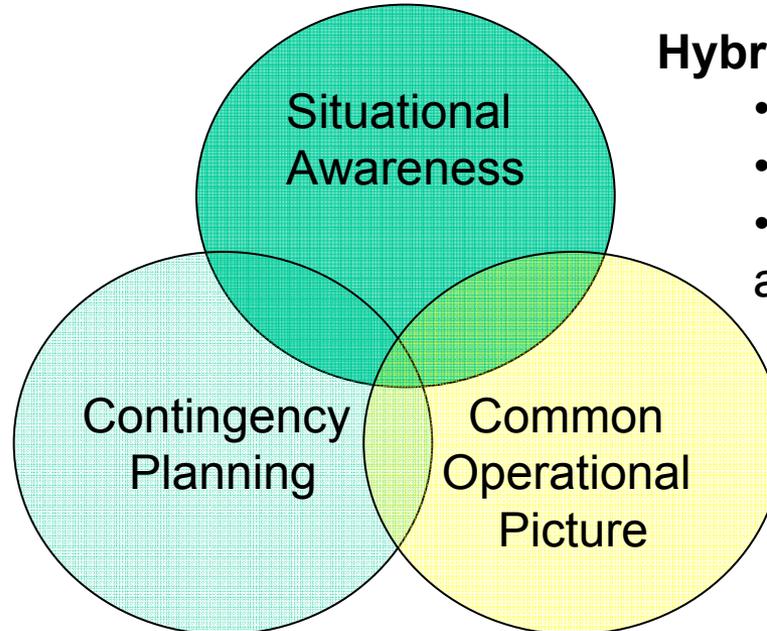
# Benefits of a Top-Down Approach

- Real-time data-driven modeling & simulation
  - CDC surveillance data, e.g. ILNet (state → age groups)
  - Google FluTrend data (state → county)



# Gryphon Platform Features

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## Hybrid agent-based modeling

- Rules/differential equations
- Continuous/discrete time
- Behaviors/interactions of agents

## Interactive what-if analysis

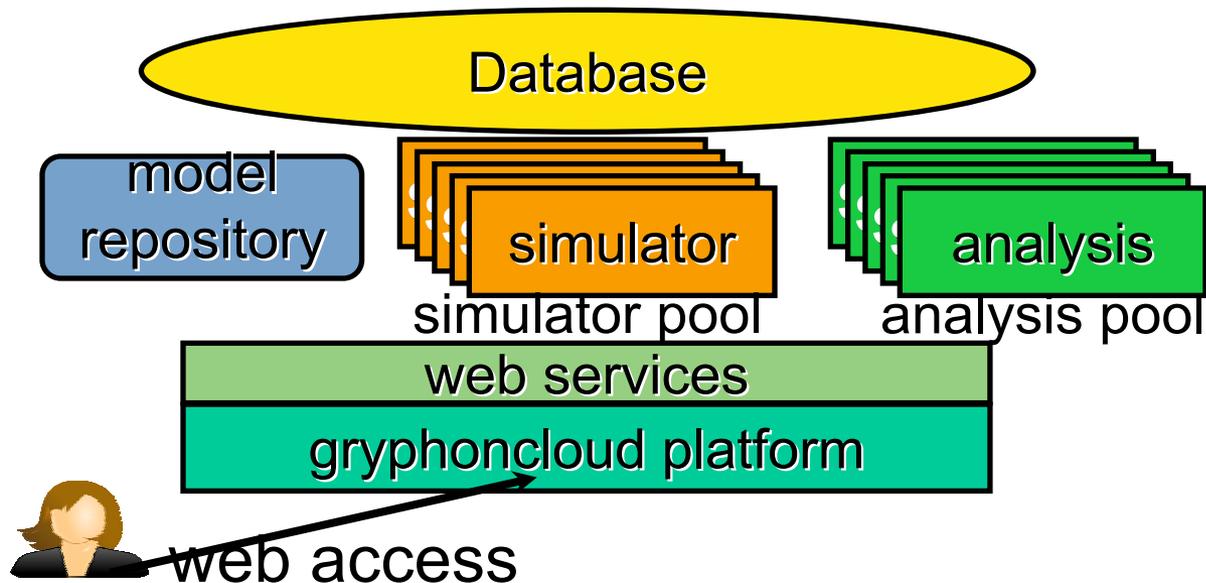
- Enabling what-if analysis of interventions or treatments under different conditions

## Generalized visualization framework

- 2D/3D GIS visualization
- Data analysis of multiple runs data over time

# Success Stories of the Gryphon Platform

- Supported by ONR since 2007
- Cobra Gold'08 (PACOM), 2008 spring
- Cyclone Nargis in Myanmar (PACOM), 2008 summer
- Swine Flu (NORTHCOM and CDC), 2009 spring
- Multi-user multi-scale M&S platform (DE HHS), 2010 spring



# Gryphon Application for Swine Flu

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Disease Simulation for the US Northern  
Command Area of Responsibility  
**(USNORTHCOM AOR)**

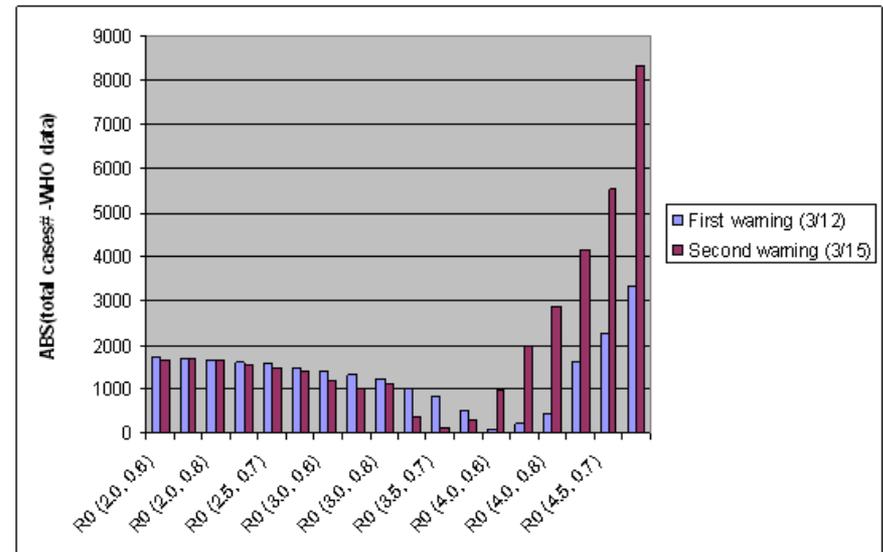
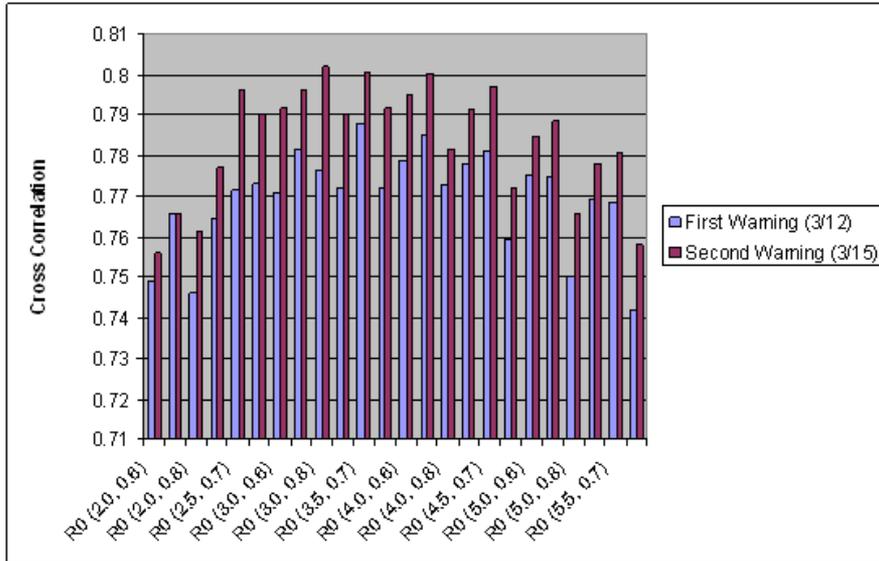
United States, Mexico, and Canada

# Data and Methods for Validation Studies

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- **Discrete time stochastic SEIR model for mixing**
- The timeline of SARS outbreak in Hong Kong and other Asian countries can be described as follows
  - February 15, 2003: Official report of a 33-year male and a 9 year old son in Hong Kong with Avian influenza (H5N1).
  - March 12, 2003: First global alert about atypical pneumonia in Vietnam and Hong Kong was issued by World Health Organization (WHO).
  - March 15, 2003: Second global alert about name of SARS and case definition was issued by WHO.
- The travel data sets are generated from the International Air Transport Association (IATA) (<http://www.iata.org>) database, which contains the number of available sets between any two given countries.
- The country data sets, including population, latitude and longitude for each country, are generated from the website (<http://www.geonames.org>).

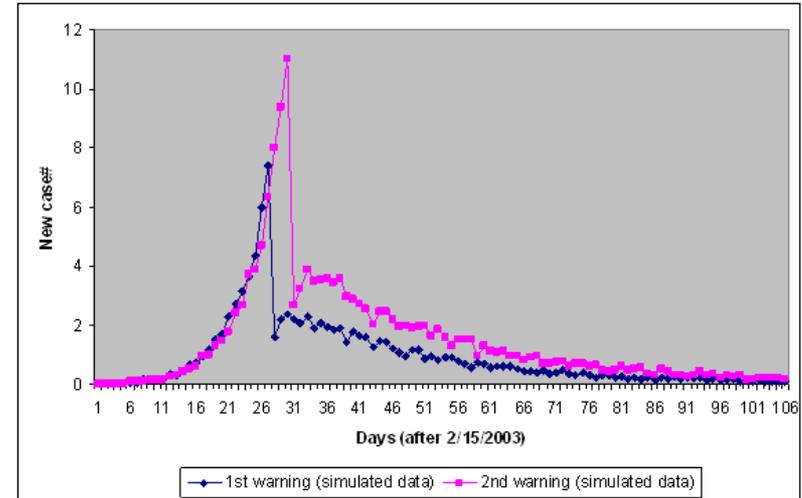
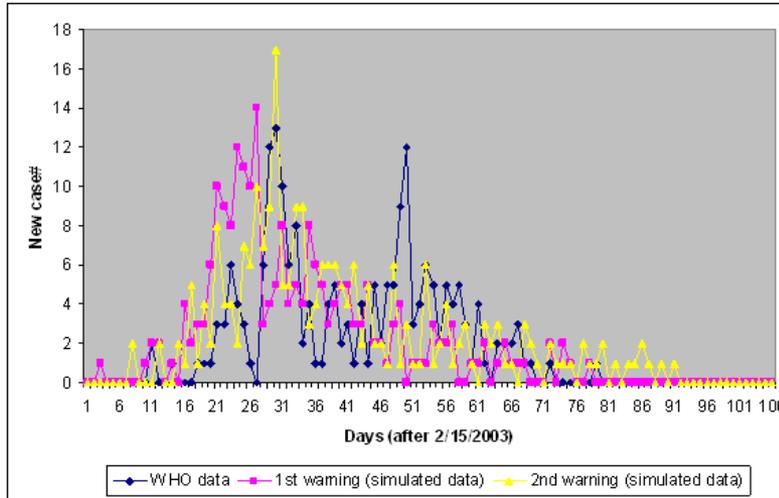
# Results for Local Transmission Dynamics



- The estimated pairwise value of  $R_0$  for Hong Kong is consistent with Wallinga and Teunis's work by assuming an exponential increase in the number of cases over time

Wallinga, J., Teunis, P.: Different epidemic curves for severe acute respiratory syndrome reveal similar impacts of control measures. *American Journal of Epidemiology* **160**(6) (2004) 509–516

# Results for non-Local Transmission Dynamics



- The predicated total case number for non-local disease transmission is close to the one given in Hufnagel's PNAS paper in 2004, in which Hufnagel et al. used a continuous-time stochastic SEIR model.

Hufnagel, L., Brockmann, D., Geisel, T.: Forecast and control of epidemics in a globalized world. PNAS 101(42) (2004) 14124–15129

# Conclusion

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## **Gryphon: A scalable, flexible and interactive simulation platform**

### **Future Work**

#### **Optimization of pharmaceutical and non-pharmaceutical interventions for infectious diseases**

##### **Pharmaceutical interventions**

- Allocation of Antiviral and Vaccines at state and county levels

##### **Non-Pharmaceutical interventions**

- Isolation, quarantine, and school closure
- Assessing the benefits of these strategies weighted against the cascading consequences that may arise from their use

#### **Modeling complex and adaptive behaviors of populations**