



# Modelling the Spread of COVID-19 with Travel Data and Incomplete Infection Data

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# COVID-19 Early Facts



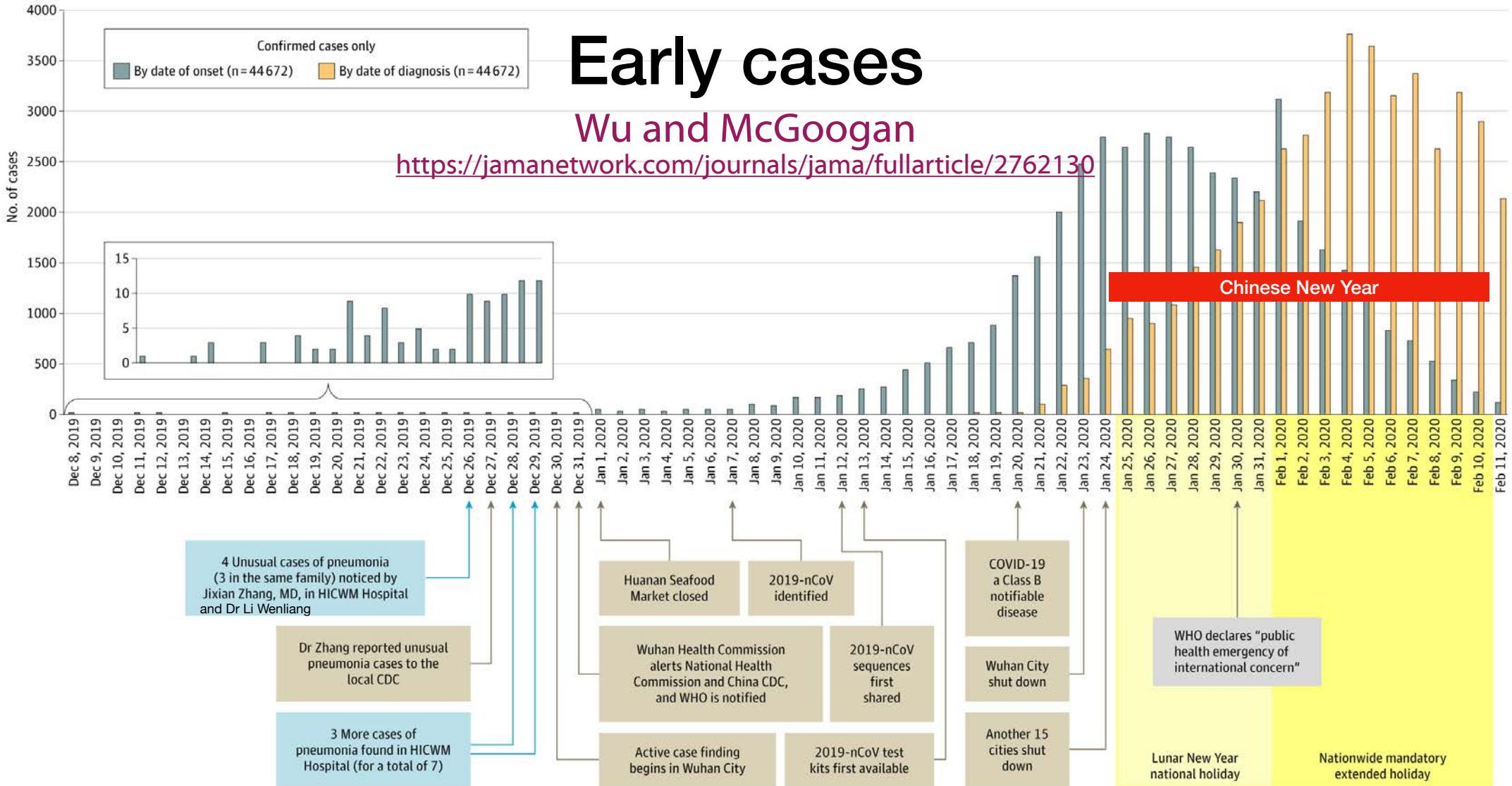
- **2019 New Coronavirus Disease (COVID-19)** began to spread from early December 2019 in **Wuhan**
- First confirmed case: December 8, 2019
- Human-to-human transmission found to occur in mid December 2019
- Active control started in late January 2020, with official instruction from central sent down on January 7, 2020.\*
- Wuhan was locked down on January 24, 2020. Other cities began to impose stringent travel control.
- Italy reported first case on January 31, 2020
- France also traced the first case back in December 2019

\*State media Qiu Shi Magazine reported President Xi's speech at the Standing Committee of the Central Political Bureau of CCP on Feb 3, 2020, sending instruction down on Jan 7, 2020 regarding control of the new coronavirus outbreak in Wuhan.

# Early cases

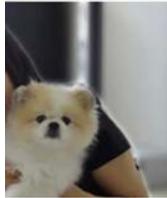
Wu and McGoogan

<https://jamanetwork.com/journals/jama/fullarticle/2762130>



# Hong Kong (second wave)

 South China Morning Post



Pet dog of Covid-19 patient infected, Hong Kong health officials confirm

- Pomeranian has low-level infection and experts unanimously conclude human-to-animal transmission
- But experts also cite previous experience with SARS suggesting cats and dogs will not fall ill or transmit virus to people

## Cancellation of events

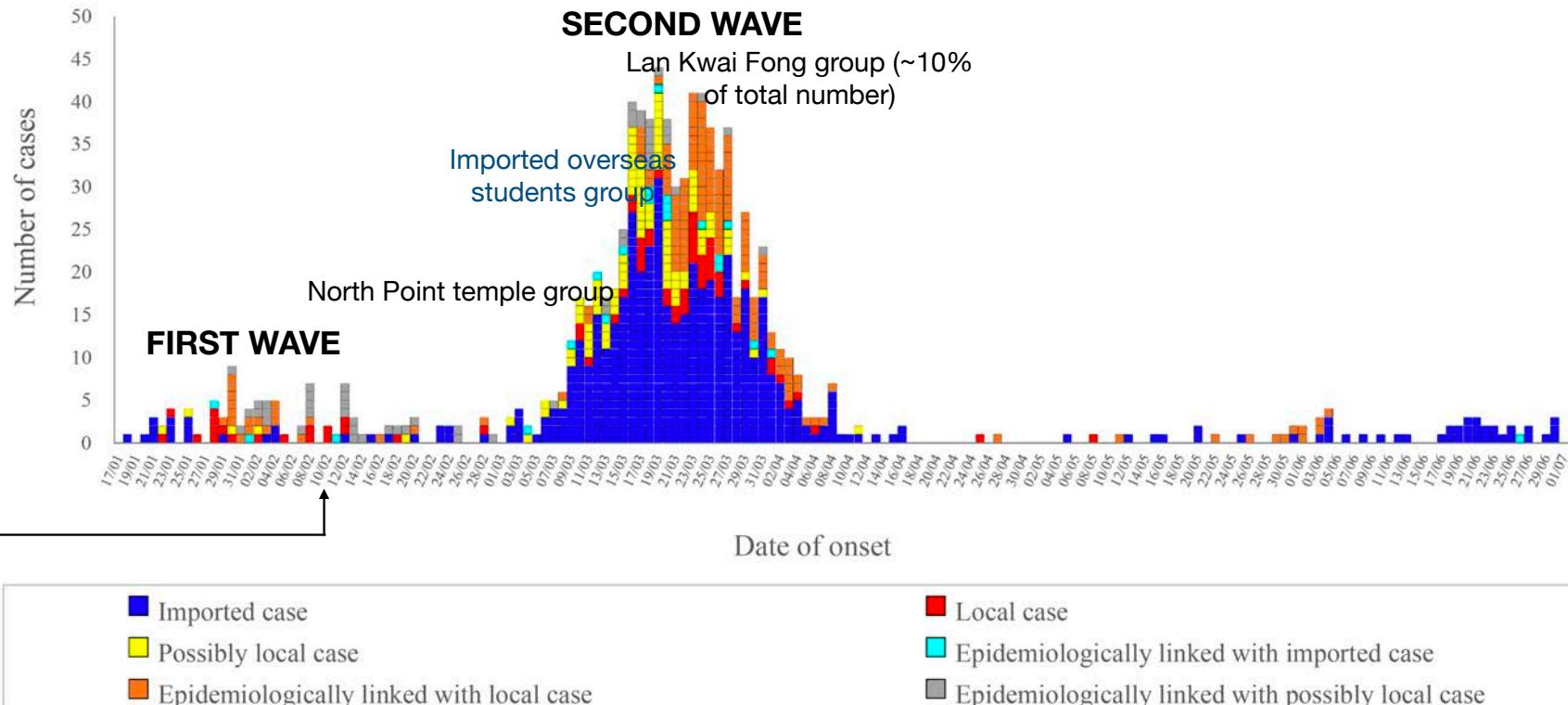
25 Eason Chan shows (Dec 2019)  
6 Zhou Jie Lun shows (Jan 2020)  
23 Andy Lau shows (Jan-Feb 2020)  
numerous concerts, theatres, ...

⇒ ~1 million audience  
1 CNY Firework show  
1 CNY Parade  
⇒ Several thousands

...  
had prevented mass gatherings and massive intercity travel during the peak travel time.

Epidemic curve of confirmed and probable cases of COVID-19 in Hong Kong (as of 1 Jul 2020)

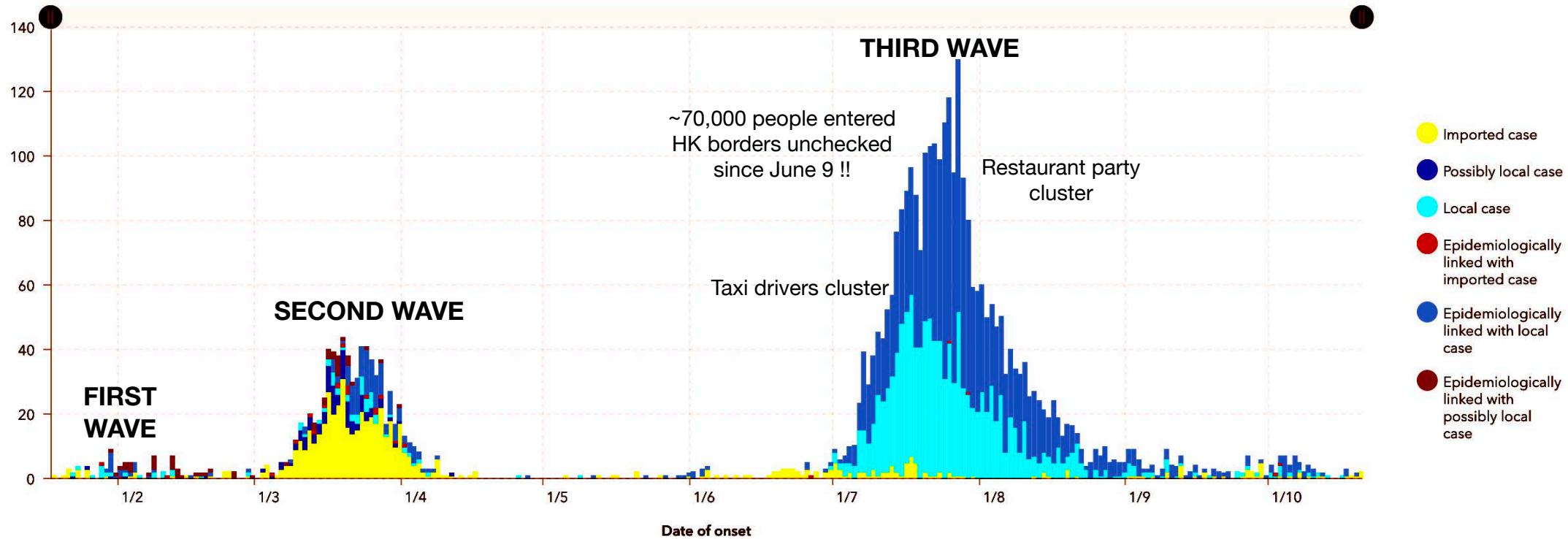
Number of confirmed and probable cases = 1234



Data: Department of Health, Hong Kong

# Hong Kong (third wave)

Epidemic curve of confirmed/probable cases of COVID-19 in Hong Kong



# Spreading escalated by the *spring travel rush*



<https://www.storm.mg/article/2212024>

- 30 days surrounding the Lunar New Year
- Migrant workers and students travel from major cities to country towns for family reunions, and return to the cities at the end of the holiday period. Holiday goers also travel to and from tourist cities.
- Ministry of Transport estimates ~3 billion trips taken during the spring travel rush.
- Wuhan, being a major transport hub and having a large number of higher education institutions as well as manufacturing plants, is among the cities with the largest outflow and inflow traffic during the spring travel rush.

# The epicenter



## WUHAN

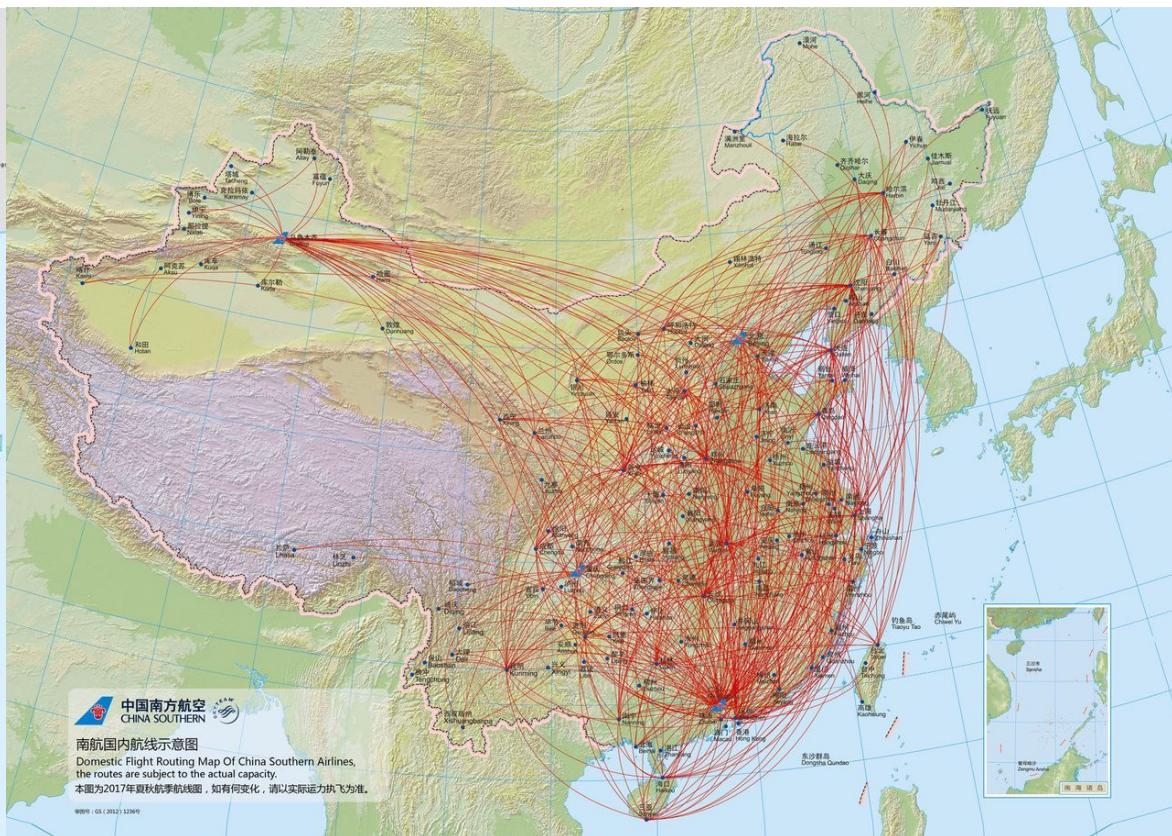
### A very important city in China

- 11 million people
- ***major traffic hub***
- center of higher education,  
manufacturing plants,  
industry and business



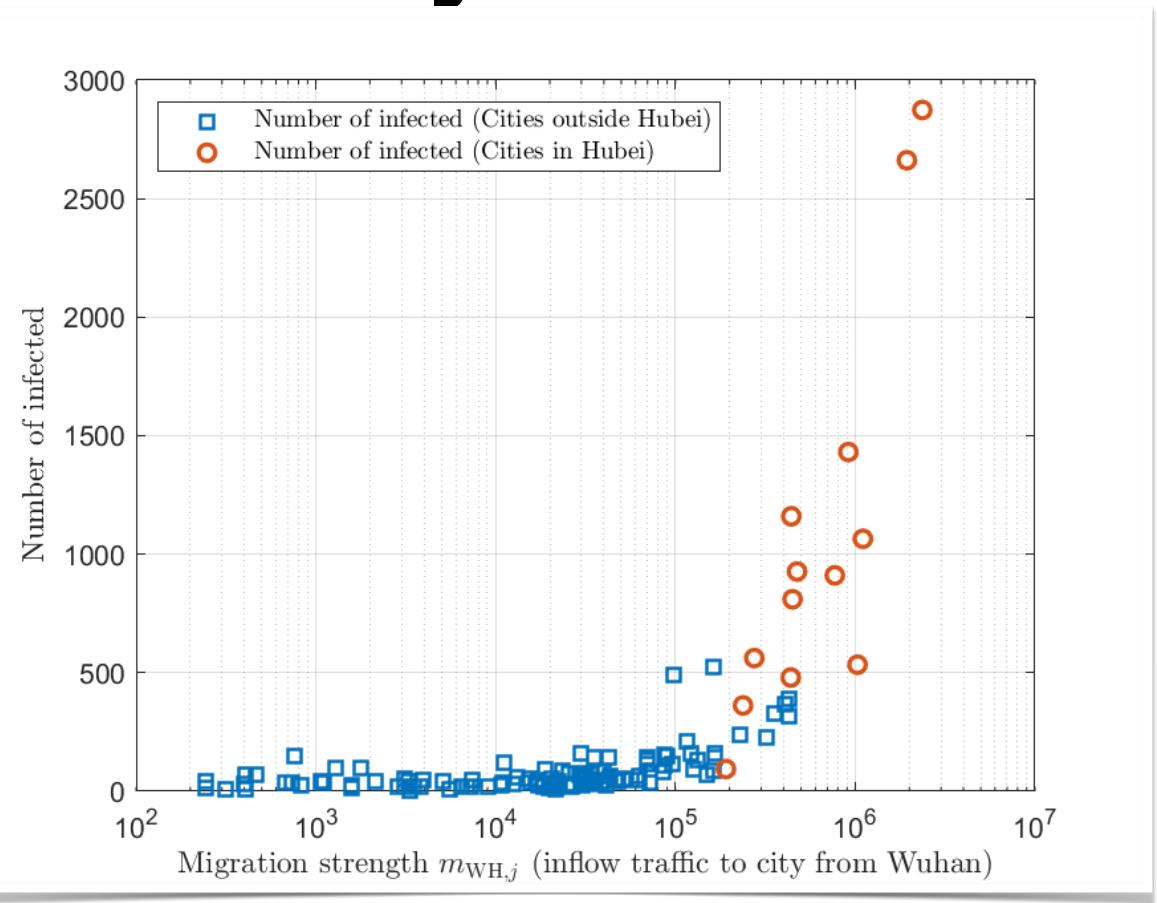
Photo by Michael Tse 2018

# Gathering of people and **intercity travel** of infected and exposed individuals within China have been the main drives that escalated the spreading of the virus. *How to get indicative travel volume data?*



# Impact of Intercity Travel

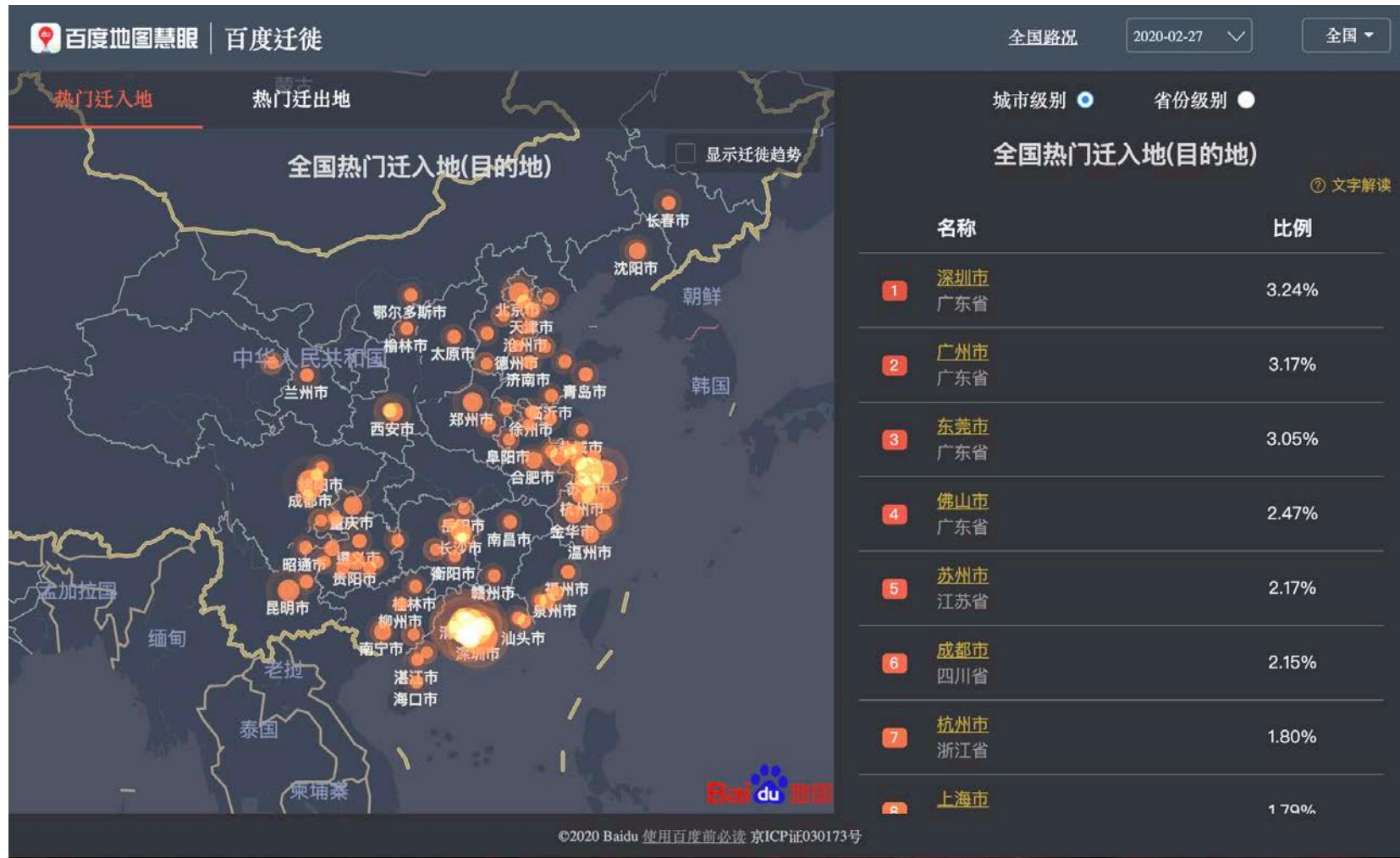
- Number of infections in a city farther from Wuhan increases almost *linearly* with the *Baidu Migration Strength* (inflow strength) of the city from Wuhan



# Data

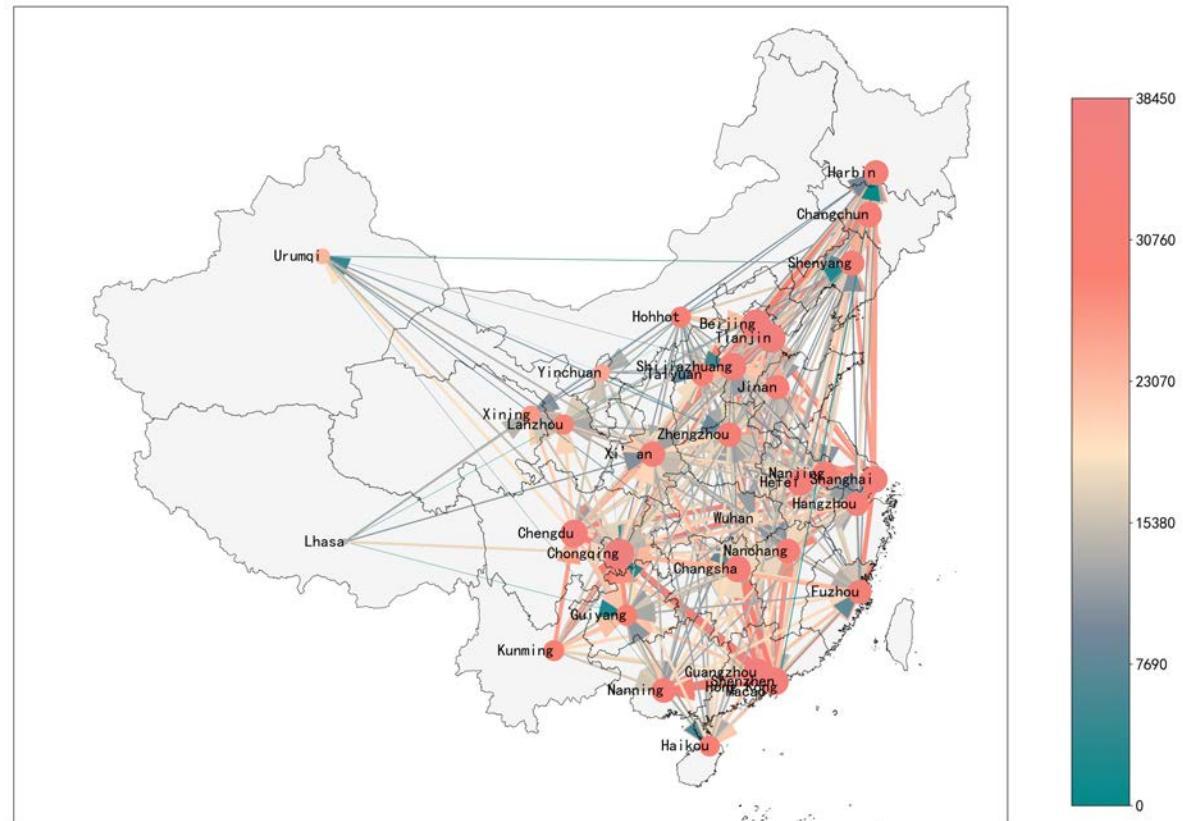
Mobile-app  
based human  
migration  
tracking data  
system  
provides  
indicative inflow  
and outflow  
volume of  
travellers  
between cities

# Baidu Migration Data

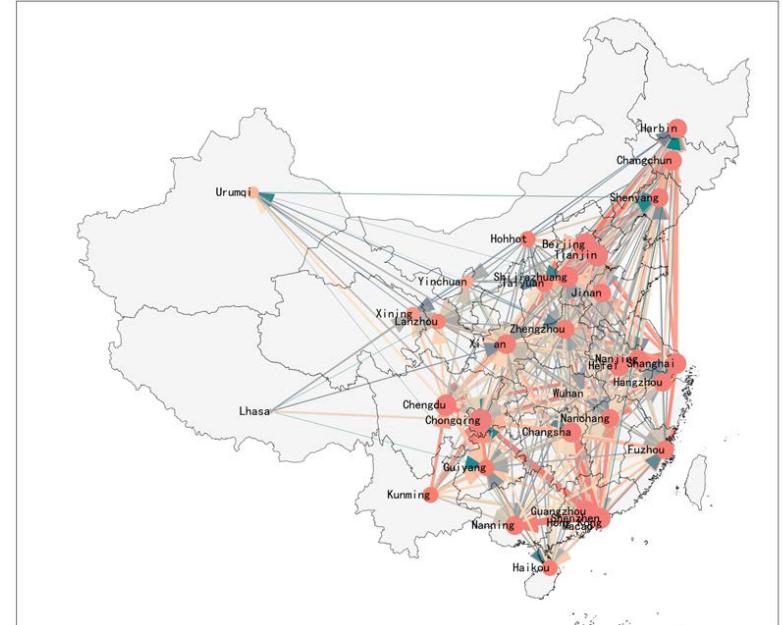
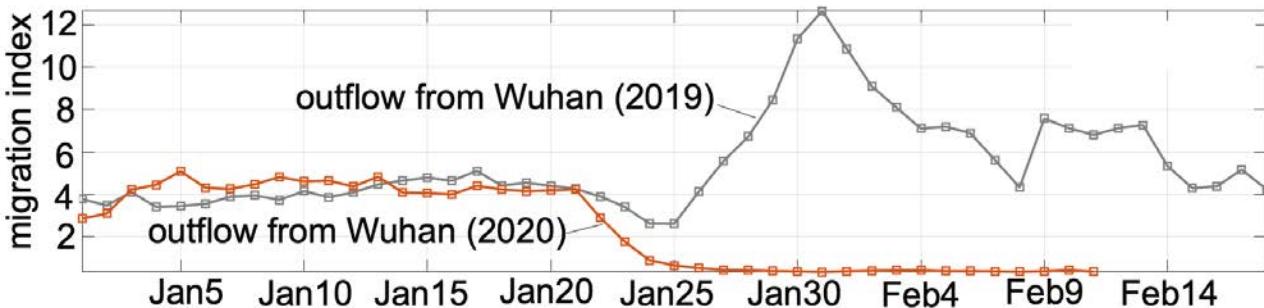
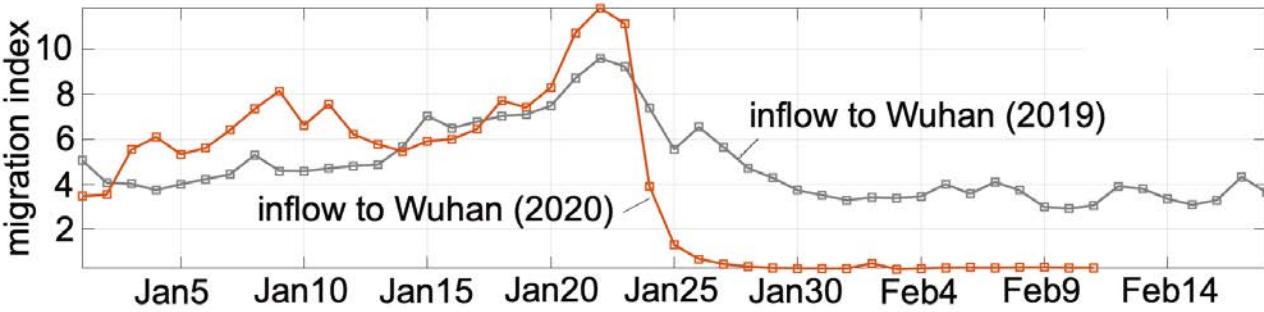


# Data collection

- Baidu Migration data for 367 cities (or administrative regions) in China.
- The data provide the migration strengths of cities which are indicative measures of the human traffic volume moving in and out of 367 cities and administrative regions.

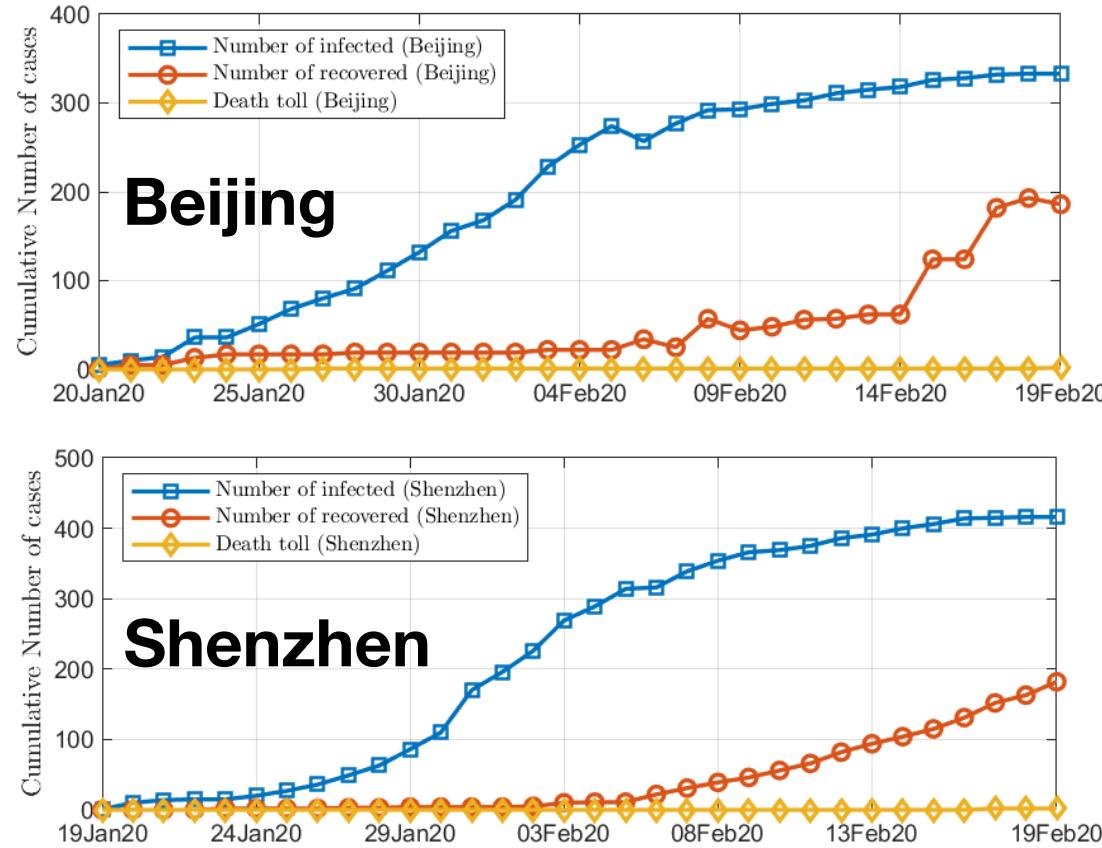


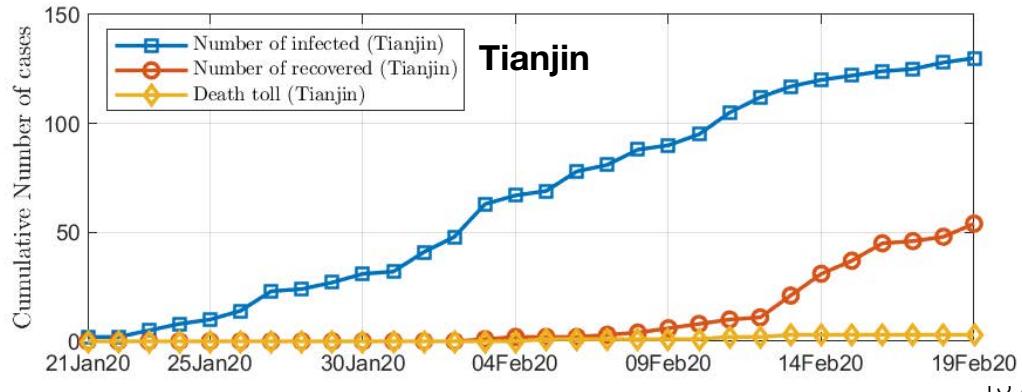
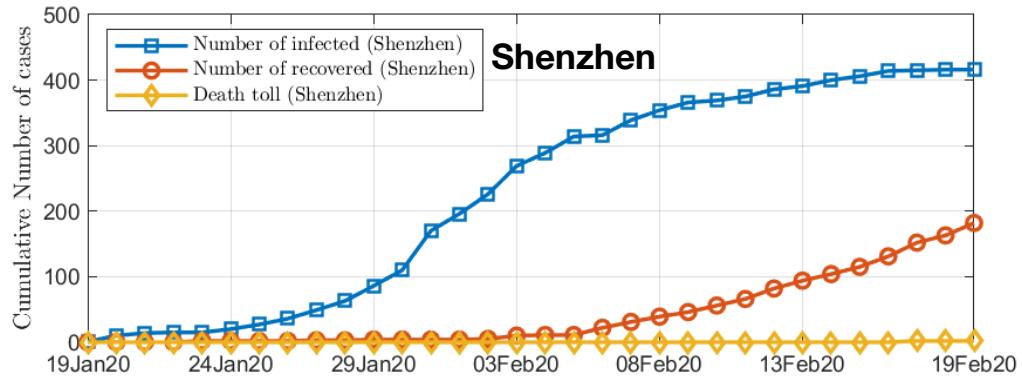
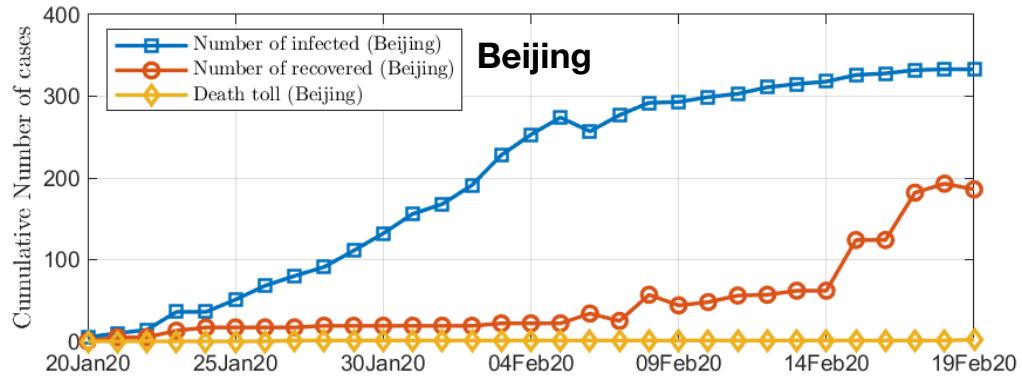
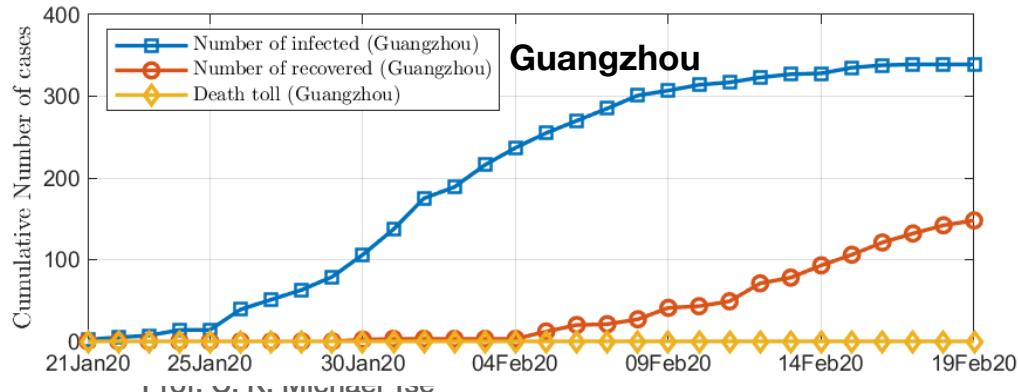
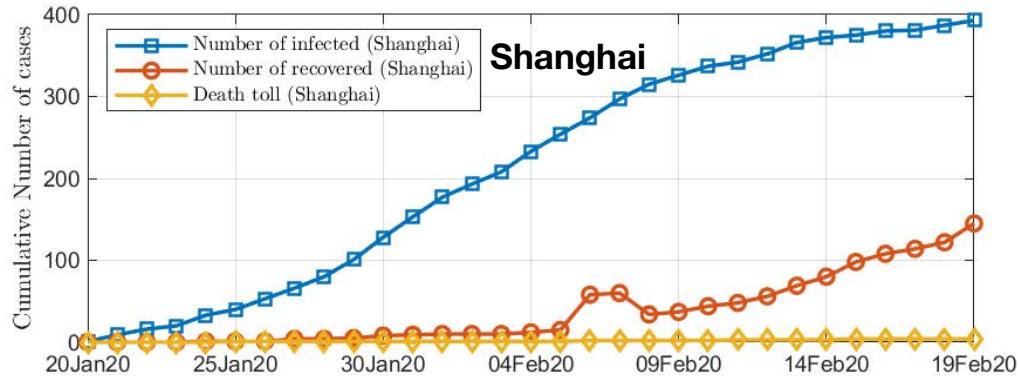
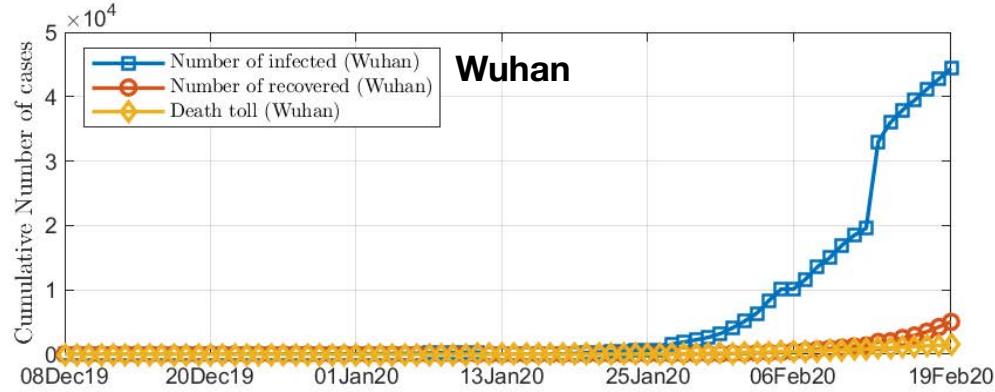
# Wuhan inbound and outbound traffic



# Data of Infection Cases

- National Health Commission of China
- Daily data, including the daily total number of confirmed cases in each city, daily total cumulative number of confirmed cases in each city, daily cumulative number of recovered cases in each city, and daily cumulative death toll in each city.
- Data collected for **367** cities





# Other countries

South Korea

Italy

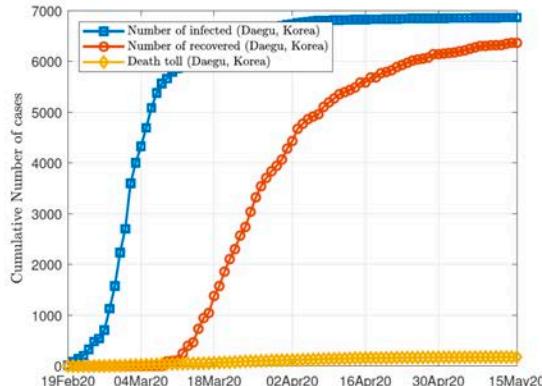
Iran

Japan

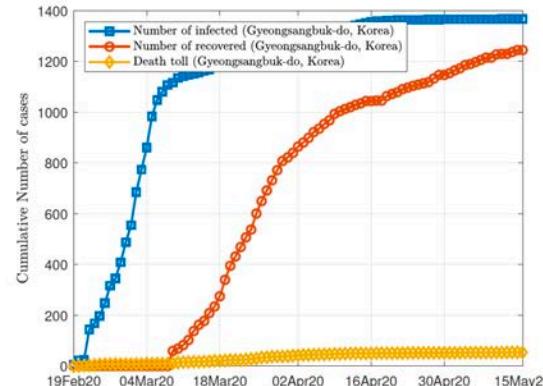
USA

UK

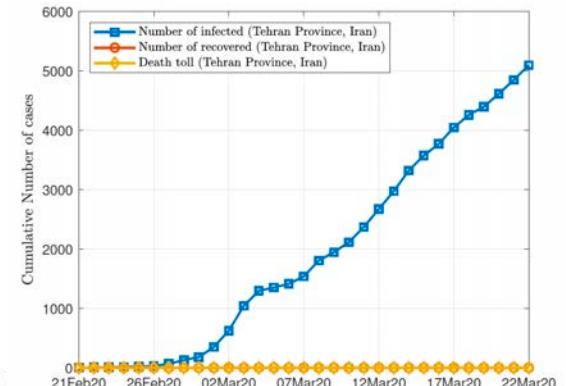
Over 500 cities



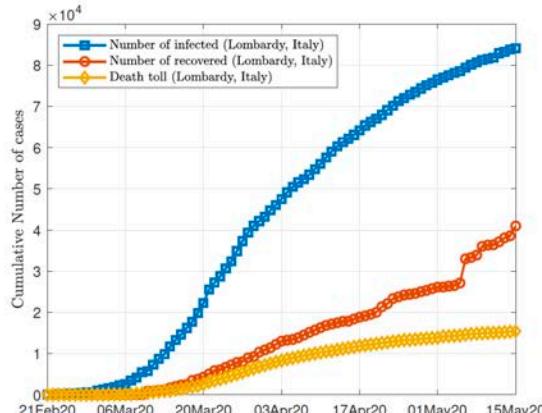
(a) Daegu



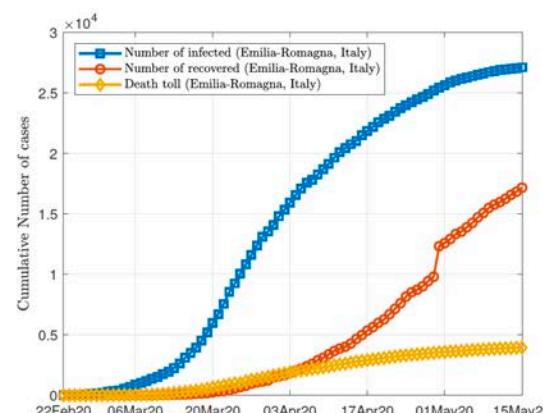
(b) Gyeongsangbuk-do



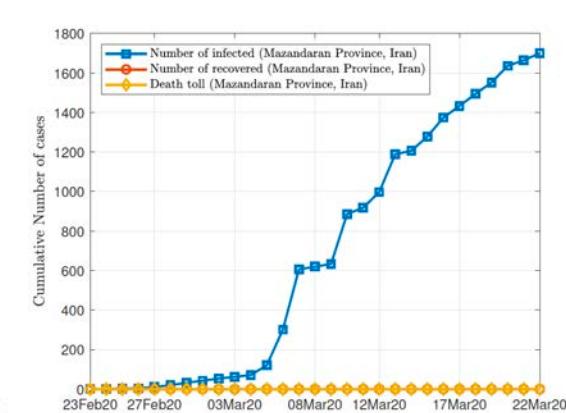
(e) Tehran



(c) Lombardy



(d) Emilia-Romagna



(f) Mazandaran

# Problem of Information Latency and Loss

- One reason for Rapid Spread of Virus:
  - *Delayed information — unawareness of the real situation* in a population
  - Most notable information latency lies in the number of confirmed cases reported, which depends on the ability of the particular country or city to perform tests as well as the possible bureaucracy in the local system of reporting.

$$I > C \text{ or } I \gg C$$

- The number of confirmed cases is certainly not the true number of infected individuals at any given time

# Model

**Issue 1: Travel Data Integration**

**Issue 2: Information Latency**

# Review of SEIR Model

On day  $t$ , there are

- **S**usceptible  $S_i(t)$
- **E**xposed  $E_i(t)$
- **I**nfected  $I_i(t)$
- **R**ecovered/ removed  $R_i(t)$

City  $i$  with population =  $P_i$



Eventual percentage of infection =  $\delta_i$

Total susceptible population  $N_i^s = \delta_i P_i$

$$N_i^s(t) = S_i(t) + E_i(t) + I_i(t) + R_i(t)$$

# Basic SEIR Dynamics

For a single population (network), we can write the state equations as

$$\begin{aligned}\dot{S}(t) &= -\beta S(t)I(t) \\ \dot{E}(t) &= \beta S(t)I(t) - kE(t) \\ \dot{I}(t) &= \kappa E(t) - \gamma I(t) \\ \dot{R}(t) &= \gamma I(t)\end{aligned}$$

$$\Rightarrow \begin{cases} \Delta S(t) &= -\beta S(t-1)I(t-1) \\ \Delta E(t) &= \beta S(t-1)I(t-1) - \kappa E(t-1) \\ \Delta I(t) &= \kappa E(t-1) - \gamma I(t-1) \\ \Delta R(t) &= \gamma I(t-1) \end{cases}$$

# Parameters

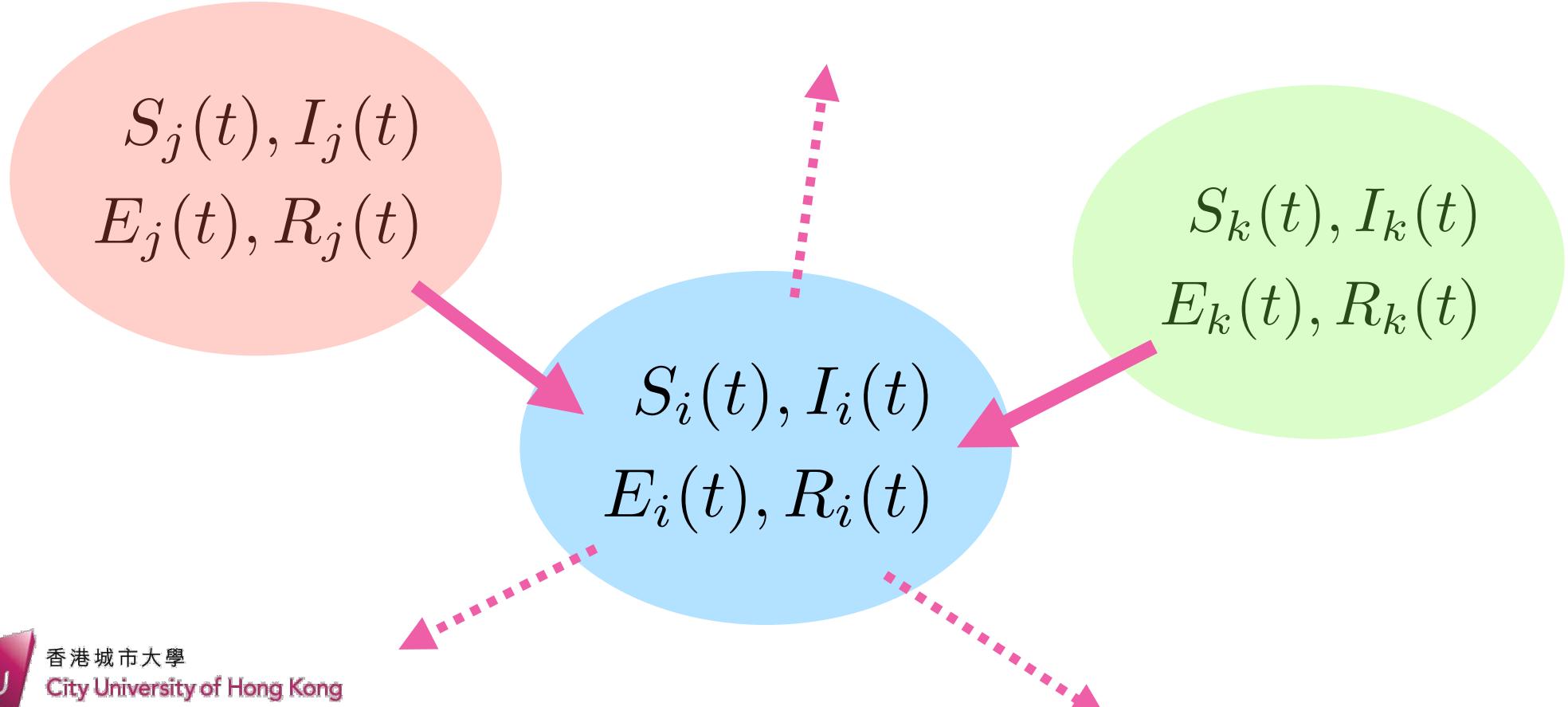
- Infection rates:  $S \rightarrow E$  and  $E \rightarrow I$ , i.e.,  $\beta_i$  and  $\kappa_i$  for each city
- Recovery rate:  $I \rightarrow R$ , i.e.,  $\gamma_i$  for each city
- Eventual infection percentage:  $\delta_i$  for each city

# Data

- Population:  $P_i$  for each city
- Infected:  $I_i, R_i$  for each city

# With intercity travel

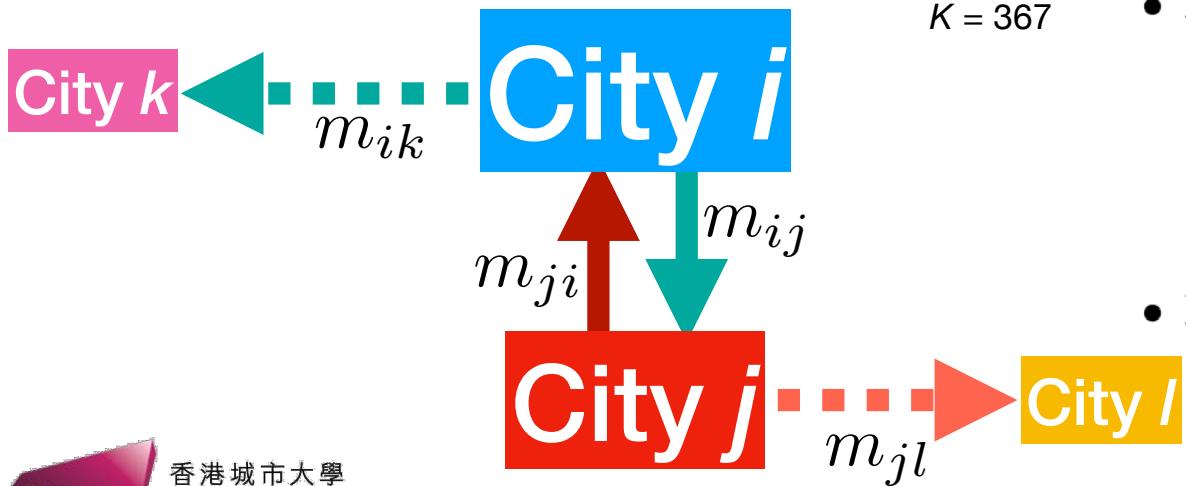
e.g., 267 Chinese cities are interacting via inflow and outflow traffic



# Intercity migration matrix

$$M(t) = \begin{bmatrix} m_{11}(t) & m_{12}(t) & \cdots & m_{1K}(t) \\ m_{21}(t) & m_{22}(t) & \cdots & m_{2K}(t) \\ \vdots & \vdots & \ddots & \vdots \\ m_{N1}(t) & m_{N2}(t) & \cdots & m_{KK}(t) \end{bmatrix}$$

/   
  $K = 367$



- $M$  records migration from one city to another. Movement within a city is not counted, i.e.,  $m_{ii}(t) = 0$  for all  $i$ .
- $M$  is non-symmetric as traffic from one city to another is not necessarily reciprocal at any given time, i.e.,  $m_{ij}(t) \neq m_{ji}(t)$ .
- Number of outflow migrants of city  $i$  at time  $t$  is

$$m_i^{(\text{out})}(t) = \sum_{i=j}^K m_{ij}(t).$$

- Number of inflow migrants of city  $i$  at time  $t$  is

$$m_i^{(\text{in})}(t) = \sum_{j=1}^K m_{ji}(t).$$

# From city $i$ to city $j$

$$\Delta I_{ij}^{\text{in}}(t) = \frac{I_i(t)m_{ij}(t)}{P_i}$$

# Leaving city $j$

$$\Delta I_j^{\text{out}}(t) = \frac{I_j(t) \sum_{i=1}^N m_{ji}(t)}{P_j}$$

**Likewise for the exposed populations**

# Assumptions / Conditions

- Susceptible and Exposed individuals can move from one city to another.
- Infected individuals, once confirmed, do NOT move to another city.
- Recovered individuals also do NOT move to another city.
- All parameters are theoretically different in different cities.
  - For computational convenience, we only assume recovery rates and infection percentages to be different in different cities.

# Travel-Data Integrated SEIR Model

MODEL	PARAMETERS
Set of Equations	(unknown)

daily changes	
$\Delta S = \dots$	$\alpha_j$
$\Delta E = \dots$	$\beta_j$
$\Delta I = \dots$	$\gamma_j$
$\Delta R = \dots$	$\kappa_j$
$\Delta P = \dots$	$\delta_j$

travel data

$$M(t) = \begin{bmatrix} m_{11}(t) & m_{12}(t) & \cdots & m_{1K}(t) \\ m_{21}(t) & m_{22}(t) & \cdots & m_{2K}(t) \\ \vdots & \vdots & \ddots & \vdots \\ m_{N1}(t) & m_{N2}(t) & \cdots & m_{KK}(t) \end{bmatrix}$$

Here we do not discuss  
the contact topology.

**S**

$$\Delta S_j(t) = -\frac{\beta_j(t)}{N_j^s(t)} I_j(t) S_j(t) - \frac{\alpha_j(t)}{N_j^s(t)} E_j(t) S_j(t) + \sum_{i=1}^N \left( \frac{S_i(t) m_{ij}(t)}{P_i(t)} \right) - \frac{S_j(t) \sum_{i=1}^N m_{ji}(t)}{P_j(t)}$$

**E**

$$\Delta E_j(t) = \frac{\beta_j(t)}{N_j^s(t)} I_j(t) S_j(t) + \frac{\alpha_j(t)}{N_j^s(t)} E_j(t) S_j(t) - \kappa_j(t) E_i(t) + \sum_{i=1}^N \left( \frac{E_i(t) m_{ij}(t)}{P_i(t)} \right) - \frac{E_j(t) * \sum_{i=1}^N m_{ji}(t)}{P_j(t)},$$

**I**

$$\Delta I_j(t) = \kappa_j(t) E_i(t) - \gamma_j(t) I_j(t) + k_I \left( \sum_{i=1}^N \left( \frac{I_i(t) m_{ij}(t)}{P_i(t)} \right) - \frac{I_j(t) * \sum_{i=1}^N m_{ji}(t)}{P_j(t)} \right)$$

**R**

$$\Delta R_j(t) = \gamma_j(t) I_j(t)$$

Population change:

$$\Delta P_j(t) = \sum_{i=1}^N m_{ij}(t) - \sum_{i=1}^N m_{ji}(t)$$

$$\Delta N_j^s(t) = k_I \left( \sum_{i=1}^N \left( \frac{I_i(t) m_{ij}(t)}{P_i(t)} \right) - \frac{I_j(t) * \sum_{i=1}^N m_{ji}(t)}{P_j(t)} \right) + \sum_{i=1}^N \left( \frac{E_i(t) m_{ij}(t)}{P_i(t)} \right) - \frac{E_j(t) * \sum_{i=1}^N m_{ji}(t)}{P_j(t)}$$

$$+ \sum_{i=1}^N \left( \frac{S_i(t) m_{ij}(t)}{P_i(t)} \right) - \frac{S_j(t) \sum_{i=1}^N m_{ji}(t)}{P_j(t)}$$

# Parameter Extraction

- Set of parameters:  $\theta_j = \{\alpha_j, \beta_j, \gamma_j, \kappa_j, \delta_j, I_{j,0}\}$  for city  $j$
- Initial infected and exposed numbers:  $\lambda_I I_j(t_0)$  and  $\lambda_E I_j(t_0)$

Modeling to be completed by finding all parameters that match the historical data.

The number of exposed individuals,  $E_j(t)$ , is also unknown.

# Method

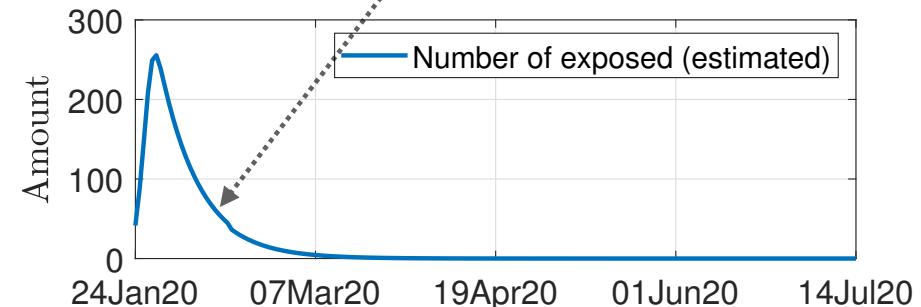
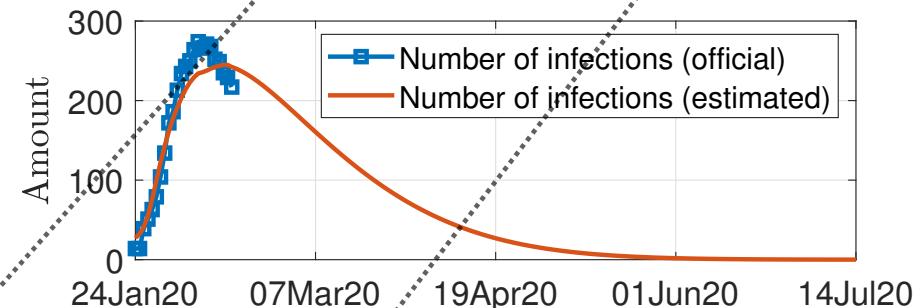
- *Constrained nonlinear optimisation* problem

$$\begin{aligned} P_0: \min_{\Theta} & \sum_{j=0}^N \|w_j(I(t_j) - \hat{I}(t_j))\|_l \\ \text{wh} \quad \text{s.t. } & \left\{ \begin{array}{l} (i) \quad \hat{x}(t+1) = \hat{x}(t) + F(\hat{x}(t)), \\ (ii) \quad \Theta_U \geq \Theta \geq \Theta_L, \end{array} \right. \end{aligned}$$

where  $\hat{x}$  is the set of estimated parameters, and  $F( \cdot )$  is the model.

- $\hat{x}(t) = [\hat{I}(t), \hat{R}(t), E(t), S(t), P(t), N^s(t)]$

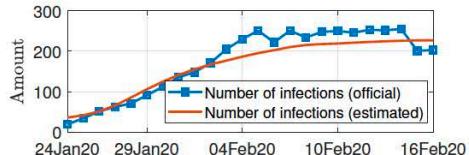
DATA FITTING with MIN ERROR  
to extract all parameters and unknown variables



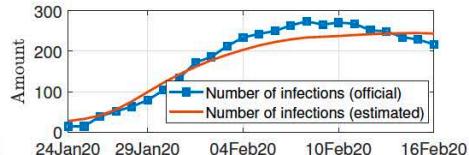
In other words, the model generates the solution that is *as close to the data as possible*.

# Fitting Results

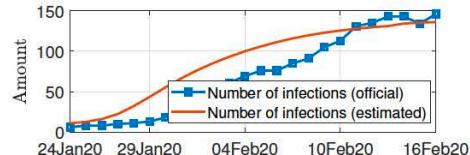
No of infected cases  
||  
Not the total cumulated  
(i.e., less the recovered)



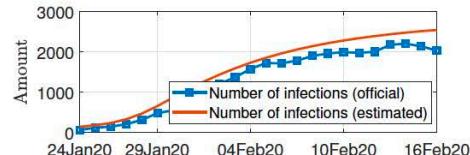
(a) Beijing



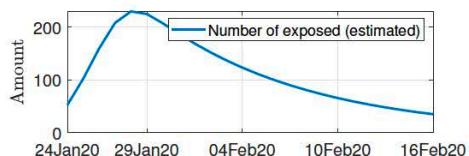
(b) Guangzhou



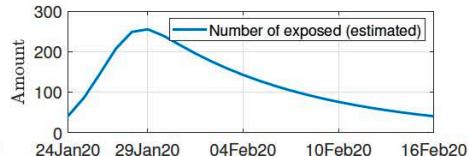
(c) Haerbin



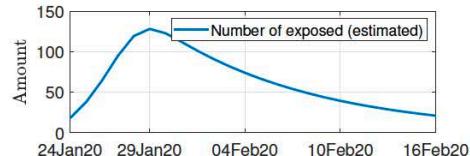
(d) Huanggang



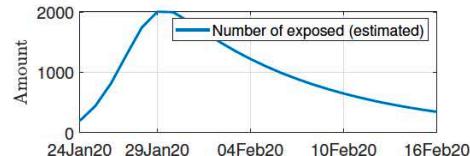
(e) Jingmen



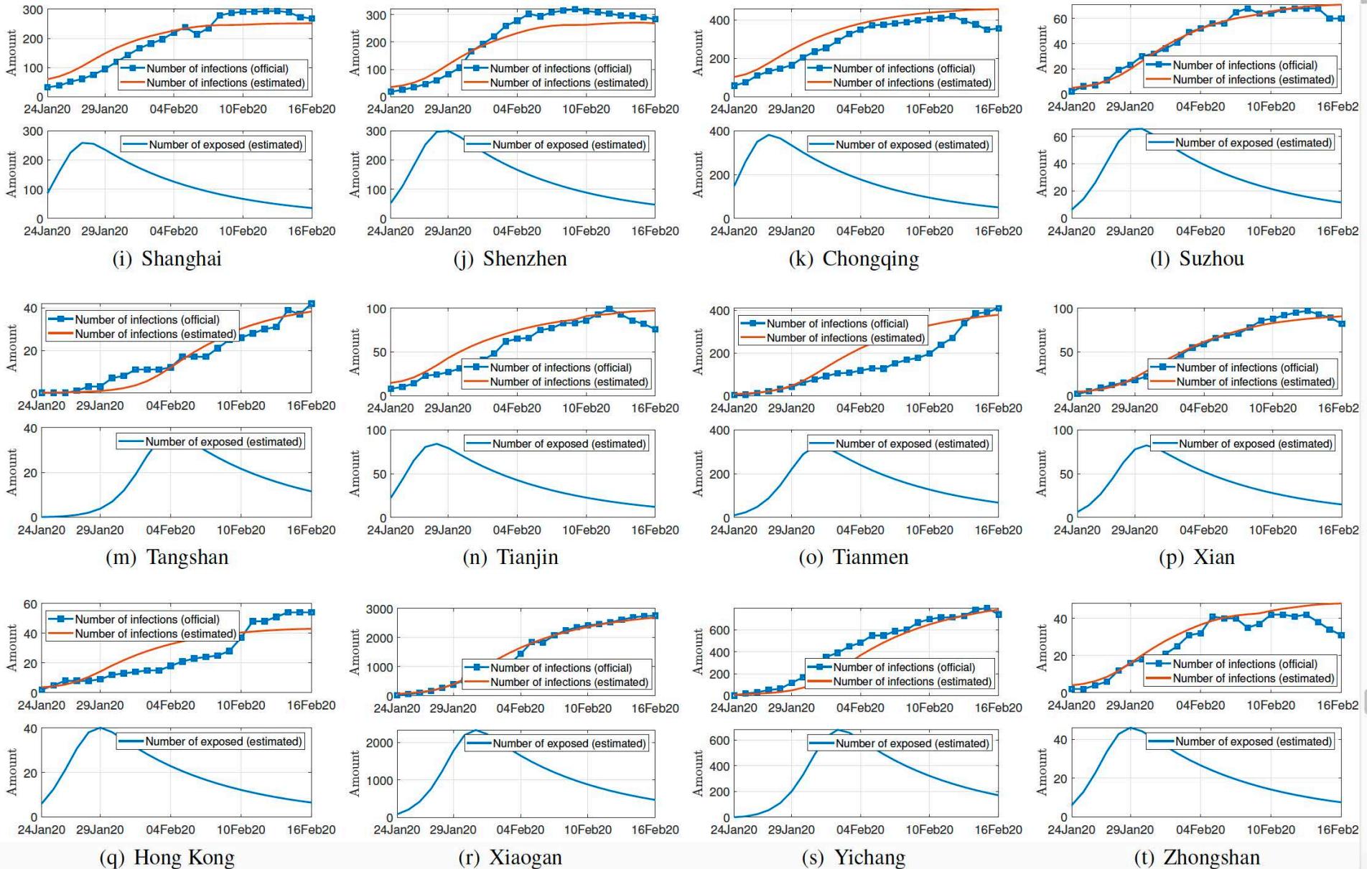
(f) Jingzhou



(g) Lianyungang



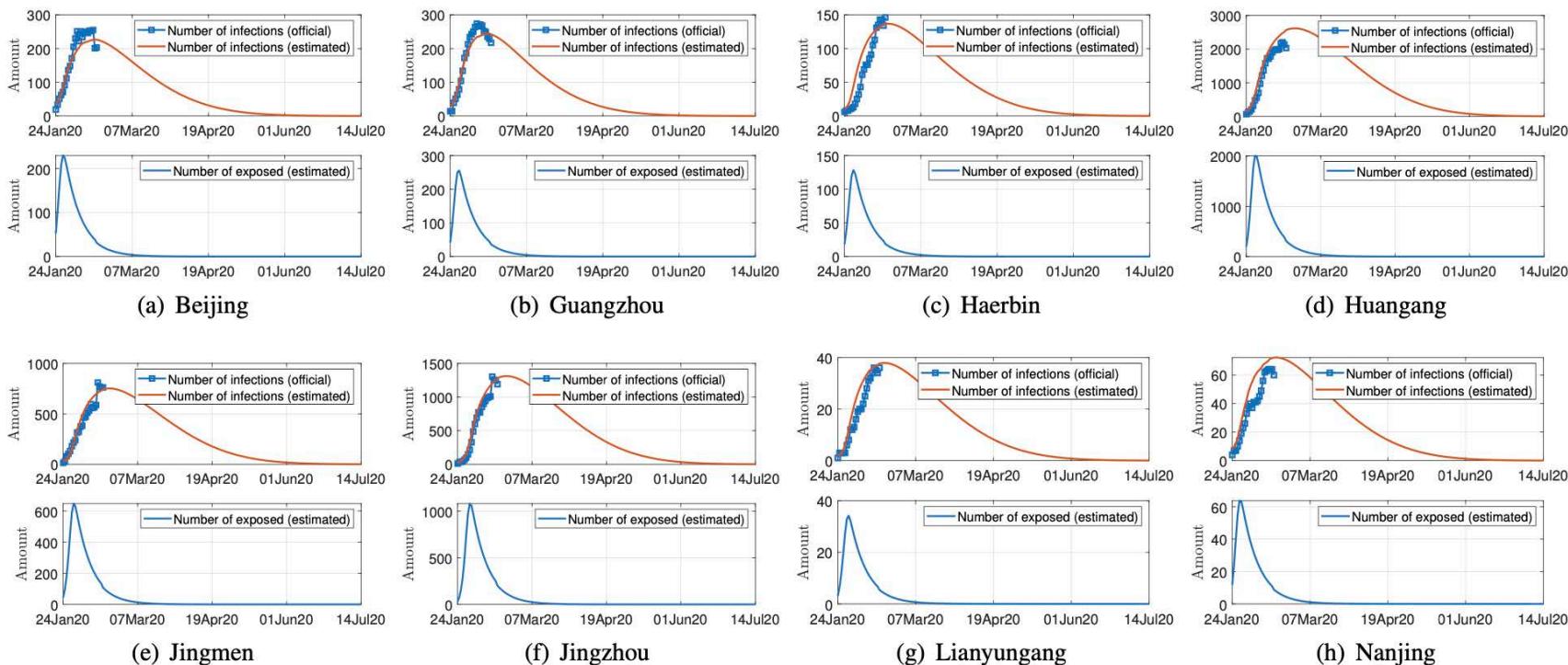
(h) Nanjing

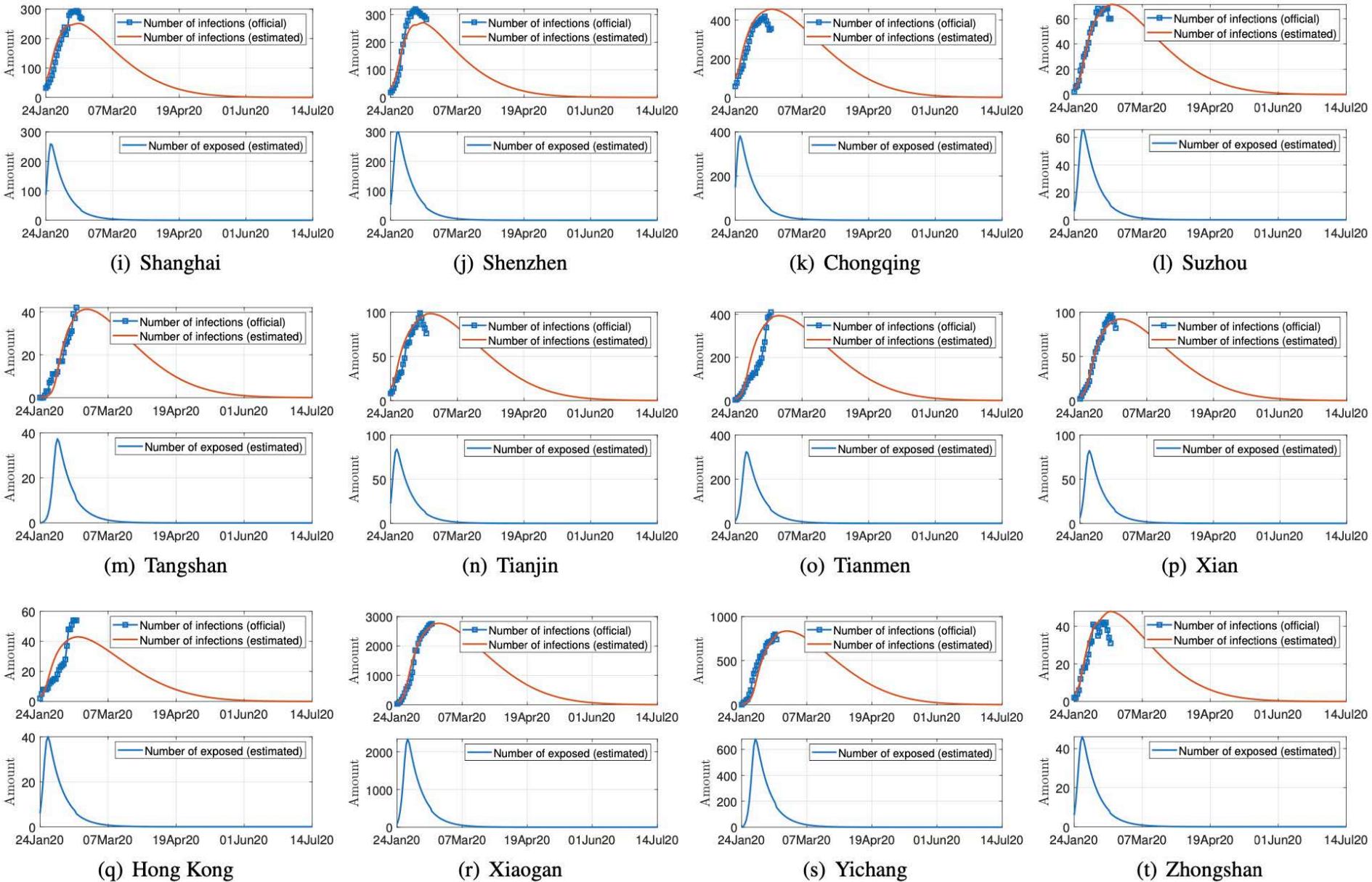


# Predictions

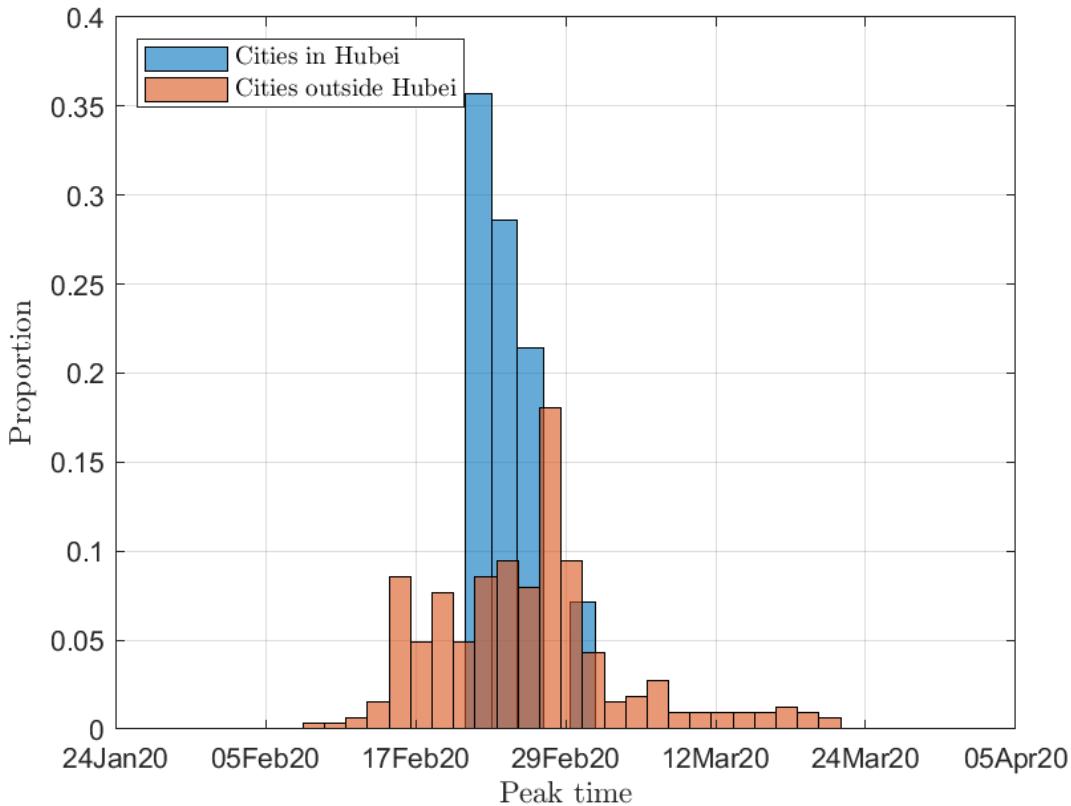
- Once we have the parameters, the model can be used for prediction assuming the validity of the same set of parameters.

**No of infected cases**  
||  
**Not the total cumulated  
(i.e., less the recovered)**



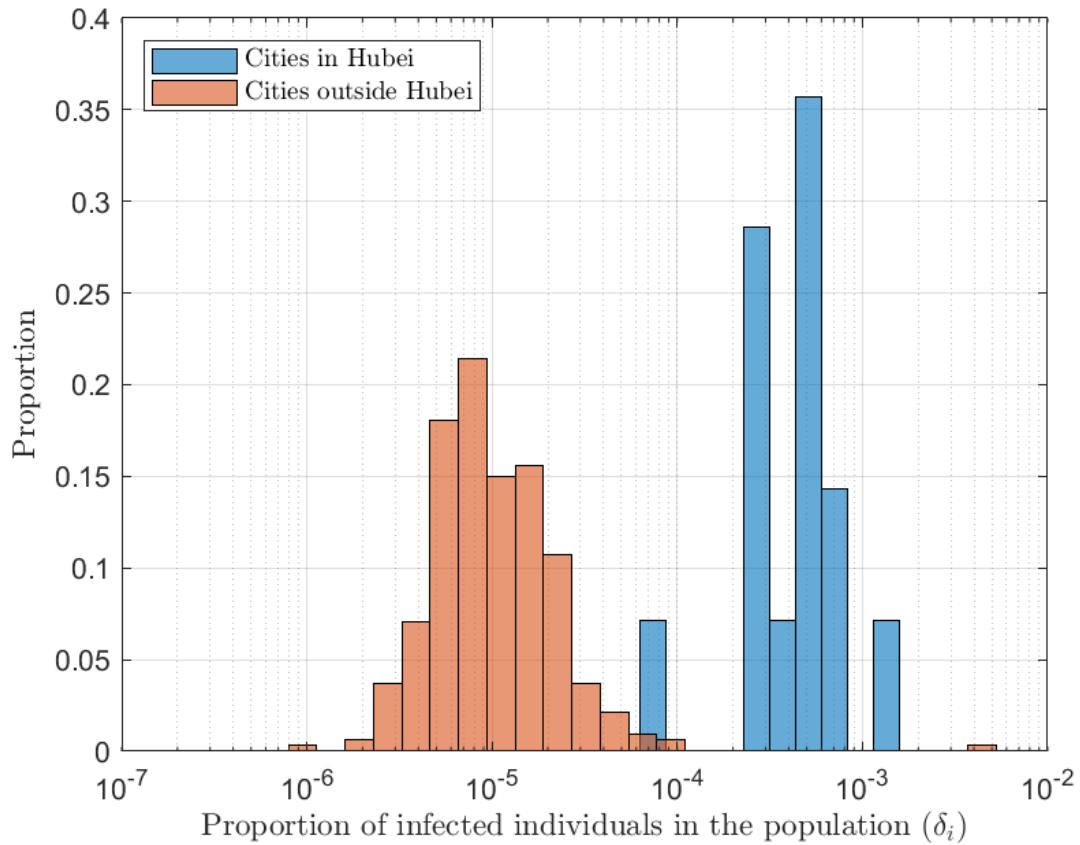


# Findings



## PEAK TIME

For most cities, the infection numbers would peak between mid February to early March 2020



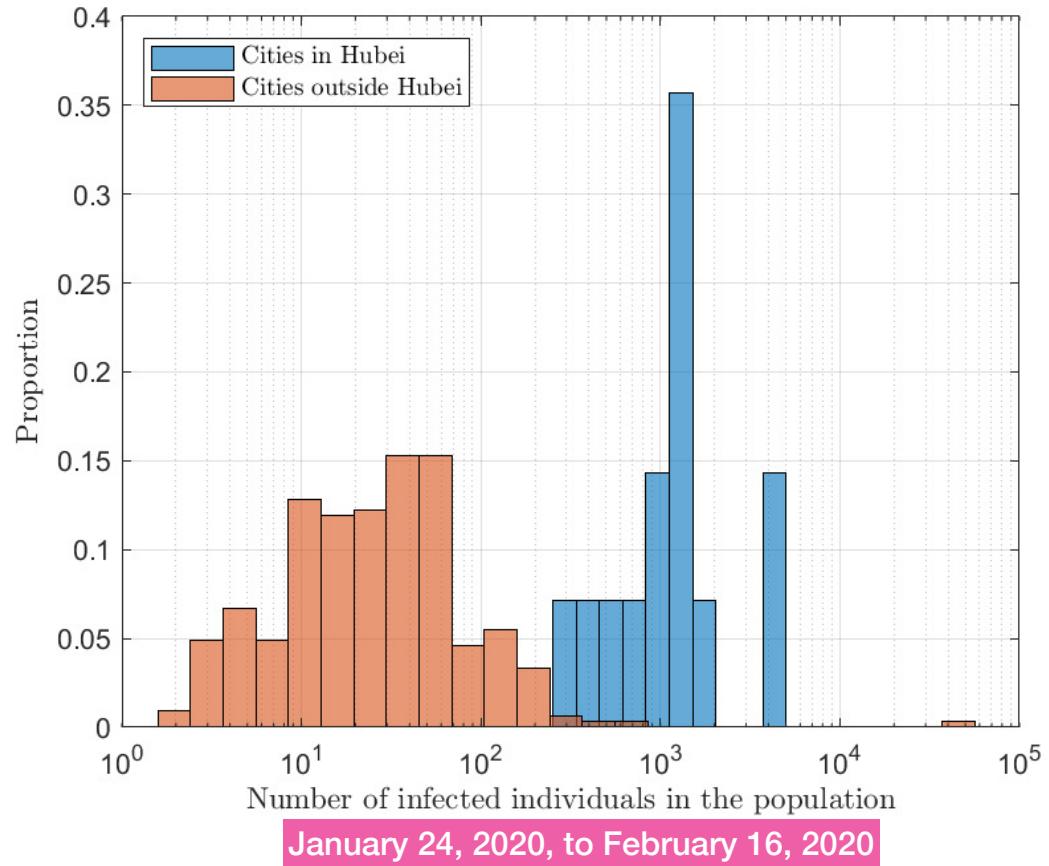
January 24, 2020, to February 16, 2020

## PROPORTION OF INFECTED IN EACH CITY $\delta_i$

At the end, about 0.8%, less than 0.1% and less than 0.01% of the population would get infected in Wuhan, Hubei Province and the rest of China, respectively.

## NUMBER OF INFECTED IN EACH CITY

Translating to actual figures, for most cities outside and within Hubei Province (except Wuhan), the total number of infected individuals was expected to be fewer than 300 and 4000, respectively



# Other countries by data-driven prediction

South Korea

Italy

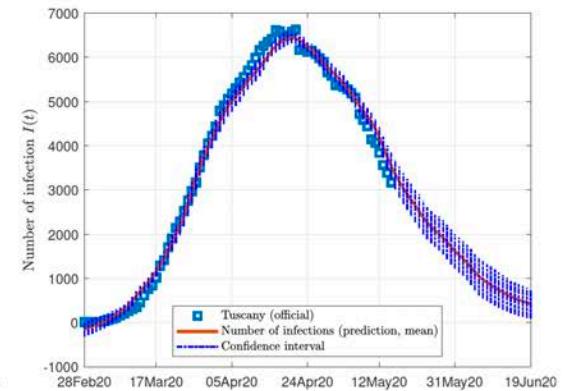
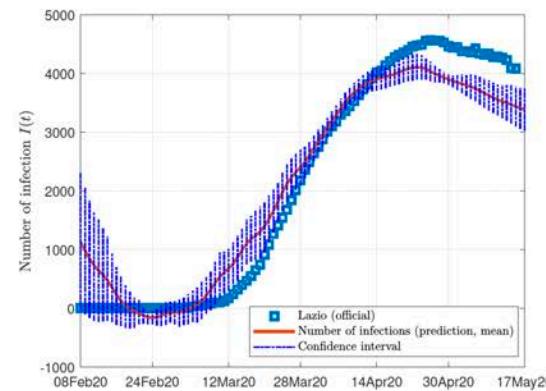
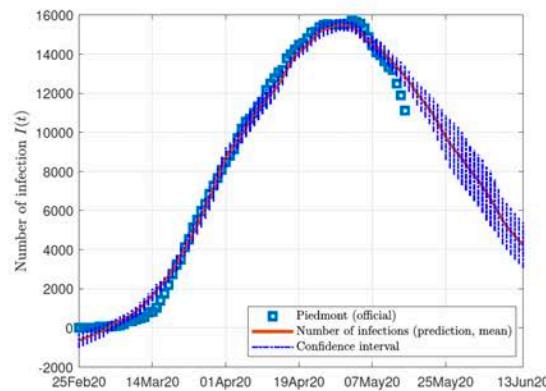
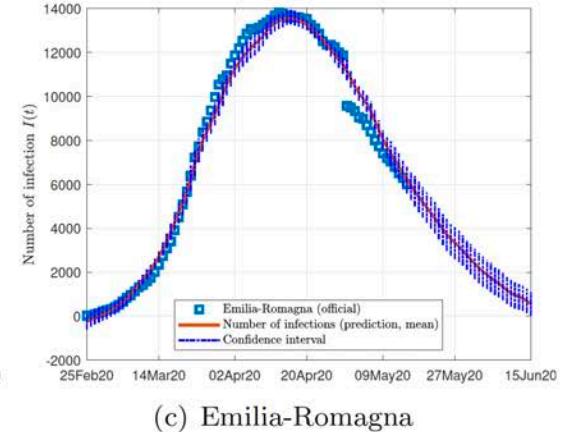
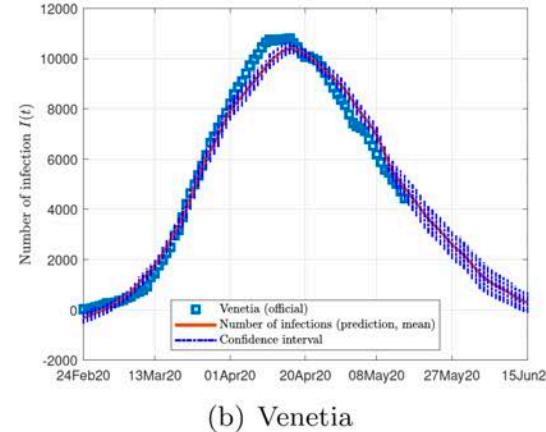
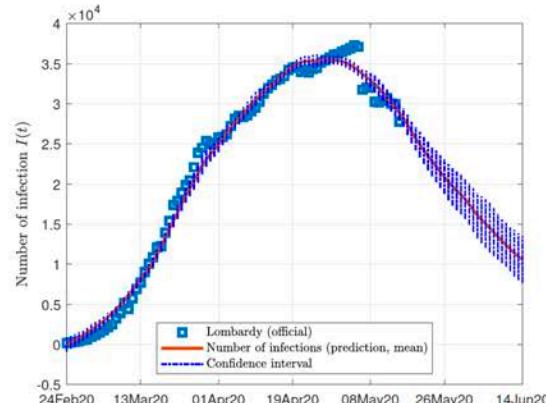
Iran

Japan

USA

UK

Over 500 cities



# Other countries by data-driven prediction

South Korea

Italy

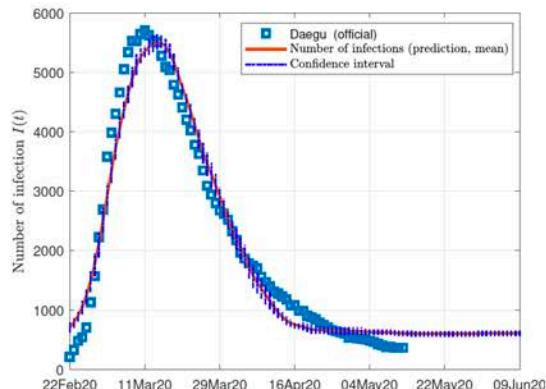
Iran

Japan

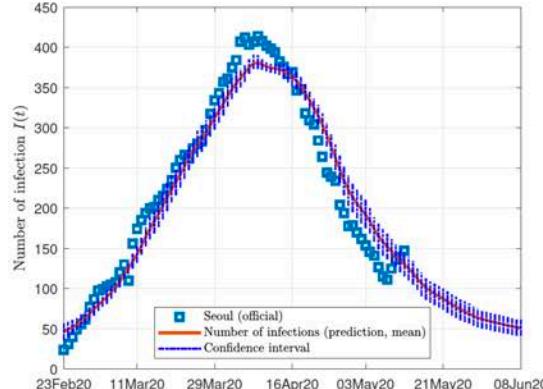
USA

UK

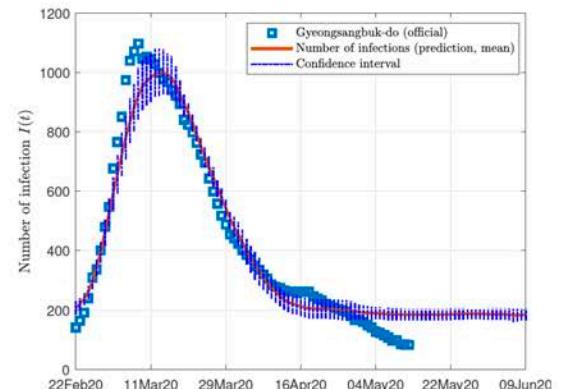
Over 500 cities



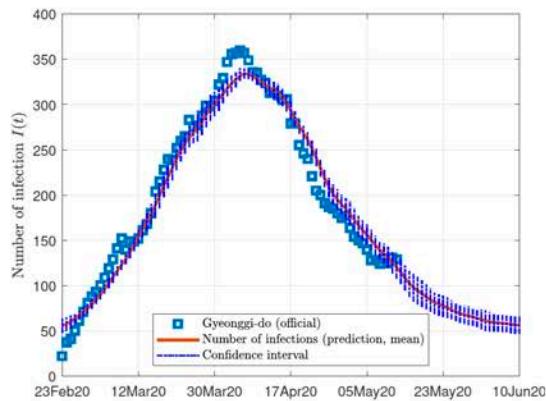
(a) Daegu



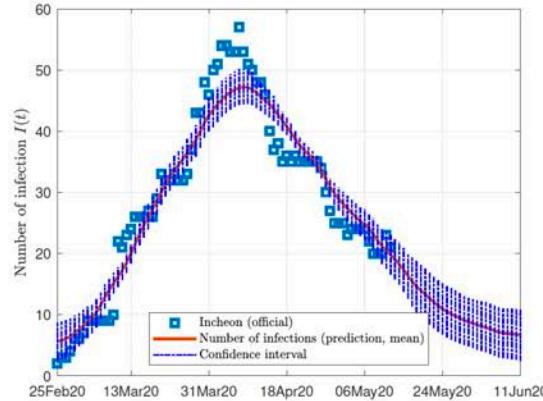
(b) Seoul



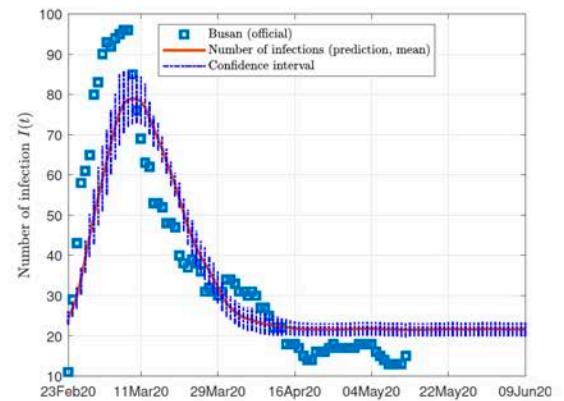
(c) Gyeongsangbuk-do



(d) Gyeonggi-do



(e) Incheon



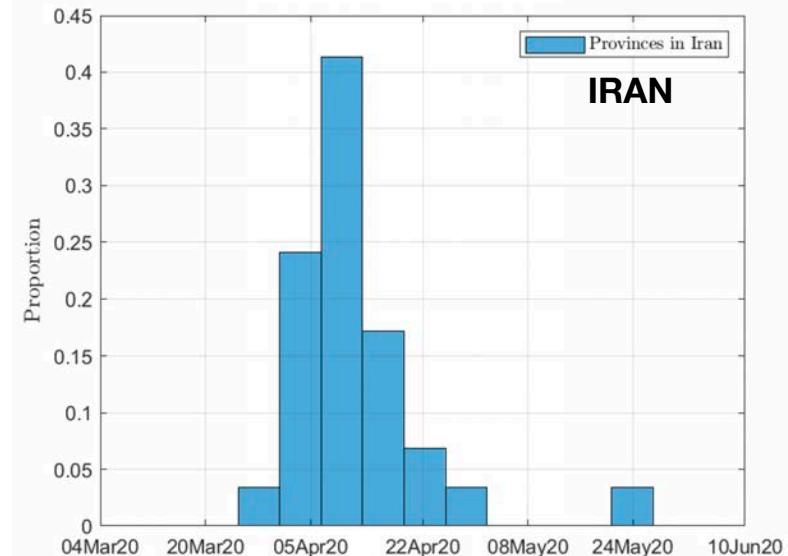
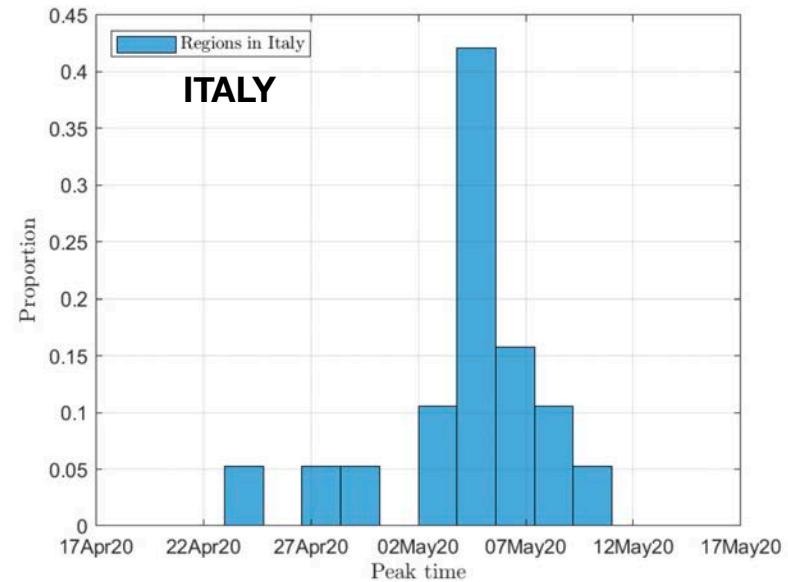
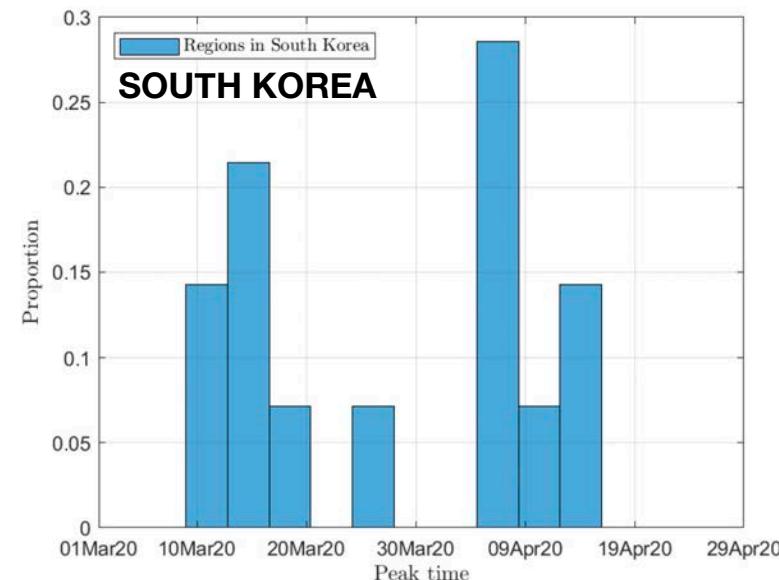
(f) Busan

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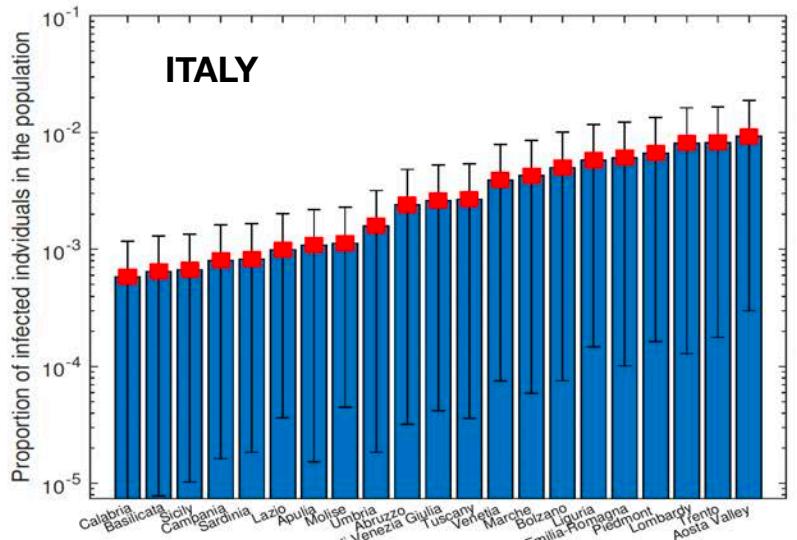
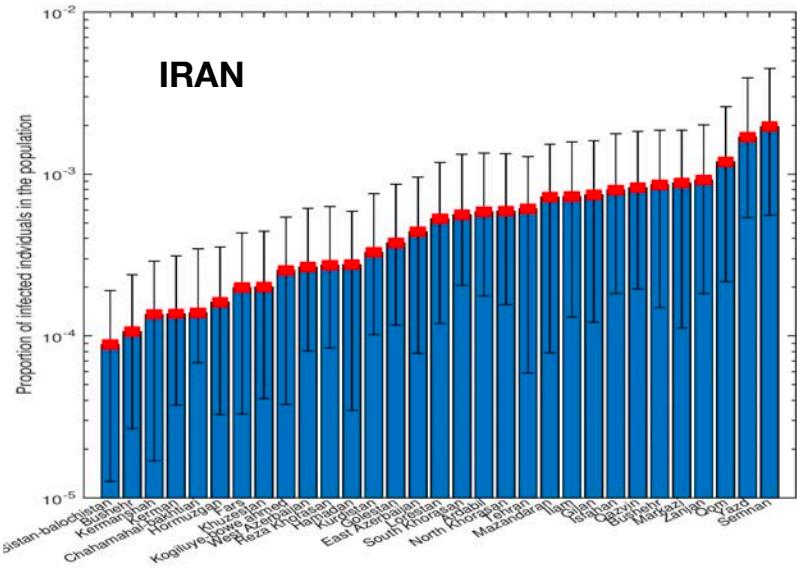
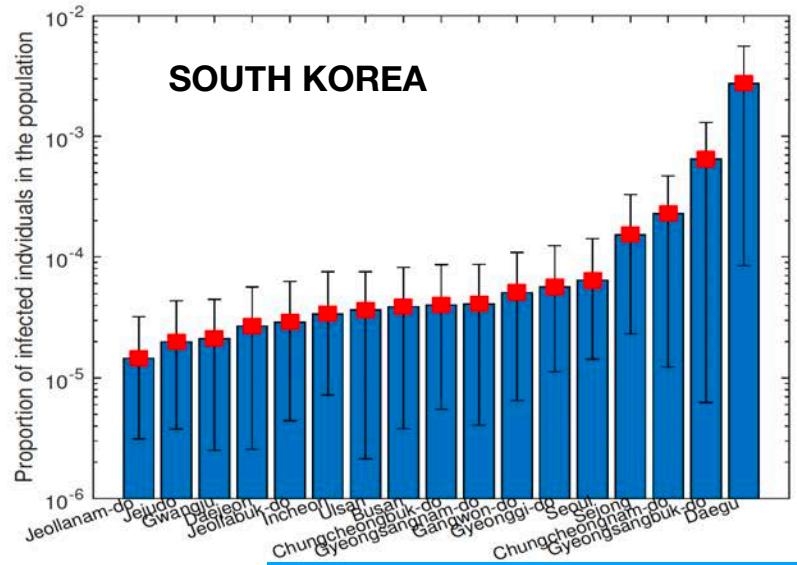
## PEAK TIME

South Korea – end by April

Italy, Iran – end by May



## PROPORTION OF INFECTED INDIVIDUALS IN THE POPULATION



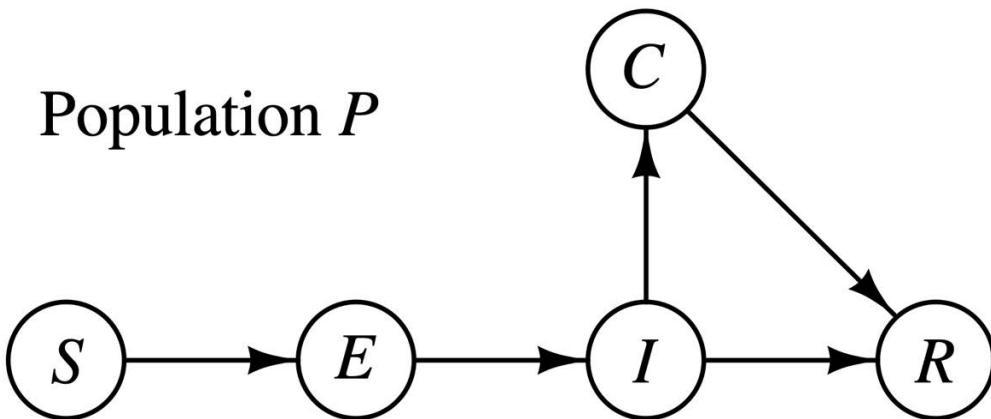
PERFORMED MARCH 8, 2020; UPDATED MAY 17, 2020

# Next Challenge: Latency / Loss of Information

Most notable information latency – the number of confirmed cases reported, which depends on

- the ability of the particular country or city to perform tests
- the possible bureaucracy in the local system of reporting

$$I \gg C$$



OUR LATEST WORK (COMPLETED JULY 27) SHOWED THAT  $C/I \approx 0.23$ , i.e.,  
**Unconfirmed : confirmed  $\approx 3.309 : 1$**   
based on data from Australia, Canada, Italy, Japan, Spain, UK and USA

# SEICR MODEL

Assume: the number of individuals eventually infected is set initially at  $N_j^s(t_0) = \delta_j P_j$  ( $\delta_j$  being constant), implying effective measures in place to limit the upper bound of the infected number.

But the model may consider inactive or less active intervention: each infected case adds to the eventual infected number. The additional  $S_j$  and  $N_j$  are:

$$k_j^{(c)} (\Delta I_j(t) + \Delta E_j(t)) \left( 1 - \frac{N_j^s(t)}{k_h P_j(t)} \right)$$

$k_j^{(c)}$  is an inverse indicator of the level of active intervention implemented, and corresponds quantitatively to an increase in the number of eventual infected individuals for each additional infected or exposed individual in region  $j$

$$\Delta I_j(t) = (1 - \lambda_j - \gamma_j^{(I)}) I_j(t) + \kappa E_i(t) + k_I \left( \sum_{i=1}^N \left( \frac{I_i(t)m_{ij}(t)}{P_i(t)} \right) - \frac{I_j(t) \sum_{i=1}^N m_{ji}(t)}{P_j(t)} \right)$$

$$\begin{aligned} \Delta E_j(t) = & \frac{\beta_j I_j(t) + \rho_j C_j(t)}{N_j^s(t)} S_j(t) + \frac{\alpha_j E_j(t)}{N_j^s(t)} S_j(t) - \kappa E_i(t) + \sum_{i=1}^N \left( \frac{E_i(t)m_{ij}(t)}{P_i(t)} \right) \\ & - \frac{E_j(t) * \sum_{i=1}^N m_{ji}(t)}{P_j(t)} \end{aligned}$$

$$\begin{aligned} \Delta S_j(t) = & -\frac{\beta_j I_j(t) + \rho_j C_j(t)}{N_j^s(t)} S_j(t) - \frac{\alpha_j E_j(t)}{N_j^s(t)} S_j(t) + \sum_{i=1}^N \left( \frac{S_i(t)m_{ij}(t)}{P_i(t)} \right) \\ & - \frac{S_j(t) \sum_{i=1}^N m_{ji}(t)}{P_j(t)} \\ & + \boxed{k_j^{(c)} (\Delta I_j(t) + \Delta E_j(t)) \left( 1 - \frac{N_j^s(t)}{k_h P_j(t)} \right)} \end{aligned}$$

$$\Delta C_j(t) = \lambda_j I_j(t) - \gamma_j^{(C)} C_j(t)$$

$$\Delta R_j(t) = \gamma_j^{(I)} I_j(t) + \gamma_j^{(C)} C_j(t)$$

$$\Delta P_j(t) = \sum_{i=1}^N m_{ij}(t) - \sum_{i=1}^N m_{ji}(t)$$

$$\begin{aligned} \Delta N_j^s(t) = & k_I \left( \sum_{i=1}^N \left( \frac{I_i(t)m_{ij}(t)}{P_i(t)} \right) - \frac{I_j(t) \sum_{i=1}^N m_{ji}(t)}{P_j(t)} \right) + \sum_{i=1}^N \left( \frac{E_i(t)m_{ij}(t)}{P_i(t)} \right) \\ & - \frac{E_j(t) \sum_{i=1}^N m_{ji}(t)}{P_j(t)} + \sum_{i=1}^N \left( \frac{S_i(t)m_{ij}(t)}{P_i(t)} \right) - \frac{S_j(t) \sum_{i=1}^N m_{ji}(t)}{P_j(t)} \end{aligned}$$

$$+ \boxed{k_j^{(c)} (\Delta I_j(t) + \Delta E_j(t)) \left( 1 - \frac{N_j^s(t)}{k_h P_j(t)} \right)}$$

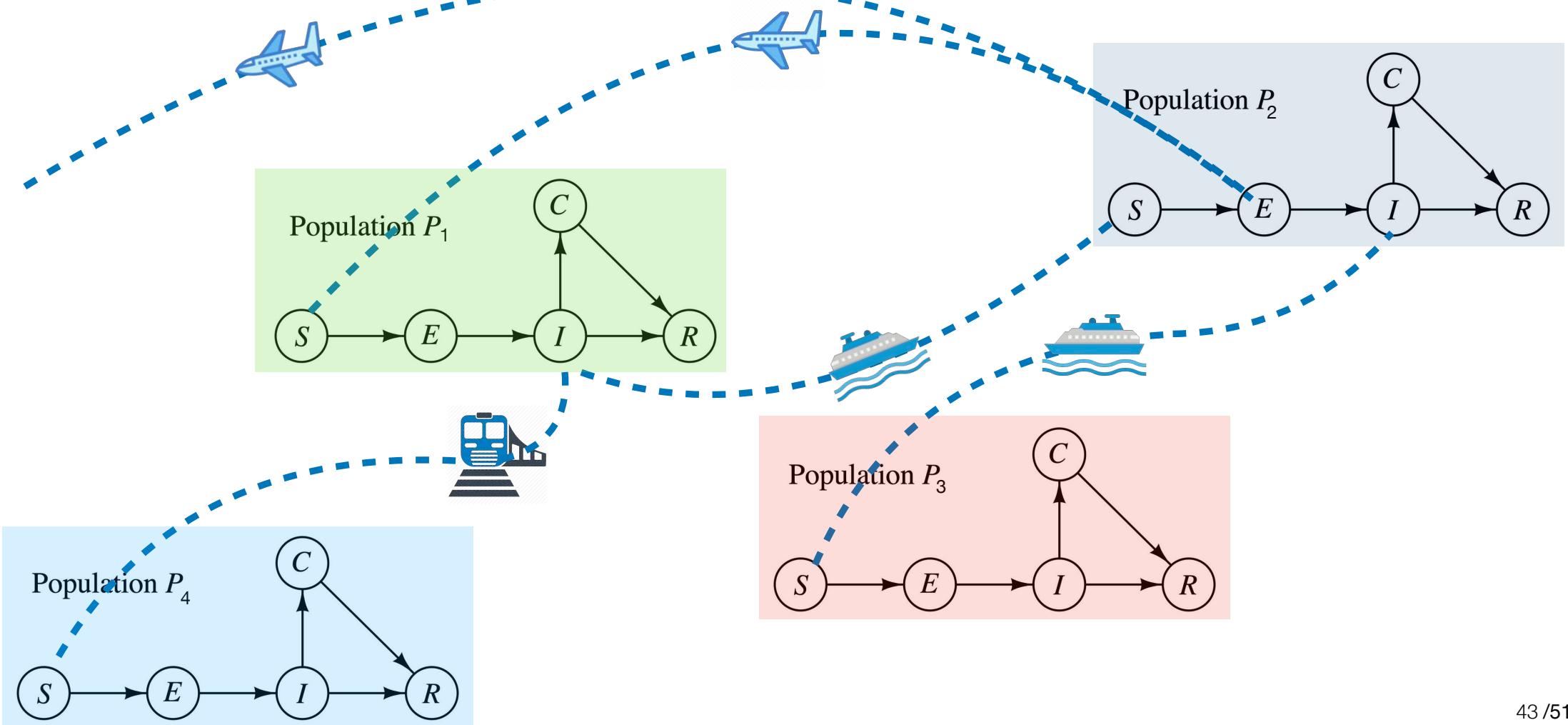
# Full Parameter Set

Symbol	Meaning
$\alpha_j$	rate of infecting a susceptible individual by an exposed individual in region $j$
$\beta_j$	rate of infecting a susceptible individual by an infected individual in region $j$
$\rho_j$	rate of infecting a susceptible individual by a confirmed individual in region $j$
$\lambda_j$	confirmed rate of infected individuals in region $j$
$\kappa_j$	rate of an exposed individual becoming infected
$\gamma^{(I)}$	recovery rate of an infected but not confirmed individual in region $j$
$\gamma^{(C)}$	recovery rate of a confirmed individual in region $j$
$k_I$	possibility of an infected individual moving from one region to another
$k_j^{(c)}$	increase in number of individuals eventually infected for each additional infected or exposed individual in region $j$
$\psi_j$	level of active intervention, $\psi_j = 1/k_j^{(c)}$
$k_h$	proportion of population infected achieving no further spreading, i.e., absolute upper bound for $N_j^s$ for all $j$
$\delta_j$	initial percentage of eventual infected individuals in region $j$
$I_{j0}$	initial number of infected individuals in region $j$
$E_{j0}$	initial number of individuals in region $j$
$C_{j0}$	initial number of confirmed infected individuals in region $j$

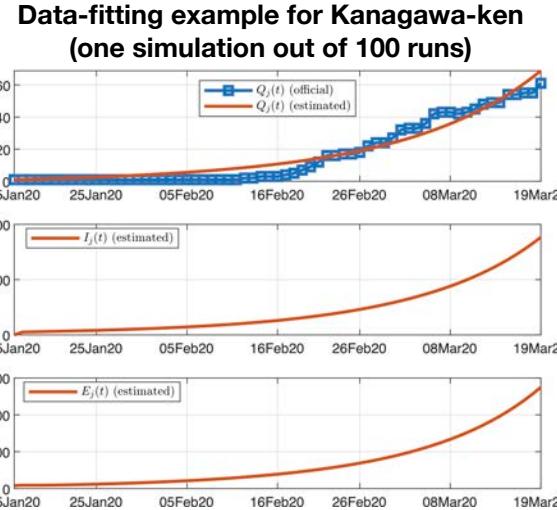
**inherent rates:**  
 social behavior  
 personal hygiene  
 mask wearing

**level of intervention:**  
 quarantine  
 border closure  
 cordon sanitaire

# Integrating Travel Data w SEICR

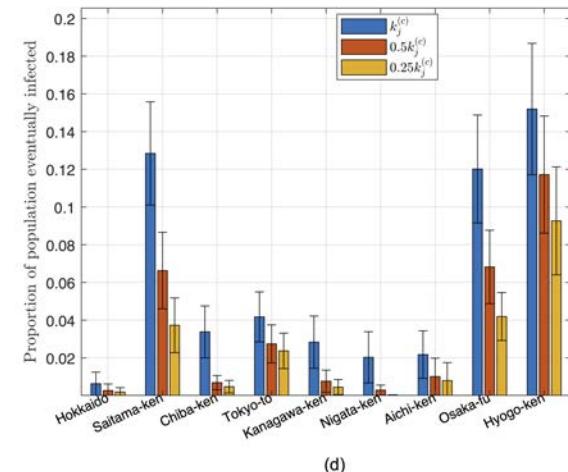
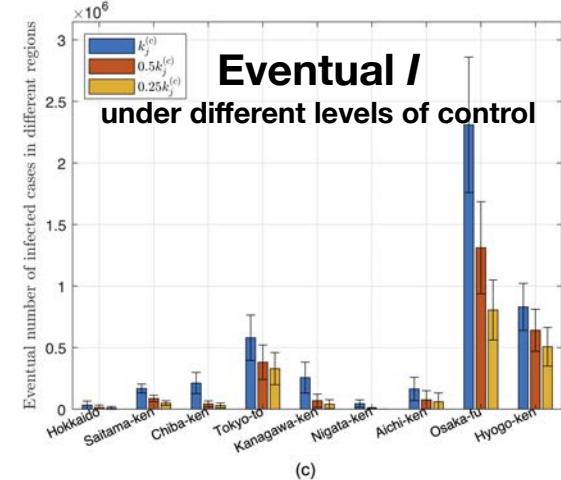
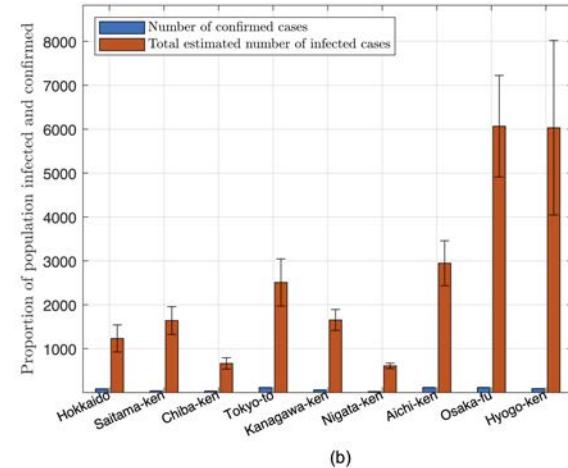
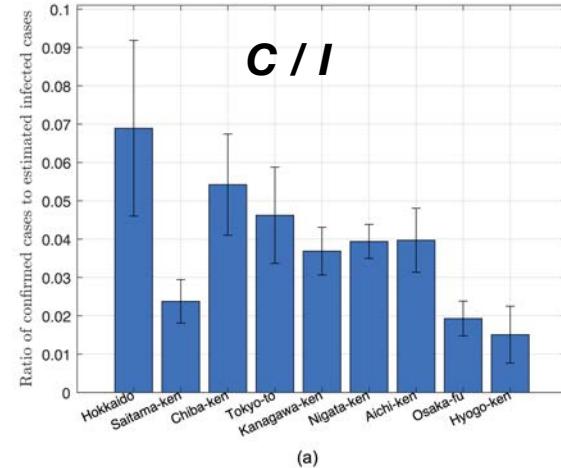


# Prediction for 46 Regions (Prefectures) in Japan



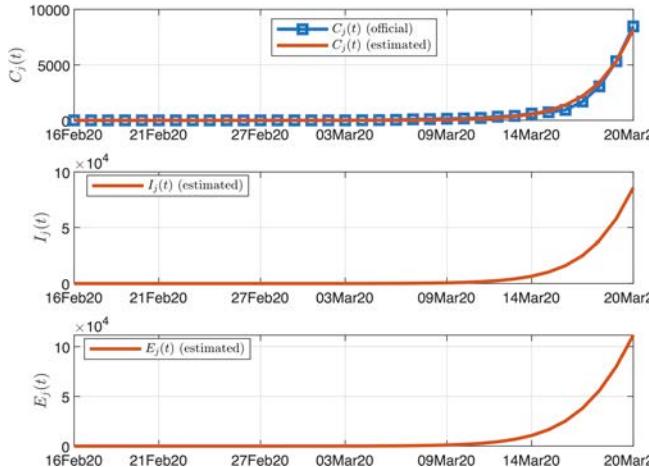
Actual infected cases can be 20-fold  
as many as the confirmed number.

$$I > 20C$$



# Prediction for 50 states (incl. DC) in USA

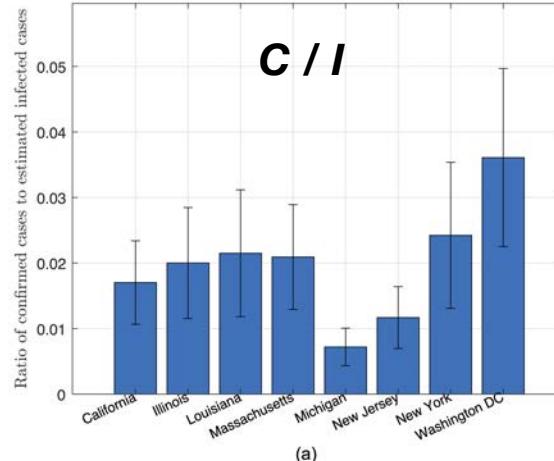
Data-fitting example for New York State  
(one simulation out of 100 runs)



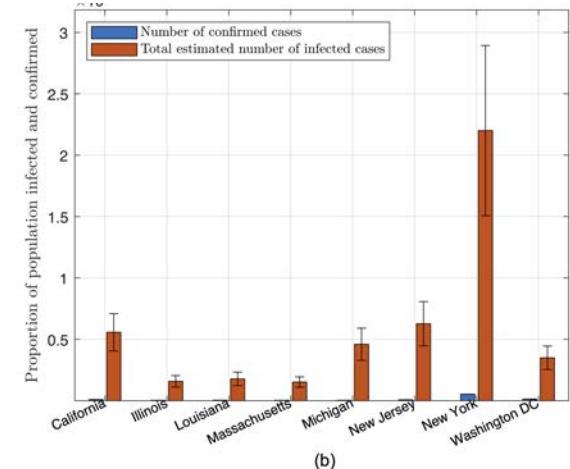
Actual infected cases can be 5-fold  
as many as the confirmed number.

$$I > 5C$$

**C / I**



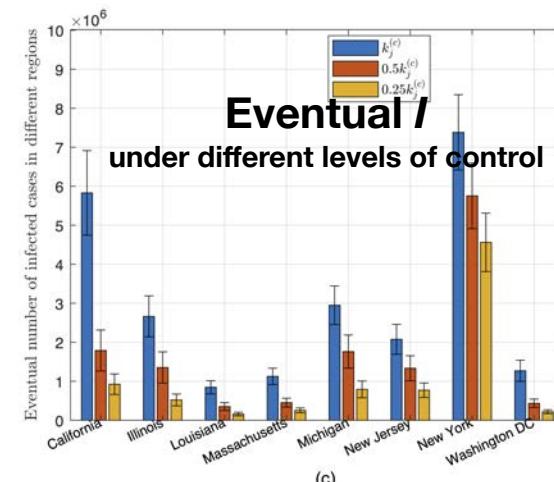
(a)



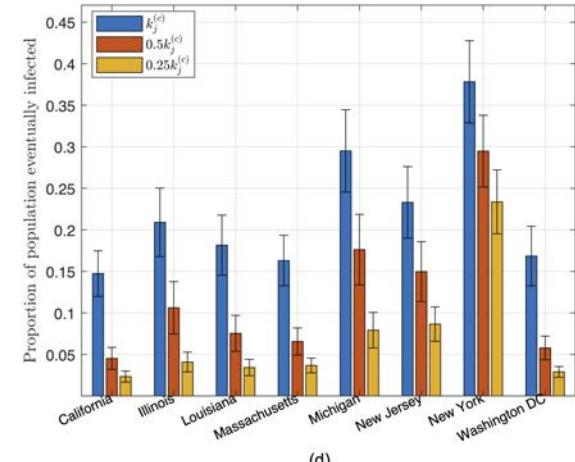
(b)

**Eventual I**

under different levels of control

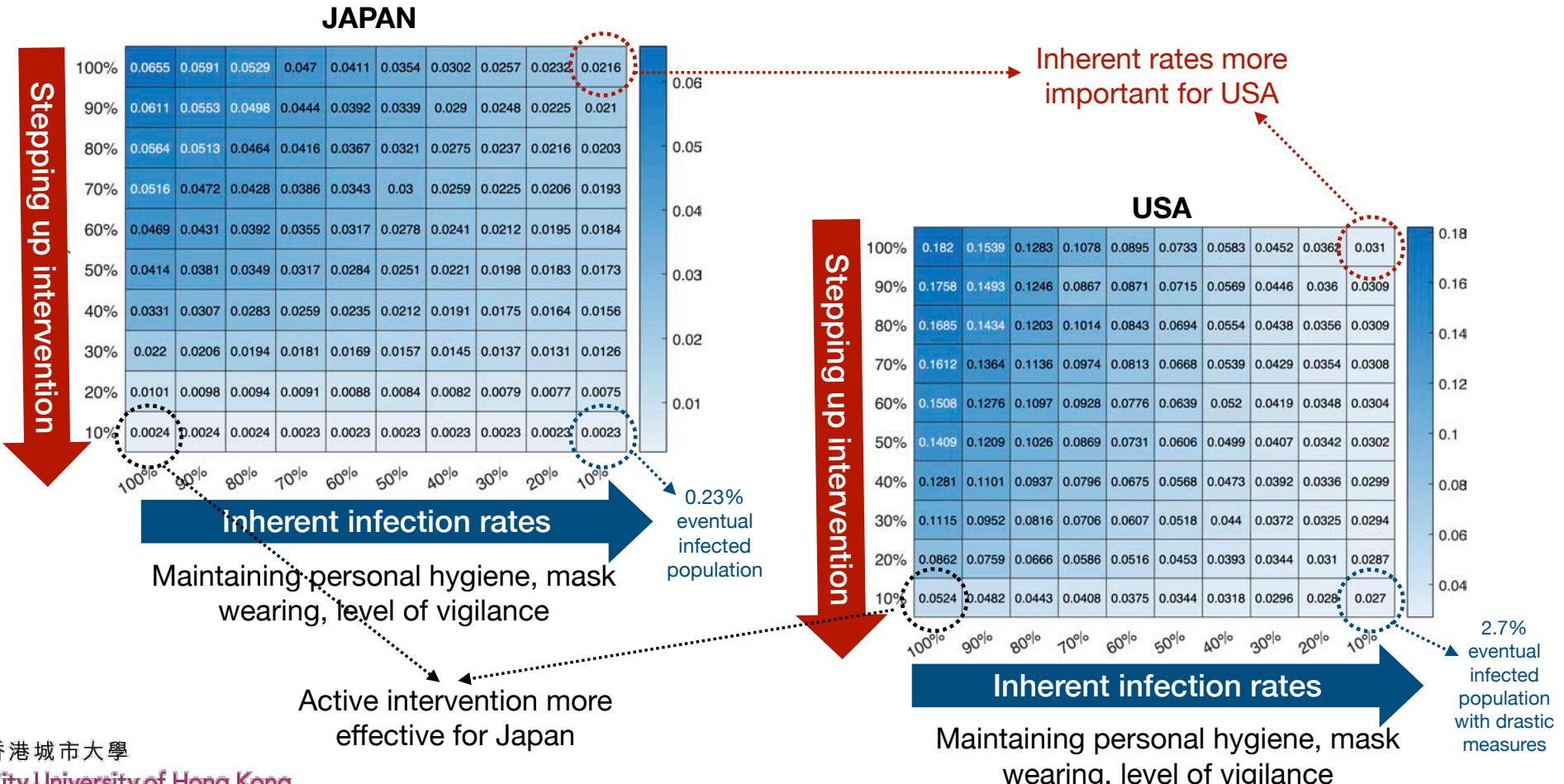


(c)



(d)

# Inherent rates vs intervention

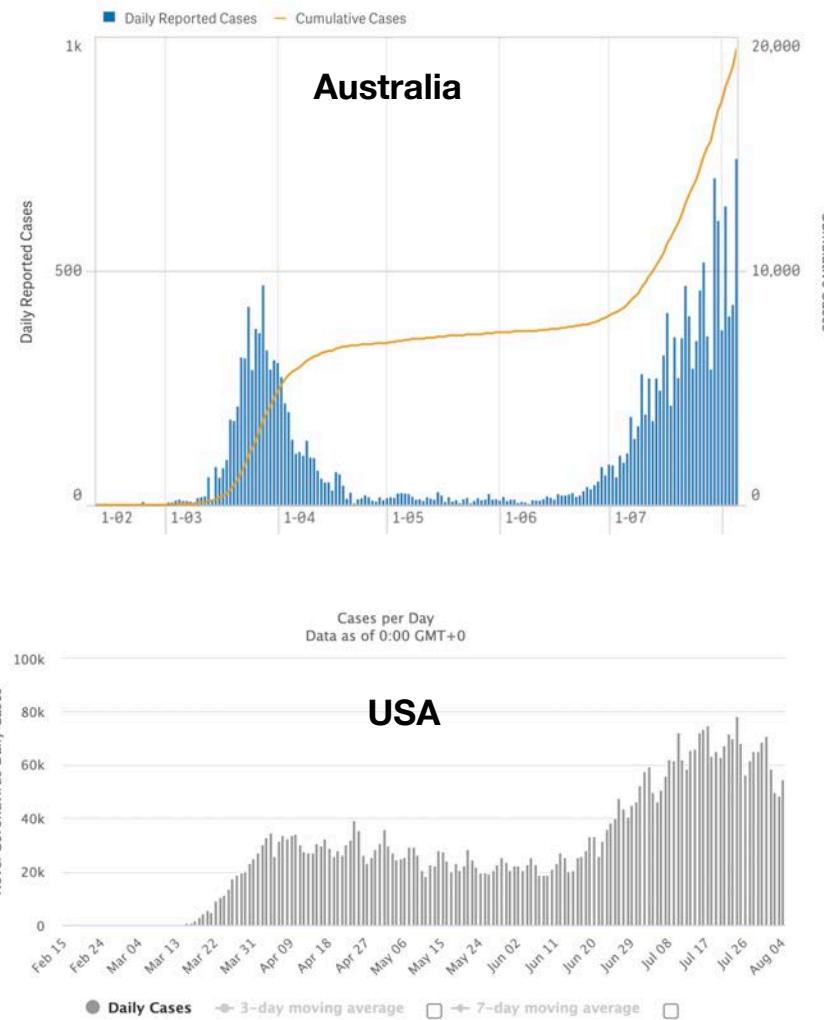


# Trends

Good News:

The **effectiveness of medical treatment** is expected to improve and the recovery rate will increase in time, making the decline rate more rapid.

Predictions of first waves are highly consistent so far,



Bad News

Borders remain practically open, and intercity travel is still active, though small. Imported cases will persist. Matter of time!!!!

Asymptomatic and presymptomatic transmissions mean that “unidentified” patients are the main infectious hosts.

# Conclusions

- Single population:
  - Gatherings of people - superspread events
- Multiple populations:
  - Intercity migration
- Value of model prediction:
  - Provides essential information on the expected severity and duration of the epidemic for illuminating social and non-pharmaceutical preventive interventions.

Drastically increase TESTING CAPACITY so as to identify the “invisible” infectious hosts.

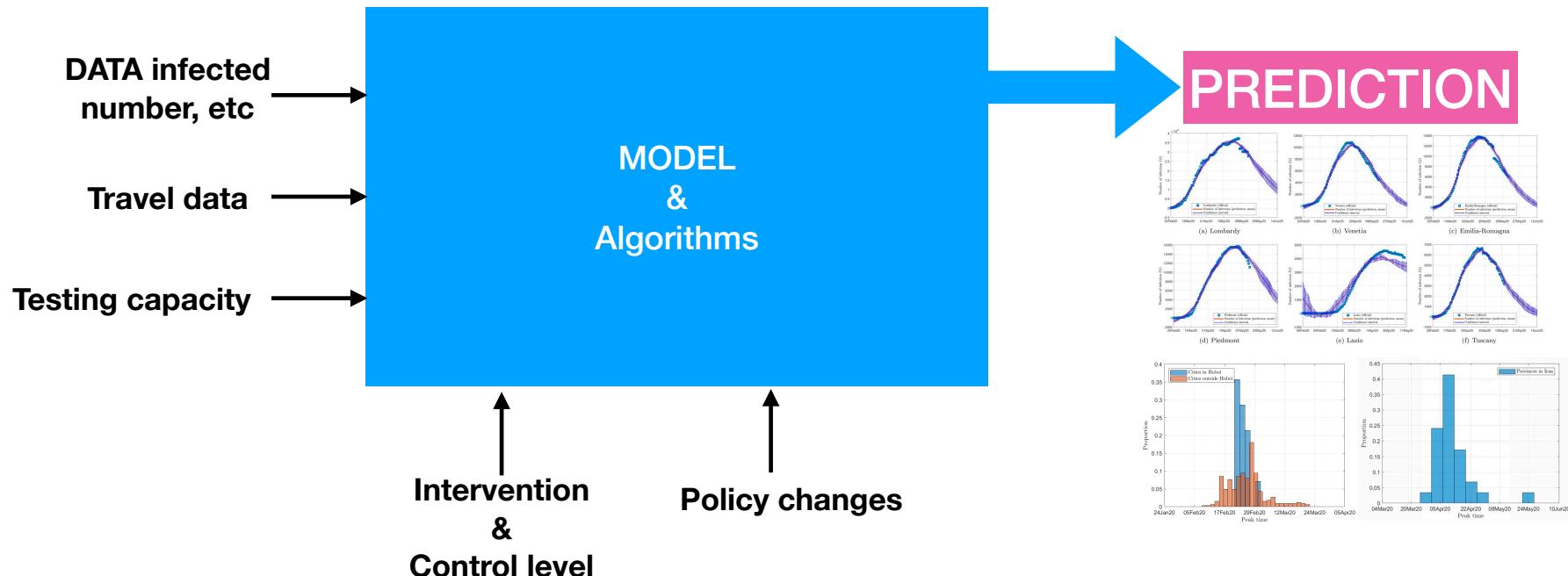
Our latest study shows that if testing capacity is raised 10-fold, the number of infected cases will reduce by 50%.

Right now, the confirmed cases are only 23% of the total infected cases.

Mass testing may be the way to go.

# Further Development

- Computerised System for ANALYSIS and PREDICTION



# References

## ON COVID-19

- C. Zhan, C. K. Tse, Y. Fu, Z. Lai, and H. Zhang, "Modeling and Prediction of the 2019 Coronavirus Disease Spreading in China Incorporating Human Migration Data," PLOS ONE, 2020. doi: <https://doi.org/10.1101/2020.02.18.20024570> [completed February 18, 2020]
- C. Zhan, C. K. Tse, Z. Lai, T. Hao, and J. Su, "Prediction of COVID-19 Spreading Profiles in South Korea, Italy and Iran by Data-Driven Coding," PLOS ONE, July 2020. doi: <https://doi.org/10.1101/2020.03.08.20032847> [completed March 8, 2020; revised on May 17, 2020]
- C. Zhan, C. K. Tse, Z. Lai, X. Chen, and M. Mo, "General Model for COVID-19 Spreading with Consideration of Intercity Migration, Insufficient Testing and Active Intervention: Modeling Study of Pandemic Progression in Japan and USA," JMIR Public Health and Surveillance, July 2020. doi: 10.2196/preprints.18880 URL: <https://preprints.jmir.org/preprint/18880> [completed March 25, 2020]

## OUR WEBSITE

<https://www.ee.cityu.edu.hk/~chitse/COVID-19/>



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### Background

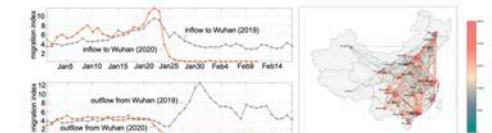
The Coronavirus Disease 2019 (SARS-CoV-2 or COVID-19) began to spread since mid December 2019 from Wuhan, which has been widely regarded as the epicentre of the epidemic, to almost all provinces throughout China and 80+ other countries. The SARS-CoV-2 outbreak began to occur and escalate in a special holiday period in China (about 20 days surrounding the Lunar New Year), during which a huge volume of intercity travel took place, resulting in outbreaks in multiple regions connected by an active transportation network. Thus, in order to understand the SARS-CoV-2 spreading process in China, it is essential to examine the human migration dynamics, especially between the epicentre Wuhan and other Chinese cities.

### Our Team

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Tianyong Hao  
Zhikang Lai  
Mingshan Mo  
Haijun Zhang



Baidu Migration Data of Inflow and Outflow Migration Strengths of Wuhan

### Model and Prediction for China (February 18, 2020)

We utilise the human migration data collected from Baidu Migration, which is a mobile-app based human migration tracking data system providing historical indicative daily volume of travellers to/from and between 367 cities in China. We combine, in this study, intercity travel data collected from Baidu Migration with the SEIR model to build a new dynamic model for the spreading of COVID-19 in China. Using official historical data of infected, recovered and death cases in 367 cities, we perform fitting of the data to estimate the best set of model parameters, which are then used to estimate the number of individuals exposed to the virus in each city and to predict the extent of spreading in the coming months. Our study shows that provided such migration control and other stringent measures continue to be in place, the number of infected cases in various Chinese cities will peak between mid February to early March 2020, with about 0.8%, less than 0.1% and less than 0.01% of the population eventually infected in Wuhan, Hubei Province, and the rest of China, respectively, and no new cases to be expected from mid March. Moreover, for most cities in and outside Hubei Province (except Wuhan), the total number of infected individuals will be less than 4000 and 300, respectively. Since January 24, 2020, very strict migration control has been imposed in

# Thank you!