

Corona Datathon

Team: C.F.R.S.S.

Member:

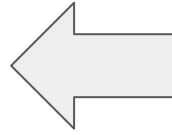
Giona SALA
Antoine SUTER
Philipp RUPP
Leonardo FOSSATI
ManHin CHENG

Overlook

- Growth Models
- Demographics
- Mobility
- A more advanced model
- Linear Regression

Logistic Growth Model

$$Q_t = \frac{a}{1 + e^{b - c(t - t_0)}}$$



$$\frac{df(t)}{dt} \propto f(t)(1 - f(t))$$

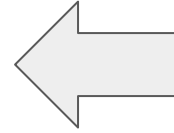
a = max. cases

b = initial condition

c = rate of growth

Gompertz Growth Model

$$Q_t = ae^{-be^{-c(t-t_0)}}$$



$$k \frac{f'(t)}{f(t)} \propto \frac{1}{f(t)}$$

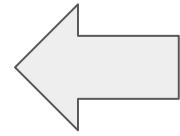
a = max. cases

b = displacement

c = rate of growth

Bertalanffy Growth Model

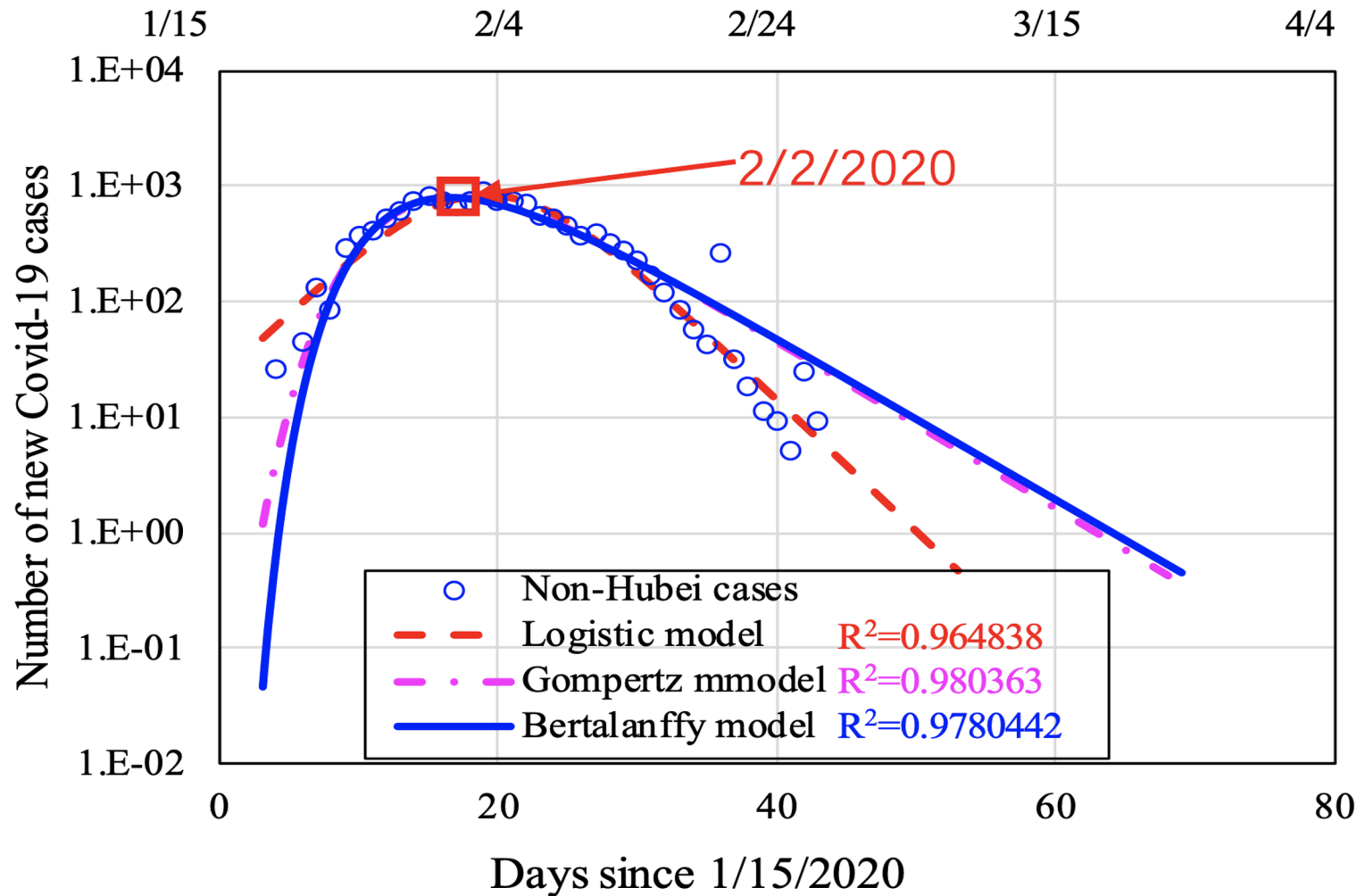
$$Q_t = a(1 - e^{-b(t-t_0)})^c$$


$$\frac{df(t)}{dt} \propto f_{t \rightarrow \infty} - f(t)$$

a = max. cases

b = asymptotic growth

c = rate of growth



Demographic influence

Assumption:

- Countries with older population are more susceptible to the virus and should present a larger mortality rate
 - Mortality rate is different depending on age group

Main Problem:

- Different countries have similar cases distribution by age group

Demographic influence

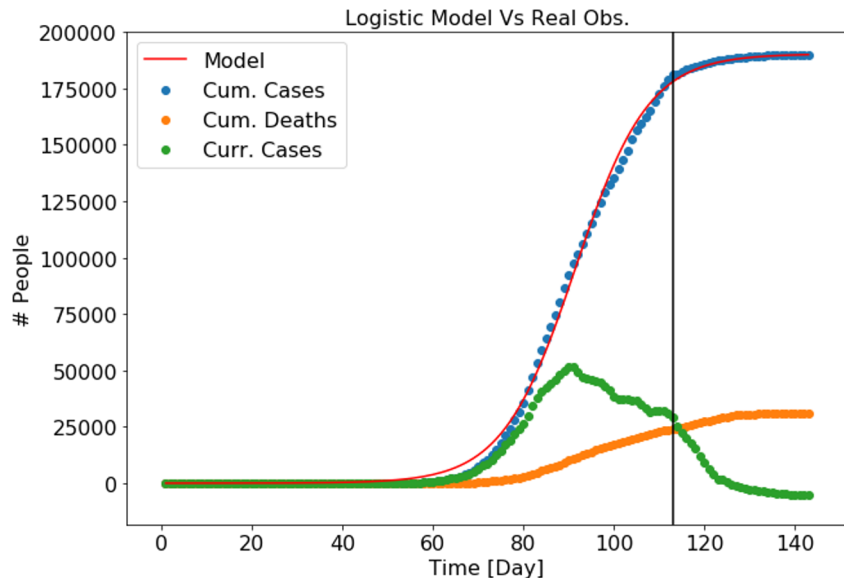
1. Calculate total number of cases [1]:
 - Everyone has the same probability to be infected
 - Mortality rate by age is based on confirmed cases
1. Dynamics of populations [2]:
 - Contact between different age groups
 - Family structures and cultures
 - Lack of data

[1] Lachmann, Alexander & Jagodnik, Kathleen M. & Giorgi, Federico Manuel & A. Ray, Forest. *Correcting under-reported COVID-19 case numbers: estimating the true scale of the pandemic*, 2020. medRxiv. doi: 10.1101/2020.03.14.20036178. <https://www.medrxiv.org/content/medrxiv/early/2020/04/05/2020.03.14.20036178.full.pdf>

[2] Wilder, Bryan & Charpignon, Marie & Killian, Jackson & Ou, Han-Ching & Mate, Aditya & Jabbari, Shahin & Perrault, Andrew & Desai, Angel & Tambe, Milind & Majumder, Maimuna. *The Role of Age Distribution and Family Structure on COVID-19 Dynamics: A Preliminary Modeling Assessment for Hubei and Lombardy*, 2020. SSRN Electronic Journal. doi: 10.2139/ssrn.3564800.

Demographic influence

Simple model which predicts the number of deaths based on confirmed cases and the average of mortality rate works better



Health system capacity

Assumption:

- Mortality rate increases drastically when the health system is at capacity

Problems:

- Lack of data
- Not really relevant until a second wave

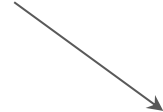
Mobility model

1. Understand and reproduce the spatial spread of an infectious disease epidemic → more reliable predictions: Pivotal Point
1. Already implemented in other studies (spread of previous SARS, Seasonal Influenza-like-illness -ILI, malaria diffusion) with outstanding results

1. Radiation model or mobile phone data model?



Mobility pattern:
Residence place - Workplace



Mobility pattern:
All movements of a single individual
recorded by mobile phone

Radiation model: a simple, but powerful tool

1

goal: reproduce commuter movements

2

based on **stochastic decision** process:

assigns work location to each potential commuter → daily commuting fluxes in a country

3

being **free-parameter:**
absence of empirical data to be fitted or regression analysis

in details : networks is generated by creating a fully connected topology between country's locations, where the weight between nodes i-j:

$$w_{ij}^r = \frac{N_i N_j}{(N_i + P_{ij})(N_i + N_j + P_{ij})} \sum_{j \neq i} w_{ij},$$

- N_i (N_j) population of origin (destination)
- P_{ij} : total population living between i and j
- E: w_{ij} : total number of commuters leaving their home in location i.

It requires: 1. knowlegde of population data (N_i , N_j , P)

2. total number of residents who commute in each administrative unit

Mobility model - Radiation model

Problems:

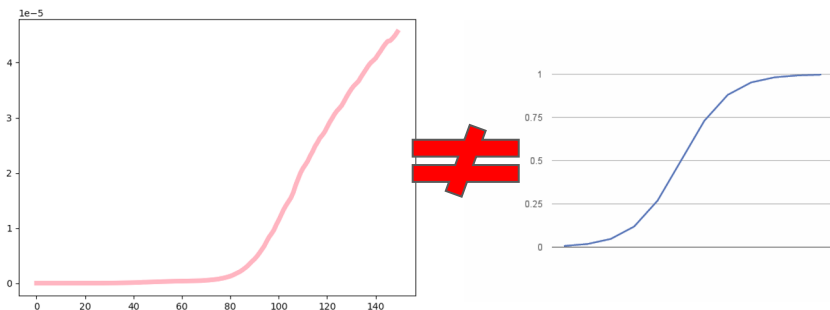
1. Only house-work pattern taken into account → **LIMITATION?**
1. Data for calibration: data of the census → accessibility? Lack of data

But very promising!

Our Complex Model

Aspects:

- Logistic Model
- Takes into isolation/mobility into account
 - With prediction of political decisions: could even model a second wave
- Based on recorded deaths to predict future



```
mod.addPopChange([np.random.uniform(0, 1, 1), np.random.normal(0, 0.003)])
return mod

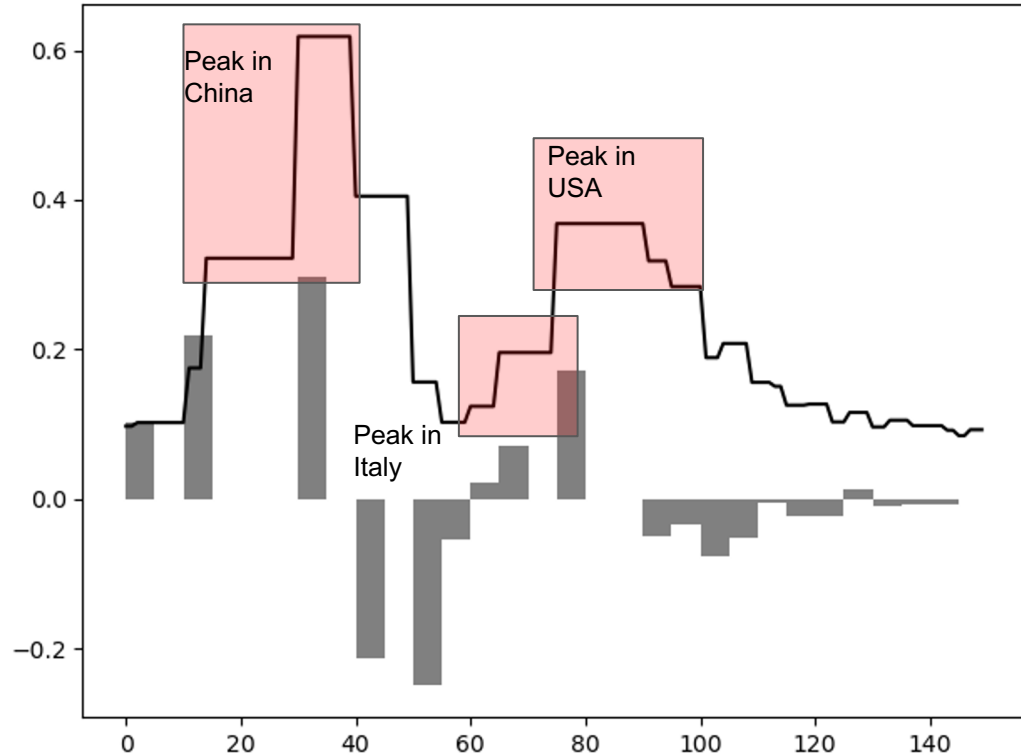
# generates a child that has similar traits(parameters) like parent
# =====
def mutation(self):
    # change parameters of model
    child = copy.deepcopy(self)
    rand = np.random.normal(0, 0.01,
                            len(child.evalParam)) # mu = 0 and sigma = 0.01 so that the change is roughly 4 %
    child.evalParam += np.multiply(rand, child.evalParam)

    attempts = np.random.uniform(0, 1, min(np.random.lognormal(0.5, 0.5), 10))
    for a in attempts:
        if a < 0.5:
            if np.random.uniform(0, 1) < 0.5:
                child.addPopChange([np.random.uniform(0, child.maxIter), np.random.normal(0.0, 0.003)])
            else:
                child.mulPopChange([np.random.uniform(0, len(child.popChangeList)), np.random.normal(1.0, 0.007)])
        else:
            child.addPopChange([np.random.uniform(0, 1), np.random.normal(0.0, 0.003)])

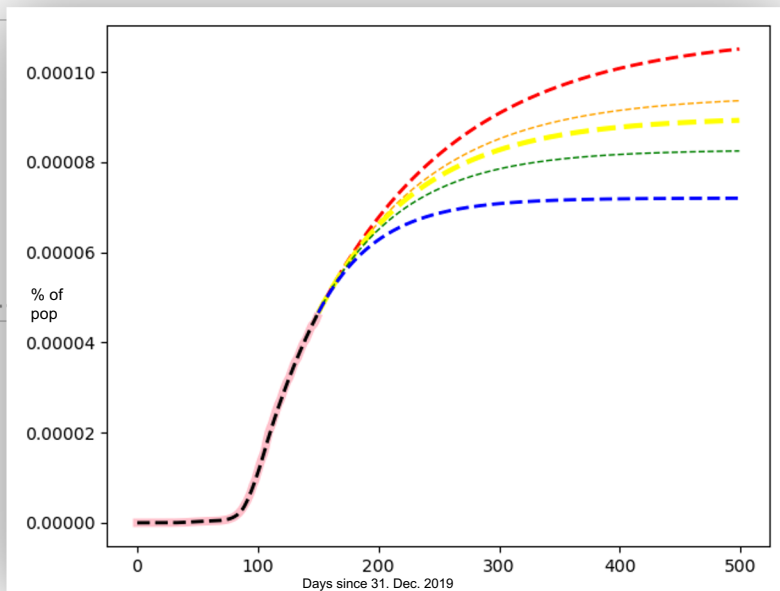
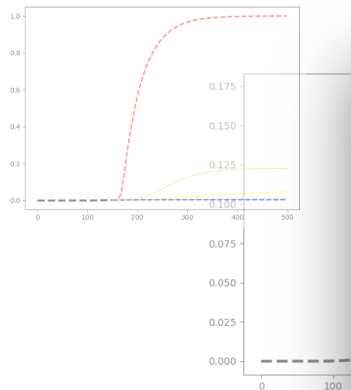
# =====
# change to the list that contains daily changes
# =====
def mulPopChange(self, change):
    day = int(change[0])
    factor = change[1]
    if day > self.maxIter:
        return False
    factor <= 0.0:
        self.delPopChange(day)
        return True
    if self.popChangeTotal * factor >= 1.0:
        factor = 1.0 / self.popChangeTotal
    pChange = self.popChangeTotal * (factor - 1.0)
    if __addPopChange__([day, popChange])
    if popChangeList.sort(key=lambda x: x[0])
    turn True

@PopChange(self, change):
    y = int(change[0])
    pChange = change[1]
    # ...
```

Susceptibility; Summarized in numbers



Predictions are sensitive



Linear Regression

- K.I.S.S: Keep It Simple Stupid
- **Short memory Markov chain** process (5 day)
- $y = mx + C$

Day 1 to Day 5 \rightarrow Day 6

Day 2 to Day 6 \rightarrow Day 7

Day 3 to Day 7 \rightarrow Day 8

Day 4 to Day 8 \rightarrow Day 9

Day 5 to Day 9 \rightarrow Day 10

Linear Regression

$$\text{Confirmed} \geq \text{Recovered} + \text{Death}$$

- Random noise to constrain it

Weekly Leaderboard

Leaderboards are displaying weekly scores (higher is better). A new competition starts every Sunday. More information [here](#). Teams with excellent prediction performances (winners of a 2day competition receive 🏆, winners of a 7day competition receive 🏆, winners of a 30day competition receives 🏆) will be selected for feature stories in our Epidemic Datathon [blog](#).

SCORE	TEAM
1530.7537906116497	C.F.R.S.S.
1500.9833290323852	GNTM_team
1239.7495485032903	stayhome
86.31297300098026	Quarenteam
5.960541171537382	ValenciaSpain

2 Day

Weekly Leaderboard

Leaderboards are displaying weekly scores (higher is better). A new competition starts every Sunday. More information [here](#). Teams with excellent prediction performances (winners of a 2day competition receive 🏆, winners of a 7day competition receive 🏆, winners of a 30day competition receives 🏆) will be selected for feature stories in our Epidemic Datathon [blog](#).

SCORE	TEAM	WINS
1443.5828241598147	C.F.R.S.S.	4x🏆,3x🏆,1x🏆
1403.3621904160848	GNTM_team	2x🏆,3x🏆,1x🏆
1350.3324663589149	stayhome	2x🏆,2x🏆
236.83130580111398	Quarenteam	
4.740510622861562	ValenciaSpain	1x🏆

7 Day

Weekly Leaderboard

Leaderboards are displaying weekly scores (higher is better). A new competition starts every Sunday. More information [here](#). Teams with excellent prediction performances (winners of a 2day competition receive 🏆, winners of a 7day competition receive 🏆, winners of a 30day competition receives 🏆) will be selected for feature stories in our Epidemic Datathon [blog](#).

SCORE	TEAM	WINS
1129.6288197582976	C.F.R.S.S.	4x🏆,3x🏆,1x🏆
1033.8505508083854	GNTM_team	2x🏆,3x🏆,1x🏆
857.9139149953027	stayhome	2x🏆,2x🏆
92.01576321396347	Quarenteam	
12.098753122499492	BuZhunBieXin	
2.2958006003576887	ValenciaSpain	1x🏆

30 Day

Thank you for your attention