

Final Presentation – stayhome

Epidemic Datathon



1. Introduction

2. Data

- a) Preprocessing
- b) Processing
- c) Post-processing

3. Analysis

- a) General remarks
- b) Model
 - Confirmed / Recovered
 - ii. Deaths
- c) Performance

4. Outcome



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General

Global Virus → Pandemic

Time delayed

Geography

Tackled in regions, but mostly in countries

Data

Collected from health institutes

 Datathon based on J.H. University

Importance regional

Adapted decision making

Importance global

Foresee trends, international studies, international solutions



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- a) Collection
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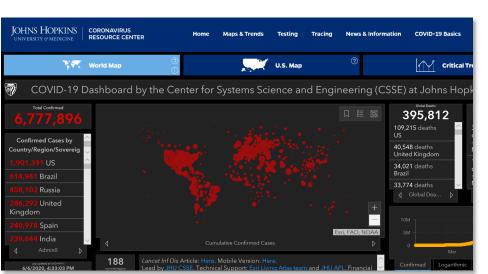
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Data - Collection





https://coronavirus.jhu.edu/map.html, last visited 06.06.2020

Collection

Accessability

Cleaning

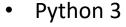
Sorted?

- Data is collected by J.H. University
- Free
- In .csv format
- Assumption of "Correctness"
- Not really necessary
- Except for adaptions in publication format of J.H. University
- Alphabetically sorted
- Country/region wise → 266 different regions in total











Read In

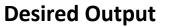
```
class Data:
    def __init__(self, confirmedFile = None, deathsFile = None, recoveredFile = None,
        self.confirmedFile = str(confirmedFile)
        self.deathsFile = str(deathsFile)
        self.recoveredFile = str(recoveredFile)
        #confirmed cases.
        self.df = self.initialize(self.confirmedFile)
        #death cases.
        self.df_deaths = self.initialize(self.deathsFile)
        #recovered cases.
        self.df_recovered = self.initialize(self.recoveredFile)
```

- Pandas Data Frames → similar to «dictionary»
- Conversion to floats





Data - Postprocessing







Read Out

- Specific naming
- Pandas Data Frames → similar to «dictionary»



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a) Future remarks

MSc ETH H. Löbner, MSc ETH S. Strub - Epidemic Datathon - Final Presentation







Not Perfect Data

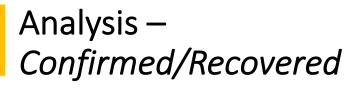
- Insufficient testing
- Time delay
- New method of counting
- Dishonesty
- ...

Decision for model

- Overview: Polyfit
- → Not so good results
- Base Model: Exponential Fit
- Advanced Model for D

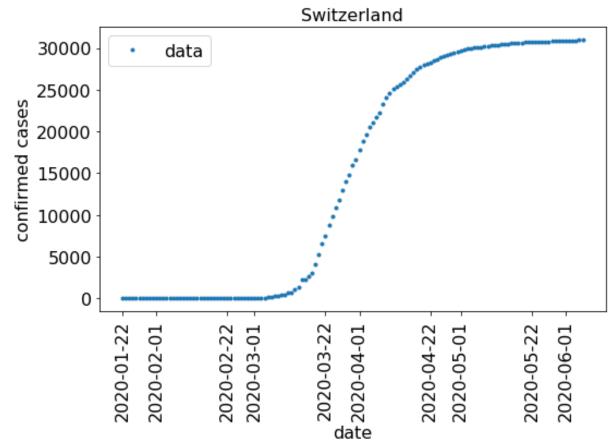
```
class Prediction(Data):
    def __init__(self,confirmedFile, deathsFile, recoveredFile, PredictionLength):
        super(Prediction, self).__init__(confirmedFile, deathsFile, recoveredFile, PredictionLength)
```





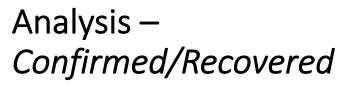






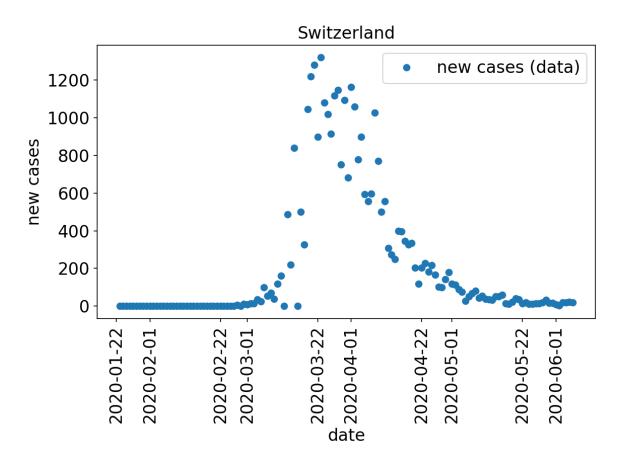
 Compute daily new cases from number of confirmed cases





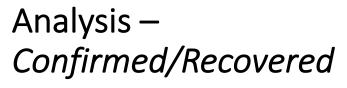






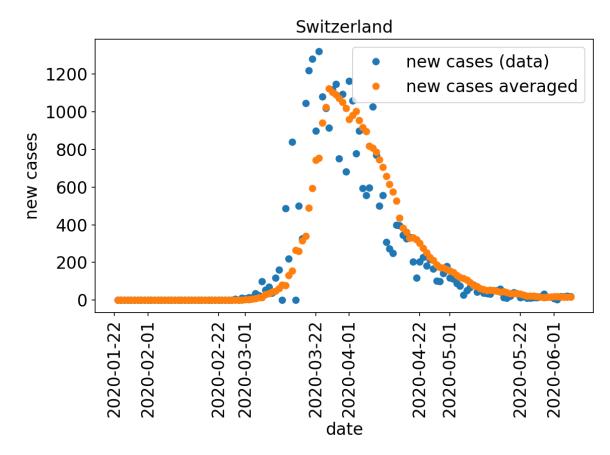
 Compute daily new cases from number of confirmed cases





6 Step plan





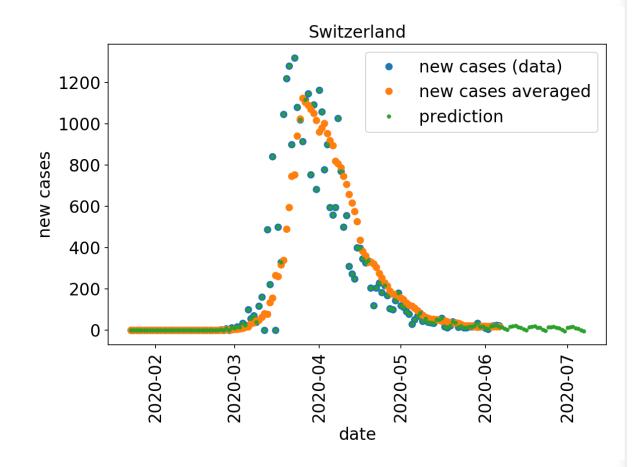
- Compute daily new cases from number of confirmed cases
- 2. Calculate average of new cases
- 3. Predict following 30 days with an exponential fit of the 5 most recent data points averaged





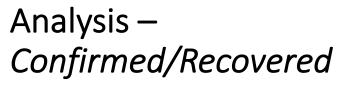






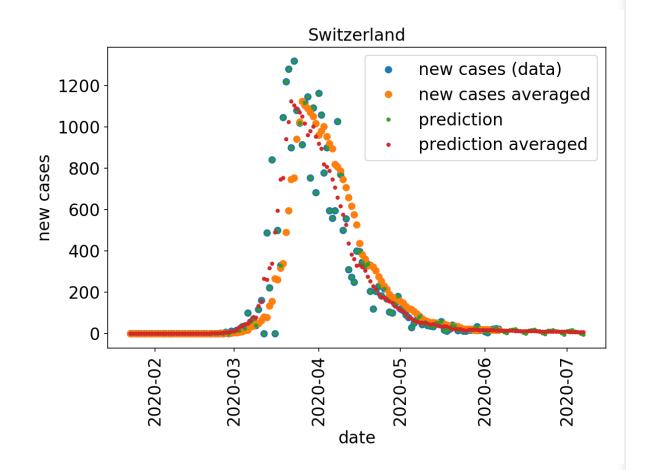
- Compute daily new cases from number of confirmed cases
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- Predict following 30 days with an exponential fit of the 5 most recent data points averaged
- Calculate backwards from the average to daily numbers





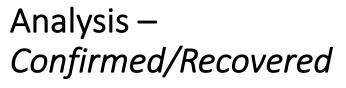
6 Step plan





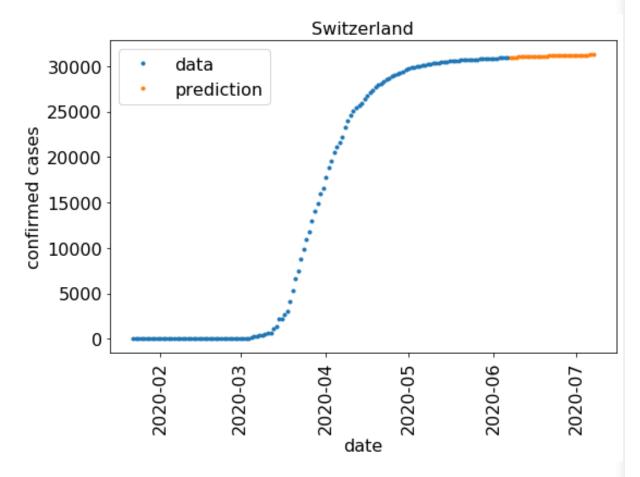
- Compute daily new cases from number of confirmed cases
- 2. Calculate average of new cases
- 3. Predict following 30 days with an exponential fit of the 5 most recent data points averaged
- Calculate backwards from the average to daily numbers
- Compute a center average of the new cases and the predicted new cases











- Compute daily new cases from number of confirmed cases
- 2. Calculate average of new cases
- 3. Predict following 30 days with an exponential fit of the 5 most recent data points averaged
- Calculate backwards from the average to daily numbers
- Compute a center average of the new cases and the predicted new cases
- 6. Sum the predicted new cases center averaged numbers to the confirmed predicted numbers

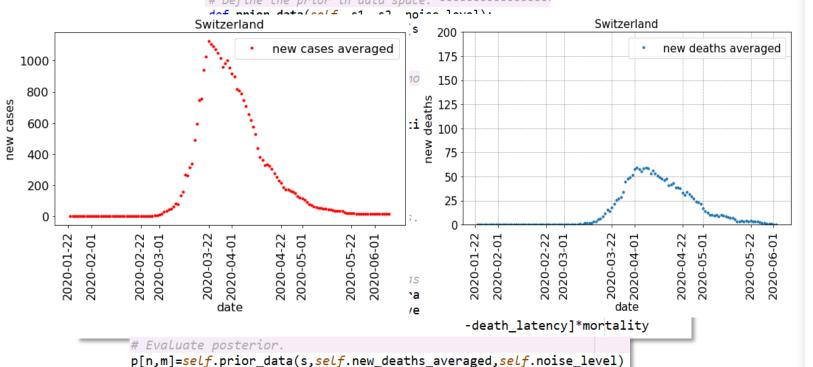


Analysis – *Deaths*



```
# Set the noise level.
self.noise_level=2
# Test time shifts in days.
self.delta=np.arange(1.0,20.0,1)
# Test mortailities.
self.mortalities=np.arange(0.001,0.15,0.001)
```

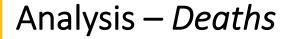
death_latency_index, mortality_index = np.where(p == np.amax(p))



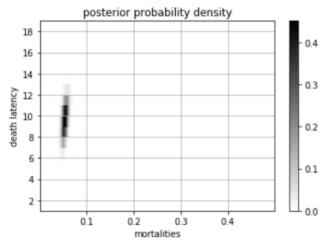
self.death latency, self.mortality = int(self.delta[death latency index]), self.mortalities[mortality index]

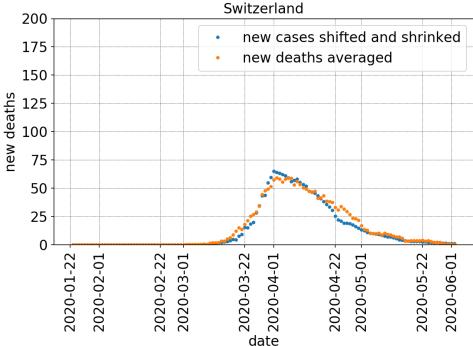
- Assumption: Connection between
 C D
- Grid search (inverse theory tool)
 - Optimize death latency and mortality
- Time shift and shrink averaged new C cases in order to match the averaged new D
- Advantage: New development in C
 will translate in the future to D











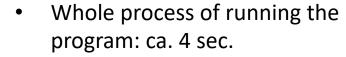
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Analysis – *Performance*

Time



Data read in: 0.5 sec
C/R: 1.0 sec
D: 1.5 sec

• Data output: 0.5 sec



Accuracy

- Good Accuracy for short term C
- Good long term accuracy for D and R
- Topping the leaderboad several times

Possible optimization

- Use gathered data to optimize parameters such as average length and fit length
- Test if our death prediction method is better than an exponential fit



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Outcome – Future Remarks



Quality of Data

Poor Data Quality

- 1. Different counting methods
- 2. Dishonesty
- 3. No complete knowledge
- 4.

Leading to

- → No modelling of disease
- → Modelling of datapoints

Thank you for your attention. Questions?

For future datathons

→ Maybe problems with *Clean* and *Correct* Data