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AN INITIATIVE OF
THE NETHERLANDS
RED CROSS

FORECASTING COVID-19

(10)(2e)

20-mar-2020



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- Goal: forecast the number of cases of covid-19
- Subgoals:
 - when will the peak of infection be?
 - How many people will be hospitalized?
- Possible approaches that we tried:
 1. **Fits** to reported number of cases with **exponential** and **logistic** functions:
 - Pros: very few free parameters, low model variance
 - Cons: valid only for short time scales (≤ 1 week)
 2. **Compartmental models** (SIR, SEIR...)
 - Pros: more realistic picture on long time scales (> 1 week)
 - Pros: possible to estimate effects of interventions
 - Cons: many unknown parameters to be estimated
- Methodology reviewed by (10)(2e), epidemiologist at Wageningen University

N.B. Predictions are **ball park estimates** and the following is **work-in-progress**

Fits with exponential and logistic (1/2)

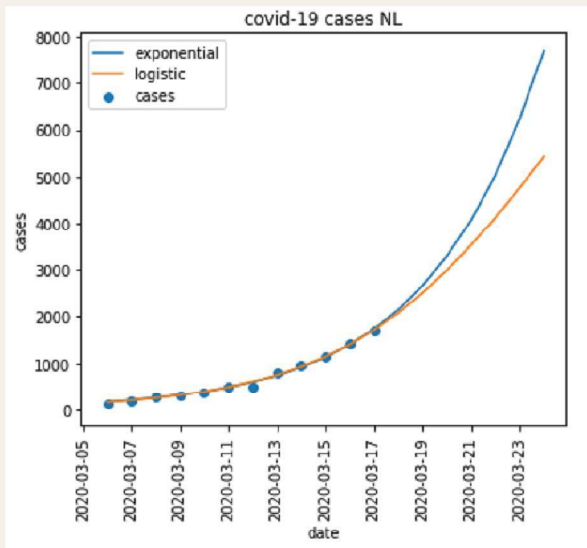


exponential

$$f(x) = ab^x$$

logistic

$$f(x) = \frac{L}{1 + e^{-k(x-x_0)}}$$



- Exponential and logistic function are biologically **motivated** descriptions of population dynamics (e.g. see [Nature education](#))
- They both describe data quite well, both at national (below) and local level (next slide)
- Predictions diverge with time: taking the **average** of exponential and logistic is probably a safe bet
- In order to account for smooth changes in growth as a function of time (e.g. effects of government interventions), **better to fit over the last ~10 days** (enough for 2 free parameters)

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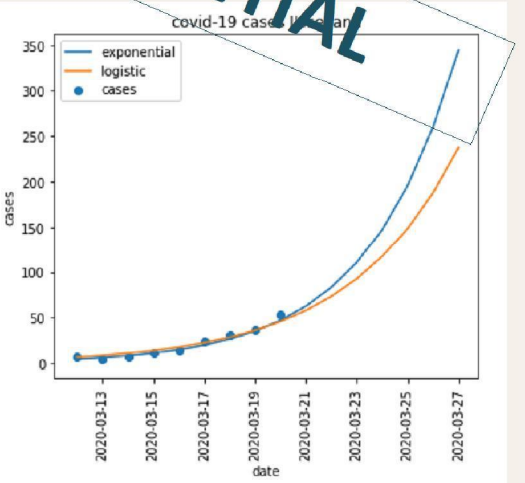
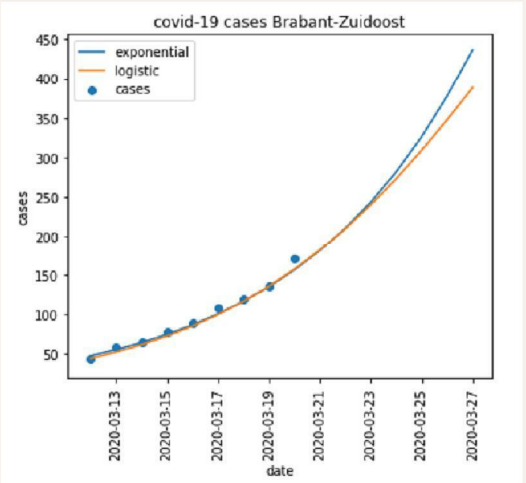
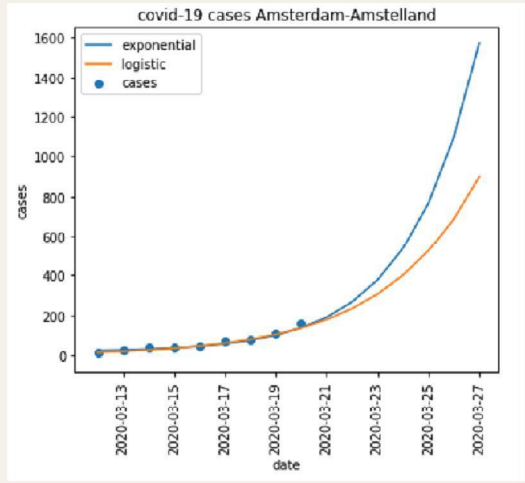
Fits with exponential and logistic (2/2)



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Good fit also at a local level (e.g. regions in the Netherlands)

Be aware of statistics if not much data is available at a local level: small numbers → big uncertainties



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- Introduction:
*“**Compartmental models** are a technique used to simplify the mathematical modeling of infectious disease. The population is divided into compartments, with the assumption that every individual in the same compartment has the same characteristics.” (source: [Wikipedia](#))*
- For covid-19, the most used ones are variations of a **SEIR model**:
 - Population is divided into
 - susceptible (S): can get the disease
 - exposed (E): got the disease, but still incubating
 - mild infections (I1): mild symptoms
 - severe infections (I1): severe symptoms → hospitalization required
 - critical infections (I1): critical symptoms → intensive care (IC) required
 - recovered (R)
 - dead (D)
 - Dynamics are governed by a series of parameters (transmission rate, death rate, etc.)
 - See more details on [Wikipedia](#) or in [this presentation](#)

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- **Many unknown parameters**, here's a collection of best estimates:
 1. basic reproduction number: 2.3 [\[source\]](#)
 2. basic reproduction number with full lock-down: 1 [\[source\]](#)
 3. incubation period: 5 days [\[source\]](#)
 4. duration of mild infections: 7 days [\[missing source\]](#)
 5. % of severe infections: 14% [\[source\]](#)
 6. % of critical infections: 5% [\[source\]](#)
 7. compartment size: number of people in country/region
- Notes:
 - It's in many cases a '**worst-case-scenario**' because it assumes random mixing (everyone has an equal chance of being in contact with each other). From a virus point of view, that is the most efficient way of spreading around.
 - **Time of the peak** will strongly depend on the compartment size: make sure you have good **population estimates**
 - Simplest implementations of SEIR **do not include age** compartments/classes: important for covid-19, work in progress

Compartemental models: preliminary results

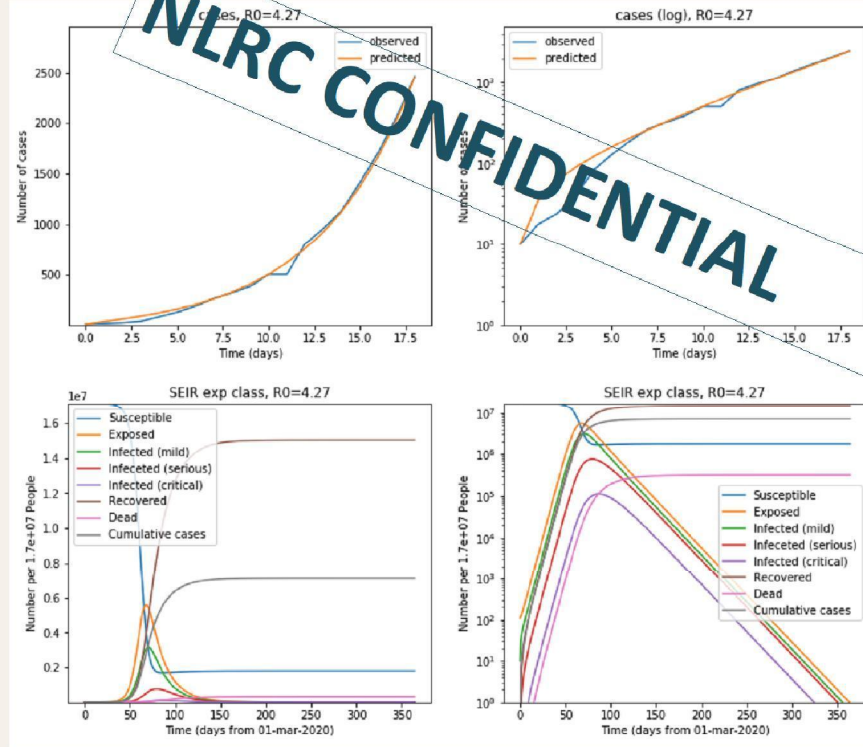


Preliminary results:

- **Decent description** of early-time evolution of cumulative number of cases in NL
- Evidence for **higher-than-expected transmission rate** (effective $R_0 \sim 4$)
- **Assuming no interventions**, for a population of about 17M **peak** is expected at **day ~60**, with about 3-5% of people hospitalized

Note:

- Full lock-down in China achieved $R_0 \sim 1$, unclear what other interventions will do



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- Python implementation of exponential and logistic fits:
 - <https://towardsdatascience.com/covid-19-infection-in-italy-mathematical-models-and-predictions-7784b4d7dd8d>
- Web-based tools for SEIR models:
 - Harvard model: <https://alhill.shinyapps.io/COVID19seir/>
 - ExploSYS model: <http://covidsim.eu/>
- Python implementation of a SEIR model:
 - https://colab.research.google.com/github/alsnhll/SEIR_COVID19/blob/master/SEIR_COVID19.ipynb
- **ALWAYS CROSS-CHECK WHAT YOU DO WITH AN EPIDEMIOLOGIST**