## SIRSi-Vaccine Dynamical Model for the Covid-19 Pandemic repository

## **URL**

https://github.com/dferruzzo/Covid-19-version2

## Description

This repository includes databases of confirmed Covid-19 cases and data on social distancing obtained from the official website of SEADE. Additionally, the repository houses Python programs that process, prepare, and analyze the raw data using least squares fit, model prediction validation, and figure generation.

In this revision, I aimed to provide a more concise and clear overview of what the repository contains and what the Python programs do. I also tried to use more parallel structure to make the text easier to read.

The files that are part of the program include the functions:

- show\_data , presents in a user-friendly way the data of confirmed cases and isolation rate used for processing.
- loaddata, loads the data corresponding to the chosen time window.
- fit isol, produces the least squares fit for the isolation index.
- fitting, produces the least squares adjustment for data from confirmed cases of Covid-19.
- validation, performs a validation of the predictions generated by the model with adjusted parameters taking data in a later time window than the data used for the adjustment.
- save\_all, saves tuning parameters and other statistical data in csv files for later use.
- load all, loads the information previously saved in csv format for use.
- produce\_figs\_param, generates most of the figures used in the article. Not all figures produced are used in the article.
- simulations, produces simulations and figures and allows changing parameters such as the vaccination rate  $\omega$  and the isolation index  $\theta$ . It is used to simulate different scenarios as proposed in the article.
- bif\_map , produces a transcritical bifurcation map in parameter spaces  $(\omega,\theta)$  that shows the trade-off of stability between disease-free and endemic equilibria, as well as the contours of peak peaks of infected associates.

## **Functionality**

This notebook can be run directly in a Codespace here on Github.

Run this notebook to:

- 1. Load the data from the Covid-19 reported confirmed cases,
- 2. Load the Isolation index data,
- 3. Produce parameter adjustment for both the isolation data and the model paramters,
- 4. Save the data produced in files for distribution and reproducibility,
- 5. Produce the plots,
- 6. For testing different scenarios.

```
In [1]: # Load the libraries
        from loaddata import *
        from fit isol import *
        from fitting import *
        from validation import *
        from save all import *
        from load all import *
        from produce figs param import *
        from omega gamma import *
        from simulations import *
        from show data import *
        from bif map import *
        import pandas as pd
In [2]: # Show the raw data used
        #show data()
In [3]: # Load the data for fitting and validation
        data for fit, data for val = loaddata()
        Loading data...
        Data loaded...OK!
In []: # Run the fitting for the isolation index
        data fit isol = fit isol(data for fit, order=1, save figs=False)
In [ ]: # Run the fit of the confirmed cases to get the model parameters
        data fit dyn = fitting(data fit isol,\
                               data for fit,\
                               save figs=False)
        # Validates the model
        validation(data fit isol,\
                   data fit dyn,\
                   data for fit,\
                   data for val,\
                   save figs=False)
```

In [ ]: # If output of the previous cell is satisfactory then saved all parameters d

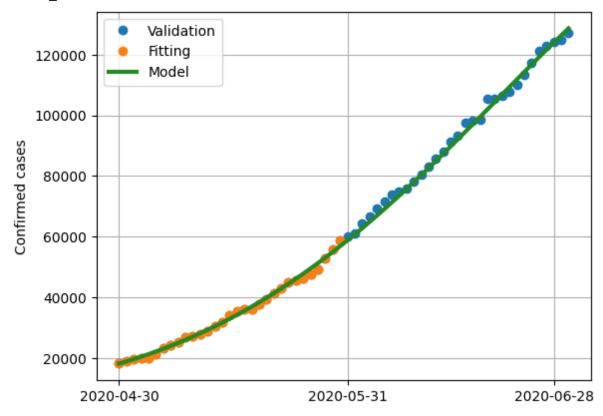
```
In [3]: # Load the data ans parameters we use in the manuscript
    csv1 = 'data_iso-2023-03-06-19-38-55.csv'
    csv2 = 'data_dyn-2023-03-06-19-38-55.csv'
    data_fit_isol_saved, data_fit_dyn_saved = load_all(csv1, csv2)
```

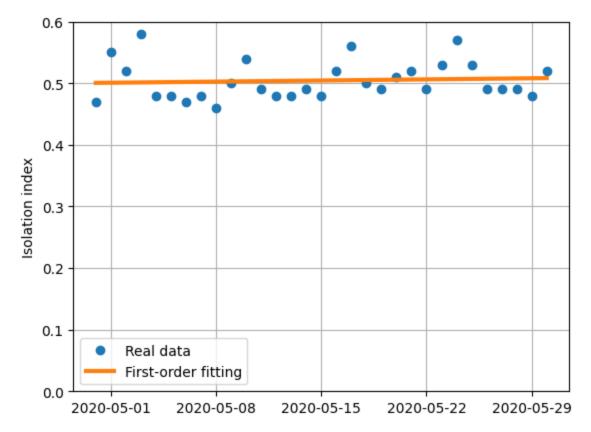
Loading data from file: data\_iso-2023-03-06-19-38-55.csv

Loading data from file: data dyn-2023-03-06-19-38-55.csv

Data loaded successfully...OK!

```
N = 62
tot_pop = 11869660
theta t = 0.50064516 + 0.00025806·x
```

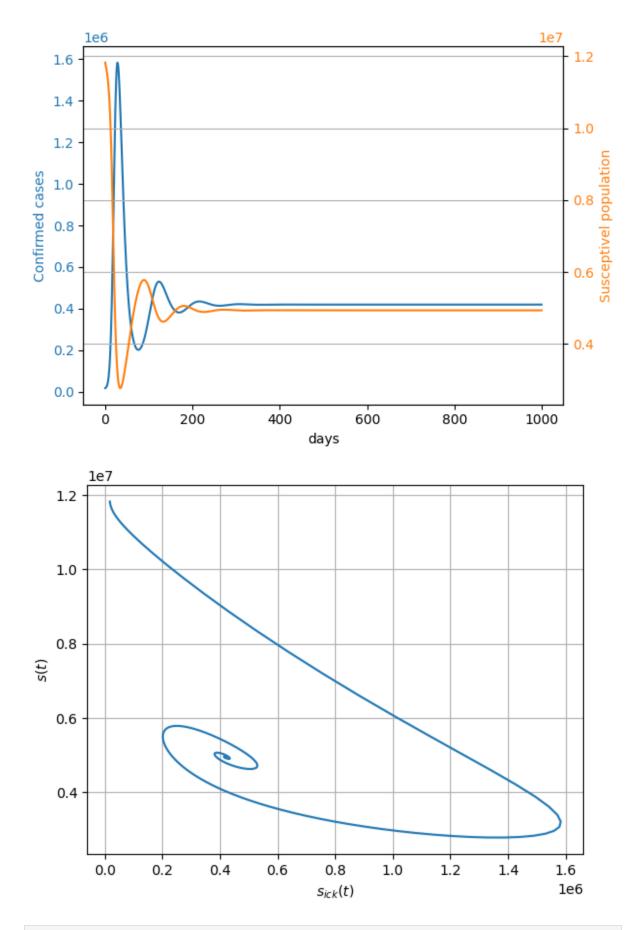




RMSE validation = 0.00010612109338681192

```
omega = -m*theta+b
theta critical = 0.5839603654983293
omega min = 0.028074066113463084
m = 0.048075293756461975
b = 0.028074066113463084
theta_test = 0.6
omega(theta)= -0.000771110140414099
```

<Figure size 640x480 with 0 Axes>



data\_fit\_dyn\_saved,\
save\_figs=False)

