



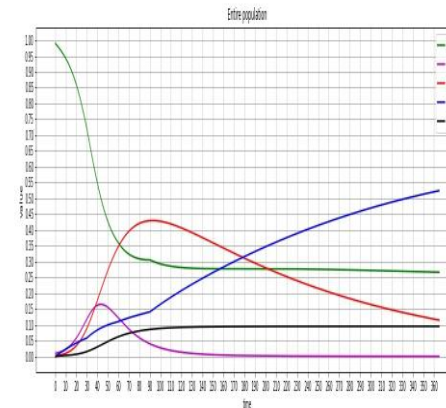
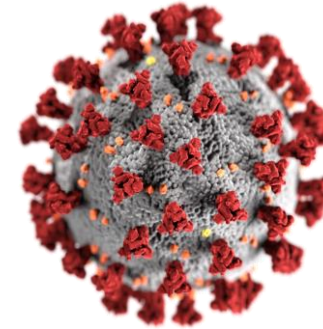
A multi-age structured SIRVSD model

Implementation of a SIRVSD model
with different age groups and
heterogeneous contacts in Python

PAOLO MURGIA

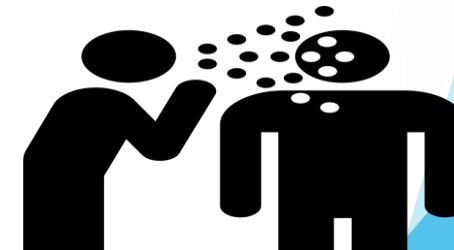
Introduction to the problem

- Covid-19 has focused on forecasting models to be able to monitor and control the number of infections and deaths within a country
- Need to evaluate different vaccination strategies to defeat the infection
- Need to predict the effect of vaccinations to organize a plan to ease restrictions, especially for the country's economy
- Compartmental models are often applied to the mathematical modelling of infectious disease using ODEs
- Population partitioned into different age groups, considering the impact of heterogeneity in susceptibility, mortality and infectivity within the population on the disease transmission



Model Assumptions

- $S+I+R+V+D = 1$
- Some coefficients are the same for all age group (only transmission, vaccination and mortality coefficients are different)
- We don't consider vital dynamics (reproduction, 'natural' death, migration)
- Heterogeneous contacts and symmetric contact matrix
- Just one injection of the vaccine is considered
- Both recovery and vaccination immunity are not forever, but they ensure 100% protection from infection
- Contacts between individuals are random, the number of infections is proportional to both I and S
- Horizontal transmission



Model and Parameters

$$\begin{cases} \frac{dS_j}{dt} = \phi R_j - \eta_j(t)S_j + \rho V_j - S_j \sum_{k=1}^M \beta_{j,k} I_k \\ \frac{dI_j}{dt} = -\gamma I_j - \mu_j I_j + S_j \sum_{k=1}^M \beta_{j,k} I_k \\ \frac{dR_j}{dt} = \gamma I_j - \phi R_j \\ \frac{dV_j}{dt} = \eta_j(t)S_j - \rho V_j \\ \frac{dD_j}{dt} = \mu_j I_j \end{cases} \quad \forall j \in (0, M]$$

with $M = 4$ age groups:

- Children (0-9 years)
- Teenagers (10-19 years)
- Adults (20-69 years)
- Senior Citizens (70+ years)

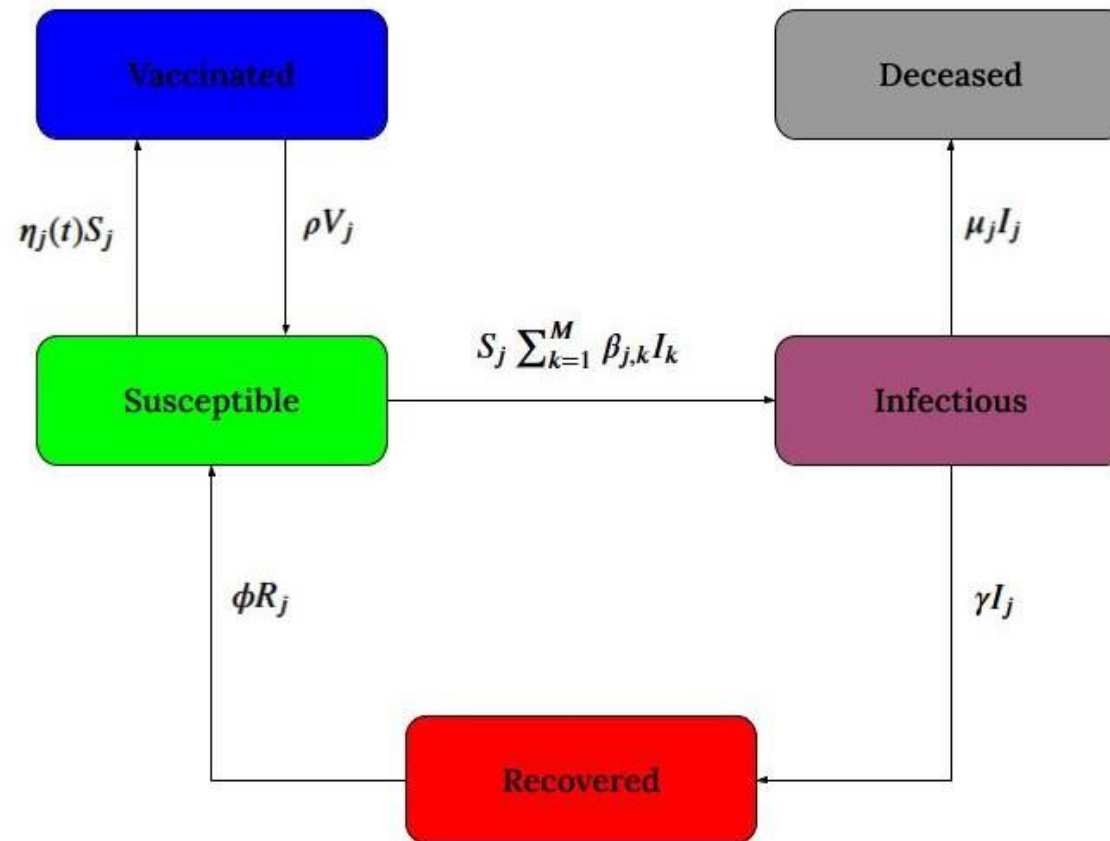
- ϕ is the *transfer coefficient* for loss of immunity from Recovered
- ρ is the *transfer coefficient* for loss of immunity from Vaccinated
- $\beta_{j,k}$ is the *infection coefficient*. We define also the entire *contact matrix*:
- γ is the *recovery coefficient* of each infected subject
- μ_j is the *mortality coefficient*, different for each age group
- $\eta_j(t)$ is a time-dependent *vaccination coefficient*, defined as follows:

$$\beta = \begin{bmatrix} \beta_{1,1} & \cdots & \beta_{1,M} \\ \vdots & \ddots & \vdots \\ \beta_{M,1} & \cdots & \beta_{M,M} \end{bmatrix}$$

$$\eta_j(t) = \begin{cases} 0 & \text{if } t < t_{vacc_j} \\ \eta_j & \text{otherwise} \end{cases}$$

where t_{vacc_j} defines the starting day of the vaccination period

Transition schema



Qualitative Analysis

- Case Study: COVID-19 pandemic with different vaccination strategies
 - Initial Configuration

Some clarifications

- fifteen days for recovering
- six months of immunity for recovered
- nine months of immunity for vaccinated
- mortality coefficient computed from [ISS covid-19 report](#)

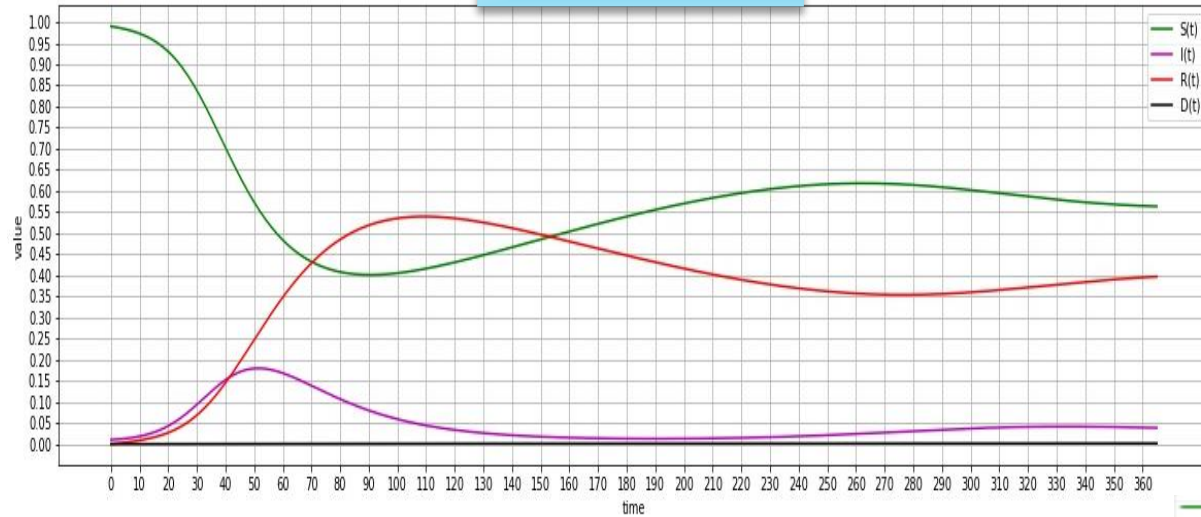
	Children	Teenagers	Adults	Senior
$S(0)$	99%	99%	99%	99%
$I(0)$	1%	1%	1%	1%
$R(0)$	0%	0%	0%	0%
$V(0)$	0%	0%	0%	0%
$D(0)$	0%	0%	0%	0%
γ	1/15	1/15	1/15	1/15
μ	0.00009	0.00005	0.00688	0.15987
ϕ	1/180	1/180	1/180	1/180
ρ	1/270	1/270	1/270	1/270

β	Children	Teenagers	Adults	Senior
Children	0.05	0.01	0.04	0.008
Teenagers	0.01	0.09	0.08	0.008
Adults	0.04	0.08	0.1	0.02
Senior	0.008	0.008	0.02	0.03

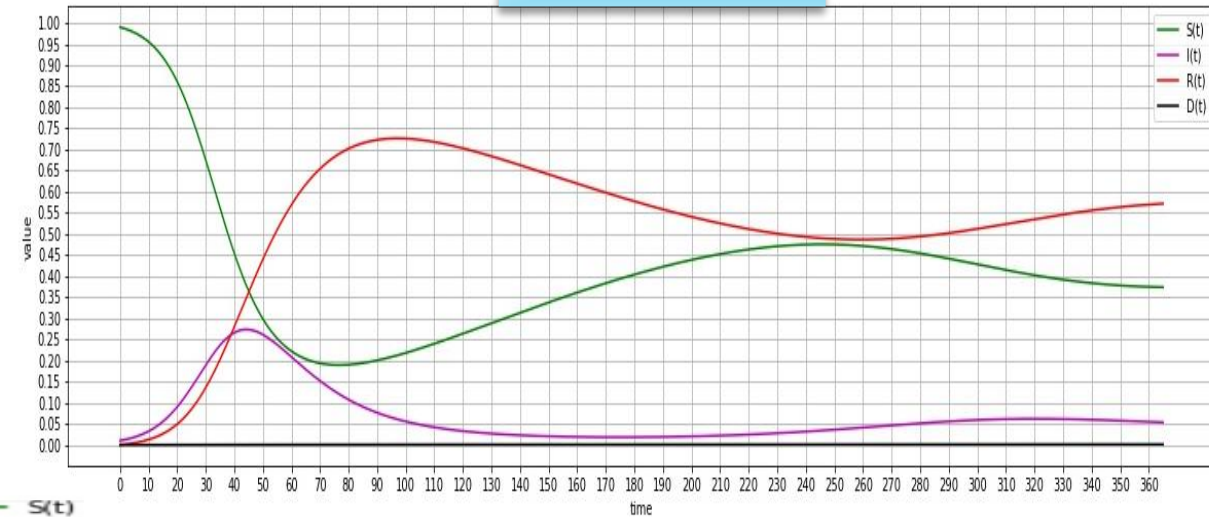
No Vaccination

Fine-grained analysis

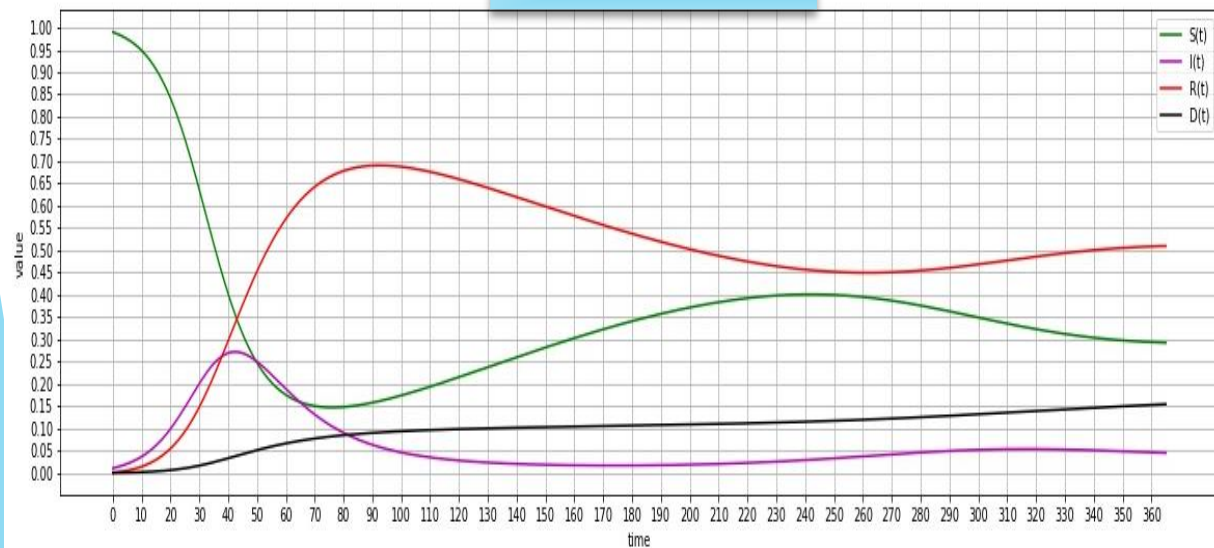
Children



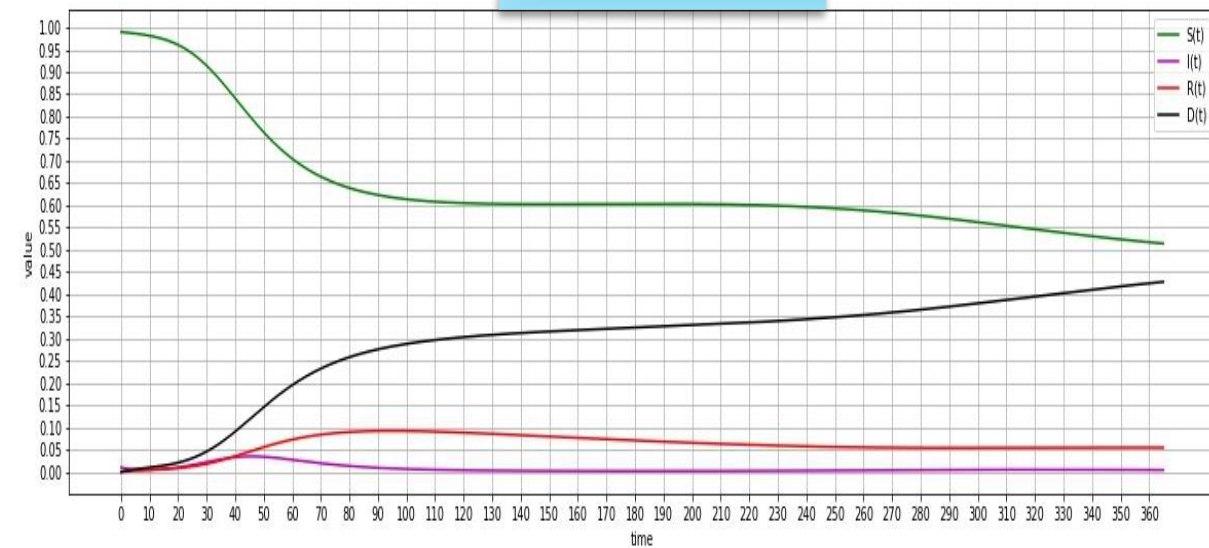
Teenagers



Adults

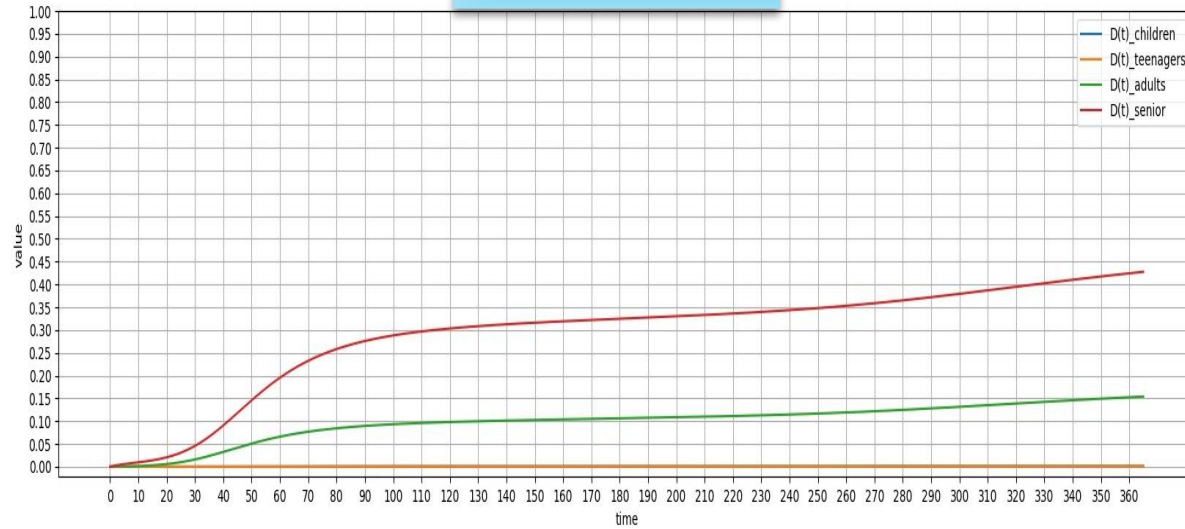


Senior

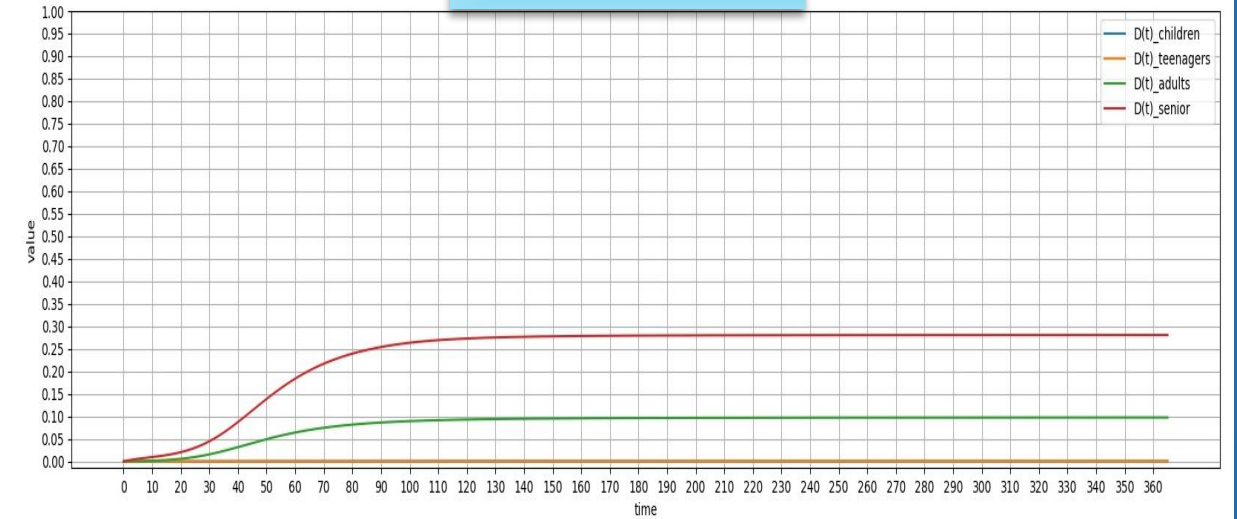


Mortality comparison for each vaccination strategy

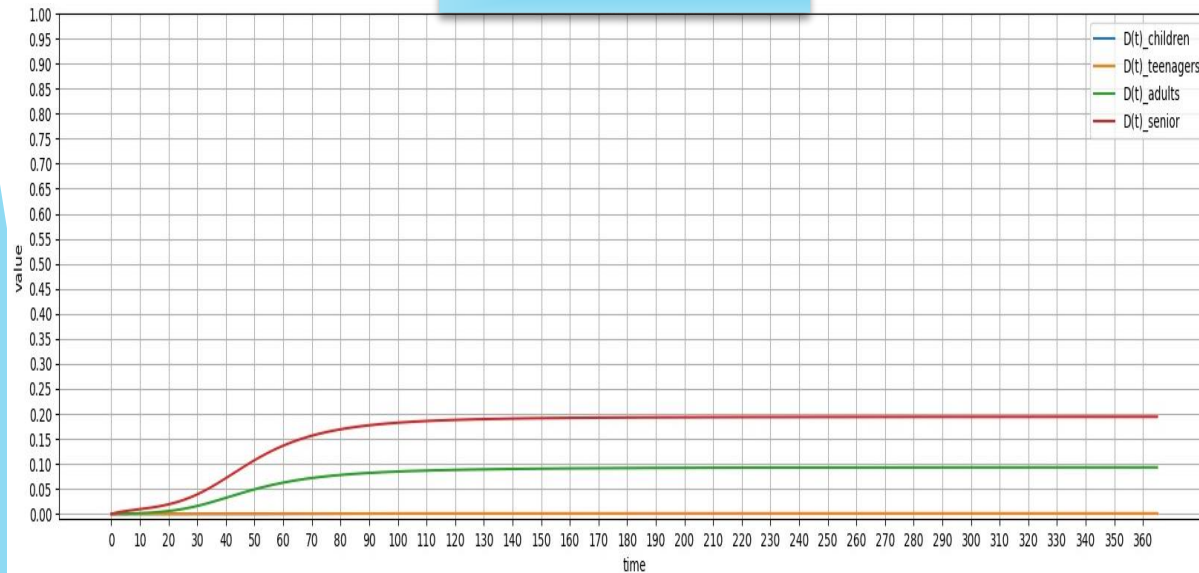
No vaccination



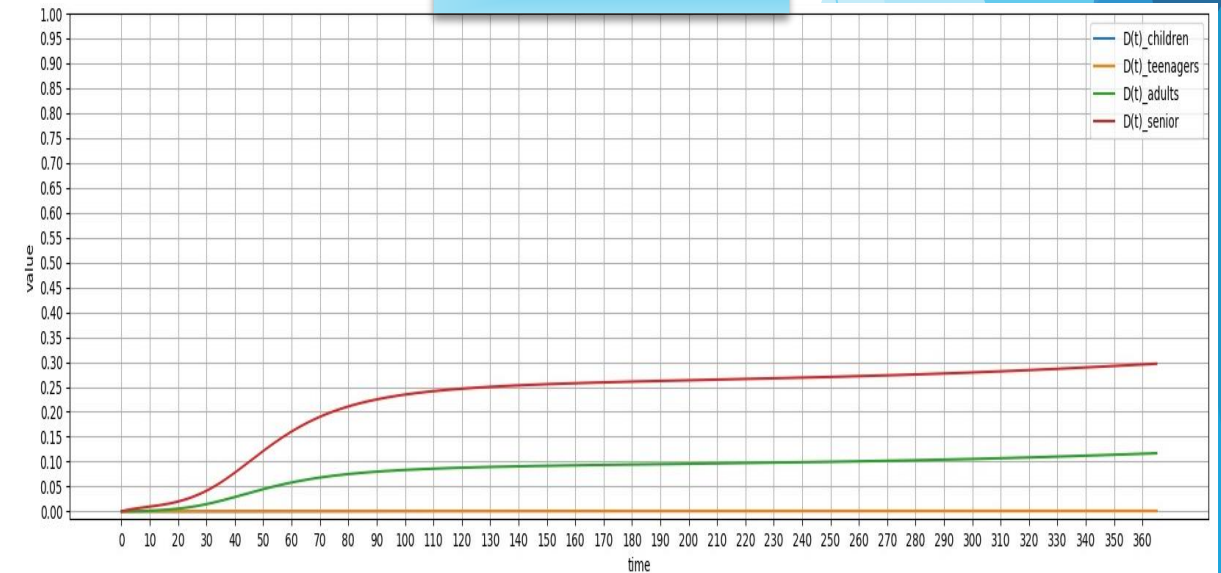
Ascending order



Descending order

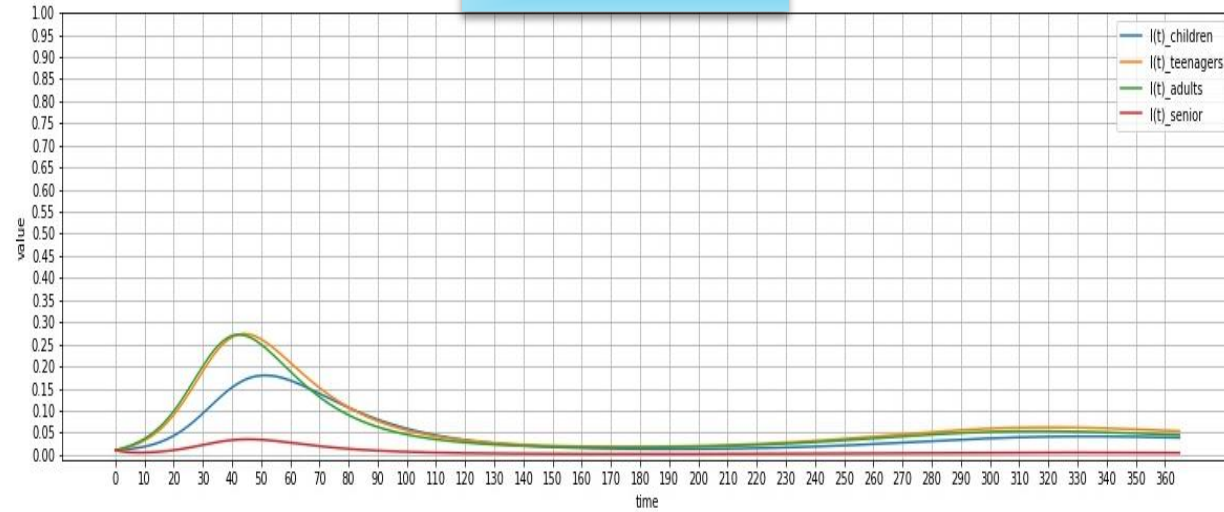


Same Time

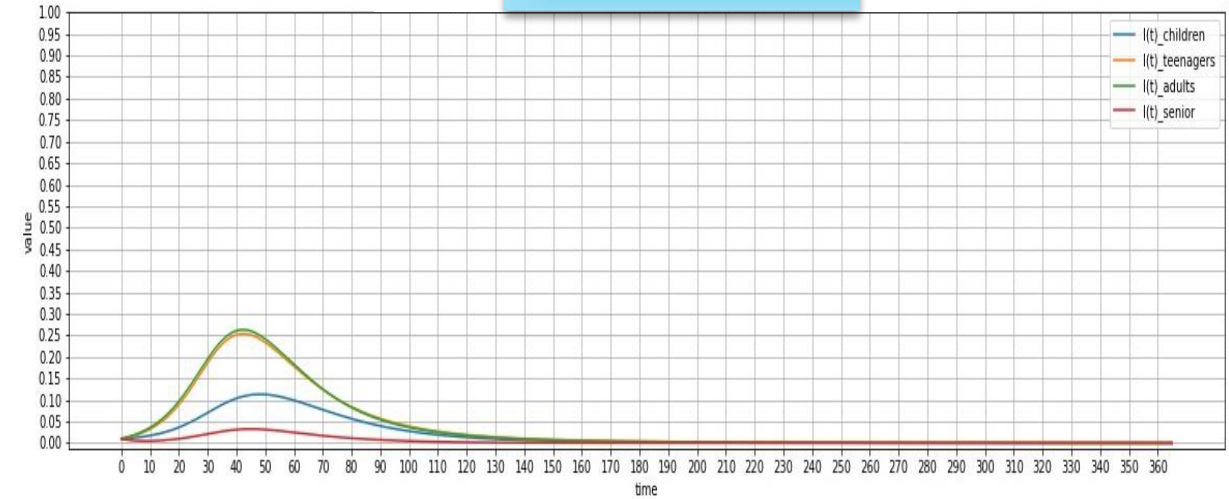


Infections comparison for each vaccination strategy

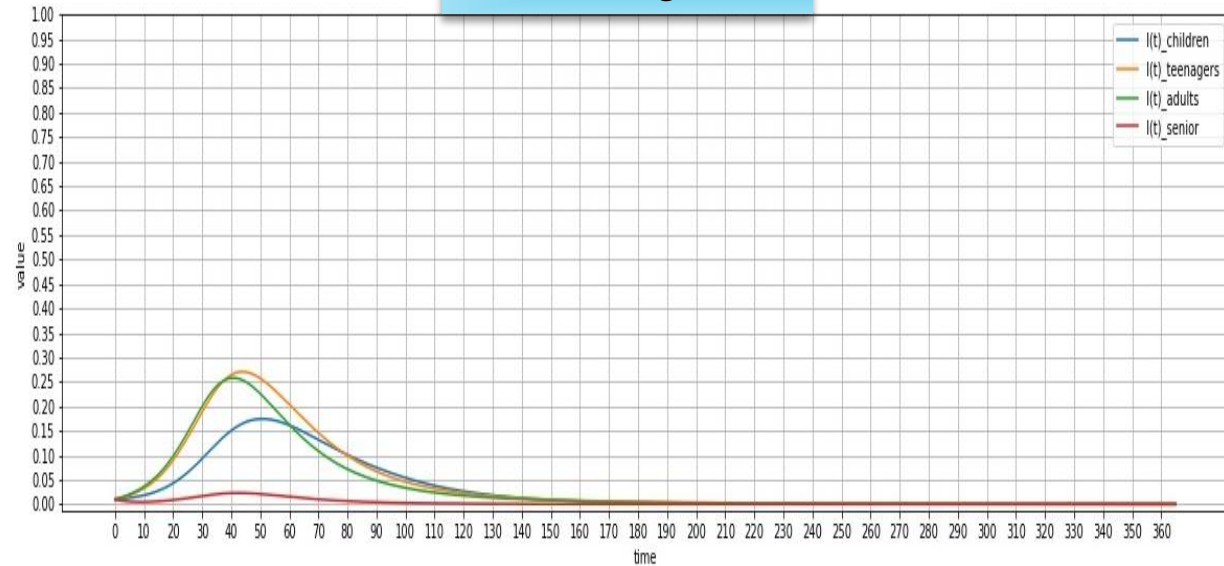
No vaccination



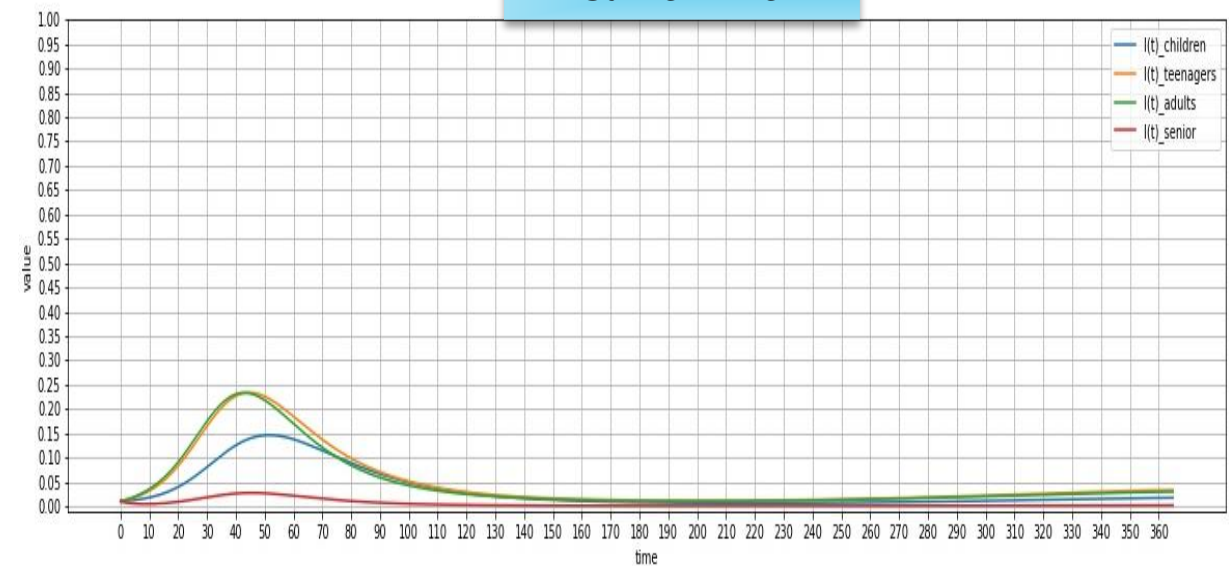
Ascending order



Descending order



Same Time



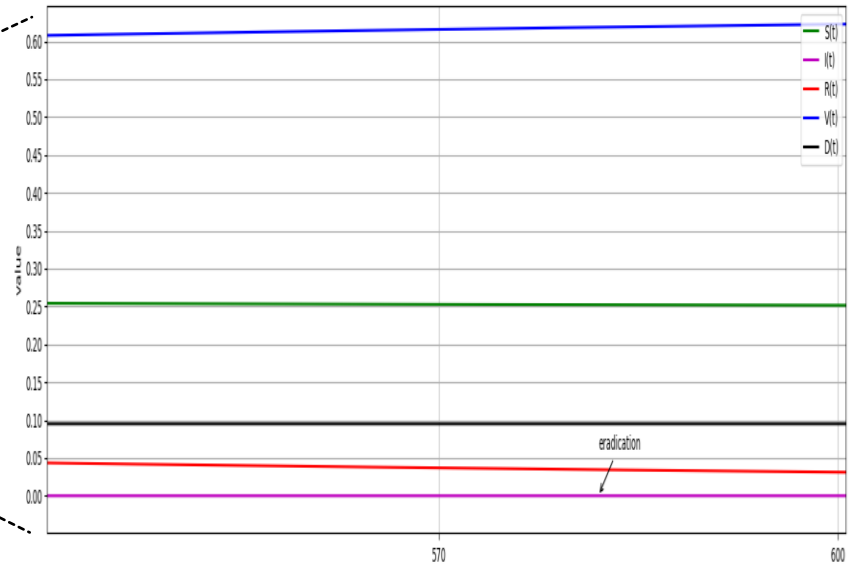
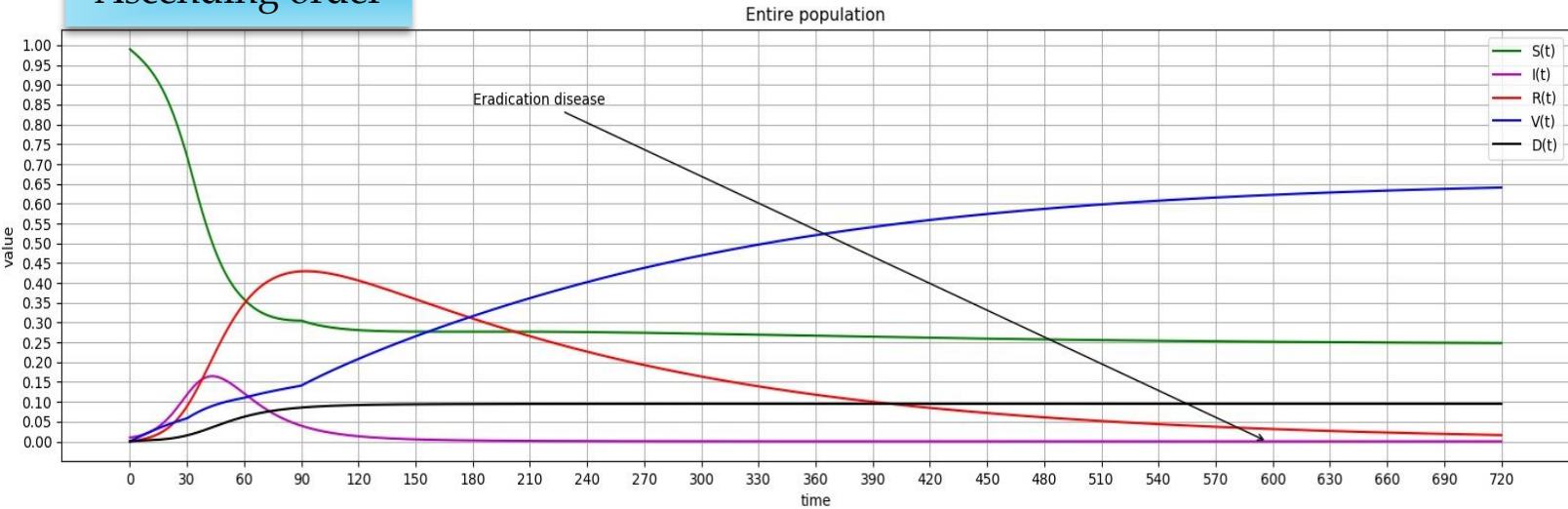
Eradication disease

Coarse-grained analysis

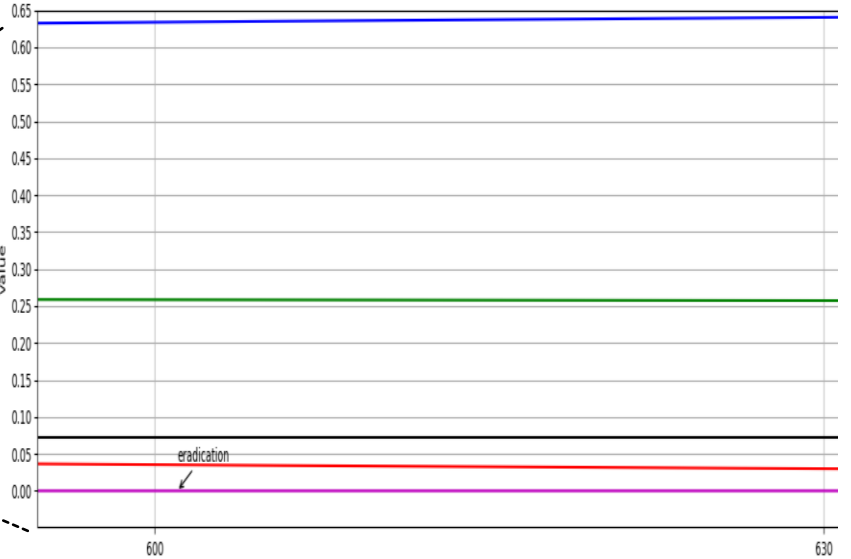
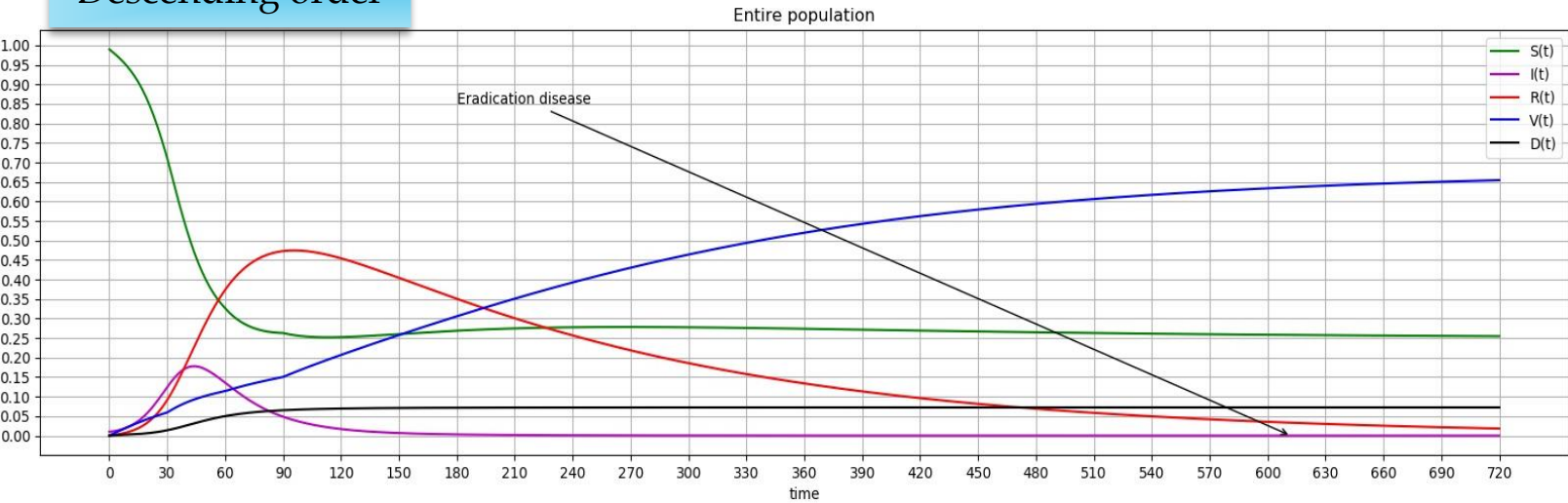
Ascending	Children	Teenagers	Adults	Senior
η	0.01	0.01	0.01	0.01
VACC-DAY	0	30	60	90

Descending equal only with reverse order for VACC_DAY

Ascending order



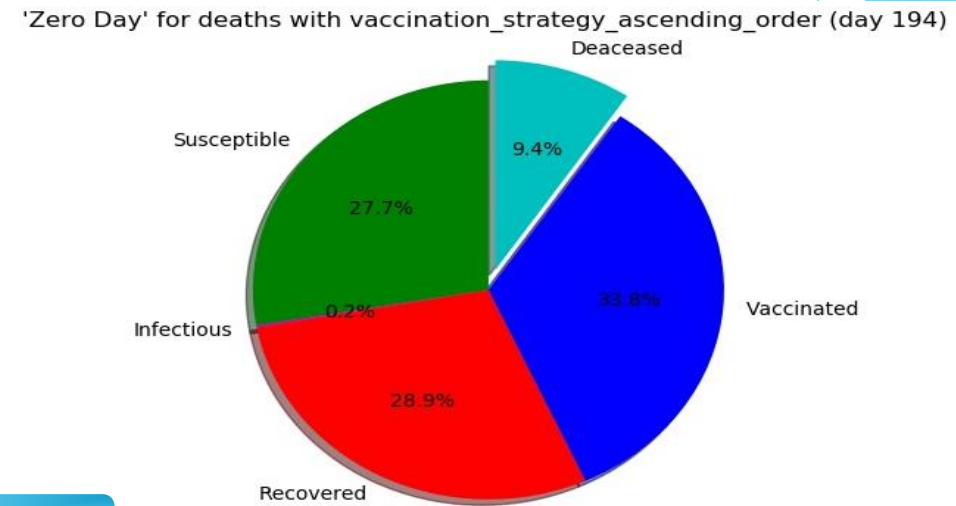
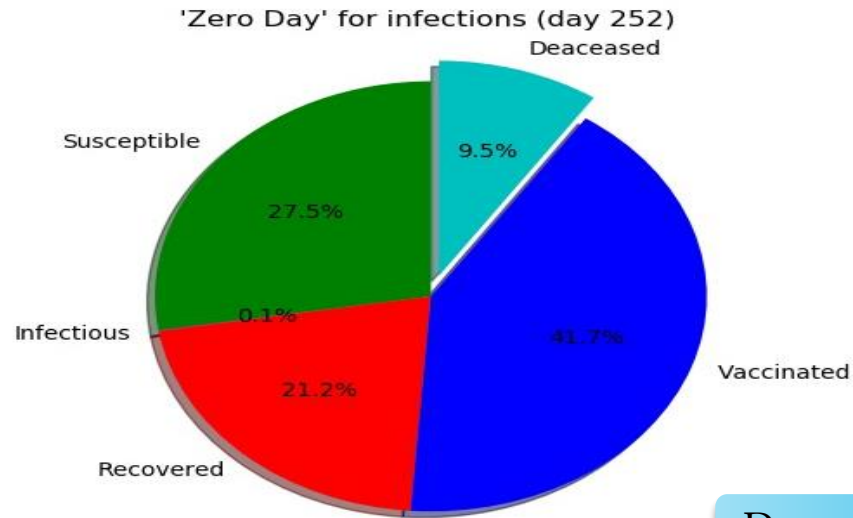
Descending order



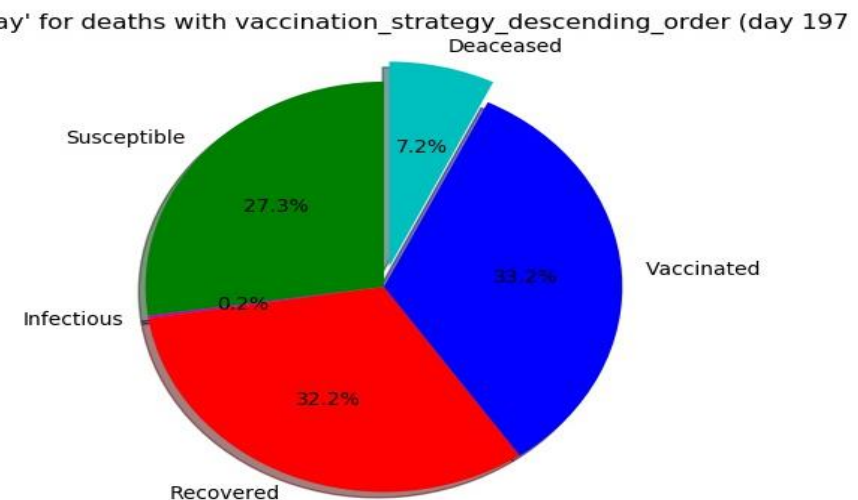
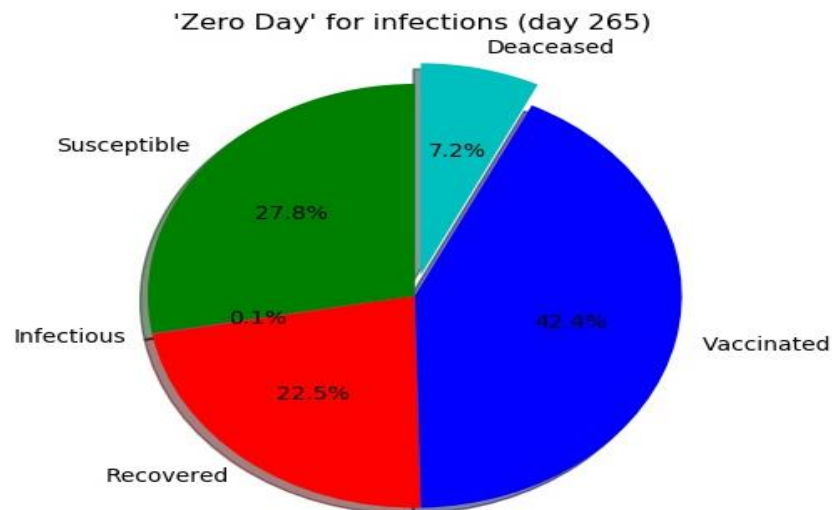
'Zero Days'

i.e. first day without new infections (or deaths)

Ascending order



Descending order

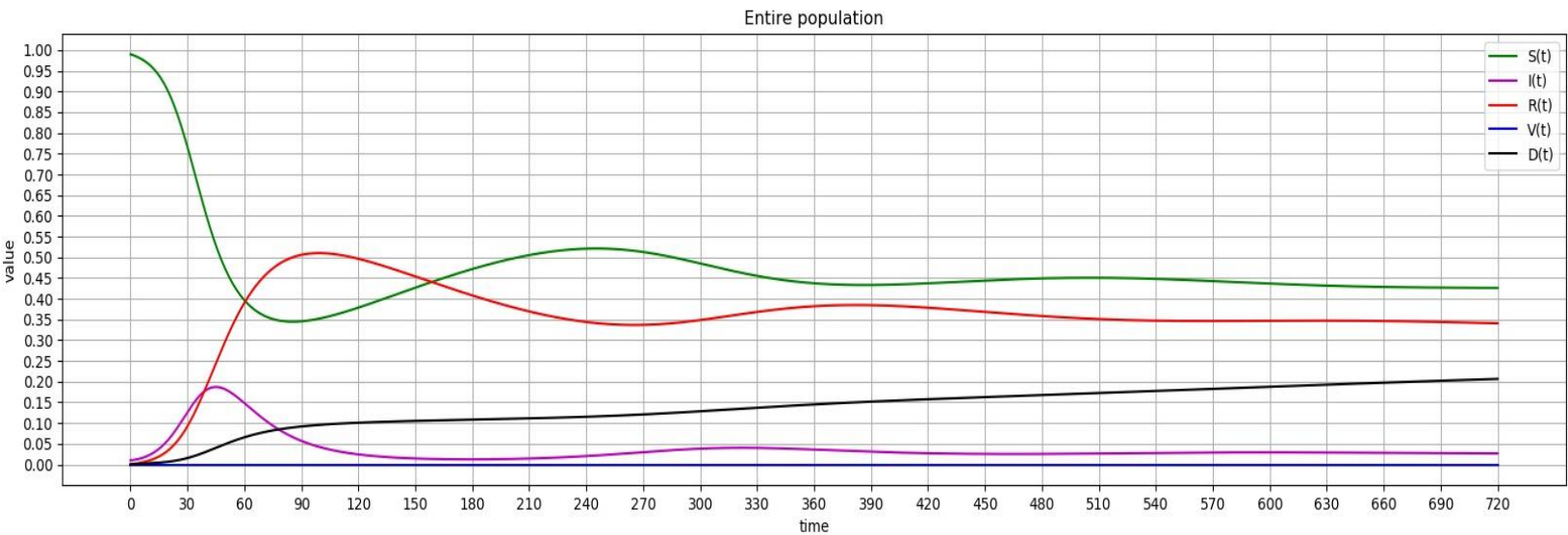
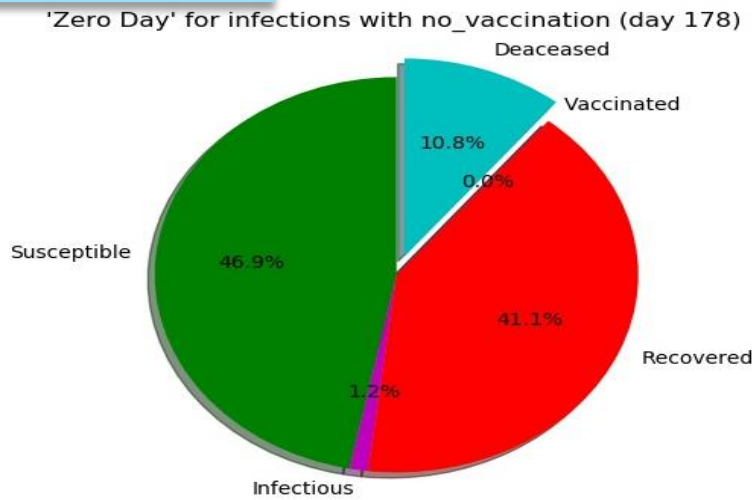


Endemic infection

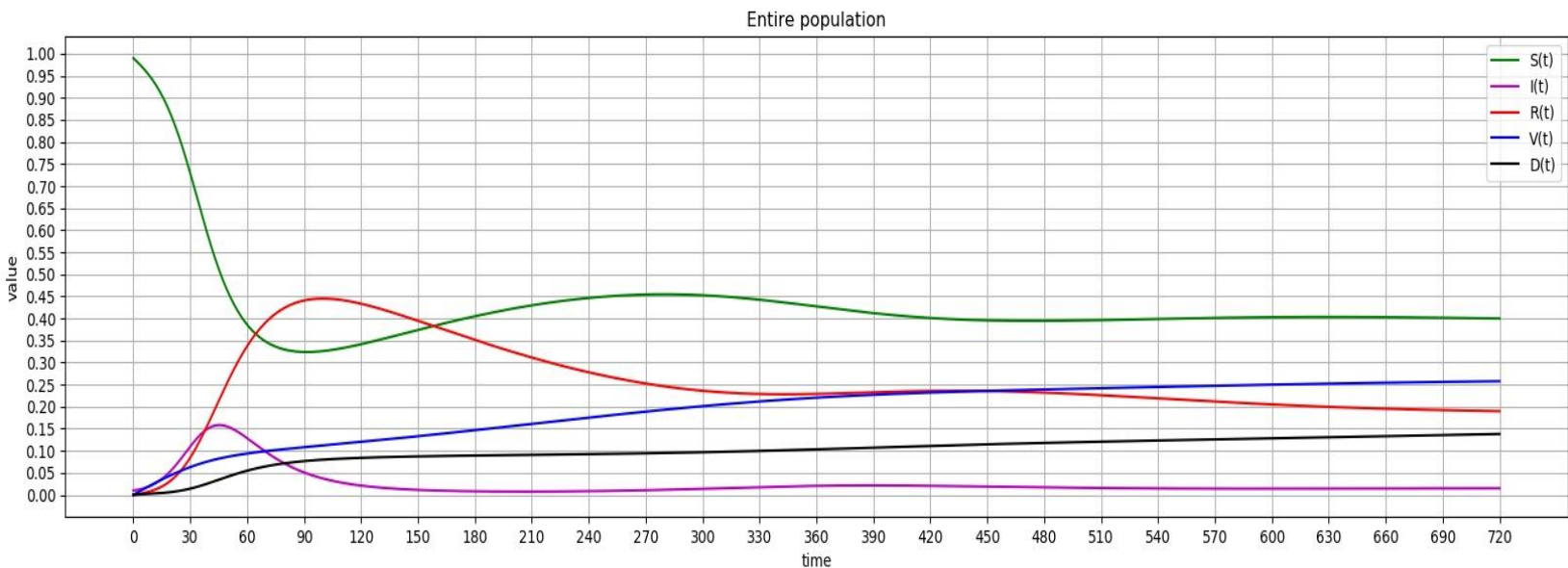
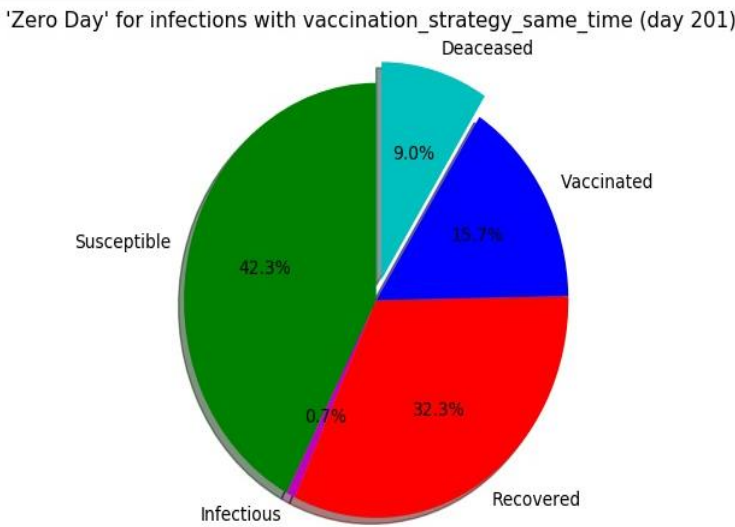
Coarse-grained analysis

Same Time	Children	Teenagers	Adults	Senior
η	0.0025	0.0025	0.0025	0.0025
VACC-DAY	0	0	0	0

No vaccination

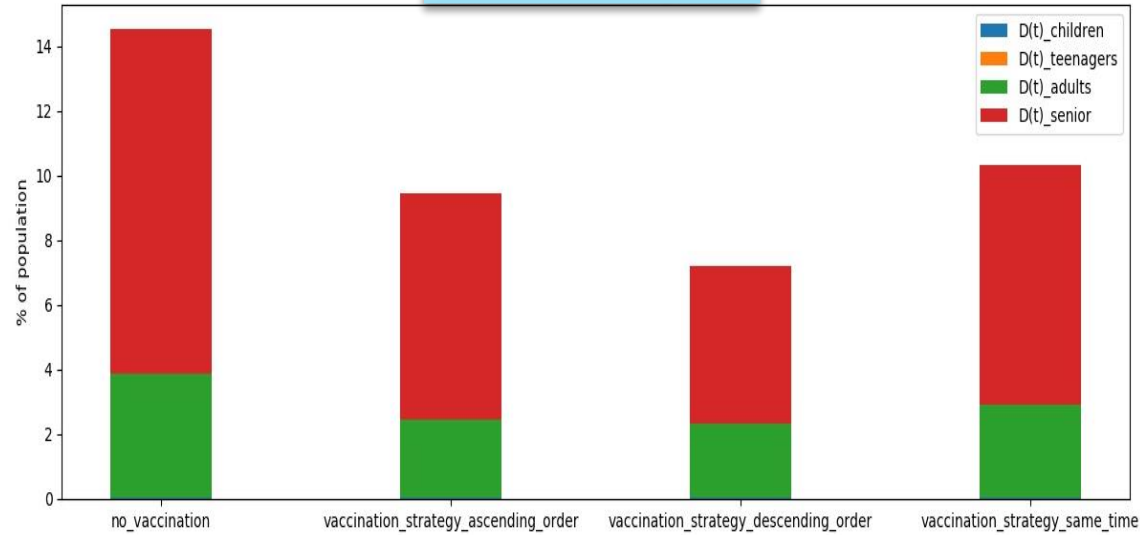


Same Time

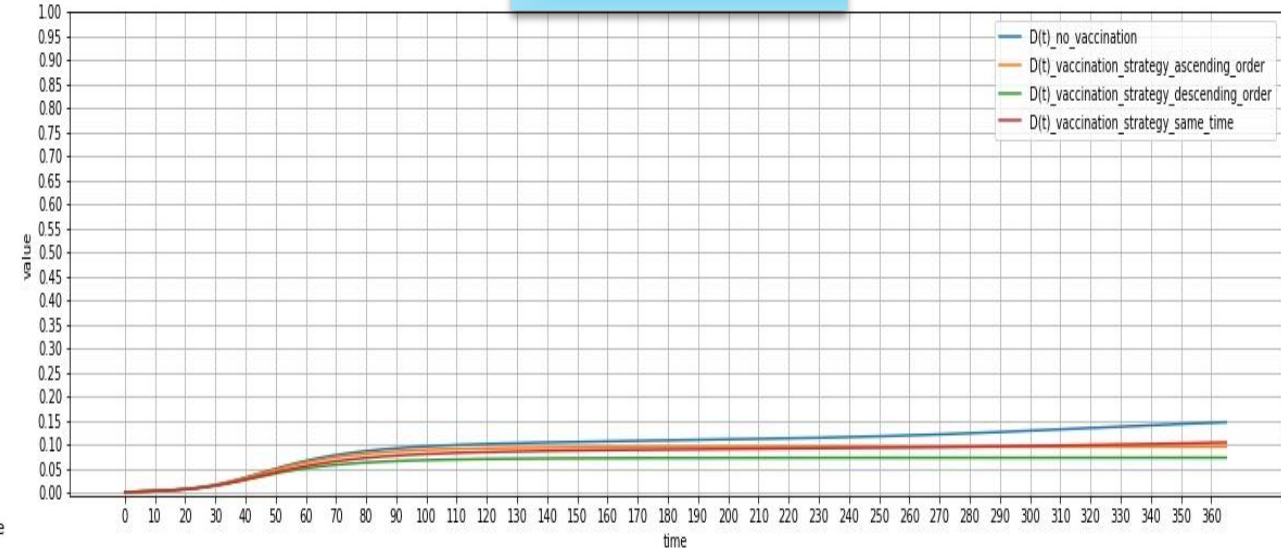


Final vaccination strategies comparison

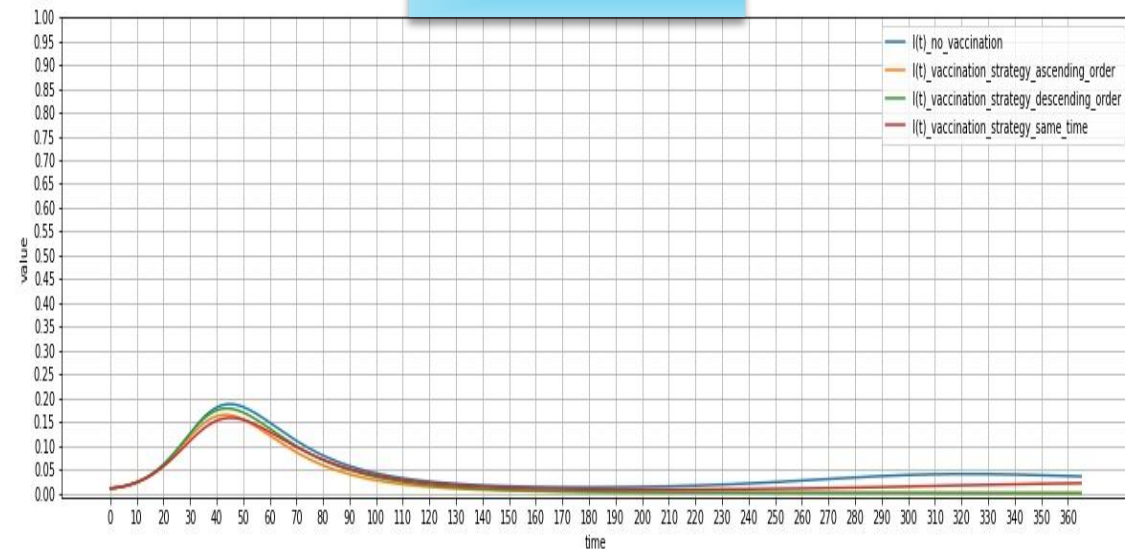
Mortality



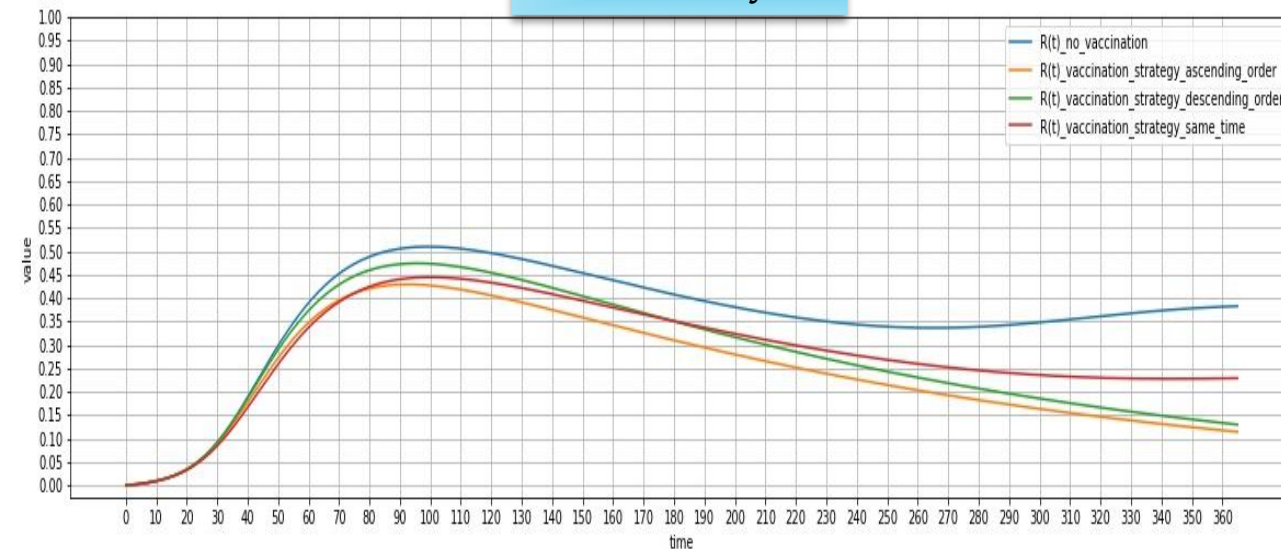
Mortality



Infections

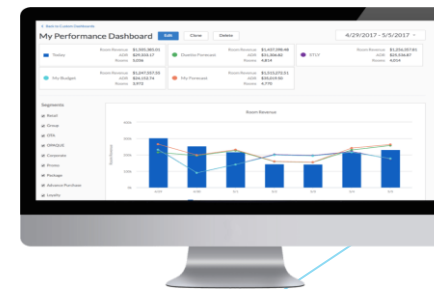
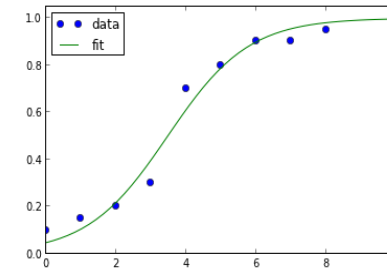
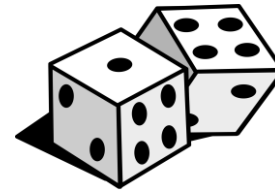


Recovery



Final Considerations and Improvements

- This model with ODEs is simple and intuitive and we obtain some expected experimental results
- The model could include compartments like quarantine, intensive care occupation and could take into account the preventive measures used
- Vaccination rate could be described by a logistic function
- Probabilities could be included in the model
- Implement an interactive dashboard with daily measurements
- Parameters estimation and fitting with a real (and accurate) dataset



References

- [Compartmental models in epidemiology — Wikipedia, the free encyclopedia](#)
- [Modeling Infectious Diseases in Humans and Animals](#)
- [Influence of nonlinear incidence rates upon the behavior of SIRS epidemiological models](#)
- [Analysis of COVID-19 Data with PRISM: Parameter Estimation and SIR Modelling](#)
- [Use of a Modified SIRD Model to Analyze COVID-19 Data](#)
- [Global results for an SIRS model with vaccination and isolation](#)
- [Mathematical models of contact patterns between age groups for predicting the spread of infectious diseases](#)
- [A statistical methodology for data-driven partitioning of infectious disease incidence into age-groups](#)
- [Lab24 - Coronavirus in Italia, i dati e la mappa](#)
- [SIR Modelling with data fitting in Python](#)
- [Matplotlib Documentation](#)