

Compartmental models of infectious disease dynamics

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Overview

Qualitative behaviour of the model

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$$\frac{dS}{dt} = -rSI, \quad \frac{dI}{dt} = rSI - \gamma I, \quad \frac{dR}{dt} = \gamma I$$

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 - That is, looking for conditions for $I(t) > I_0$ for some time t ;

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- Also, for $dS/dt < 0$, $S < S_0$ for all t .
- Thus, if $S_0 < \gamma/r$ then $S(t) < \gamma/r$ for all t

$$\left[\frac{dI}{dt} \right] = rSI - \gamma I = I(rS - \gamma) < 0,$$

and thus $I(t) < I_0$ and will be no epidemic.

- If $S_0 > \gamma/r$ there will be an epidemic (as $\left[\frac{dI}{dt} \right]_0 > 0$). Is $I(t)$ a **monotonic function?**

Qualitative behaviour of the model: the threshold phenomena

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- The inverse of the relative removal rate is the well known threshold parameter called the basic reproductive ratio (R_0), for this case $R_0 = \frac{rS_0}{\gamma}$; if $R_0 < 1$ no epidemic other we have an outbreak.

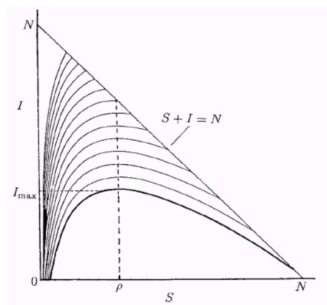
Phase portrait of the SIR model

- Let us look at the dynamics in the phase space.
- We have three variables, but as $S + I + R = N$, we can eliminate one of them. Say, R . So we now have a two-dimensional phase space S e I .

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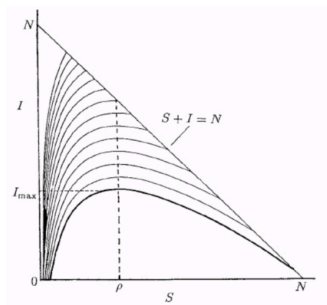
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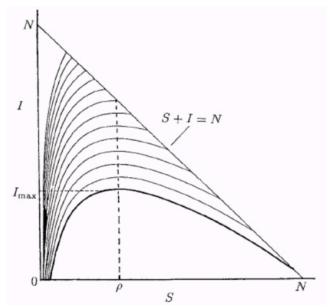


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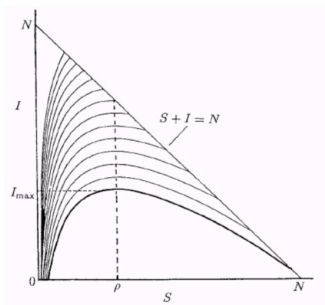


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- This will be further extended through **final epidemic size** analysis.

Find the curves governing the (S, I) phase plane. Determine when the maximum value of I on each of these orbits is attained. Give the expression of I_{max} .

Interpretation of R_0

- If we assume that $S_0 \approx N$, then $S_0 < \gamma/r$ implies $R_0 = \frac{rN}{\gamma} < 1$ while $S_0 > \gamma/r$ implies $R_0 = \frac{rN}{\gamma} > 1$.
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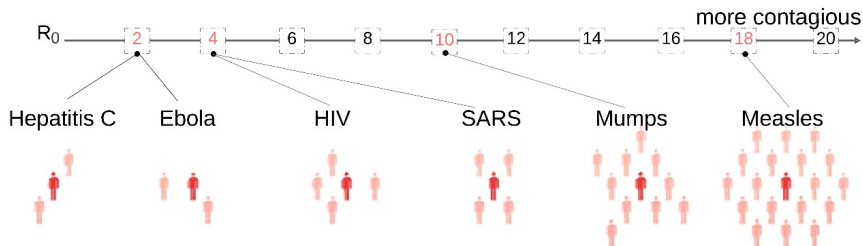
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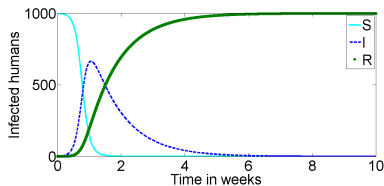
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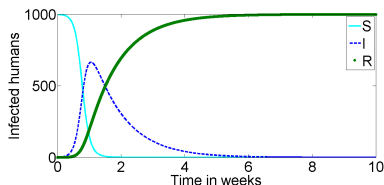
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The time-evolution of model variables



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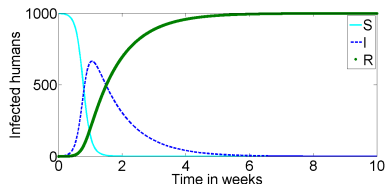
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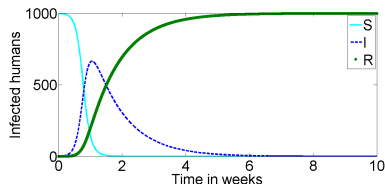
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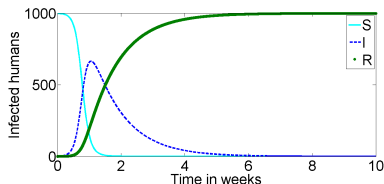
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- we can use the model as a starting point then extend it for more realistic scenarios.

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Remark

The chain of transmission eventually breaks due to the decline in infectives, not due to a complete lack of susceptibles.

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- So, there is a zero, which is the solution.

Herd immunity and Social distancing

Herd immunity

The interruption or resistance to onward infections that results if a sufficient high proportion of individuals are immune.

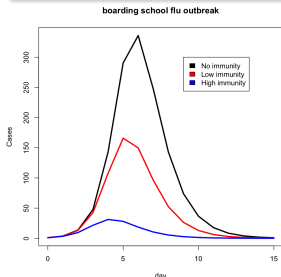
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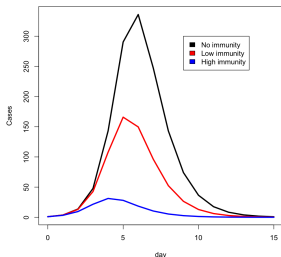
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boarding school flu outbreak

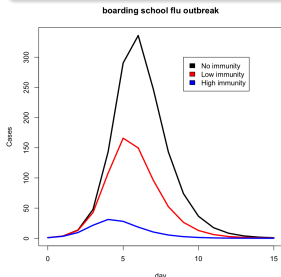


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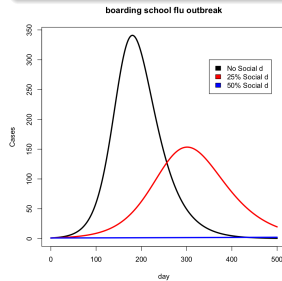
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Conditions for a Second Wave of COVID-19 Due to Interactions Between Disease Dynamics and Social Processes

Sansao A. Pedro¹, Frank T. Ndjomatchoua², Peter Jentsch^{3,4}, Jean M. Tchuente⁵, Madhur Anand³ and Chris T. Bauch^{4*}

¹ Departamento de Matemática e Informática, Universidade Eduardo Mondlane, Maputo, Mozambique, ² Sustainable Platform, Geospatial Science and Modelling Cluster, International Rice Research Institute, Metro Manila, Philippines, ³ of Environmental Sciences, University of Guelph, Guelph, ON, Canada, ⁴ Department of Applied Mathematics, University of Waterloo, Waterloo, ON, Canada, ⁵ American Health, Glastonbury, CT, United States

Summary: general properties of epidemics

- For an infectious disease with an average infectious period given by $1/\gamma$ and a transmission rate $r = \beta$, its basic reproductive ratio R_0 is determined by β/γ .
- In a closed population, an infectious disease with a specified R_0 can invade only if there is a threshold fraction of susceptibles greater than $1/R_0$.
- Vaccination can be used to reduce the proportion of susceptibles below $1/R_0$ and hence eradicate the disease;
- Social distancing can be used to delay the peak and buy time for control;
- The chain of transmission eventually breaks due to the decline in infectives, not due to a complete lack of susceptibles.

- In many infectious diseases there is an exposed period after the transmission of infection from susceptibles to potentially infective members but before these potential infectives can transmit infection. Establish the final size relation for the extended model.
- Consider an *SIR* epidemic model and extend it by assuming that there a treatment for infection once a person has been infected. Model this by supposing that a fraction γ per unit time of infectives is selected for treatment, and that treatment reduces infectivity by a fraction δ . Suppose that the rate of removal from the treated class is η . Give the corresponding flow chart and equations of the model.

End Lecturer Two!

References



M.J. Keeling and P. Rohani (2007)

Modeling Infectious Diseases in Humans and Animals

Princeton.



J.D. Murray (2002)

Mathematical Biology I

Springer.



R.M. Anderson and (1982)

Population Dynamics of Infectious Diseases: Theory and Applications

Springer.



G.F. Raggett: Modeling the Eyam plague. IMA J., 18, 221–226 (1982)

References



O. Diekmann, J.A.P. Heesterbeek, J.A.J. Metz (1990),
On the definition and the computation of the basic reproduction ratio R_0 in
models for infectious diseases in heterogeneous populations,
J. Math. Biol. 28: 365-382.



P. VAn den Driessche, J. Watmough (2002),
Reproduction numbers and sub-threshold endemic equilibria for compartmental
models of disease transmission,
Mathematical Biosciences, 180: 29-48.



Pedro SA, Ndjomatchoua FT, Jentsch P, Tchuente JM, Anand M and Bauch CT
(2020) Conditions for a Second Wave of COVID-19 Due to Interactions Between
Disease Dynamics and Social Processes. *Front. Phys.* 8:574514. doi:
10.3389/fphy.2020.574514



R. M. Anderson and R. M. May (1992),
Infectious Disease of Humans.,
Oxford University Press

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