

Nonlinear Systems

SIR Model

John Meth

April 3, 2020

$S(t)$ is the number of susceptible individuals

$I(t)$ is the number of infected individuals

$R(t)$ is the number of recovered individuals

$S(t)$ is the number of susceptible individuals

$I(t)$ is the number of infected individuals

$R(t)$ is the number of recovered individuals

A linear system in S , I , and R would look like

$$\frac{d}{dt} S = a_1 S + a_2 I + a_3 R$$

$$\frac{d}{dt} I = b_1 S + b_2 I + b_3 R$$

$$\frac{d}{dt} R = c_1 S + c_2 I + c_3 R$$

$S(t)$ is the number of susceptible individuals

$I(t)$ is the number of infected individuals

$R(t)$ is the number of recovered individuals

Our model of disease spread in S , I , and R is *nonlinear*

$$\frac{d}{dt}S = -\alpha SI$$

$$\frac{d}{dt}I = \alpha SI - \gamma I$$

$$\frac{d}{dt}R = \gamma I$$

$S(t)$ is the number of susceptible individuals

$I(t)$ is the number of infected individuals

$R(t)$ is the number of recovered individuals

Our model of disease spread in S , I , and R is *nonlinear*

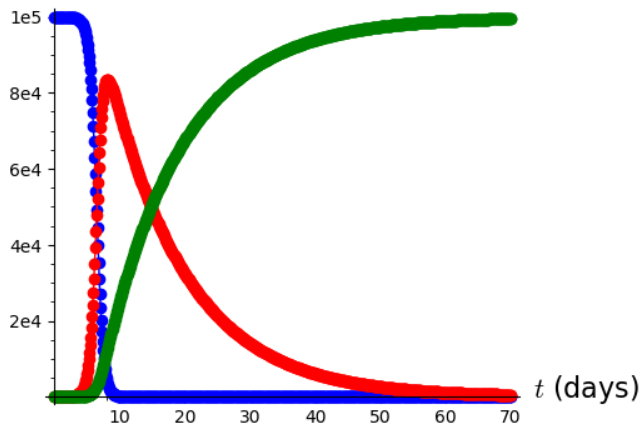
$$\frac{d}{dt} S = -\alpha SI$$

$$\frac{d}{dt} I = \alpha SI - \gamma I$$

$$\frac{d}{dt} R = \gamma I$$

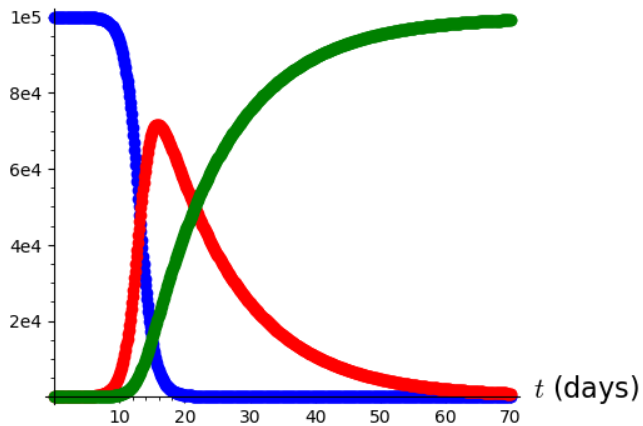
$$\alpha = \left\{ \begin{array}{l} \text{Proportion of possible} \\ \text{contacts that actually} \\ \text{occur} \end{array} \right\} \cdot \left\{ \begin{array}{l} \text{Proportion of those} \\ \text{contacts that actually} \\ \text{result in an infection} \end{array} \right\}$$

S, I, R (individuals)



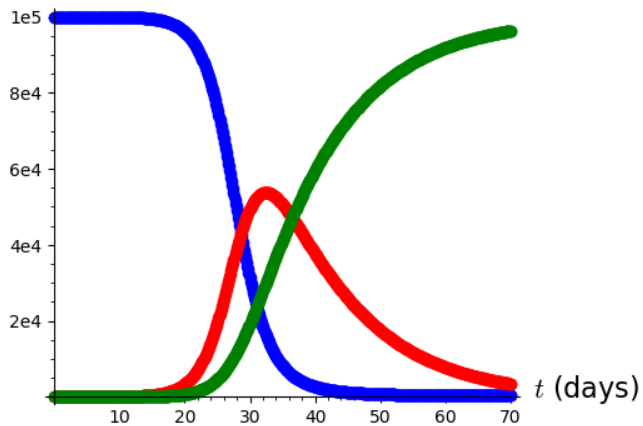
$$\alpha = .00002$$

S, I, R (individuals)



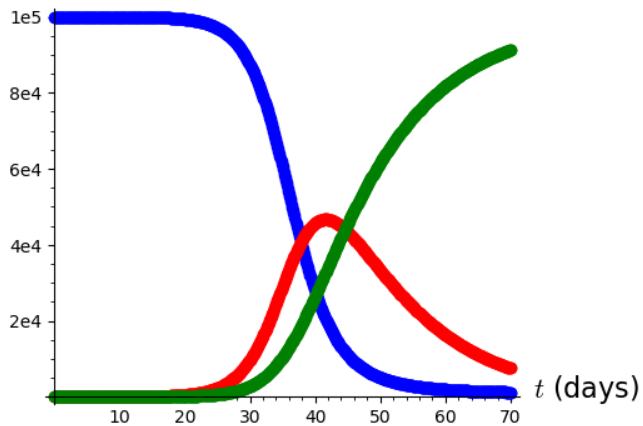
$$\alpha = .00001$$

S, I, R (individuals)



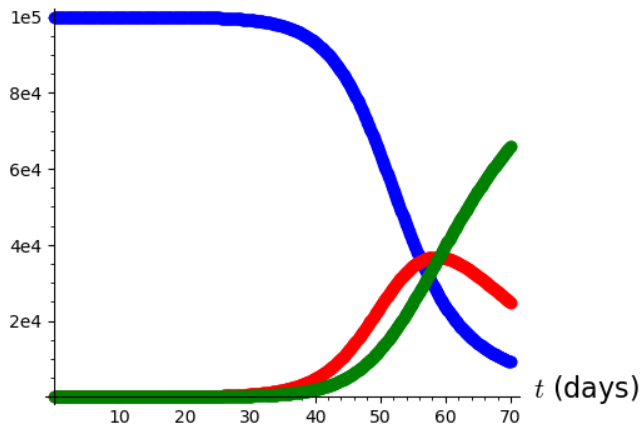
$$\alpha = .000005$$

S, I, R (individuals)



$$\alpha = .000004$$

S, I, R (individuals)



$$\alpha = .000003$$

- ▶ https://en.wikipedia.org/wiki/Compartmental_models_in_epidemiology
 - ▶ <http://math.colorado.edu/~stade/CLS/sage3.html>
 - ▶ <https://cocalc.com/>
 - ▶ <http://faculty.sfasu.edu/judsontw/ode/html-20190821/systems01.html>
- Section 2.1.3 of Judson