

Calculating Biological Quantities

CSCI 2897

Prof. Daniel Larremore

2021, Lecture 12

daniel.larremore@colorado.edu

[@danlarremore](https://twitter.com/danlarremore)

Last time on CSCI 2987: The SIR Model

$$\dot{S} = -\beta SI$$

$$\dot{I} = \beta SI - \gamma I$$

$$\dot{R} = \gamma I$$

where $S + I + R = 1$

$$\implies \dot{S} + \dot{I} + \dot{R} = 0$$

Equilibrium when:

$$I = 0$$

Epidemic peak:

$$S^* = \frac{\gamma}{\beta}$$

Herd Immunity (vaccination)

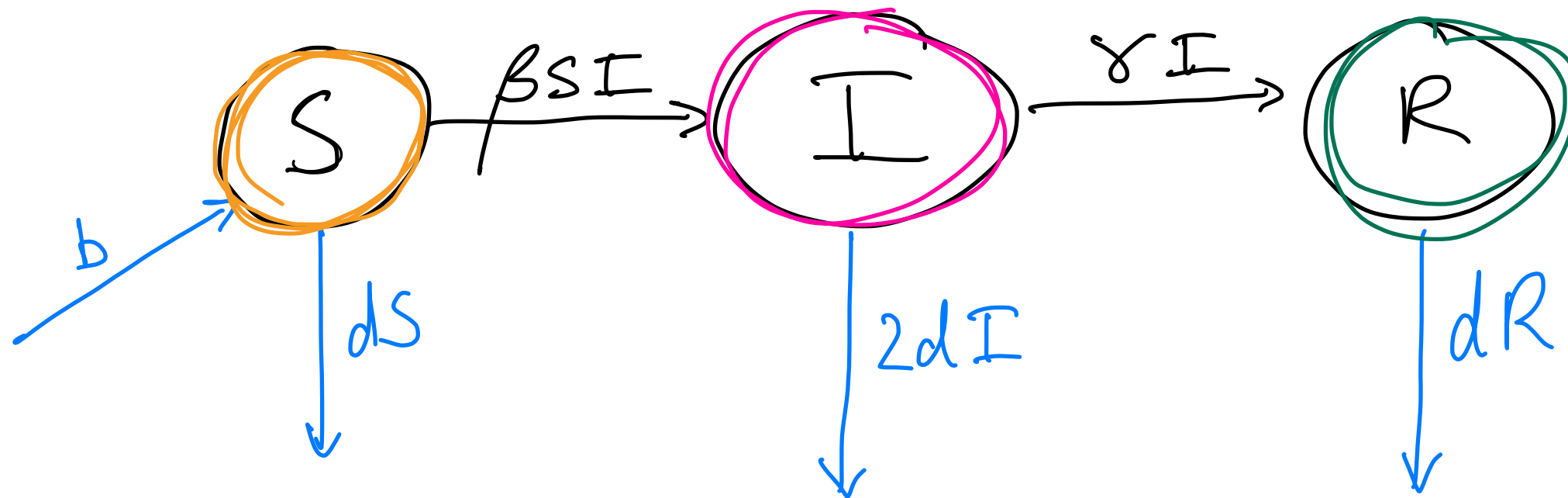
$$v > 1 - \frac{\gamma}{\beta}$$

Basic Reproduction Number

$$R_0 = \frac{\beta}{\gamma}$$

Midterm: SIR model with birth and death

$$\frac{dN}{dt} = b - dS - 2dI - dR = b - dI - d(S + I + R) = b - d(I + N)$$



$$\frac{dS}{dt} = b - dS - \beta SI$$

$$\frac{dI}{dt} = \beta SI - \gamma I - 2dI$$

$$\frac{dR}{dt} = \gamma I - dR$$

The SEIR model—exposure without infectiousness

Some diseases have a **latent period** in which a person is infected but not yet infectious to others.

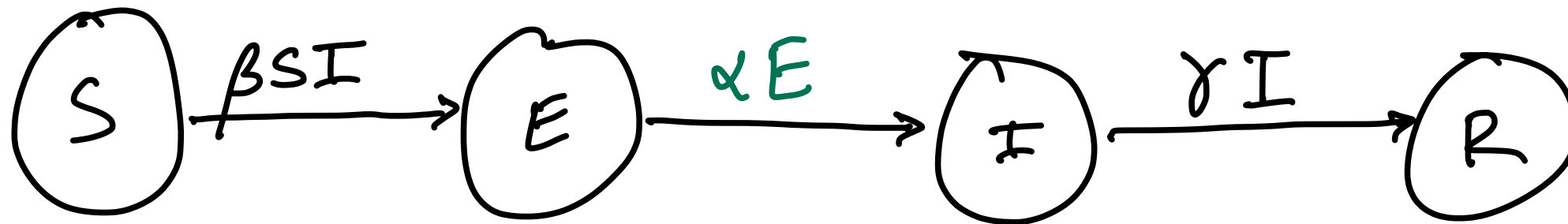
Let's consider a **new** compartment: **E**xposed, with a transition rate α .

$$\dot{S} = -\beta SI$$

$$\dot{E} = \beta SI - \alpha E$$

$$\dot{I} = \alpha E - \gamma I$$

$$\dot{R} = \gamma I$$



E puts a delay in the model!

$$\dot{S} + \dot{E} + \dot{I} + \dot{R} = 0$$

(constant pop. size)

SEIR Model — Equilibrium

$$\dot{S} = -\beta SI = 0$$

$$\textcircled{4} -\beta SI = 0 \quad (by \textcircled{1})$$

$$\dot{E} = \beta SI - \alpha E = 0$$

$$\textcircled{3} \beta SI - \alpha E = 0 \quad (by \textcircled{1})_0 \quad (by \textcircled{2})$$

$$\dot{I} = \alpha E - \gamma I = 0$$

$$\textcircled{2} \alpha E - \gamma I = 0 \Rightarrow E = 0$$

$$\dot{R} = \gamma I = 0$$

$$\textcircled{1} I = 0$$

satisfied

where $S + E + I + R = 1$

equilibrium = $(S, 0, 0, 1-S)$

SEIR Model — Out of Equilibrium?

$$\dot{S} = -\beta SI$$

$$\dot{E} = \beta SI - \alpha E$$

$$\dot{I} = \alpha E - \gamma I$$

Let's hop into Jupyter Notebooks to explore a bit.

$$\dot{R} = \gamma I$$

where $S + E + I + R = 1$

SEIR Model — Out of Equilibrium?

With Social Distancing? $0 \leq \rho \leq 1$

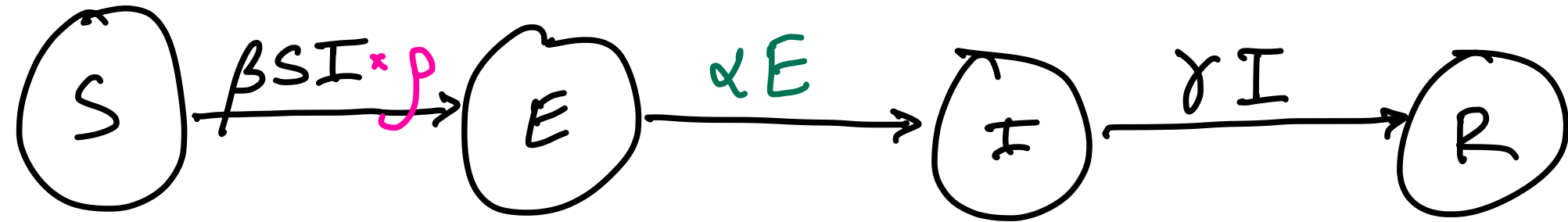
$$\dot{S} = -\beta SI \rho$$

$$\dot{E} = \rho \beta SI - \alpha E$$

$$\dot{I} = \alpha E - \gamma I$$

$$\dot{R} = \gamma I$$

where $S + E + I + R = 1$



Let's hop into Jupyter Notebooks to explore a bit.

ρ = fraction of contacts
that are still allowed

$\rho = 1 \rightarrow$ No Soc. Dist.

$\rho = 0 \rightarrow$ total isolation.

As you change ρ
from 1 downward,
you "flatten the curve."

How should we model vaccination in the SEIR model?

$$\dot{S} = -\beta SI$$

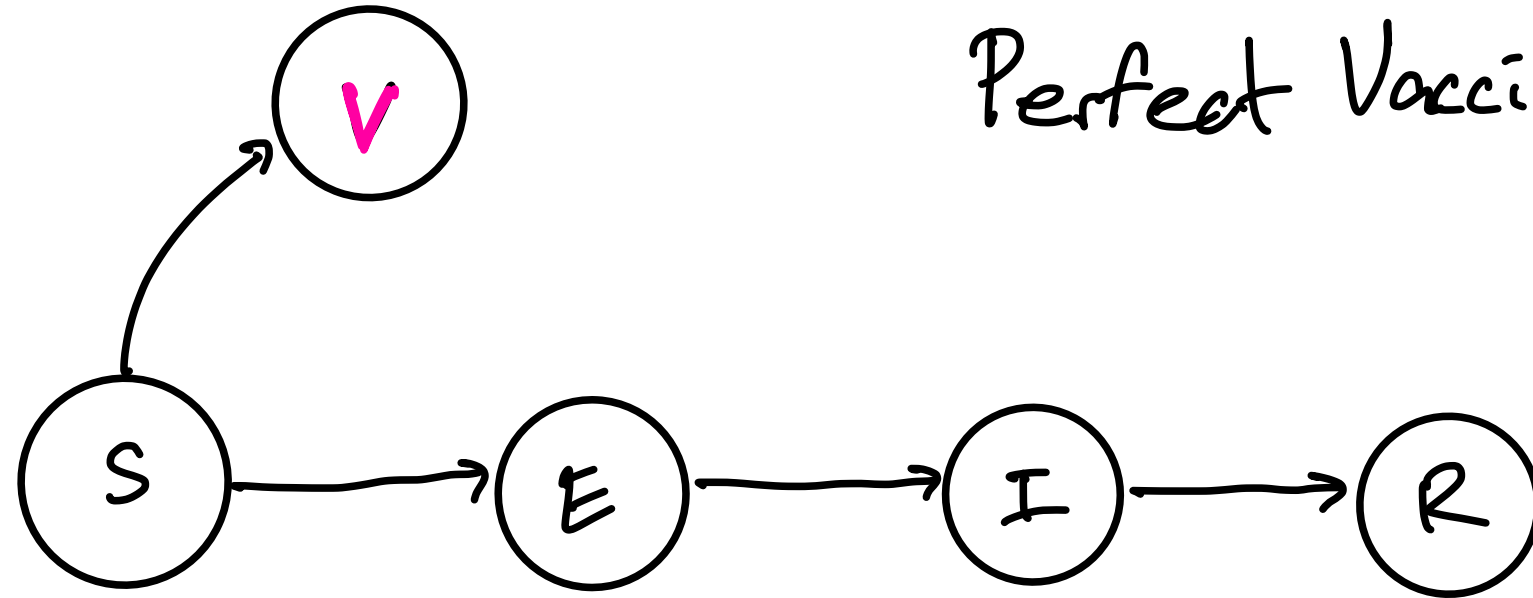
$$\dot{E} = \beta SI - \alpha E$$

$$\dot{I} = \alpha E - \gamma I$$

$$\dot{R} = \gamma I$$

$$\text{where } S + E + I + R \stackrel{+ \checkmark}{=} 1$$

Perfect Vaccine Model

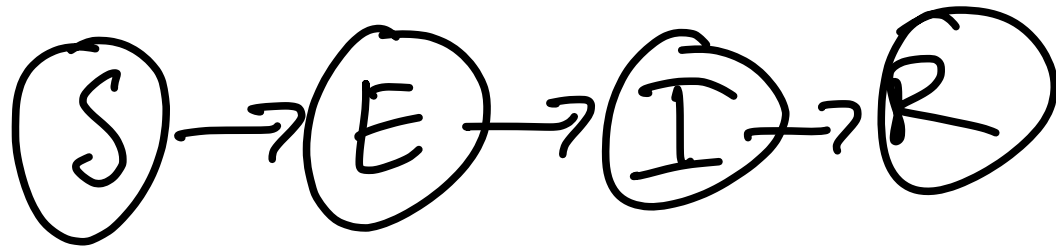
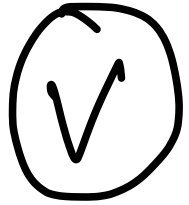


vaccination makes people not-susceptible.

- ① Prevents infection — protects you.
- ② Prevents you from spreading virus — protects others.
- ③ Prevents morbidity + mortality — protects you.

Model 1: The Perfect Vaccine model

A **perfect vaccine** provides complete protection against infection.



$$\dot{S} = -\beta SI$$

$$\dot{E} = \beta SI - \alpha E$$

$$\dot{I} = \alpha E - \gamma I$$

$$\dot{R} = \gamma I$$

$$\dot{V} = 0$$

$$S + E + I + R + V = 1$$

Model 1: The Perfect Vaccine model

A **perfect vaccine** provides complete protection against infection.

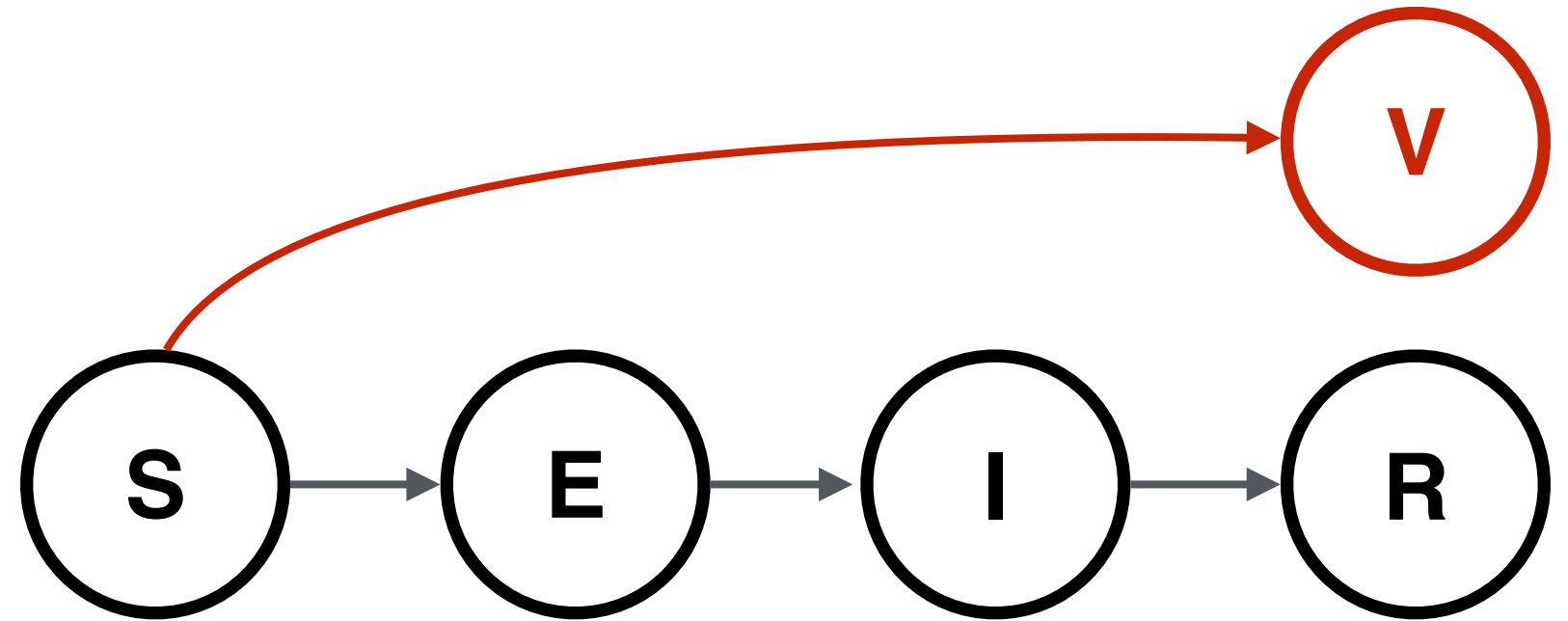
$$\dot{S} = -\beta SI$$

$$\dot{E} = \beta SI - \alpha E$$

$$\dot{I} = \alpha E - \gamma I$$

$$\dot{R} = \gamma I$$

where $S + E + I + R + V = 1$

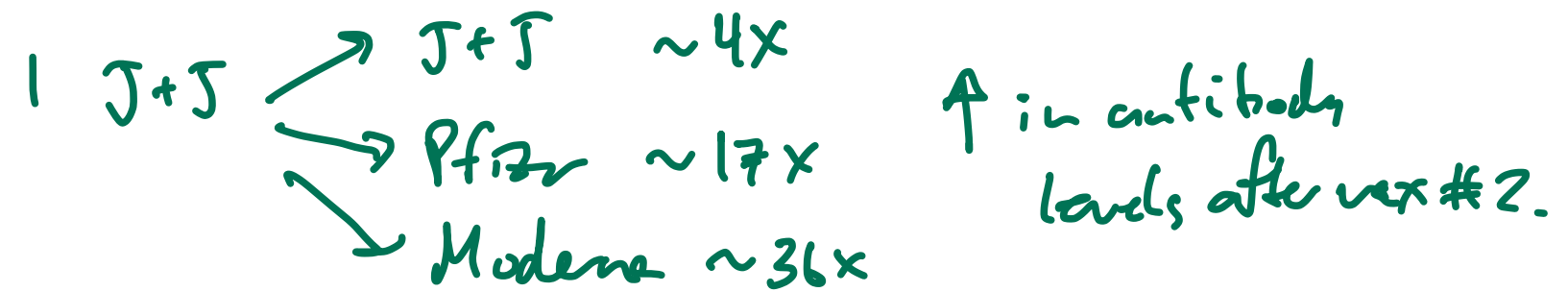


This is a model for a vaccine with $VE = 1$.
What does this mean?

What about vaccines with imperfect protection?

Vaccine efficacy (VE) is the reduction in disease outcomes in a vaccinated group compared to an unvaccinated group under trial conditions.

Vaccine effectiveness is the ability of the vaccine to prevent those disease outcomes in the real world.



What are some of the determinants of VE ?

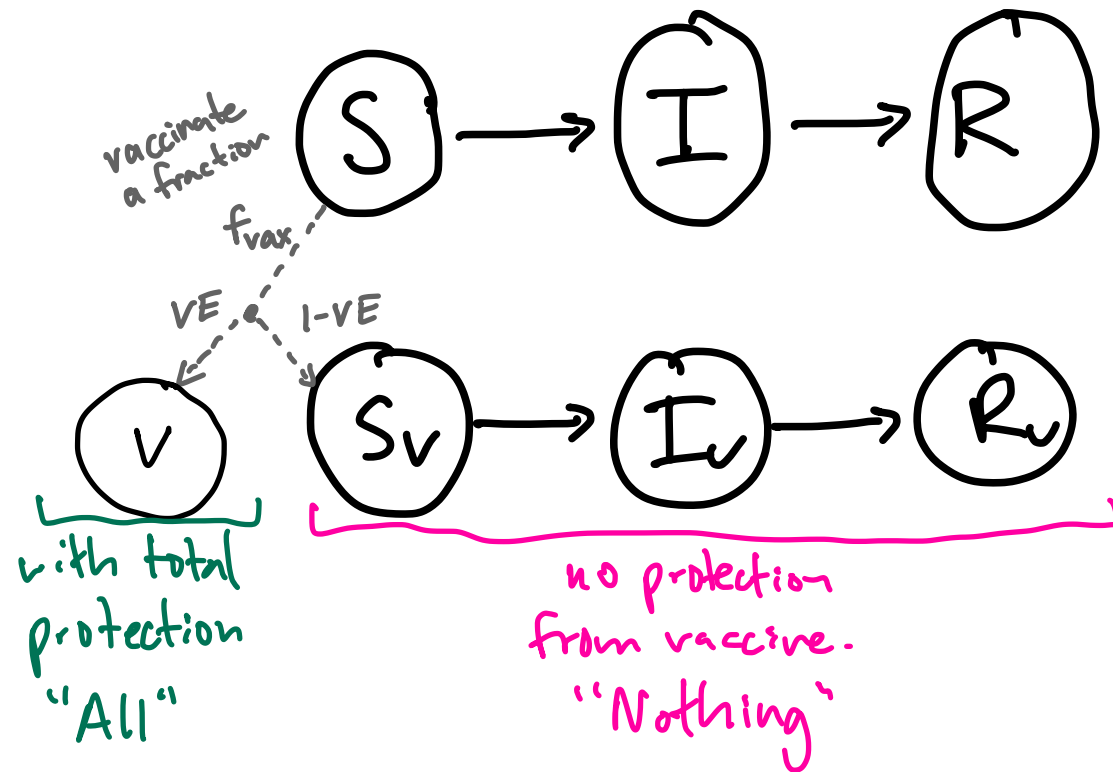
- strength of immune response to the vaccine (antibody titers, neutralization assays)
- specificity of immune response. (vac is against WildType SARS-CoV2 spike, but circulating variant is different.)
- individual effects - e.g. age - ^{immune}immunosenesence ^{steepness}

Model 2: The All-or-Nothing vaccine model

new

SIR

An **all-or-nothing** vaccine completely protects VE and leaves $1 - VE$ unprotected.



[All of the vaccination takes place as an initial condition (Assumed).]

$$\dot{S} = -\beta SI - \beta SI_v = -\beta S(I + I_v)$$

$$\dot{I} = \beta S(I + I_v) - \gamma I$$

$$\dot{R} = \gamma I$$

$$\dot{S}_v = -\beta S_v I_v - \beta S_v I = -\beta S_v(I + I_v)$$

$$\dot{I}_v = \beta S_v(I + I_v) - \gamma I_v$$

$$\dot{R}_v = \gamma I_v$$

$$\dot{V} = 0$$