

**Population Dynamics of Infectious Disease**

***Infection of Host by Microparasite***

**1. Contact with free-living pathogen stage**

***Bacillus anthrax* spores in soil**

**Nuclear polyhedrosis virus on vegetation**

**Windborne, waterborne pathogen dispersal**

**2. Contact with biotic vector**

**Parasitism preceding pathogen transmission**

**Animal macroparasites: mosquito, tick ,fly**

**Parasitic plants: *Dodder* spp**

**Aphids: persistent and non-persistent viruses**

**3. Contact with previously infected host**

**S     susceptible**

**E     exposed (latent period)**

**I     infectious**

**R     removed (dead, recover with immunity)**

**Compartment models infection by direct-contact**

**Compare two transmission dynamics**

## **Direct-contact transmission**

**Assumptions ( $S, I$ )  $\rightarrow$  ( $S - 1, I + 1$ )**

**Lotka-Volterra (density-dependent)**

**Mass action (frequency-dependent)**

**Local, probabilistic**

## ***SI Model with Immigration, Include Virulence***

**$S_t$  susceptible density, time  $t$**

**$I_t$  infectious-host density, time  $t$**

### **Density-dependent dynamics:**

$$\frac{dS}{dt} = \lambda - \mu S - \beta I S$$

$$\frac{dI}{dt} = \beta I S - (\mu + \gamma) I$$

$\lambda$  **Birth/Immigration constant**

$\mu$  **Natural mortality, density-independent**

$\gamma$  **Virulence (extra mortality from disease)**

$\beta$  **Transmission coefficient**

**Lotka-Volterra: Rate of contacts proportional to product of densities,  $IS$**

**Pr [ Transmit pathogen | contact ] =  $\beta$**

**Generalization  $\beta I^p S^q$**

## ***Equilibrium Nodes***

$$[S^*, I^*] = \left[ \frac{\lambda}{\mu}, 0 \right] \quad \text{Disease-free host population}$$

**Invasion analysis: Can rare pathogen advance?**

$$[S^*, I^*] = \left[ \frac{\mu + \gamma}{\beta}, \frac{\lambda}{\mu + \gamma} - \frac{\mu}{\beta} \right] \quad \text{Endemic disease}$$

**( Epidemic advance to final size, then fadeout )**

**Disease equilibrium (when stable)**

$$\frac{\partial S^*}{\partial \mu} > 0; \frac{\partial I^*}{\partial \mu} < 0 \quad \text{Background mortality}$$

$$\frac{\partial S^*}{\partial \gamma} > 0; \frac{\partial I^*}{\partial \gamma} < 0 \quad \text{Virulence}$$

$$\frac{\partial S^*}{\partial \lambda} = 0; \frac{\partial I^*}{\partial \lambda} > 0 \quad \text{Flow of susceptibles}$$

$$\frac{\partial S^*}{\partial \beta} < 0; \frac{\partial I^*}{\partial \beta} > 0 \quad \text{Pathogen fitness (?)}$$

## ***Stability of Disease-Free Equilibrium***

**Can pathogen invade host population?**

### **Jacobian**

***How do densities affect dynamics?***

### **Recall dynamics of *SI* model**

$$\frac{dS}{dt} = \lambda - \mu S - \beta I S$$

$$\frac{dI}{dt} = \beta I S - (\mu + \gamma) I$$

$$\frac{\partial \dot{S}}{\partial S} = -\mu - \beta I$$

$$\frac{\partial \dot{S}}{\partial I} = -\beta S$$

$$\frac{\partial \dot{I}}{\partial S} = \beta I$$

$$\frac{\partial \dot{I}}{\partial I} = \beta S - \mu - \gamma$$

$$J = \begin{bmatrix} -\mu - \beta I & -\beta S \\ \beta I & \beta S - \mu - \gamma \end{bmatrix}$$

**Jacobian Matrix**

**Evaluate  $J$  at disease-free equilibrium**

$$J\left(\frac{\lambda}{\mu}, 0\right) = \begin{bmatrix} -\mu & -\beta\left(\frac{\lambda}{\mu}\right) \\ 0 & \beta\left(\frac{\lambda}{\mu}\right) - \mu - \gamma \end{bmatrix}$$

**Characteristic equation:  $\text{Det} [J - \xi I] = 0$**

$\xi$  **Eigenvalue; stability analysis**

**Pathogen invades (Disease advances when rare):**

$$\mu + \gamma - \beta\left(\frac{\lambda}{\mu}\right) < 0 \Rightarrow \mu + \gamma < \beta\left(\frac{\lambda}{\mu}\right) = \beta S^*$$

**Pathogen invades *iff***  $S^* > \frac{(\mu + \gamma)}{\beta}$

**Critical population size**

**Promote invasion: increase  $\lambda, \beta$**

**Inhibit invasion: increase  $\mu, \gamma$**

## ***R<sub>0</sub> and Pathogen Invasion***

**Disease advances:**  $\mu + \gamma - \beta \left( \frac{\lambda}{\mu} \right) < 0$

$$\beta \left( \frac{\lambda}{\mu} \right) - \mu - \gamma > 0 \Rightarrow \frac{\beta \left( \frac{\lambda}{\mu} \right)}{\mu + \gamma} > 1 \Leftrightarrow \frac{\beta S^*}{\mu + \gamma} > 1$$

## **R<sub>0</sub> Criterion**

### **Single infective**

**S\* contacts/unit time**

**Pr [Infect | contact ] = β**

**Duration infectious period = (μ + γ)<sup>-1</sup>**

$$R_0 = \frac{\beta S^*}{\mu + \gamma}$$

### **Recover stability criterion**

### **Criterion function of host disease-free density**

## ***Frequency-Dependent Transmission***

### **Dynamics:**

$$\begin{aligned} \frac{dS}{dt} &= \lambda - \mu S - \beta I \left( \frac{S}{N} \right) \\ \frac{dI}{dt} &= \beta I \left( \frac{S}{N} \right) - (\mu + \gamma) I \end{aligned}$$

$\lambda, \mu, \gamma$  as above

**$N = S + I$  total density**

**$I_i$  infectives make  $\beta I$  total contacts/unit time**

**Fraction contacts susceptible =  $S_i/N$**

**Total transmission rate =  $\beta I \left( \frac{S}{N} \right)$**

**N.B. :  $\beta$  has different units in 2 models**

## ***R<sub>0</sub> from Invasion Analysis***

**Disease-free equilibrium**  $[S^*, I^*] = \left[ \frac{\lambda}{\mu}, 0 \right]$   
**As above**

**Rare infective, total contact rate  $\beta$**

**Susceptible frequency =  $S^*/N = S^*/S^* = 1$**

**Duration of infectious period =  $(\mu + \gamma)^{-1}$**

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

**Independent of host density; no critical pop size**

**Equilibrate dynamics, endemic densities differ**

**Different assumptions make different predictions**

**Each has limited empirical support**

**Is there a critical population size?**

**Does  $R_0$  depend on susceptible density?**

**Transients, endemic densities compare models?**

**In particular, effect of immigration on endemic  $S^*$**

**Do experiments address both density variation  
and number variation?**

**What are consequences for pathogen evolution  
via strain competition?**