

Environmentally Transmitted Pathogens

Models – Part deux :)

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January 2023

Some considerations about numerics

- The tetanus model of Cvjetanović

- The model of Capasso for ETP

- A model for zoonotic transmission of waterborne disease

- The first schistosomiasis model of Woolhouse

- The third schistosomiasis model of Woolhouse – Heterogeneous contacts

- Spatial aspects – Cholera in Haiti

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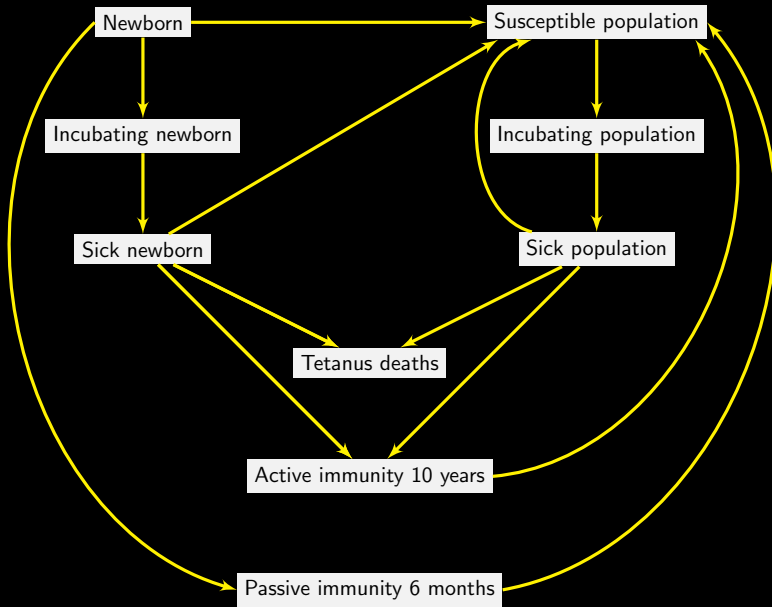
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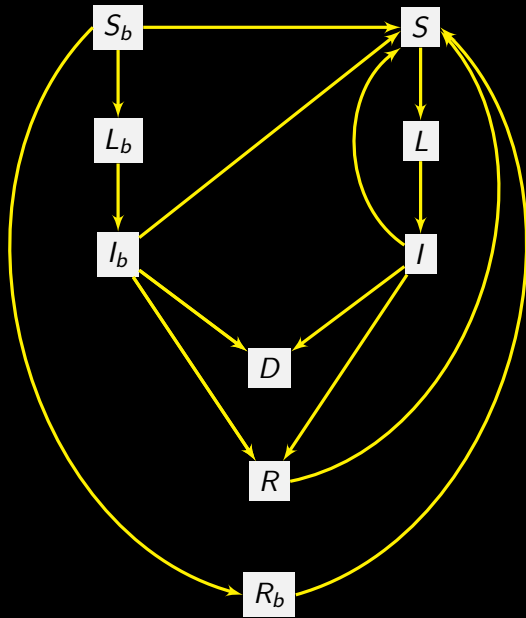
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The discrete-time tetanus model (notation mine)

$$\Delta S_b = bT \quad (1a)$$

$$\Delta S = b(1 - \lambda_b)(T - R) + \nu R + \nu_b I + R_{5,2}\gamma_b I_b + R_{6,2}\gamma I - (\lambda + d - \delta_T)S \quad (1b)$$

$$\Delta L_b = \lambda_b b(T - R) - (\varepsilon_b + d - \delta_T)L_b \quad (1c)$$

$$\Delta L = \lambda S - (\varepsilon + d - \delta_T)L \quad (1d)$$

$$\Delta I_b = \varepsilon_b L_b - (\gamma_b + d - \delta_T)I_b \quad (1e)$$

$$\Delta I = \varepsilon L - (\gamma + d - \delta_T)I \quad (1f)$$

$$\Delta R = R_{5,7}\gamma_b I_b + R_{6,7}\gamma I - (\nu + d - \delta_T)R \quad (1g)$$

$$\Delta R_b = bR - (\nu_b + d - \delta_T)R_b \quad (1h)$$

$$\Delta D = R_{5,9}\gamma_b I_b + R_{6,9}\gamma I \quad (1i)$$

where

$$T = S + L_b + L + I_b + I + R + R_b \quad \text{and} \quad \delta_T = \frac{\Delta D}{T} \quad (1j)$$

Parameter assumptions – Tetanus

- ▶ **Incubation period** – Mean duration 6 days for newborn and 8 days for general population \Rightarrow daily rate of exit $\varepsilon_b = 0.1667$ and $\varepsilon = 0.125$
- ▶ **Period of sickness** – Mean duration 3 days for newborn and 14 days for general population \Rightarrow daily rate of exit $\gamma_b = 0.3333$ per sick newborn and $\gamma = 0.0714$ for sick general in general population
- ▶ **Mortality from tetanus** – Untreated tetanus cases, fatality rate 90% for newborn S_b and 40% for general population. Treated: 80% for newborn and 30% general population
- ▶ **Immunity** – Tetanus cases do not lead to immunity to reinfection. But as a general rule, recovered people are vaccinated. Convalescents and general population effectively immunised by complete course of vaccination go to R for average 10 years, daily rate of exit is $\nu = 0.000274$ per person.
- ▶ **Immunity of newborns** – Newborn to women vaccinated during pregnancy are temporarily protected by maternal antibodies and pass through R_b for a mean duration of 6 months. Daily rate of exit $\nu_b = 0.005479$ per immunised newborn

Parameter assumptions – Demography

Live birth rate 35 per 1,000 population and annual crude death rate 15 per 1,000 population (annual rate of growth 2%) \Rightarrow daily birth and death rates $b = 0.00009889$ and $d = 0.0000411$ per person, respectively

Parameter assumptions – Force of infection

No H2H transmission \Rightarrow incidence proportional to number of susceptible individuals and force of infection, which quantifies combined effect of all variables involved in infection process:

- ▶ degree of soil contamination with *Clostridium tetani*
- ▶ climate
- ▶ frequency of lesions
- ▶ proportion of rural population
- ▶ socioeconomic conditions
- ▶ level of medical care for the wounded and during deliveries

Force of infection acting on newborn (λ_b) and susceptible population (λ) fixed at 3 different levels adequate for reproducing the following stable annual incidence rates of tetanus cases in the community

- ▶ For newborn, 200 cases, 400 cases and 600 cases per 100,000 newborn
- ▶ For general population (without newborn), 9, 18 and 27 cases

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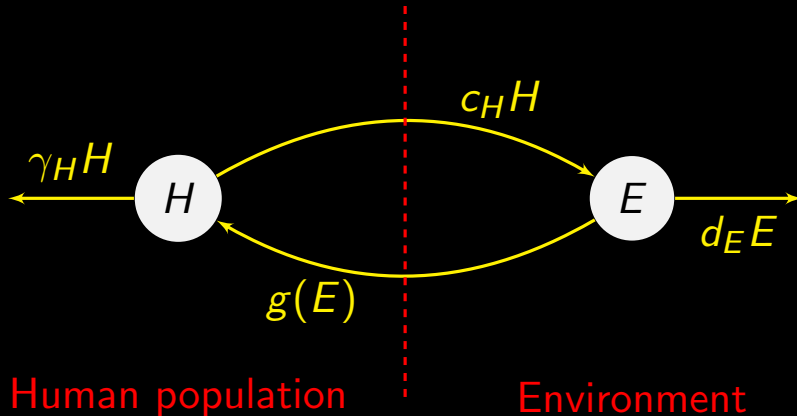
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A minimal model of V. Capasso



$1/\gamma_H$ mean infectious period, $1/d_E$ mean lifetime of the agent in the environment, c_H growth rate of the agent due to the human population, $g(E)$ “force of infection” (I would say “incidence”) of the agent on human population

Incidence function

$$g(E) = h(E)N\beta p \quad (2)$$

where

- ▶ $h(E)$ probability for an exposed susceptible to get the infection
- ▶ N total human population
- ▶ β fraction of susceptible individuals in N
- ▶ p fraction exposed to contaminated environment per unit time (“probability per unit time to have a “snack” of contaminated food”)

Typically, we would assume p and β independent of E and H and h to be saturating

To ensure (2) satisfies these conditions, we can assume

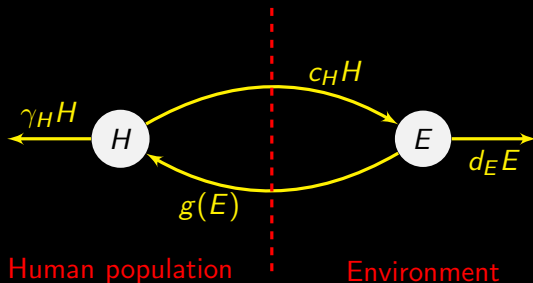
- ▶ $0 < g(e_1) < g(e_2)$ for $0 < e_1 < e_2$
- ▶ $g(0) = 0$
- ▶ $g''(z) < 0$ for all $z > 0$
- ▶ $0 < g'_+(0) < \infty$ (right derivative)
- ▶ $\lim_{z \rightarrow \infty} \frac{g(z)}{z} < \frac{d_E \gamma_H}{c_H}$

Of course, we also assume $d_E, c_H, \gamma_H > 0$

The model

$$E' = c_H H - d_E E \quad (3a)$$

$$H' = g(E) - \gamma_H H \quad (3b)$$



Pay attention to the flows...! E' does not have a $-g(E)$ and H' does not have $-c_H H$. Why?

Let

$$\mathcal{R}_0 = \frac{g'_+(0)c_H}{d_E\gamma_H} \quad (4)$$

Theorem 1

- ▶ *If $0 < \mathcal{R}_0 < 1$, then (3) admits only the trivial equilibrium in the positive orthant, which is GAS*
- ▶ *If $\mathcal{R}_0 > 1$, then two EP exist: $(0,0)$, which is unstable, and $z^* = (E^*, H^*)$ with $E^*, H^* > 0$, GAS in $\mathbb{R}_+^2 \setminus \{0,0\}$*

Adding a periodic component

Assume p in (2) takes the form

$$p(t) = p(t + \omega) > 0, \quad t \in \mathbb{R} \quad (5)$$

i.e., p has period ω . So we now consider the incidence

$$g(t, E) = p(t)h(E) \quad (6)$$

with h having the properties prescribed earlier. Letting

$$p_{min} := \min_{0 \leq t \leq \omega} p(t), \quad p_{max} := \max_{0 \leq t \leq \omega} p(t) \quad (7)$$

then we require that

$$\lim_{z \rightarrow \infty} \frac{g(z)}{z} < \frac{d_E \gamma_H}{c_H p_{max}} \quad (8)$$

Let

$$\mathcal{R}_0^{min} = \frac{c_H p_{min} h'_+(0)}{d_E \gamma_H}, \quad \mathcal{R}_0^{max} = \frac{c_H p_{max} h'_+(0)}{d_E \gamma_H} \quad (9)$$

Theorem 2

- ▶ *If $0 < \mathcal{R}_0^{max} < 1$, then (3) with incidence (6) always goes to extinction*
- ▶ *If $\mathcal{R}_0^{min} > 1$, then a unique nontrivial periodic endemic state exists for (3) with incidence (6)*

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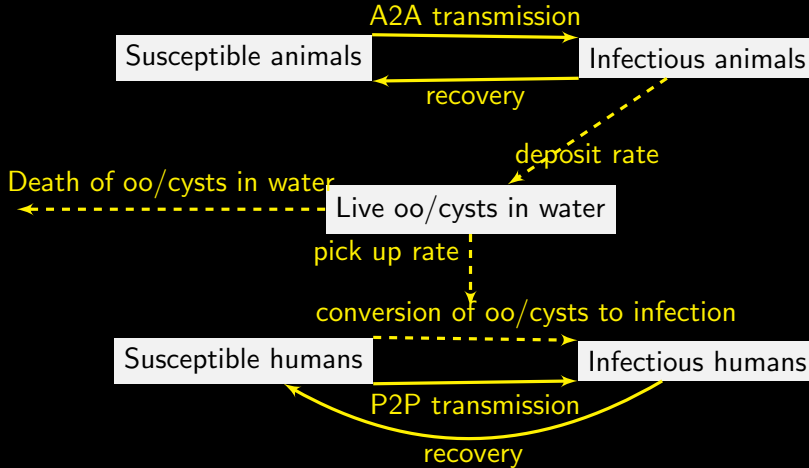
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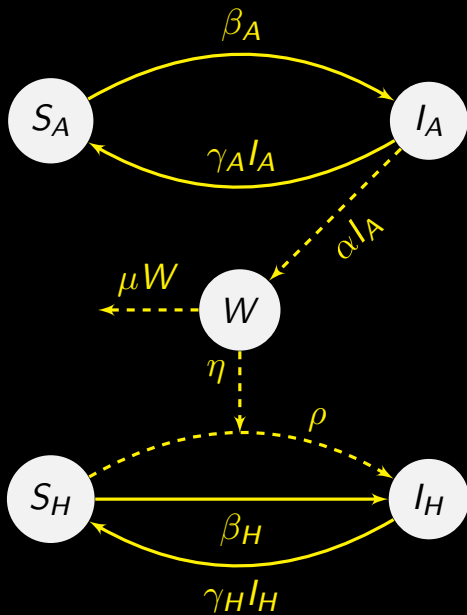
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Zoonotic transmission of waterborne disease

Waters, Hamilton, Sidhu, Sidhu, Dunbar. Zoonotic transmission of waterborne disease: a mathematical model. *Bull Math Biol* (2016)

Used for instance to model *Giardia* transmission from possums to humans





The full model

$$S'_A = -\beta_A S_A I_A + \gamma_A I_A \quad (10a)$$

$$I'_A = \beta_A S_A I_A - \gamma_A I_A \quad (10b)$$

$$W' = \alpha I_A - \eta W(S_H + I_H) - \mu W \quad (10c)$$

$$S'_H = -\rho \eta W S_H - \beta_H S_H I_H + \gamma_H I_H \quad (10d)$$

$$I'_H = \rho \eta W S_H + \beta_H S_H I_H - \gamma_H I_H \quad (10e)$$

Considered with $N_A = S_A + I_A$ and $N_H = S_H + I_H$ constant

Simplified model

Because N_A and N_H are constant, (10) can be simplified:

$$I'_A = \beta_A N_A I_A - \gamma_A I_A - \beta_A I_A^2 \quad (11a)$$

$$W' = \alpha I_A - \eta W N_H - \mu W \quad (11b)$$

$$I'_H = \rho \eta W (N_H - I_H) + \beta_H N_H I_H - \gamma_H I_H - \beta_H I_H^2 \quad (11c)$$

Three EP: DFE (0,0,0); endemic disease in humans because of H2H transmission;
endemic in both H and A because of W

Three EP: DFE $(0, 0, 0)$; endemic disease in humans because of H2H transmission; endemic in both H and A because of W

Let

$$\mathcal{R}_{0A} = \frac{\beta_A}{\gamma_A} N_A \quad \text{and} \quad \mathcal{R}_{0H} = \frac{\beta_H}{\gamma_H} N_H \quad (12)$$

- ▶ DFE LAS if $\mathcal{R}_{0A} < 1$ and $\mathcal{R}_{0H} < 1$, unstable if $\mathcal{R}_{0A} > 1$ or $\mathcal{R}_{0H} > 1$
- ▶ If $\mathcal{R}_{0H} > 1$ and $\mathcal{R}_{0A} < 1$, (11) goes to EP with endemicity only in humans
- ▶ Endemic EP with both A and H requires $\mathcal{R}_{0A} > 1$ and $\mathcal{R}_{0H} < 1$

Note that proof is **not** global

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A model of Woolhouse

Woolhouse. On the application of mathematical models of schistosome transmission dynamics. I. Natural transmission. *Acta Tropica* **49**:241-270 (1991)

The model

Population of H individuals using a body of water containing N snails

i_H mean number of schistosomes per person and i_S the proportion of patent infections in snails (prevalence)

$$i'_H = \alpha N i_S - \gamma i_H \quad (13a)$$

$$i'_S = \beta H i_H (1 - i_S) - \mu_2 i_S \quad (13b)$$

- ▶ α number of schistosomes produced per person per infected snail per unit time
- ▶ $1/\gamma$ average life expectancy of a schistosome
- ▶ $1/\mu_2$ average life expectancy of an infected snail
- ▶ β transmission parameter

Let the basic reproductive rate for schistosomes be

$$\mathcal{R}_0 = \frac{\alpha N \beta H}{\gamma \mu_2} \quad (14)$$

(13) has two EP

- ▶ $(i_H^*, i_S^*) = (0, 0)$, LAS when $\mathcal{R}_0 < 1$ and unstable when $\mathcal{R}_0 > 1$
- ▶ $(i_H^*, i_S^*) = \left(\frac{\alpha N}{\gamma} - \frac{\mu_2}{\beta H}, 1 - \frac{1}{\mathcal{R}_0} \right)$, which only “exists” when $\mathcal{R}_0 > 1$ (and is LAS then)

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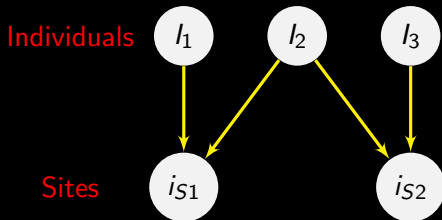
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Heterogeneities in contact rates

I_i the number of schistosomes in person $i = 1, \dots, H$ and i_{Sj} the proportion of patent infected snails in site $j = 1, \dots, L$ (L sites each supporting N snails)



l_i the number of schistosomes in person $i = 1, \dots, H$ and i_{Sj} the proportion of patent infected snails in site $j = 1, \dots, L$ (L sites each supporting N snails)

$$l'_i = \alpha \left(\sum_j \eta_{ij} N i_{Sj} \right) - \gamma l_i \quad (15a)$$

$$i'_{Sj} = \beta \left(\sum_i \eta_{ij} l_i \right) (1 - i_{Sj}) - \mu_2 i_{Sj} \quad (15b)$$

η_{ij} rate of water contact by individual i at site j

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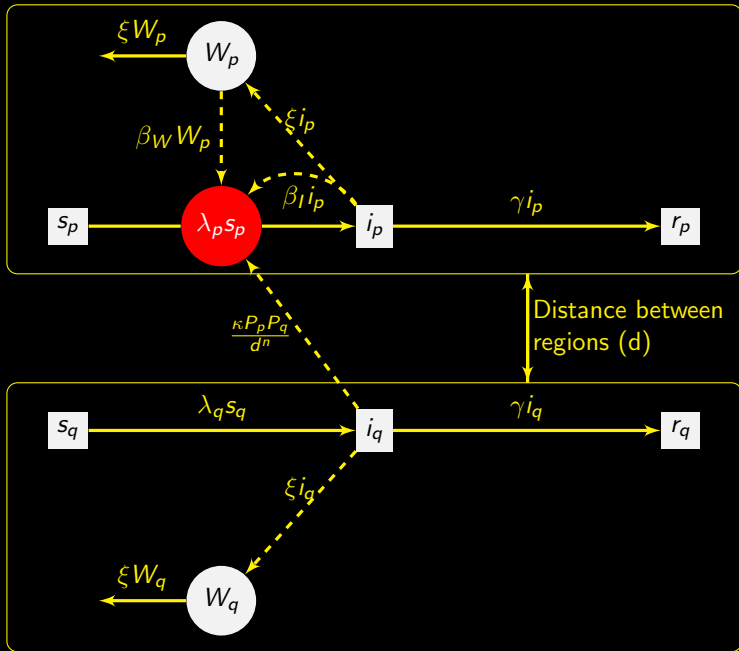
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Spatial aspects – Cholera in Haiti

Tuite, Tien, Eisenberg, Earn, Ma & Fisman. Cholera Epidemic in Haiti, 2010: Using a Transmission Model to Explain Spatial Spread of Disease and Identify Optimal Control Interventions. *Annals of Internal Medicine* **154**(9) (2011)



Metapopulation model with **implicit** movement

$$s'_p = \mu - \lambda_p s_p - \mu s_p \quad (16a)$$

$$i'_p = -\gamma i_p + \lambda_p s_p - \mu i_p \quad (16b)$$

$$r'_p = \gamma r_p - \mu r_p \quad (16c)$$

$$w'_p = \xi(i_p - w_p) \quad (16d)$$

with force of infection

$$\lambda_p = \beta_{i_p} i_p + \beta_{w_p} w_p + \sum_{q=1}^{10} \theta_{pq} i_q \quad (16e)$$

Influence of infection prevalence in q on incidence in p is gravity-type

$$\theta_{pq} = \kappa \frac{P_p P_q}{d^n}$$

