Environmentally Transmitted Pathogens Models

Julien Arino

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A few models of Cvjetanović

A tetanus model A cholera model

A model of Capasso for ETP

A model for zoonotic transmission of waterborne disease

A few models of schistosomiasis A first model of Woolhouse A second model of Woolhouse – Latency A third model of Woolhouse – Heterogeneous contacts Spatial aspects – Cholera in Haiti Cvjetanović, Grab, Uemura & World Health Organization. Dynamics of acute bacterial diseases : epidemiological models and their application in public health. World Health Organization (1978)

 Briscoe. On the use of simple analytic mathematical models of communicable diseases. *International Journal of Epidemiology* 9(3) (1980)

Models of (Branko) Cvjetanović are in discrete time and quite detailed on the epi side

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Tetanus

Bacterial infection caused by *Clostridium tetani*

Spores everywhere in the environment, including soil, dust, and manure

Spores develop into bacteria when they enter the body

Not spread P2P (so not a classic model)

Incubation period usually 3-21 days (average 8 days). Can range from 1 day to several months, depending on the kind of wound. Most cases occur within 14 days

1 to 2 in 10 cases are fatal

People of all ages need TETANUS VACCINES



DTaP for young children

- 2, 4, and 6 months
 15 through 18 months
 4 through 6 years
- Tdap for preteens
- ✓ 11 through 12 years
- for adults ✓ Every 10 years

<u>Td</u> or Tdap



www.cdc.gov/tetanus





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A cholera model



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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

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Incidence function

$$g(E) = N\beta ph(E) \tag{1}$$

where

- ► 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")

• h(E) probability for an exposed susceptible to get the infection Typically, we would assume p and β independent of E and H and h to be saturating To ensure (1) satisfies these conditions, we can assume

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

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The model



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

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Let

$$\mathcal{R}_0 = \frac{g'_+(0)c_H}{d_E\gamma_H} \tag{3}$$

Theorem 1

- If $0 < \mathcal{R}_0 < 1$, then (2) 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 (1) takes the form

$$p(t) = p(t + \omega) > 0, \quad t \in \mathbb{R}$$
 (4)

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

$$g(t,E) = p(t)h(E)$$
(5)

with h having the properties prescribed earlier. Letting

$$p_{\min} := \min_{0 \le t \le \omega} p(t), \quad p_{\max} := \max_{0 \le t \le \omega} p(t)$$
(6)

then we require that

$$\lim_{z \to \infty} \frac{g(z)}{z} < \frac{d_E \gamma_H}{c_H p_{max}}$$
(7)

Let

$$\mathcal{R}_{0}^{min} = \frac{c_{H} \rho_{min} h'_{+}(0)}{d_{E} \gamma_{H}}, \quad \mathcal{R}_{0}^{max} = \frac{c_{H} \rho_{max} h'_{+}(0)}{d_{E} \gamma_{H}}$$
(8)

Theorem 2

- If $0 < \mathcal{R}_0^{max} < 1$, then (2) with incidence (5) always goes to extinction
- If R₀^{min} > 1, then a unique nontrivial periodic endemic state exists for (2) with incidence (5)

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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



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The full model

$$S'_{A} = -\beta_{A}S_{A}I_{A} + \gamma_{A}I_{A}$$
 (9a)

$$I'_{A} = \beta_{A} S_{A} I_{A} - \gamma_{A} I_{A}$$
(9b)

$$W' = \alpha I_{\mathcal{A}} - \eta W(S_{\mathcal{H}} + I_{\mathcal{H}}) - \mu W$$
(9c)

$$S'_{H} = -\rho\eta W S_{H} - \beta_{H} S_{H} I_{H} + \gamma_{H} I_{H}$$
(9d)

$$I'_{H} = \rho \eta W S_{H} + \beta_{H} S_{H} I_{H} - \gamma_{H} I_{H}$$
(9e)

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, (9) can be simplified:

$$I'_{A} = \beta_{A} N_{A} I_{A} - \gamma_{A} I_{A} - \beta_{A} I^{2}_{A}$$
(10a)

$$W' = \alpha I_{\mathcal{A}} - \eta W N_{\mathcal{H}} - \mu W \tag{10b}$$

$$I'_{H} = \rho \eta W (N_{H} - I_{H}) + \beta_{H} N_{H} I_{H} - \gamma_{H} I_{H} - \beta_{H} I_{H}^{2}$$
(10c)

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

$$\mathcal{R}_{0A} = \frac{\beta_A}{\gamma_A} N_A$$
 and $\mathcal{R}_{0H} = \frac{\beta_H}{\gamma_H} N_H$ (11)

- ▶ DFE LAS if $R_{0A} < 1$ and $R_{0H} < 1$, unstable if $R_{0A} > 1$ or $R_{0H} > 1$
- If R_{0H} > 1 and R_{0A} < 1, (10) goes to EP with endemicity only in humans
- \blacktriangleright Endemic EP with both A and H requires $\mathcal{R}_{0\text{A}}>1$ and $\mathcal{R}_{0\text{H}}<1$

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)

$$i'_{H} = \alpha Ni_{S} - \gamma i_{H}$$
(12a)
$$i'_{S} = \beta Hi_{H}(1 - i_{S}) - \mu_{2}i_{S}$$
(12b)

α number of schistosomes produced per person per infected snail per unit time

- $\blacktriangleright~1/\gamma$ average life expectancy of a schistosome
- ▶ $1/\mu_2$ average life expectancy of an infected snail
- $\blacktriangleright \beta$ transmission parameter

Let the basic reproductive rate for schistosomes be

$$\mathcal{R}_0 = \frac{\alpha N \beta H}{\gamma \mu_2} \tag{13}$$

(12) has two EP

($i_{H}^{\star}, i_{S}^{\star}$) = (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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Extending the model

Interval between infection of a snail and onset of patency (release of cercariae) is *prepatent* or *latent* period

$$i'_{H} = \alpha N i_{S} - \gamma i_{H} \tag{14a}$$

$$\ell_{S}^{\prime} = \beta H i_{H} (1 - \ell_{S} - i_{S}) - \sigma \ell_{S} - \mu_{1} \ell_{S}$$
(14b)

$$i'_{S} = \sigma \ell_{S} - \mu_{2} i_{S} \tag{14c}$$

1/σ average duration of prepatent period
 f = σ/(σ + μ₁) fraction of infected snails surviving prepatent period

The basic reproductive rate for schistosomes is now

$$\mathcal{R}_0 = f \frac{\alpha N \beta H}{\gamma \mu_2} \tag{15}$$

(14) has endemic EP

$$(i_{H}^{\star}, i_{S}^{\star}) = \left(\frac{\alpha N\sigma}{\gamma(\sigma + \mu_{2})} - \frac{\mu_{2}(\sigma + \mu_{1})}{\beta H(\sigma + \mu_{2})}, \frac{\sigma}{\sigma + \mu_{2}}\left(1 - \frac{1}{\mathcal{R}_{0}}\right)\right)$$

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Also has models

- where snails lose infectiousness (assumed to happen sometimes)
- with larval population dynamics
- single variable models
- human immigration and emigration
- reservoir hosts

Really worth a read

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Heterogeneities in contact rates

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



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

$$I'_{i} = \alpha \left(\sum_{j} \eta_{ij} N_{i} S_{j}\right) - \gamma I_{i}$$
(16a)
$$i'_{Sj} = \beta \left(\sum_{i} \eta_{ij} I_{i}\right) (1 - i_{Sj}) - \mu_{2} i_{Sj}$$
(16b)

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

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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

$$s'_{p} = \mu - \lambda_{p} s_{p} - \mu s_{p}$$
 (17a)

$$i'_{p} = -\gamma i_{p} + \lambda_{p} s_{p} - \mu i_{p} \tag{17b}$$

$$r'_{p} = \gamma r_{p} - \mu r_{p} \tag{17c}$$

$$w_p' = \xi(i_p - w_p) \tag{17d}$$

with force of infection

$$\lambda_{\rho} = \beta_{i_{\rho}}i_{\rho} + \beta_{W_{\rho}}w_{\rho} + \sum_{q=1}^{10}\theta_{\rho q}i_{q}$$
(17e)

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

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

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Days Since 21 October 2010