

Modelling Macroparasitic Diseases

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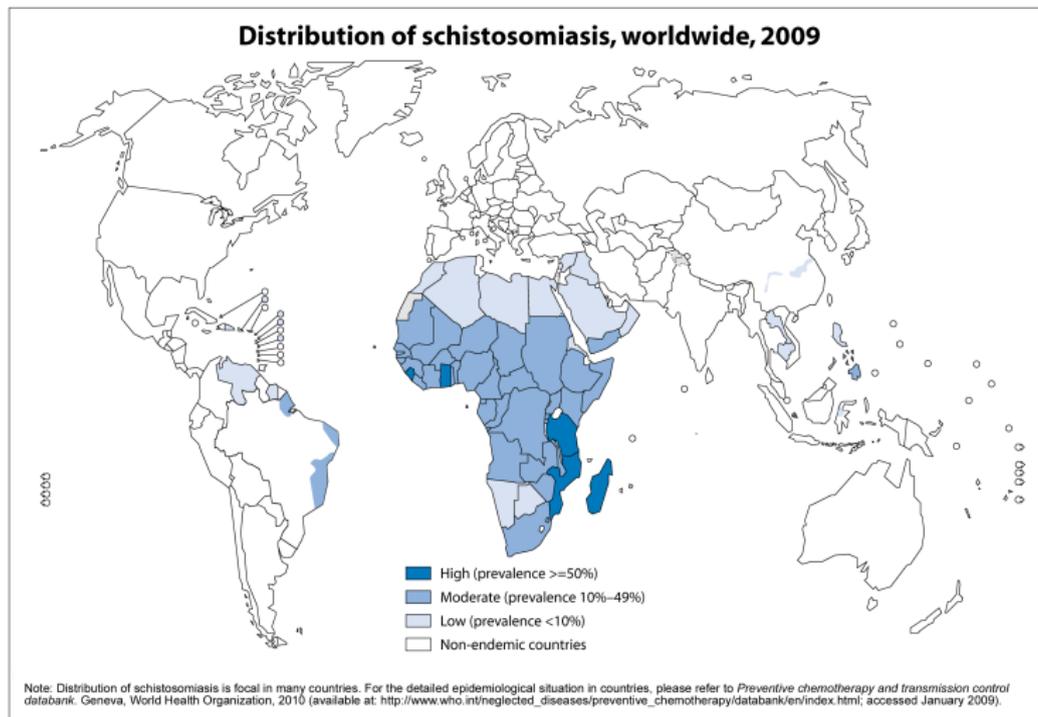
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- We will focus on schistosomiasis, which effects around 300 million people worldwide, contributing to around 200,000 deaths per year.
- Symptoms include abdominal pain, blood in urine/stool, diarrhoea. Long term it causes damage to internal organs, anaemia, stunted growth, reduced cognitive ability.

Schistosomiasis distribution



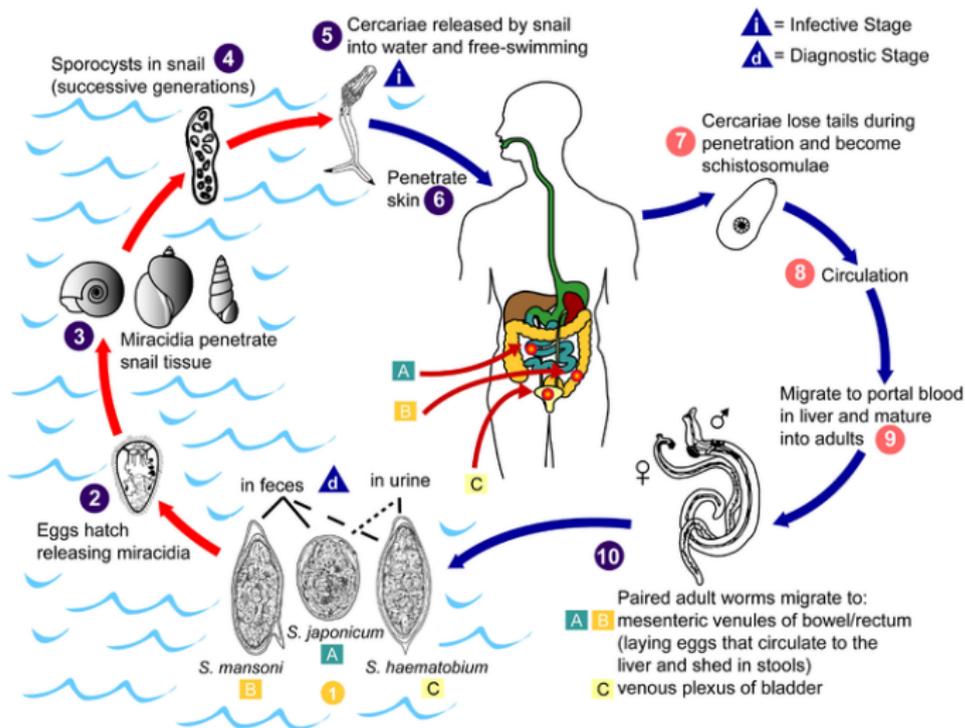
The boundaries and names shown and the designations used on this map do not imply the expression of any opinion whatsoever on the part of the World Health Organization concerning the legal status of any country, territory, city or area or of its authorities, or concerning the delimitation of its frontiers or boundaries. Dotted lines on maps represent approximate border lines for which there may not yet be full agreement. © WHO 2010. All rights reserved

Data Source: World Health Organization
Map Production: Control of Neglected
Tropical Diseases (NTD)
World Health Organization



Schistosomiasis life-cycle

Schistosomiasis



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- We cannot use standard infectious disease models (e.g. SIR). It is not enough to classify an individual as either infected or not.
- Worm *burden* as well as infection needs to be considered. How do we do this?
- Lets start off by considering a population of human hosts of constant size N .
- Let $p_i(t)$ denote the probability that a human host is carrying i adult worms at time t
- The mean adult worm burden is just $M(t) = \sum ip_i(t)$

Modelling - Initial considerations

- Humans are infected by juvenile worms. Supposing there are $L(t)$ of them, the rate of contact/infection per person is β , and they take a time τ_1 to become adults during which they die at rate d_1 , then

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- Lets assume that humans die with a rate μ (killing worms inside), and that worms die within humans with rate μ_1

$$\frac{dM}{dt} = \beta L(t - \tau_1) d_1 - \mu \sum i p_i(t) - \sum \mu_1(i) i p_i(t)$$

Modelling - Initial considerations

- Now let's consider L . If ϕ is the probability that a female worm is mated, s is the proportion of female worms in the population, $\lambda(i)$ is the per capita egg production rate of female worms, and τ_2 is the time taken for eggs to reach juvenile age during which they die at rate d_2 , then we get

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- If infective juvenile worms die at rate μ_2 , and infect humans with rate βN then

$$\frac{dL}{dt} = s\phi Nd_2 \sum \lambda(i)ip_i(t - \tau_2) - \mu_2 L - \beta NL$$

- This gives us the basic model

$$\begin{aligned}\frac{dM}{dt} &= \beta L(t - \tau_1) d_1 - \mu \sum i p_i(t) - \sum \mu(i) i p_i(t) \\ \frac{dL}{dt} &= s \phi N d_2 \sum \lambda(i) i p_i(t - \tau_2) - \mu_2 L - \beta N L\end{aligned}$$

Empirical distribution of parasites among hosts

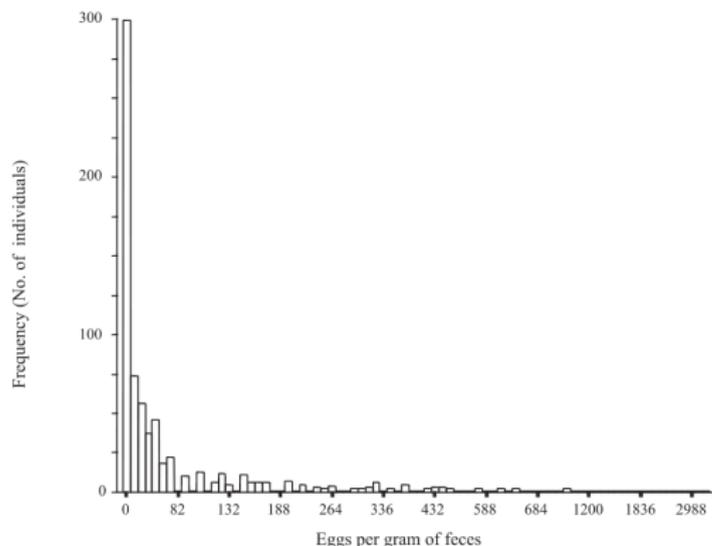


Figure 1 Distribution of *S. mansoni* eggs in the study sample expressed in eggs per gram of faeces ($n = 597$).

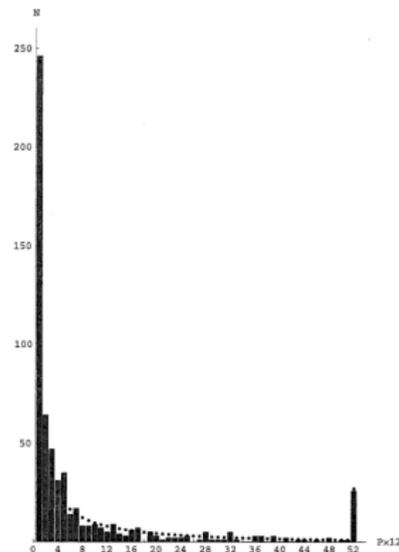
- Distribution of *Schistosoma Mansoni* burden (measured as eggs per gram of faeces) among individuals from a village in Brazil (*Bethony et al., 2001*)

Empirical distribution of parasites among hosts

- Fit of data with negative-binomial distribution (Feng et al 2004)
- Negative binomial is a discrete probability distribution, parameterised by mean m and aggregation parameter k , defined by

$$p(i) = \frac{(k + i - 1)!}{i!(k - 1)!} (1 + m/k)^{-k-i} (m/k)^i$$

For this data $\bar{x} = 9.5 \times 12$,
 $k = 0.24$.



- First, let's over-simplify our model by neglecting the density dependent effects of worm death and egg production i.e. $\mu_1(i) = \mu_1$, $\lambda(i) = \lambda$, and setting the probability of worm mating $\phi = 1$. Then we get

$$\begin{aligned}\frac{dM}{dt} &= \beta L(t - \tau_1) d_1 - (\mu + \mu_1) M(t) \\ \frac{dL}{dt} &= s N d_2 \lambda M(t - \tau_2) - (\mu_2 + \beta N) L(t)\end{aligned}$$

Basic reproduction rate, R_0

- It can be shown that this pair of equations leads to ever increasing M and L under the condition

$$R_0 := \frac{s\lambda\beta Nd_1d_2}{(\mu + \mu_2)(\mu_2 + \beta N)} > 1$$

- If $R_0 < 1$ then the worm population will die out.
- Alternatively, we can reformulate this as a threshold population size for humans above which the number of hosts is able to support a worm population

$$N_T = \frac{\mu_2(\mu + \mu_1)/\beta}{s\lambda d_1 d_2 - (\mu + \mu_1)}$$

Density dependent fecundity

- Without any density dependent effects the basic model does not produce a stable non-zero equilibrium.
- Empirical evidence suggests that there is a decline in per capita fecundity of female worms, and it can be modelled approximately by the relationship

$$\lambda(i) = \lambda_0 \exp(-\gamma(i - 1))$$

- Assuming a negative binomial distribution, and subbing this into $\sum \lambda(i)ip_i$ means that λ is replaced in the simplified model with

$$\lambda \rightarrow \lambda_0 (1 + M(1 - z)/k)^{-(k+1)}$$

where $z = e^{-\gamma}$

Some more simplifying

- The maturation times are short compared to the life of a worm, so lets set $\tau_1 = \tau_2 = 0$.
- It is reasonable to assume that the dynamics of the juvenile worms occur on much shorter time-scales than the dynamics of mature adults, so L will be very close to its equilibrium value obtained by setting $dL/dt = 0$:

$$L(t) = sd_2\lambda NM(t)/(\mu_2 + \beta N)$$

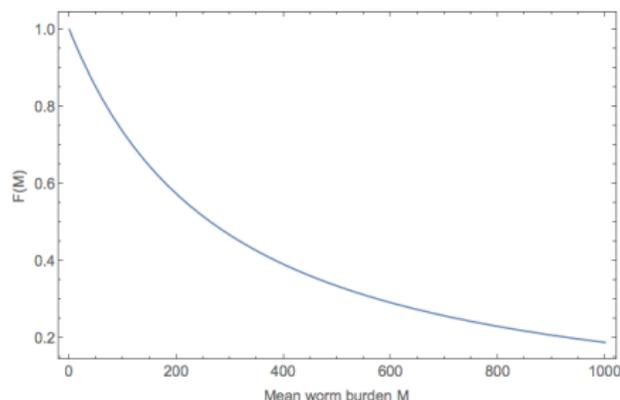
- Which results in a single ODE for $M(t)$

$$\frac{dM}{dt} = (\mu + \mu_1)(R_0 f(M) - 1)M(t)$$

- where $f(M) = (1 + M(1 - z)/k)^{-(k+1)}$

Equilibrium M^*

This is what f looks like with parameters $k = 0.25$, $\gamma = 0.0003$.



- The non-zero equilibrium mean worm burden for this model is given by $R_0 f(M^*) = 1$:

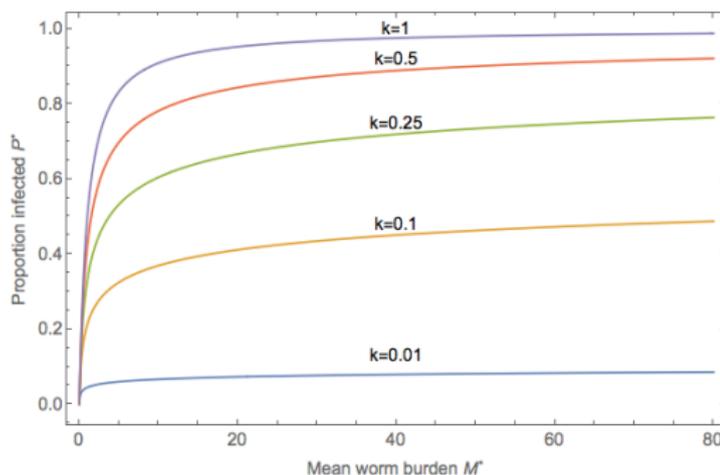
$$M^* = k(R_0^{1/(k+1)} - 1)/(1 - z)$$

which is stable iff $R_0 > 1$

Proportion of people infected

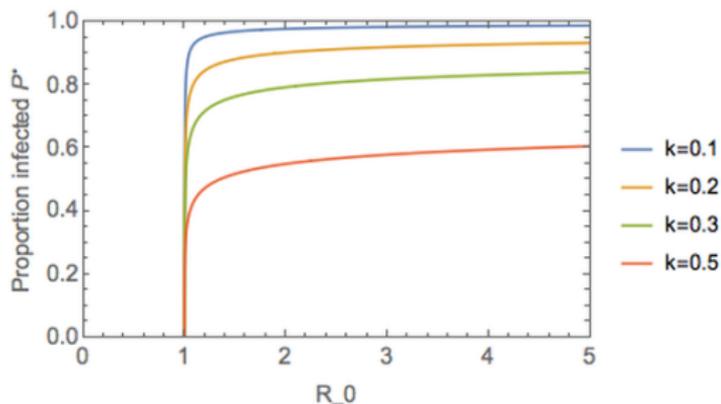
- Recalling that we are assuming a negative-binomial distribution of worms across the host population, the proportion of people infected (the prevalence) is just

$$P = 1 - (1 + M/k)^{-k}$$



Proportion of people infected

- We can also see the effect that R_0 has on the equilibrium prevalence.



In order to eradicate or control the disease, there are a few options we could look at:

- Change behaviour (hard)
- Kill snails (hard)
- Mass drug administration programs
- Vaccinate (no vaccine available yet)

The effect of chemotherapy

- There is a safe and effective drug, Praziquantel, that is able to reduce worm burden by 90%.
- The pharma company Merck is currently donating 250 million doses per year to the WHO
- Let's suppose we have a drug which we are able to administer randomly to a proportion g of the population per unit time, with efficacy h (the proportion of worm burden killed by a single dose). Then the death rate will increase by

$$c = -\ln(1 - gh)$$

The effect of chemotherapy

- With this additional death rate

$$\frac{dM}{dt} = (\mu + \mu_1)(R_0 f(M) - 1 - c/(\mu + \mu_1))M(t)$$

which changes the equilibrium to

$$M^* = k \left(\left(\frac{R_0}{A \ln(1 - gh)} \right)^{1/(k+1)} - 1 \right) / (1 - z)$$

- $A = 1/(\mu + \mu_1)$ is the life expectancy of the adult worm (without drug treatment)

The effect of chemotherapy

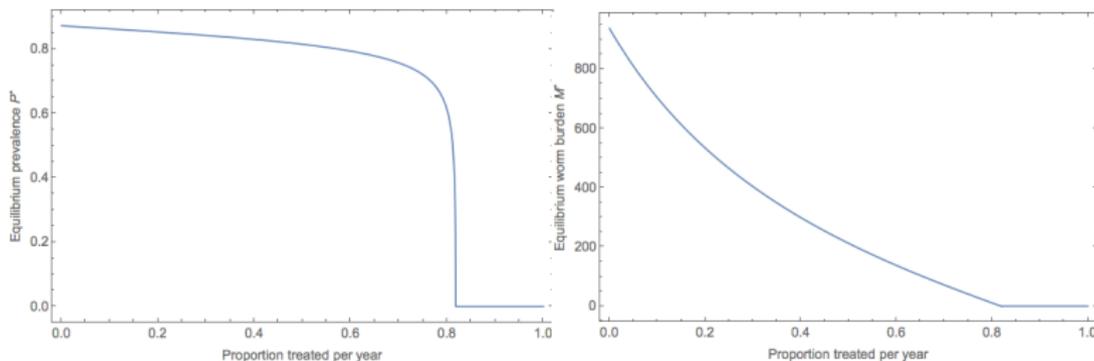
- This has effectively modified the basic reproductive rate R_0 . For the drug treatment program to eradicate the disease we need this rate to be less than one.
- Expressed in terms of a threshold proportion of the population per unit time needed to treat, this is:

$$g_c = \left(1 - e^{(1-R_0)/A}\right) / h$$

- For an area badly effected area by schistosomiasis, we might expect $R_0 = 5$, adult worm life-span $A = 3$ years, and praziquantel has an efficacy of $h = 0.9$ so we would need to treat 82% of the population.

The effect of chemotherapy

- Here we have the equilibrium prevalence and mean burden as a function of proportion treated ($R_0 = 5$, $z = 0.99$, $A = 3$, $k = 0.25$, $h = 0.9$).



Some extra stuff

- We've been assuming that the probability of a female worm being mated $\phi = 1$.
- In reality, the chance of a randomly selected female being mated will be dependent on the density of worms in the population, and also the aggregation k .
- Can you think of a reasonable functional form for ϕ ? How will this affect the long term behaviour?

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