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**School on Modelling Tools and Capacity Building in Climate and Public Health**

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**An agent-based model of schistosomiasis transmission and water temperature**

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# An agent-based model of schistosomiasis transmission and water temperature

Nicky McCreesh





# Outline

- 1. Agent-based vs population-based models.**
2. My schistosomiasis model.

# Population-based models (aka compartmental)

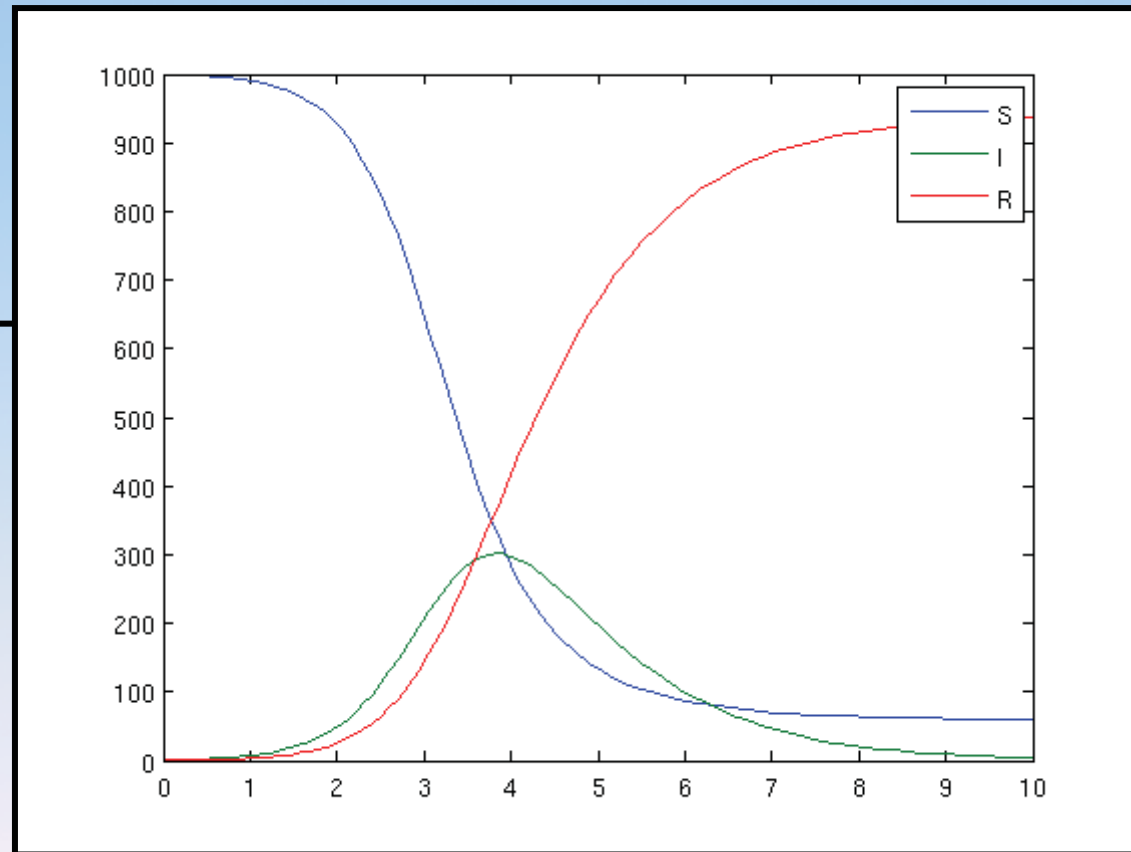
- Model populations of individuals (e.g. susceptibles, infecteds, recovered).
- Mutually exclusive groups.
- Homogeneity within groups.
- Can subdivide to create more groups:
  - Sex
  - Age
  - Location (“metapopulation” or “patch” models)
- Tend to be deterministic.



- $dS/dt = -\beta SI$

- $dI/dt = \beta SI$

- $dR/dt = \gamma I$

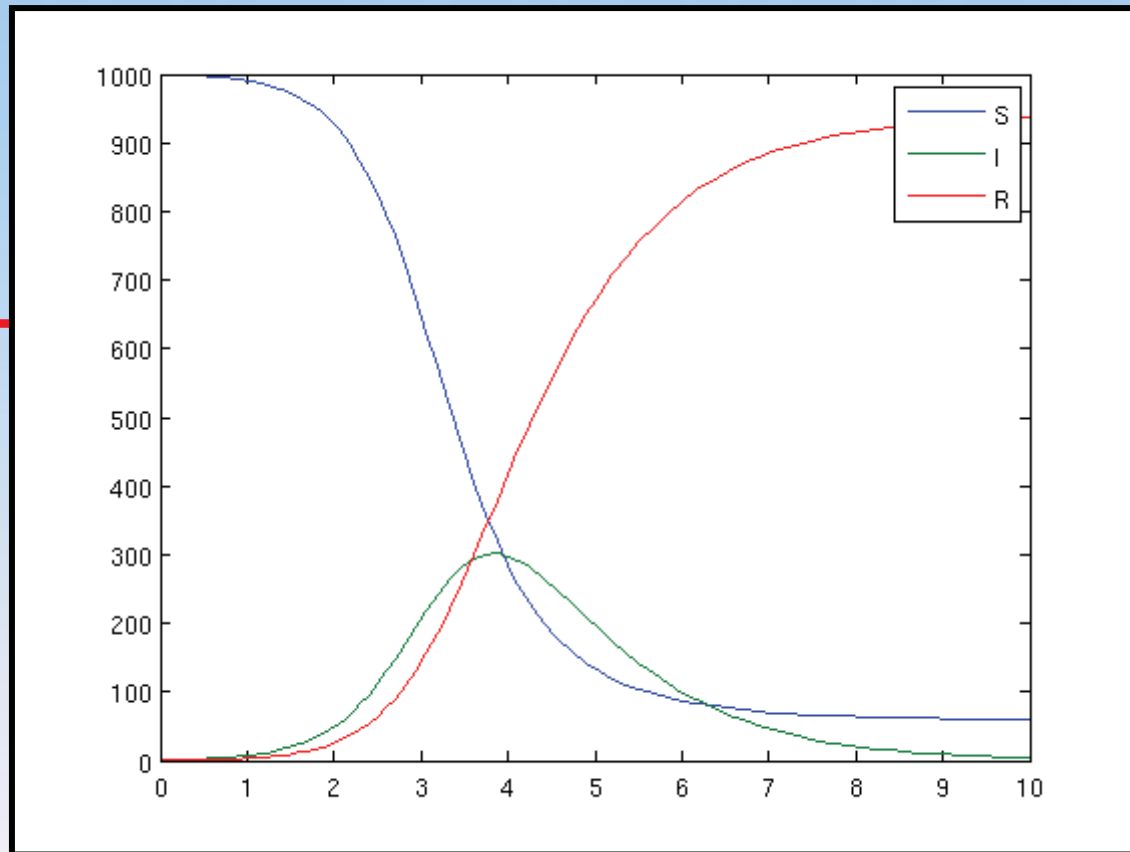




- $dS/dt = -\beta SI$

- $dI/dt = \beta SI$

- $dR/dt = \gamma I$



# Agent-based models

(aka individual-based, microsimulation)

- Models individuals in the population.
- Each individual has:
  - An ID.
  - One or more states (e.g. age, sex, infection status, immunity).
- Characteristics of each individual are tracked through time.
  - Some do not change (e.g. sex).
  - Others do (e.g. infection status).
- Tend to be stochastic.
- Can include more information at the individual level.
- Have much more control over characteristics of individuals (e.g. prepatent periods).
- Can be much slower to run and require more computing power.

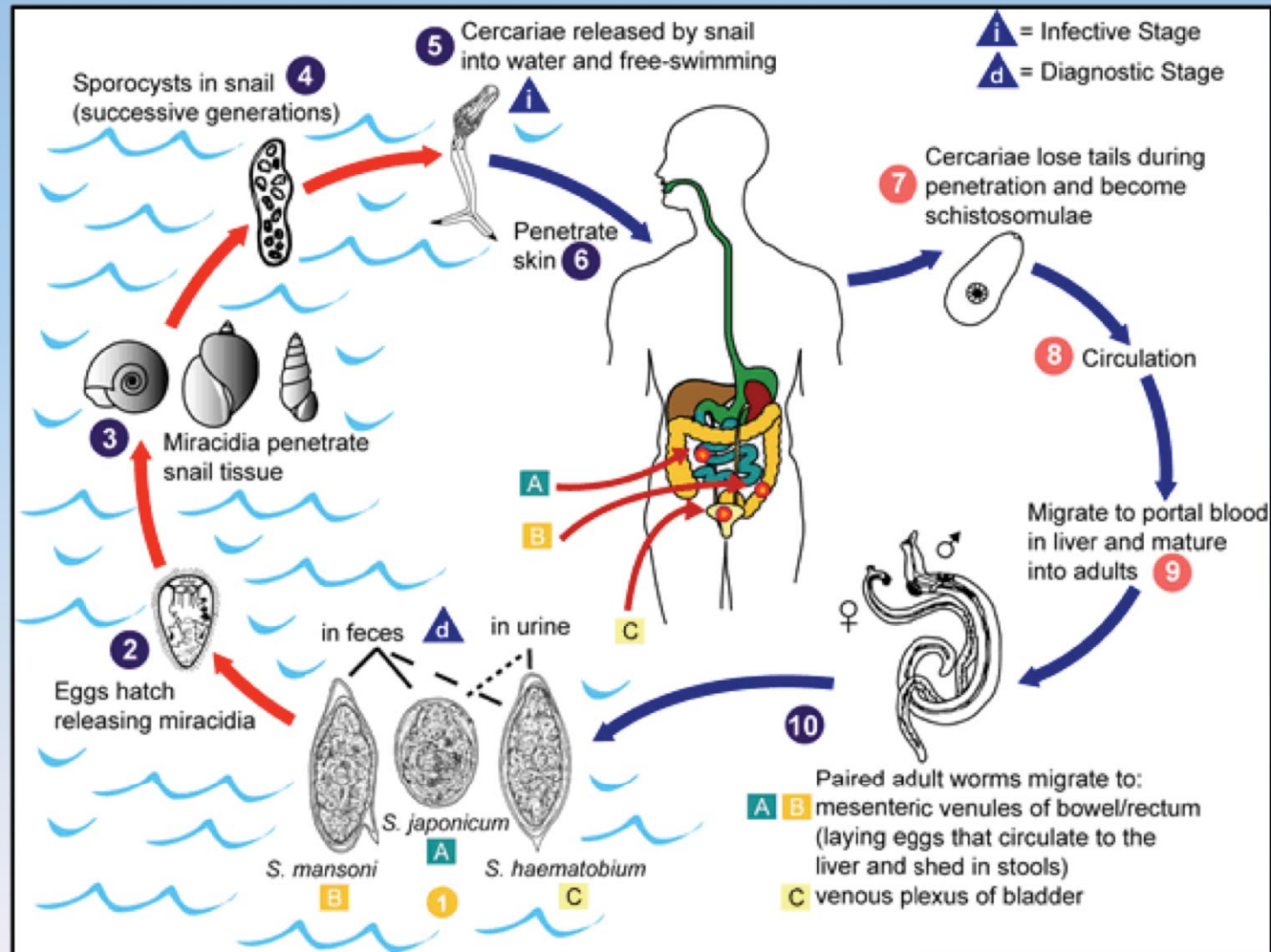


# Outline

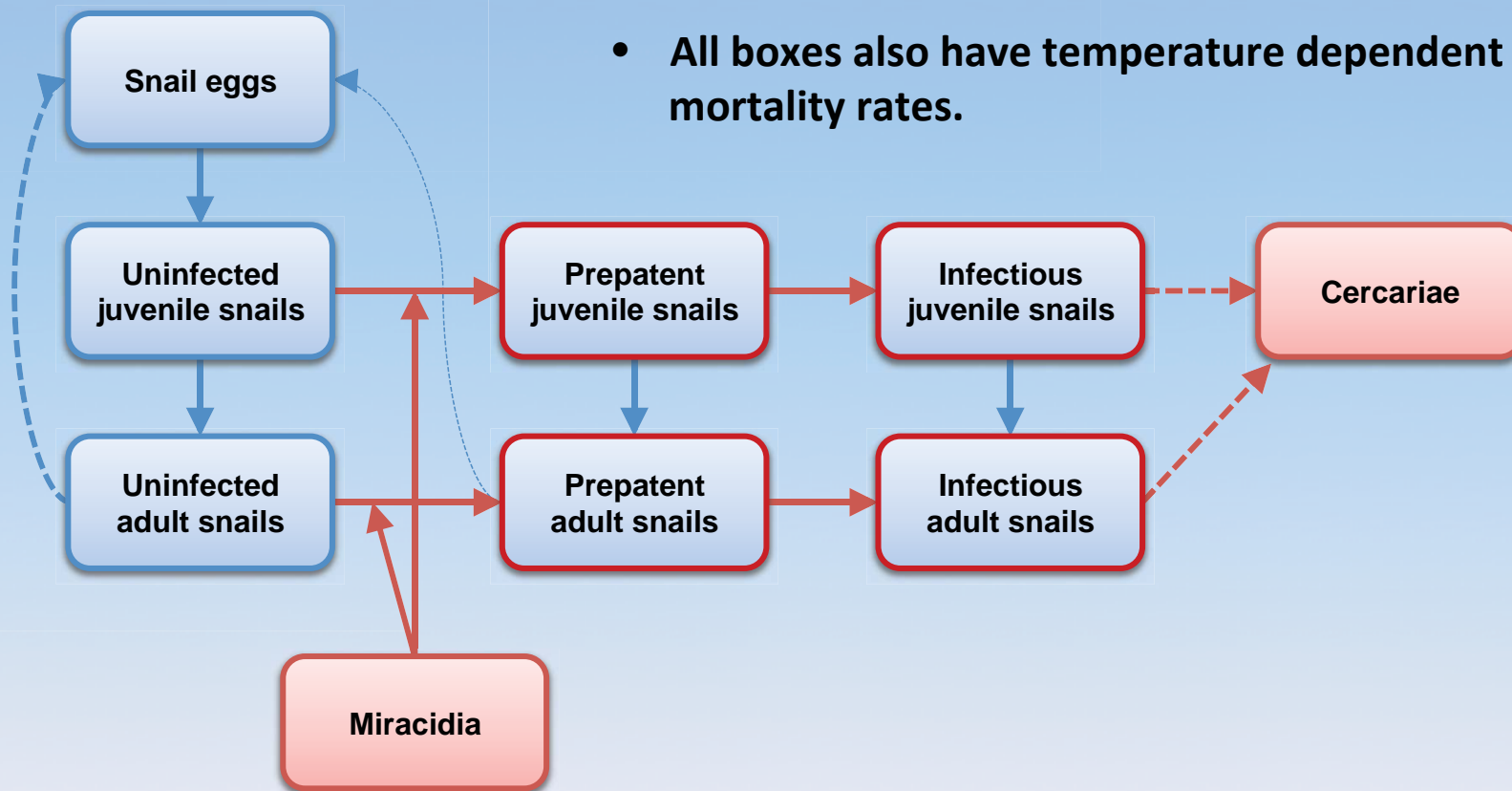
1. Agent-based vs population-based models.
- 2. My schistosomiasis model.**



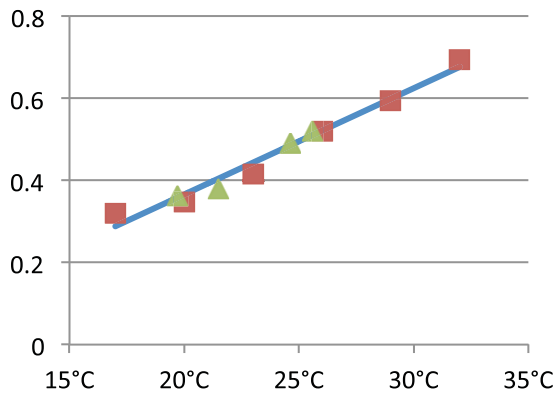
# Schistosoma lifecycle



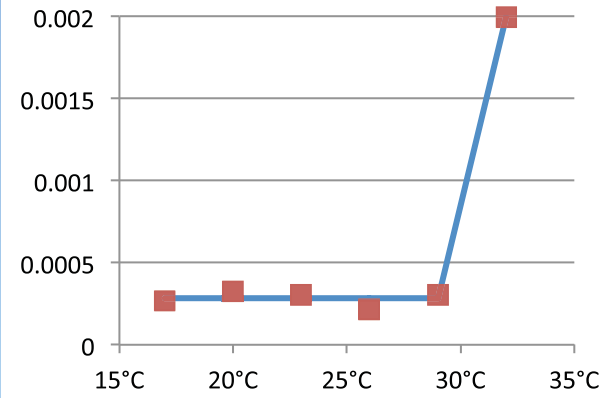
- All arrows indicate temperature dependent rates.
- All boxes also have temperature dependent mortality rates.



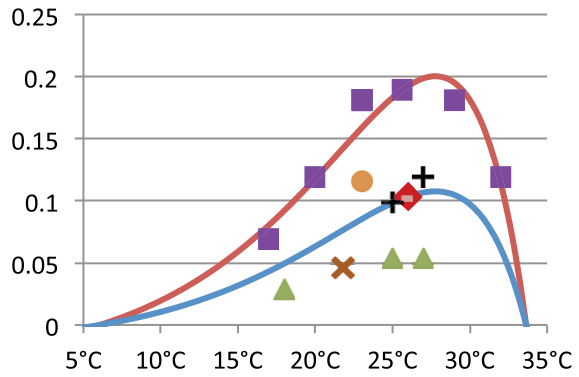
**Rate of egg development**



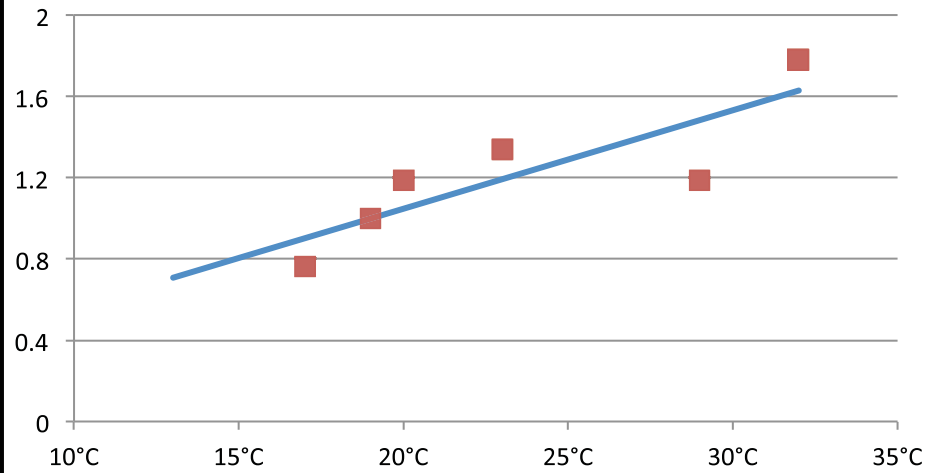
**Egg mortality rates**

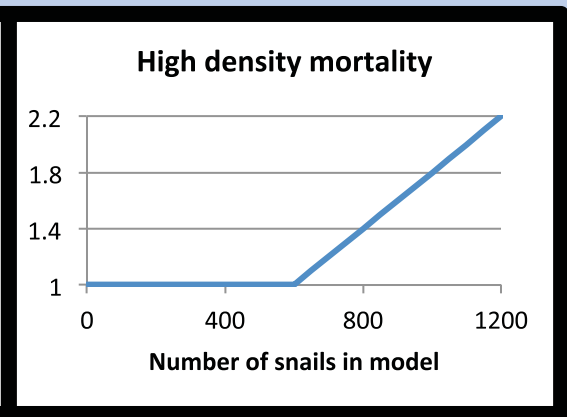
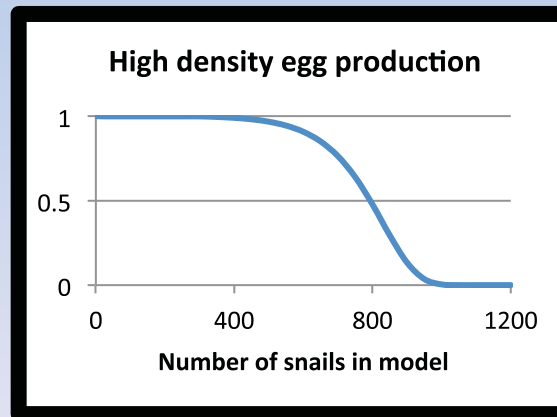
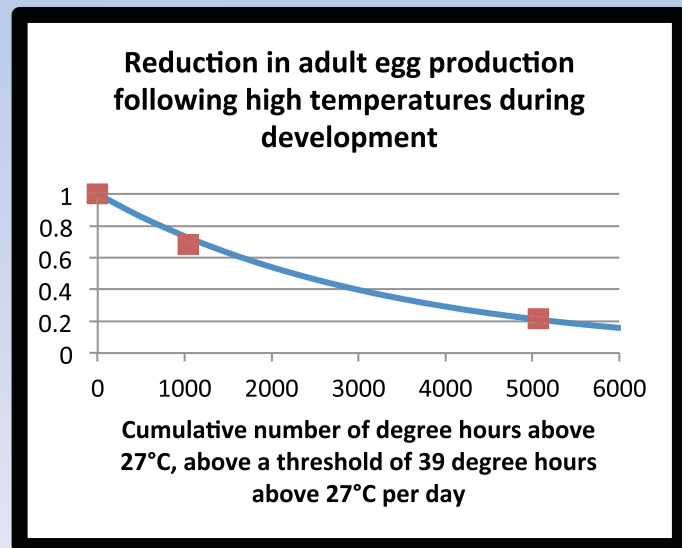
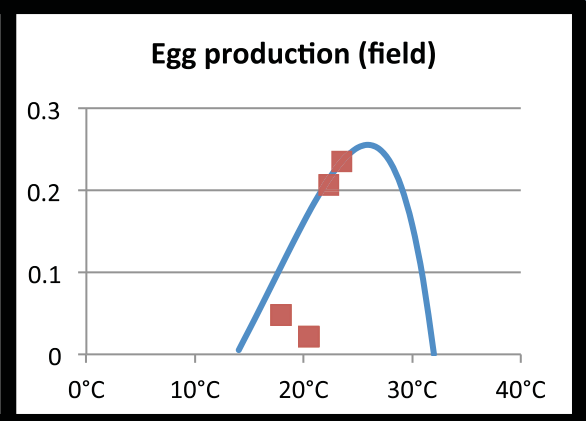
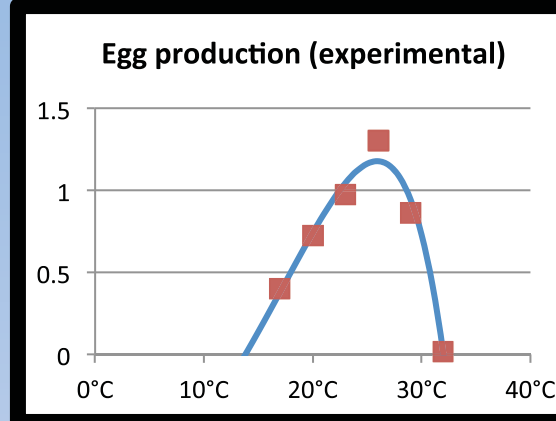
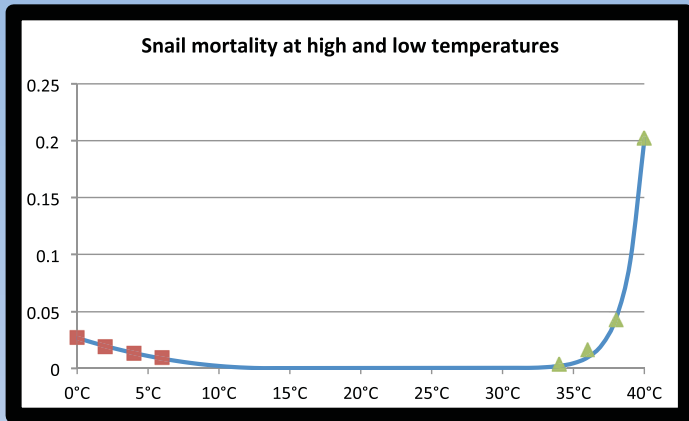


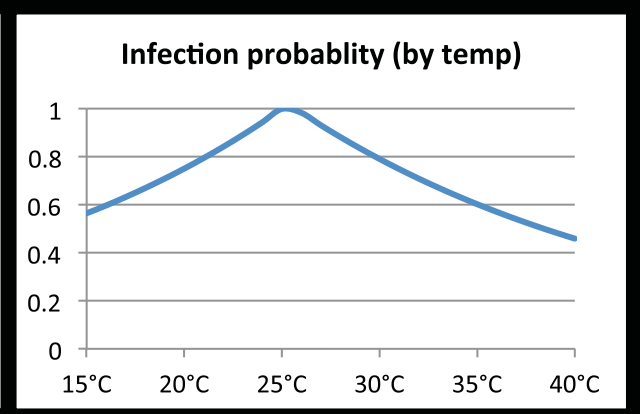
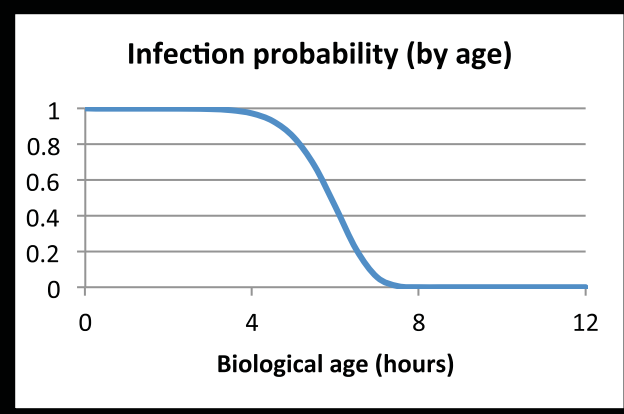
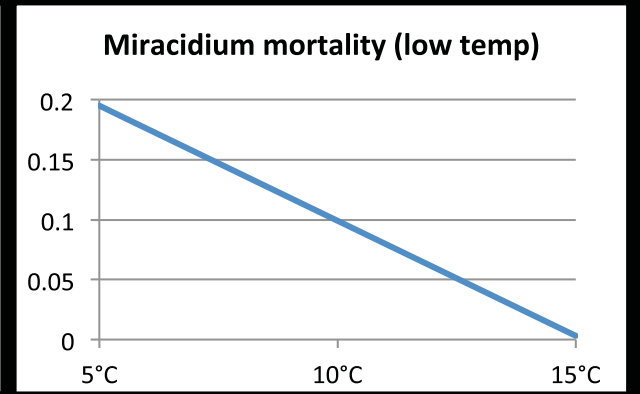
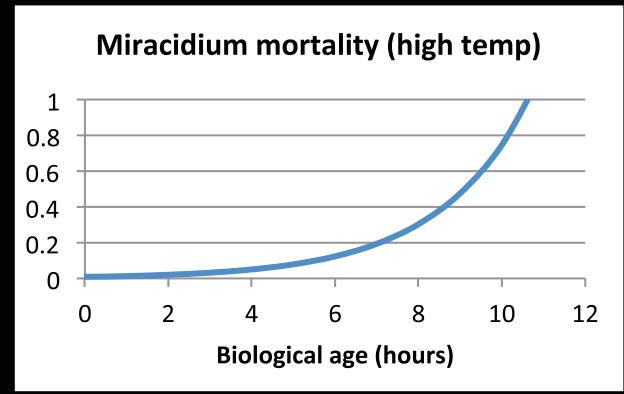
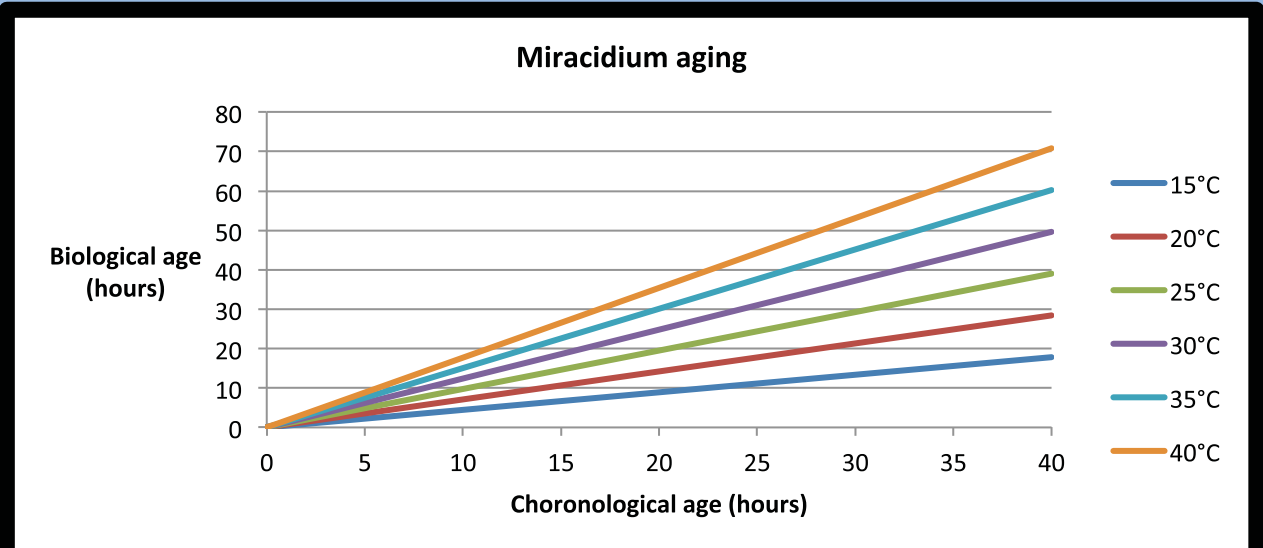
**Juvenile development rate**

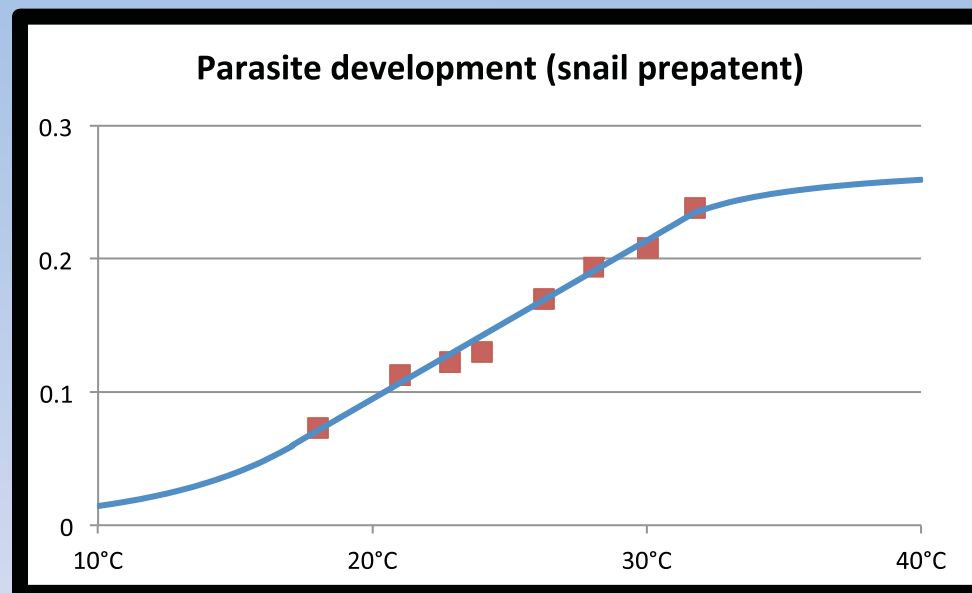


**Snail mortality relative to mortality at 19°C**

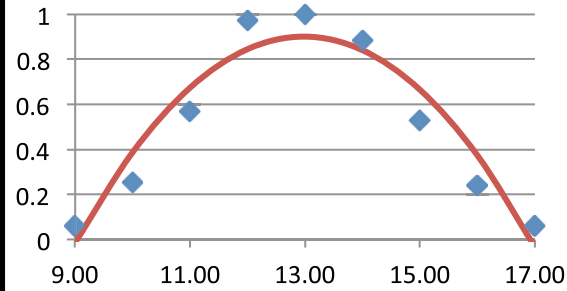




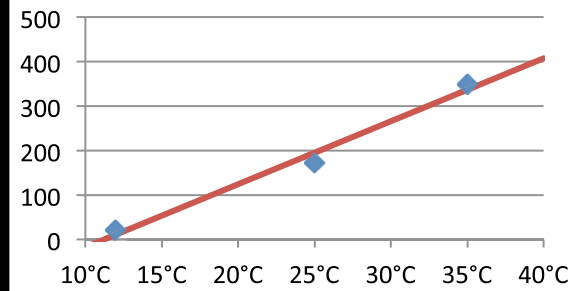




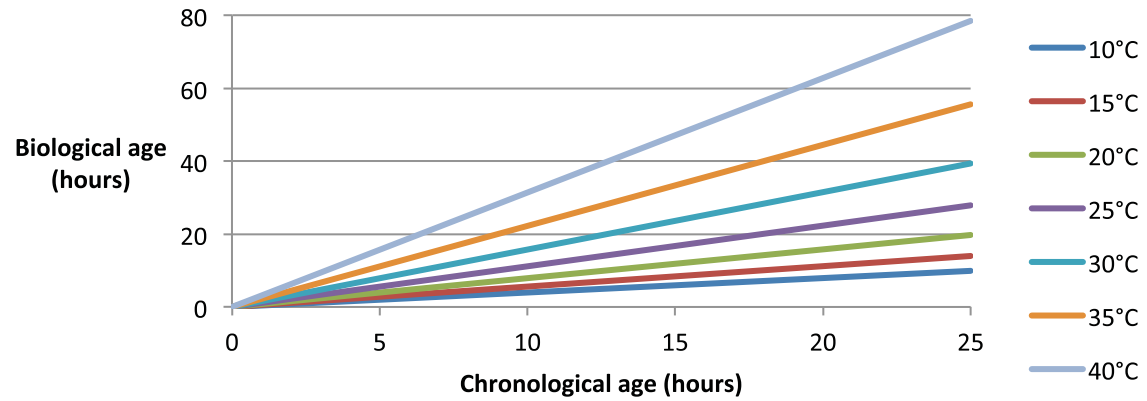
Cercaria production (time of day)



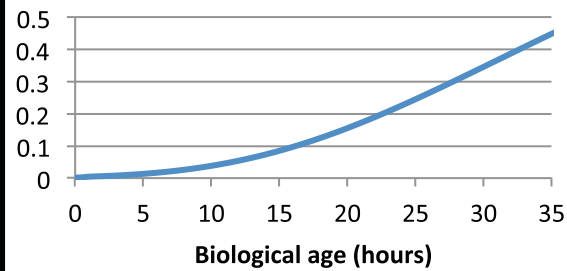
Cercaria production (temp)



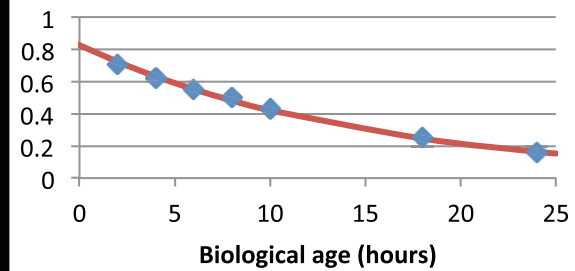
Cercaria aging

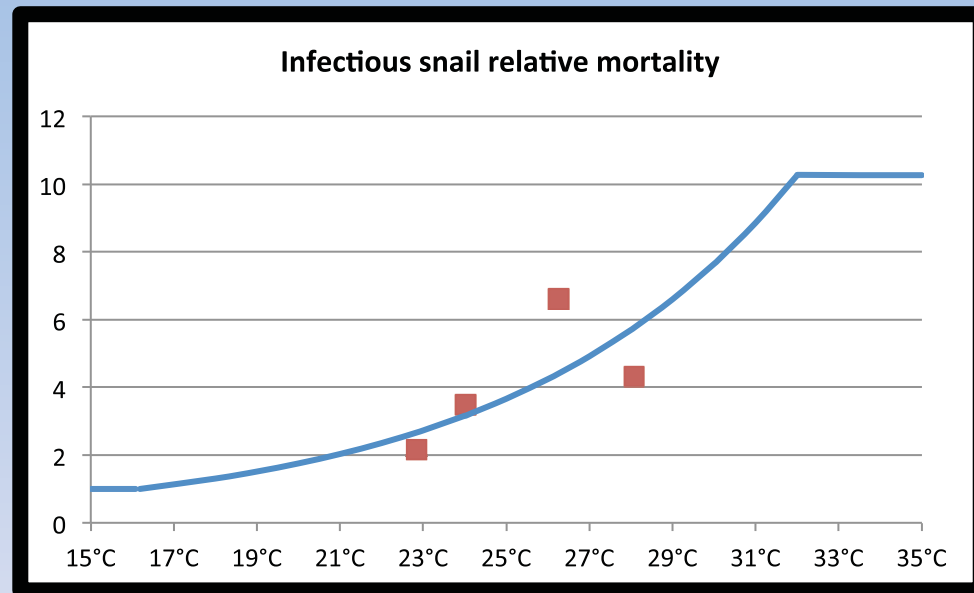


Cercaria mortality



Infection probability



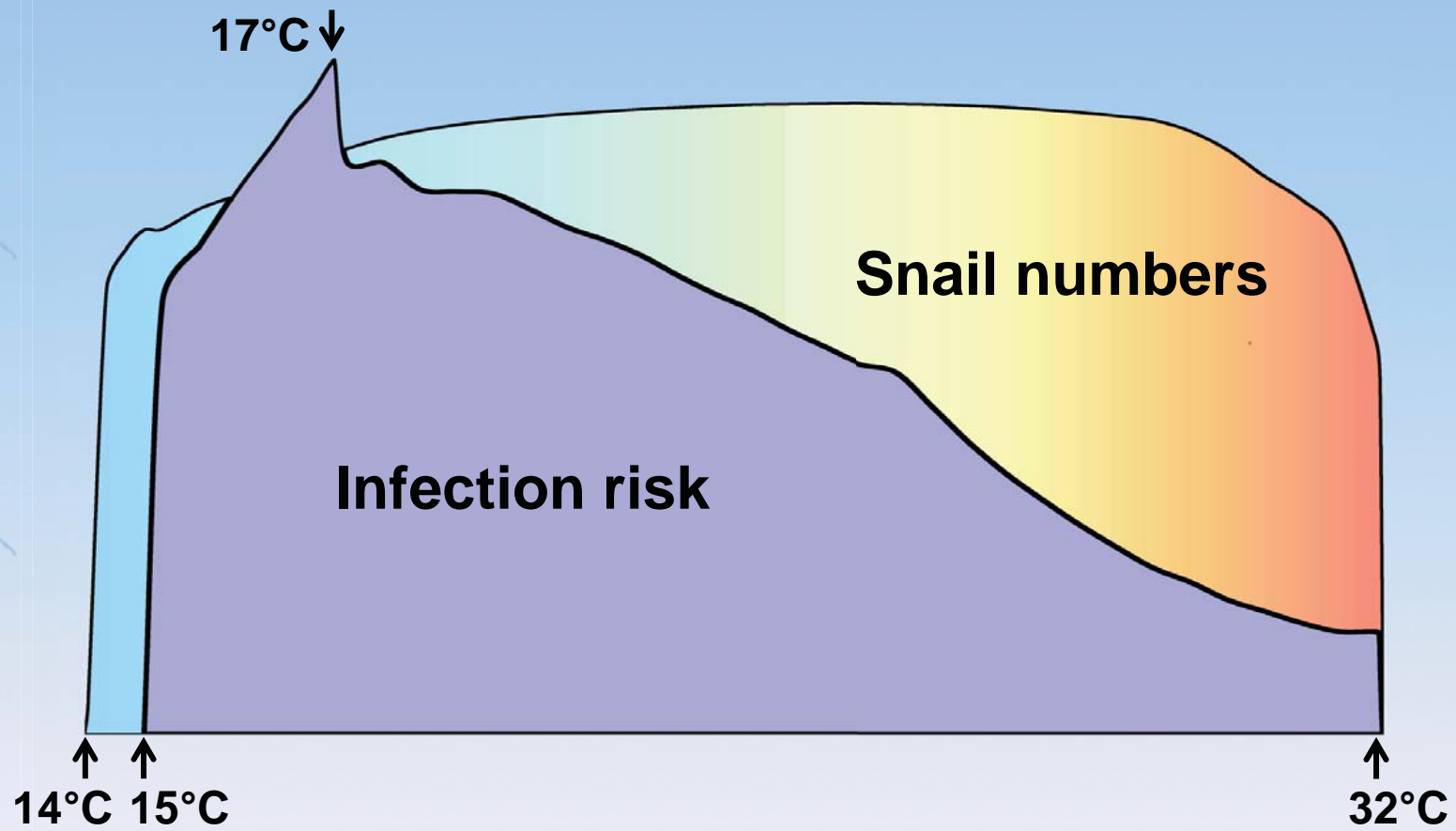




## Methods

- The model was run using constant water temperatures and results were averaged over 200 runs.
- Infection risk is measured as the mean number of cercariae in the model, adjusted for their decreasing probability of causing infection with increasing biological age.

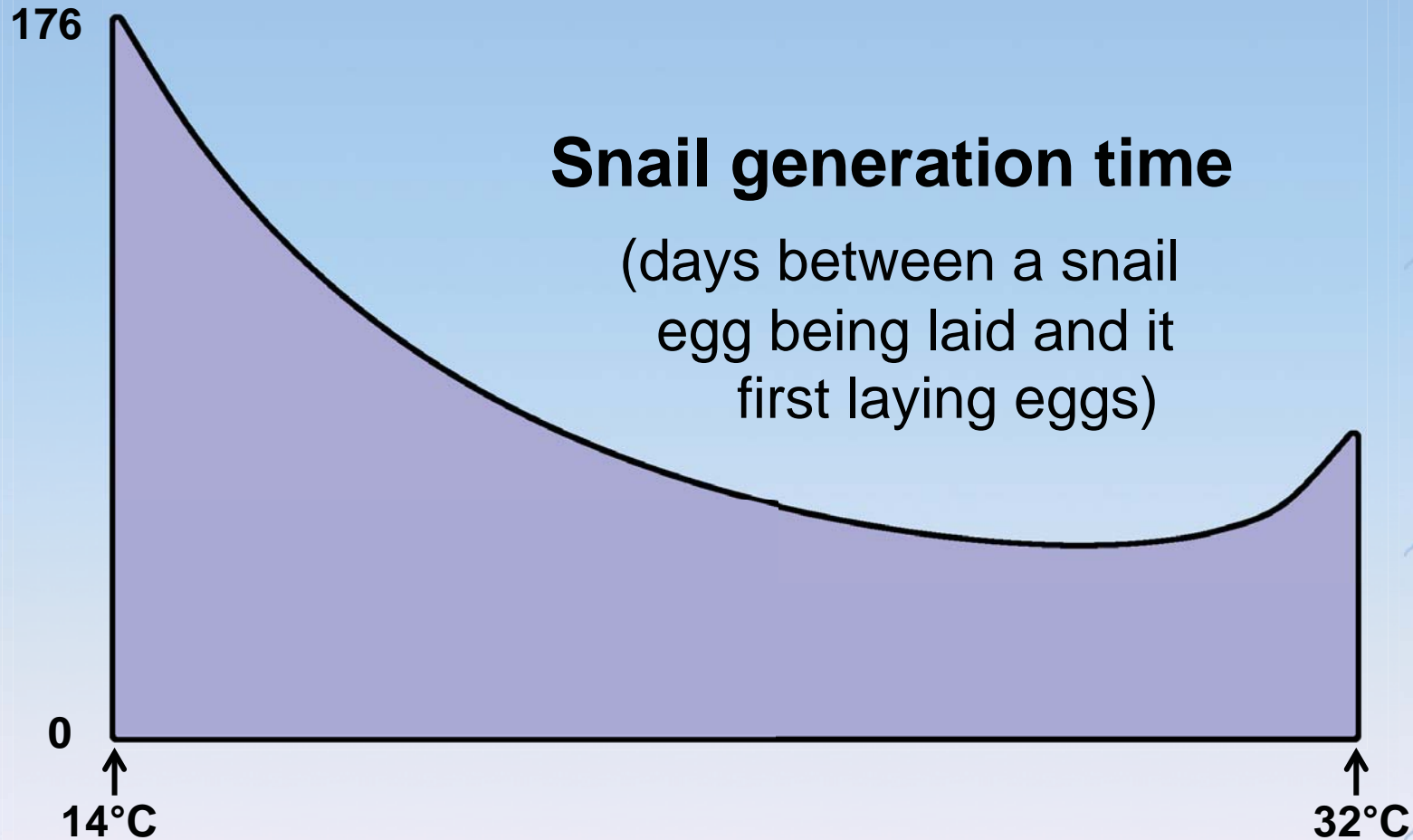
# Results



# Results

## Snail generation time

(days between a snail egg being laid and it first laying eggs)



# Conclusions

- Our results suggest that in most areas where *S. mansoni* is found, infection risk will decrease slightly as temperatures increase, but the difference will not be substantial.
- Snail generation times will decrease however, meaning that snail populations will recover faster from reductions in their numbers.
- In areas where *B. pfeifferi* snails are currently found but where low temperatures prevent sustained transmission, infection risk may increase dramatically over coming years and decades leading to epidemics of schistosomiasis.
- These areas will fall outside current control programs and people at risk may have little or no knowledge of or immunity to schistosomiasis.
- There is an urgent need for these areas to be monitored to minimise the impact of future epidemics.

## What next?

- Incorporate real current and predicted future temperature data into the model.
- Produce relative risk maps.