# Introduction to MCMC for Infectious Diseases

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# Long history of modeling infectious disease transmission





# Given data, what can we say about our model parameters?



time

#### Resource for Bayesian statistics



#### Bayesian inference

- **Observed data:** e.g., number of observed infections (D)
- **Probability model:** how is the observed data generated?

- Model is parameterized by  $\ \theta \in \Theta$ 

- Fitting the model: we want to find  $\theta$  that have high probability given our observed data
  - We want  $P(\boldsymbol{\theta}|\boldsymbol{D})$  to be large
  - i.e., we want to find "highly plausible"  $\theta$  given our data

# Bayesian updating

• Bayes' rule tells us how to compute these "plausibilities":

$$P(\boldsymbol{\theta}|D) = \frac{P(D|\boldsymbol{\theta})P(\boldsymbol{\theta})}{P(D)}$$

- $P(\boldsymbol{\theta})$  is the **prior probability**
- $P(D|\theta)$  is the **likelihood**
- ${\cal P}(D)$  is a normalizing constant
- $P(\boldsymbol{\theta}|\boldsymbol{D})\,$  is the posterior probability

## Globe tossing example

- Toss globe 9 times: observe W = 6 water and L = 3 land
- What's the proportion p of Earth that is covered in water?



- Observed data D = (W, L)
- $\bullet$  Binomial model with parameter p
  - $W \sim \operatorname{Binom}(p)$
- Question: which p are most plausible given our data?

#### Globe tossing Bayesian updating

• Apply Bayes' rule:

$$P(p|W) = \frac{P(W|p)P(p)}{P(W)} \qquad P(p|W) \propto P(W|p)P(p)$$

• Binomial likelihood: 
$$P(W|p) = {W+L \choose W} p^W (1-p)^L$$

• Uniform prior: P(p) = 1

#### Globe tossing Bayesian updating



# Analyzing the Bayesian estimate

- The **entire** posterior distribution is the Bayesian estimate!
- Extract **point estimates** 
  - Find the parameter value with highest probability (MAP estimate)
  - Find the median parameter value (50% quantile)
- Extract credible intervals
  - e.g., find the 2.5% and 97.5% quantiles of the parameter values
- Make posterior predictions
  - Simulate new data taking into account all of the uncertainty embodied in the posterior distribution





#### Sampling from the Bayesian estimate

• Usually easier to work with samples from the posterior



#### What can we learn from this data?



time

# Need to add demographic processes to obtain periodic case trajectories



$$\begin{aligned} \frac{dS}{dt} &= \mu N - \beta S I / N - \mu S \\ \frac{dI}{dt} &= \beta S I / N - \gamma I - \mu I \\ \frac{dR}{dt} &= \gamma I - \mu R \end{aligned}$$

SIR schematic source: https://sineadmorris.github.io/post/the-sir-model/

## Key epidemiological parameter: $R_0$

- Average number of secondary cases that an infectious individual would generate in a completely susceptible population
  - $R_0 < 1$ : no outbreak
  - $R_0 > 1$ : yes outbreak

$$\frac{dS}{dt} = \mu N - \beta SI/N - \mu S$$

$$\frac{dI}{dt} = \beta SI/N - \gamma I - \mu I$$

$$\frac{dR}{dt} = \gamma I - \mu R$$

# Perform inference on $R_0$



## Grid approximation can fail

- Globe tossing example: choose a discrete set of *p* values and compute the posterior probability for each *p* value
- Can we do something similar with  $R_0$  values instead?
  - Problem:  $P(D|R_0)$  values are extremely small
  - Could be numerical issues with trying to sample  $R_0$  values from a grid approximated posterior
- Grid approximation computationally infeasible with high-dimensional parameter spaces

## Markov Chain Monte Carlo (MCMC)

• Algorithm for obtaining **samples** from the posterior distribution without ever computing the actual posterior distribution



#### Markov chains (weather space)



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sunny

Markov chains (proportion of water p space)



#### Metropolis-Hastings Algorithm

 Constructs a Markov chain that gives us parameter samples in proportion to their posterior probability





Nicholas Metropolis

Arianna Rosenbluth



Marshall Rosenbluth



Augusta Teller



Edward Teller



Wilfred Hastings

### Metropolis Algorithm

- 1. Start the chain at a parameter value  $\theta_0$
- 2. Choose a new parameter value ( $\theta^*$ ), based on the current parameter value ( $\theta$ )
- 3. If the  $P(D|\theta^*)P(\theta^*) > P(D|\theta)P(\theta)$ , move the chain to  $\theta^*$ . Otherwise, move the chain to  $\theta^*$  with probability  $P(D|\theta^*)P(\theta^*)/P(D|\theta)P(\theta)$
- 4. Repeat steps 2-3 until the distribution of parameter values converges

#### Burn-in samples need to be removed



#### Grid approximation versus MCMC

0.30 R0\_posterior 0.20 300 Frequency 0.10 100 0.00 վլլլլե 0 1.0 1.5 1.5 2.0 2.5 3.0 1.0 2.0 2.5 3.0

R0\_grid

Histogram of MCMC\_samples

R0

#### Prior and posterior mean trajectories



## Thanks for listening!