# Introduction to MCMC for Infectious Diseases

CCB Skills Seminar – 10/10/2024

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# Long history of modeling infectious disease transmission



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#### 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 *θ* that have high probability given our observed data
	- We want  $P(\theta|D)$  to be large
	- i.e., we want to find "highly plausible" *θ* given our data

## Bayesian updating

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

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

- $P(\theta)$  is the **prior probability**
- $\bullet$   $P(D|\theta)$  is the **likelihood**
- $P(D)$  is a normalizing constant
- $P(\theta|D)$  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)*
- Binomial model with parameter *p*
	- $W \sim \text{Binom}(p)$
- **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



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

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

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

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

$$
\begin{aligned}\n\frac{dS}{dt} &= \mu N - \beta SI/N - \mu S \\
\frac{dI}{dt} &= \beta SI/N - \gamma I - \mu I\n\end{aligned}\n\qquad R_0 = \frac{\beta}{\gamma + \mu}
$$
\n
$$
\frac{dR}{dt} = \gamma I - \mu R
$$

# 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
- $\bullet$  Can we do something similar with  $R_0$  values instead?
	- Problem:  $P(D|R_0)$  values are extremely small
	- $\bullet$  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)





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



Rosenbluth

Marshall 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 (*θ*)
- 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

 $0.30$ R<sub>0</sub>\_posterior 0.20 300 Frequency  $0.10$ 100  $0.00$  $\circ$  $1.5$  $1.0$  $2.0$  $2.5$  $3.0\,$  $1.0$  $1.5$  $2.0$  $2.5$  $3.0$ R0\_grid R<sub>0</sub>

**Histogram of MCMC\_samples** 

#### Prior and posterior mean trajectories



## Thanks for listening!