

# **SOC542 Statistical Methods in Sociology II**

## **Introduction to Bayesian statistics**

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# Plan

- ▶ Probability review
- ▶ Bayes' theorem and its applications
- ▶ Comparing Bayesian and Frequentist approaches
- ▶ Bayesian estimation
- ▶ Lab: Bayesian regression in R

# Probability review

## Simple probability

- ▶  $P(A)$  refers to the probability of an event  $A$ 
  - ▶ e.g.  $P(A) = 0.5$  when referring to the probability of receiving a heads on a fair coin toss.
  - ▶ e.g.  $P(B) = \frac{1}{6}$  is the probability of rolling six with a fair die.
- ▶ In each case, we have a *random process* with a set of possible outcomes (e.g. heads or tails) referred to as the *sample space*.

# Probability review

## Simple probability

- ▶ What is the probability of tossing a coin twice and getting two heads?

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- ▶ What is the probability of a sequence of  $N$  heads?
  - ▶  $P(A)^N$
- ▶ In this case,  $P(A)$  becomes vanishingly small as  $n \rightarrow \infty$ 
  - ▶  $0.5^{10} = 0.00098 = \frac{1}{1024}$



# Probability review

## Simple probability

- ▶ We can easily use simulations to verify our calculation. In this case, I use the `rbinom` function to simulate 1024 sequences of 10 tosses of a fair coin.

```
sims <- rbinom(1024, 10, 0.5)
print(length(sims[sims >= 10]))
```

```
## [1] 0
```

# Probability review

## Independence

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$$P(A, B) = P(A)P(B) = \frac{1}{2} * \frac{1}{6} = \frac{1}{12}$$

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## Independence

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$$P(A, B) = P(A)P(B) = \frac{1}{2} * \frac{1}{6} = \frac{1}{12}$$

- ▶ The two events are independent of one another, so the *joint probability* is simply the product of the probabilities of the two events.

# Probability review

## Conditional probability and independence

- ▶  $P(A)$  and  $P(B)$  are independent *if and only if*  $P(A|B) = P(A)$ .
  - ▶ e.g. The number we rolled on the die has no effect on the outcome of the coin toss.

# Probability review

## Conditional probability and independence

- ▶ Consider a deck of 52 standard playing cards. What is the probability of randomly drawing an Ace?<sup>1</sup>

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<sup>1</sup>Example from Cunningham 2021, p. 17.

# Probability review

## Conditional probability and independence

- ▶ Consider a deck of 52 standard playing cards. What is the probability of randomly drawing an Ace?

$$P(\text{Ace}) = 4/52 = 1/13$$

- ▶ Let's assume we pick an Ace and put it to the side. What's the probability we get another Ace?

# Probability review

## Conditional probability and independence

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$$P(\text{Ace}) = \frac{4}{52} = \frac{1}{13}$$

- ▶ Let's assume we pick an Ace and put it to the side. What's the probability we get another Ace?
- ▶ Wrong answer:  $P(\text{Ace}_2) = \frac{4}{52} = \frac{1}{13}$ .



# Probability review

## Conditional probability and independence

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- ▶ Let's assume we pick an Ace and put it to the side. What's the probability we get another Ace?
- ▶ Wrong answer:  $P(\text{Ace}_2) = \frac{4}{52} = \frac{1}{13}$ .
- ▶ Correct answer:  $P(\text{Ace}_2) = P(\text{Ace}_2|\text{Ace}_1) = 3/51 = 0.059$ .
- ▶ This is an example of *conditional probability* since  $P(\text{Ace}_2|\text{Ace}_1) \neq P(\text{Ace}_1)$ .

# Probability review

## Conditional probability and independence

- ▶ We can express a conditional probability as:

$$P(A|B) = \frac{P(B, A)}{P(B)}$$

- ▶ The probability of  $A$  conditional on  $B$  is the **joint probability** of  $A$  and  $B$ , divided by the **marginal probability** of  $B$ .
- ▶ The denominator is the sum of over possible joint probabilities of  $B$  and  $A$ ,  $\sum_{A^*} P(B, A^*)$ .
  - ▶ The  $*$  denotes that  $A^*$  may take multiple values.

# Probability review

## Conditional probability and independence

- ▶ If two events are independent, then  $P(A|B) = P(A)$ .
- ▶ To reject independence, we need to show that  $P(A, B) \neq P(A)P(B)$

# Probability review

## Bayes' theorem

$$P(A|B) = \frac{P(B|A)P(A)}{P(B)}$$

# Probability review

## Bayes' theorem

- ▶ What's the probability it is going to rain given that we can see clouds?

$$P(Rain|Cloud) = \frac{P(Cloud|Rain)P(Rain)}{P(Cloud)}$$

# Probability review

## Bayes' theorem

- ▶ Let's say we live in England...
  - ▶  $P(\text{Cloud}) = 0.7$
  - ▶  $P(\text{Rain}) = 0.3$
  - ▶  $P(\text{Cloud}|\text{Rain}) = 1$

$$P(\text{Rain}|\text{Cloud}) = \frac{P(\text{Cloud}|\text{Rain})P(\text{Rain})}{P(\text{Cloud})} = \frac{1 * 0.3}{0.7} = \frac{0.3}{0.7} \approx 0.429$$

# Probability review

## Deriving Bayes' theorem

- ▶ Start with the definition of conditional probability:

$$P(A|B) = \frac{P(B, A)}{P(B)}$$

- ▶ Multiply each side by  $P(B)$ :

$$P(A|B)P(B) = P(B, A)$$

- ▶ Analogously, if we start with  $P(B|A)$  we can get:

$$P(B|A)P(A) = P(B, A)$$

# Probability review

## Deriving Bayes' theorem

- ▶ The previous example shows that the following quantities are equal:

$$P(A|B)P(B) = P(B|A)P(A)$$

- ▶ Divide both sides by  $P(B)$  to get Bayes' theorem:

$$P(A|B) = \frac{P(B|A)P(A)}{P(B)}$$



# Bayes' theorem

## COVID-19 tests

$$P(A|B) = \frac{P(B|A)P(A)}{P(B)}$$

$$P(C19|+) = \frac{P(+|C19)P(C19)}{P(+)}$$

# Bayes' theorem

## COVID-19 tests

$$P(C19|+) = \frac{P(+|C19)P(C19)}{P(+)}$$

- ▶  $P(C19|+)$ : Probability you have COVID-19 given that you test positive.
- ▶  $P(+|C19)$ : Probability you test positive given that you have COVID-19.
- ▶  $P(C19)$ : Probability you have COVID-19 given population infection rates.
- ▶  $P(+)$ : Probability a test returns a positive result.

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# Bayes' theorem

## COVID-19 tests

- ▶ Assume there is a 1% chance you have COVID-19.
- ▶ Assume a test has a false negative rate of 2%.
  - ▶ 98% of the time it correctly diagnoses COVID-19, 2% of the time it fails to detect it.
- ▶ Assume the same test has a false positive rate of 5%.
  - ▶ 95% of the time it correctly rejects COVID-19 when a person is negative, 5% of the time it falsely diagnoses COVID-19.
- ▶ What is the probability you really have COVID-19 following a positive test?

# Bayes' theorem

## COVID-19 tests: $P(+|C19)$

$$P(C19|+) = \frac{P(+|C19)P(C19)}{P(+)}$$

- ▶ If we assume a false negative rate of 2%. Then the probability of a positive test given COVID-19 is  
 $P(+|C19) = 1 - 0.02 = 0.98$ .



# Bayes' theorem

## COVID-19 tests: $P(C19)$

$$P(C19|+) = \frac{P(+|C19)P(C19)}{P(+)}$$

- ▶ Assume 1% of the population has COVID-19, then  $P(C19) = 0.01$ .

# Bayes' theorem

## COVID-19 tests: $P(+)$

- ▶ To calculate the proportion of positive tests we need to count all the positive tests, irrespective of whether someone is positive.
- ▶ To obtain this, we can reformulate Bayes rule as

$$\frac{P(B|A)P(A)}{P(B|A)P(A) + P(B|A^*)P(A^*)}$$
$$\frac{P(+|C19)P(C19)}{P(+|C19)P(C19) + P(+|C19-)P(C19-)}$$

# Bayes' theorem

## COVID-19 tests: $P(+)$

$$P(C19|+) = \frac{P(+|C19)P(C19)}{P(+)}$$

- ▶ We already know the first part of the denominator,  
 $P(+|C19)P(C19) = 0.98 * 0.01$ .
- ▶ If the test has a false positive rate of 5%,  
 $P(+|C19-) = 0.05 * (1 - 0.01)$
- ▶ Thus, we take the sum of these probabilities to get the marginal probability of a positive test:  
 $P(+) = (0.98 * 0.01) + (0.05 * (1 - 0.01))$

# Bayes' theorem

## COVID-19 tests: Calculating $P(C19|+)$

- If we plug the numbers into Bayes' theorem we get

$$P(C19|+) = \frac{0.98 * 0.01}{0.98 * 0.01 + 0.05 * 0.99}$$

- We can use R to do the calculation for us

```
(0.98*0.01) / ((0.98*0.01) + (0.05*(1-0.01)))
```

```
## [1] 0.1652614
```

# Bayes' theorem

## Terminology

**Posterior**  $\propto$  **Likelihood**  $\times$  **Prior**

- ▶ In the previous example,
  - ▶  $P(C19|+)$  is the **posterior**.
  - ▶  $P(+|C19)$  is the **likelihood of the data**.
  - ▶  $P(C19)$  is the **prior**.
- ▶ The denominator  $P(+)$  ensures the result is a probability. It is sometimes referred to as the **marginal likelihood** or **normalizing constant**.

# Bayes' theorem

## COVID-19 tests: Tabular explanation

- ▶ The four cells in the middle of the table represent the *joint probabilities* of two events.
- ▶ The row and column totals represent the *marginal probabilities* of each event.
- ▶  $\theta$  is used to denote the parameters we are estimating.

Test result	$\theta = C19+$	$\theta = C19-$	Marginal (Test)
+	$P(+ C19)P(C19)$	$P(+ C19-)P(C19-)$	$\sum_{\theta} P(+ \theta)P(\theta)$
-	$P(- C19)P(C19)$	$P(- C19-)P(C19-)$	$\sum_{\theta} P(- \theta)P(\theta)$
Marginal C19	$P(C19+)$	$P(C19-)$	1.0

# Bayes' theorem

## COVID-19 tests: Tabular explanation

- ▶ To calculate  $P(C19|+)$  we can take the *joint probability* of C19 and a positive test and divide it by the *marginal probability* of a positive test.
- ▶ We can get the relevant values directly from the table:  
 $0.98 * 0.01 / 0.06$ .

Test result	$\theta = C19+$	$\theta = C19-$	Marginal (Test)
+	$0.98 * 0.01$	$0.05 * (1 - 0.01)$	0.06
-	$(1 - 0.98) * 0.01$	$(1 - 0.05) * (1 - 0.01)$	0.94
Marginal C19	0.01	$(1 - 0.01)$	1.0

# Bayes' theorem

## Changing our priors

- ▶ If there is a new surge we could update our prior and recalculate
- ▶ Let's change our prior to assume 10% COVID-19 prevalence in the population

$$(0.98 * 0.1) / ( (0.98 * 0.1) + (0.05 * 0.9) )$$

```
## [1] 0.6853147
```

- ▶ Now we get a much higher posterior probability.
- ▶ We could also extend this analysis by incorporating other prior information, e.g. symptoms, exposure



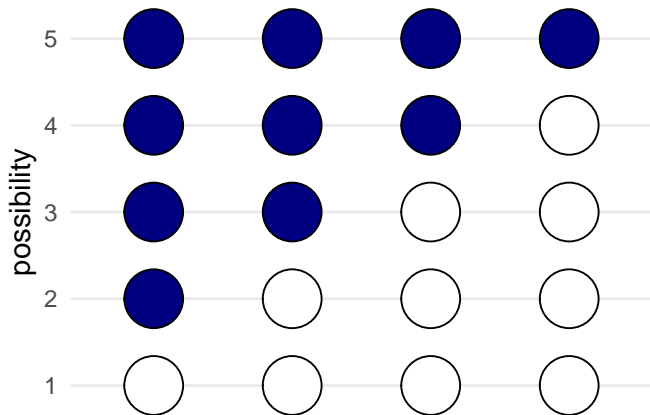
# Bayesian inference as counting

## McElreath's marble counting example

- ▶ Consider a bag containing four marbles
- ▶ The marbles can be white or blue
- ▶ We draw a sample of marbles from the bag (with replacement)

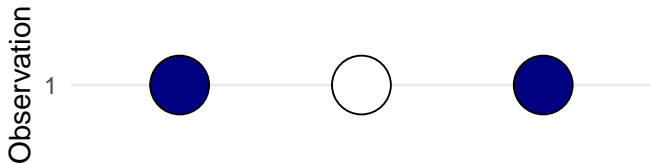
# Bayesian inference as counting

Conjecture: Five possibilities



# Bayesian inference as counting

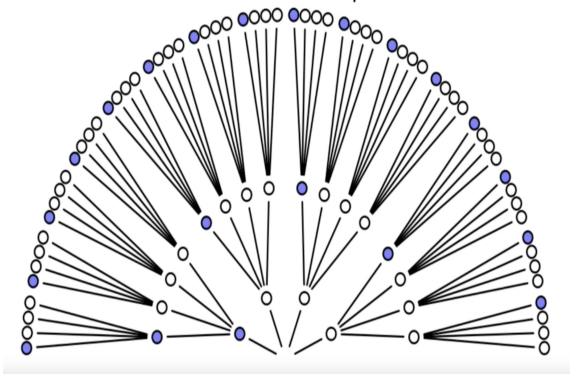
A sample from the bag produces



# Bayesian inference as counting

## Sampling and possibilities

How many ways can we get this sample if the bag contains 3 white and 1 blue?



McElreath 2020, Fig. 2.2 (p. 22)

# Bayesian inference as counting

## Counting the possibilities

Conjecture	Ways to produce [B,W,B]
[W,W,W,W]	$0 \times 4 \times 0 = 0$
[B,W,W,W]	$1 \times 3 \times 1 = 3$
[B,B,W,W]	$2 \times 2 \times 2 = 8$
[B,B,B,W]	$3 \times 1 \times 3 = 9$
[B,B,B,B]	$4 \times 0 \times 4 = 0$

# Bayesian inference as counting

## From counts to probability

Conjecture	Proportion B	Ways [B,W,B]	Plausibility
[W,W,W,W]	0.00	0	0.00
[B,W,W,W]	0.25	3	0.15
[B,B,W,W]	0.50	8	0.40
[B,B,B,W]	0.75	9	0.45
[B,B,B,B]	1.00	0	0.00

# Bayesian inference as counting

## Summary

- ▶ We enumerated the set of plausible data generating processes  $p$
- ▶ We counted the ways we could produce the data given each value of  $p$ . This is known as the *likelihood*.
- ▶ We normalized these counts to get *posterior* probabilities, which indicate the relative plausibility of each option  $p$ .
- ▶ The most plausible value is the one that has the most ways of generating the data.

# Bayesian inference as counting

## Incorporating prior information

- Now let's say we pick another marble and it's blue. We can use the prior information to update our counts.

Conjecture	Ways to produce [B]	Prior counts	New counts
[W,W,W,W]	0	0	$0 \times 0 = 0$
[B,W,W,W]	1	3	$3 \times 1 = 3$
[B,B,W,W]	2	8	$8 \times 2 = 16$
[B,B,B,W]	3	9	$9 \times 3 = 27$
[B,B,B,B]	4	0	$0 \times 4 = 0$



# Bayesian inference as counting

## Bayes' theorem and data analysis

- In a general sense, we can think about Bayesian inference as calculating the posterior distribution in the following way:

$$\textit{Posterior} = \frac{\textit{Probability of the data} * \textit{Prior}}{\textit{Average probability of the data}}$$

# Bayesian inference

**"Bayesian inference is reallocation of credibility across possibilities" - John Kruscke<sup>2</sup>**

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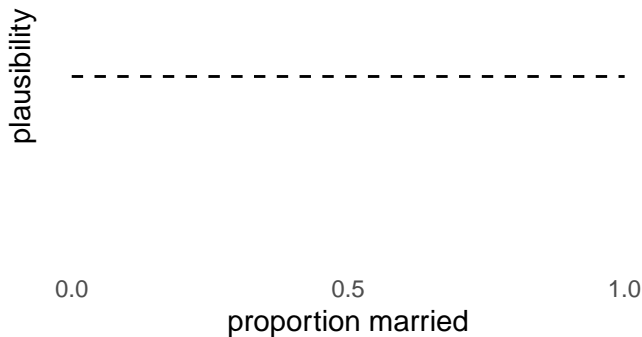
<sup>2</sup>Chapter 2 of Kruschke's 2015 book *Doing Bayesian Data Analysis* provides an outline of his argument and is [available online](#).

# Bayesian inference for a continuous parameter

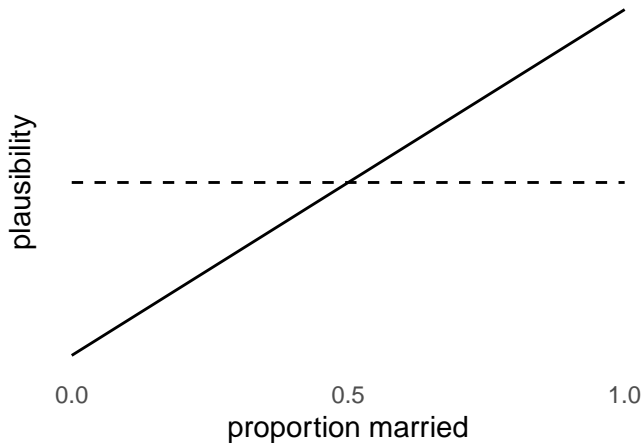
## Estimating the married population

- ▶ Assume a demographer is interested in estimating the probability that someone is married
- ▶ The demographer starts out with a “flat” prior
  - ▶ The marriage rate could be anywhere from 0 (nobody is married) to 1 (everybody is married).
- ▶ The demographer samples people at random and asks them their marital status.

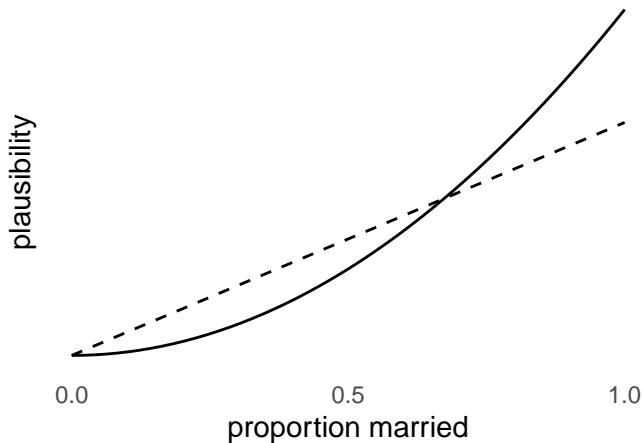
## Assume zero knowledge with a flat (uniform) prior



## First observation: Married



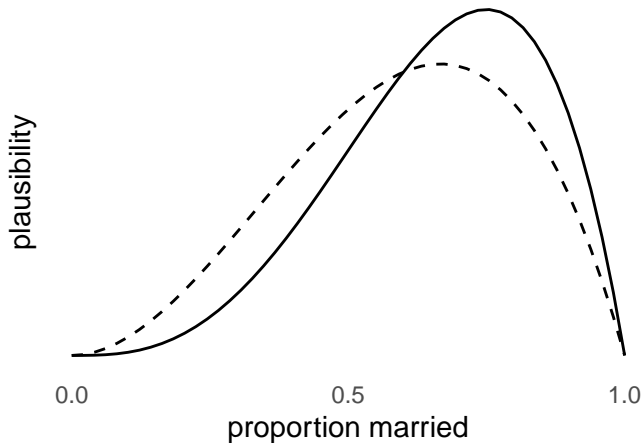
## Second observation: Married



## Third observation: Single

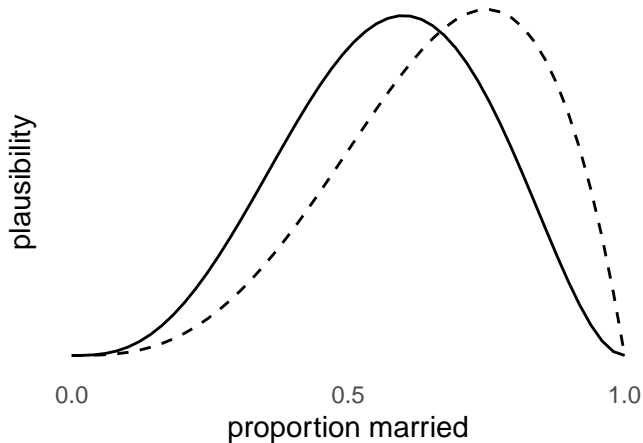


## Fourth observation: Married





## Fifth observation: Single



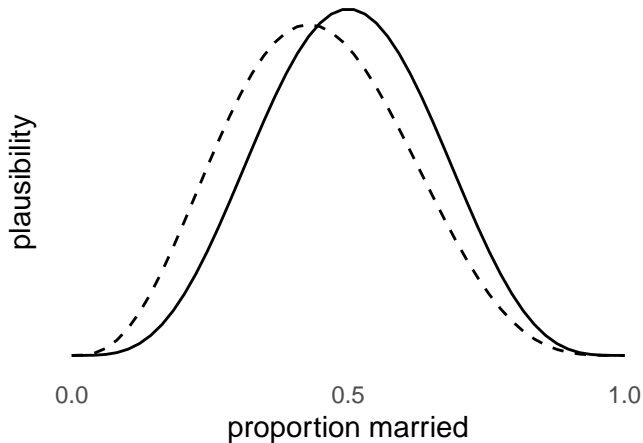
## Sixth observation: Single



## Seventh observation: Single



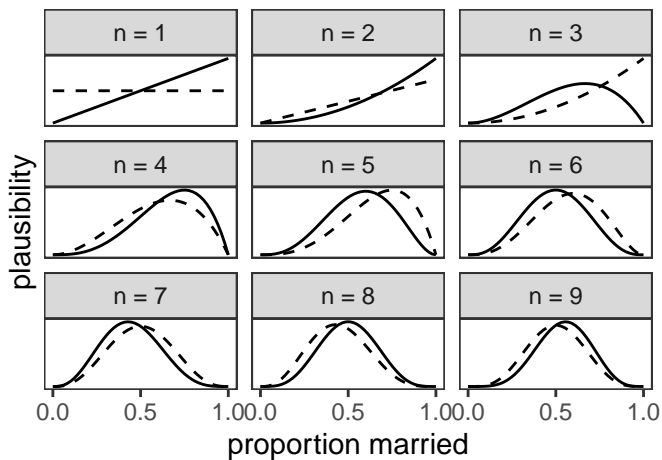
## Eighth observation: Married



## Nineth observation: Single



# Priors and Posteriors



# Bayesian Updating

- ▶ This example demonstrates the concept of **Bayesian updating**
  - ▶ We use new information to update our beliefs
- ▶ Each time we update we use the previous **posterior** as the new **prior**!

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# Bayesian Updating

- ▶ This example demonstrates the concept of **Bayesian updating**
  - ▶ We use new information to update our beliefs
- ▶ Each time we update we use the previous **posterior** as the new **prior**!
- ▶ Most of the time we use all our data at once to get the final posterior rather than iteratively updating.
- ▶ Bayesian updating is order invariant: we will get the same result regardless of the way observations are ordered.

# Formalizing the marriage model

## Writing down a model

- ▶ This problem can be represented using the Beta-Binomial model.<sup>3</sup>

$$\textit{Marriage} \sim \textit{Binomial}(N, p)$$

$$p \sim \textit{Beta}(a, b)$$

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<sup>3</sup>Chapter 3 of *Bayes Rules!* for an extended discussion.

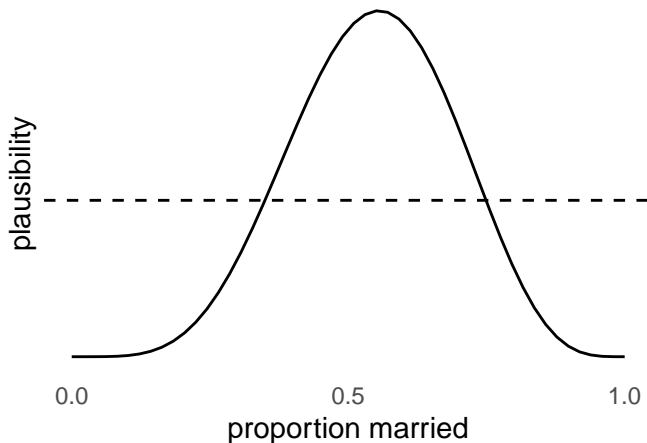
# Formalizing the marriage model

## Writing down a model

- ▶ The goal of this analysis is to produce an estimate of  $p$ , the probability of marriage.
- ▶ Our prior for  $p$  is represented by  $Beta(a, b)$ 
  - ▶ The Beta distribution is bounded to  $[0,1]$
  - ▶ It is equivalent to a *Uniform* distribution when  $a = b = 1$ , representing complete uncertainty

# Formalizing the marriage model

## Prior and posterior distributions



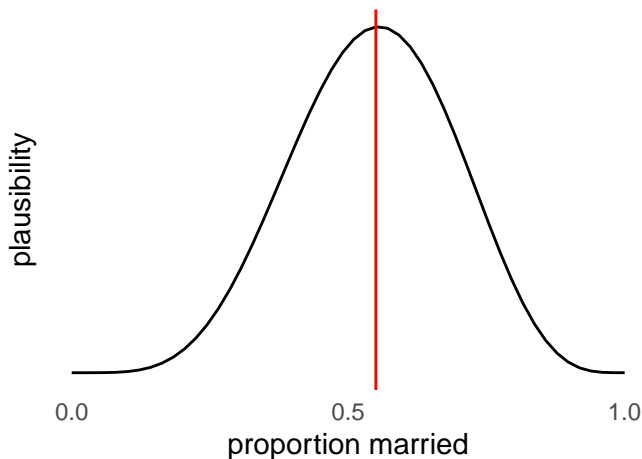
# Formalizing the marriage model

## Understanding the posterior distribution

- ▶ Unlike previous discrete examples, our estimate of  $p$  is represented by the entire posterior distribution
- ▶ In this case, the posterior distribution is simple to calculate
  - ▶  $Beta(1, 1) \rightarrow Beta(1 + married, 1 + N) \rightarrow Beta(6, 10)$
  - ▶ This is due to **conjugacy**, as the prior and posterior belong to the same family of distributions
- ▶ As our models get more complex, we need to use more sophisticated approaches to estimate posterior distributions

# Formalizing the marriage model

## Summarizing the posterior distribution



# Bayesian Regression

- ▶ We can extend this approach to linear regression, where outcomes are modeled using the Normal distribution.

$$y_i \sim \text{Normal}(\mu_i, \sigma)$$

$$\mu_i = \beta_0 + \beta_1 x_i$$

$$\beta_0 \sim \text{Normal}(0, 1)$$

$$\beta_1 \sim \text{Normal}(0, 1)$$

$$\sigma \sim \text{Uniform}(0, 1)$$

# Bayesian Regression

- ▶ In this case, we make the *assumption* that  $y_i$  is normally distributed and that we can express its mean in terms of  $x$  (recall that  $E[y|x] = \beta_0 + \beta_1 x_i$ )
- ▶ After estimating a model using the data we get the *posterior* distribution for each parameter
  - ▶ We can then make statistical inferences regarding  $\hat{\beta}_1$



# Comparing Bayesian and Frequentist approaches

Thomas Bayes (1701-1761)



Source: [Wikipedia](#).

# Comparing Bayesian and Frequentist approaches

Pierre-Simon Laplace (1749-1827)



Source: [Wikipedia](#).

# Comparing Bayesian and Frequentist approaches

Ronald Fisher (1890-1962)



Source: [Wikipedia](#).

# Comparing Bayesian and Frequentist approaches

## Historical developments

- ▶ Frequentist (or “Fisherian”) statistics dominated for most of the 20th century.
- ▶ Bayesian inference critiqued as too subjective and difficult to implement
- ▶ Reversal over the past couple of decades as critiques of Bayesianism debunked, cheap compute power makes it tractable, and key tenets of Frequentist statistics are questioned (e.g. controversy over p-hacking).
- ▶ The Bayesian approach is now mainstream in statistics and much of the natural sciences, but the social sciences have been slower to adopt.<sup>4</sup>

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<sup>4</sup>See Scott and Bartlett 2019.

# Comparing Bayesian and Frequentist approaches

## Theoretical foundations

- ▶ Frequentist
  - ▶ Long-run probabilities
  - ▶ Sampling distributions
- ▶ Bayesian
  - ▶ Probability theory

# Comparing Bayesian and Frequentist approaches

## Sample size

- ▶ Frequentist
  - ▶ Properties of estimators depend on minimal sample size
- ▶ Bayesian
  - ▶ No minimum sample size
  - ▶ But larger samples improve precision of estimates

# Comparing Bayesian and Frequentist approaches

## Point estimates

- ▶ Frequentist
  - ▶ Models produce point estimates
- ▶ Bayesian
  - ▶ No singular point estimates
    - ▶ Many different summaries of the posterior distribution are possible (e.g. mean, median, mode)

# Comparing Bayesian and Frequentist approaches

## P-values

- ▶ Frequentist
  - ▶ p-values used to communicate statistical significance
- ▶ Bayesian
  - ▶ Critique: p-values are based on arbitrary distributional assumptions
  - ▶ Uncertainty is captured by entire posterior distribution
  - ▶ *Bayes' Factor* is a Bayesian version of a p-value<sup>5</sup>

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<sup>5</sup>See Kruschke and Liddell 2018.



# Comparing Bayesian and Frequentist approaches

## Confidence intervals

- ▶ Frequentist
  - ▶ Confidence intervals defined using test statistics and conventions
  - ▶ Assumption that a parameter is fixed and that interval is derived from a sample
- ▶ Bayesian
  - ▶ Critique: Frequentist conventions are arbitrary
  - ▶ Assumption that a parameter has a distribution
  - ▶ *Credible intervals* or *compatibility intervals* can be used to summarize the posterior distribution

# Comparing Bayesian and Frequentist approaches

## Confidence intervals: Interpretation of a 95% interval

- ▶ Frequentist
  - ▶ Over many repeat samples, 95% of calculated confidence intervals would contain the true value of the parameter
- ▶ Bayesian (assume an interval over 95% of the posterior distribution)
  - ▶ There is a 95% probability that the estimated parameter lies within the defined range, given the model and the data.
  - ▶ “What the interval indicates is a range of parameter values compatible with the model and the data.” McElreath, p. 54.

# Computation and Bayesian Estimation

## Bayesian Estimation

- ▶ Three methods for estimating the posterior distribution
  - ▶ Analytical calculations
  - ▶ Grid and quadratic approximation
  - ▶ Markov Chain Monte Carlo

# Computation and Bayesian Estimation

## Analytical calculations

- ▶ For simple problems we can use calculus to provide an analytical solution for the posterior distribution
- ▶ But this approach does not scale well beyond simple problems like the marriage example

# Computation and Bayesian Estimation

## Grid and quadratic approximation

- ▶ Grid approximation (see McElreath 2.4.3)
  - ▶ We can approximate continuous spaces by using grids
    - ▶ But scales very poorly to complex examples
- ▶ Quadratic approximation (see McElreath 2.4.4)
  - ▶ A more robust approach that involves using distributions to approximate the posterior
  - ▶ Flexible for many regression problems but also has trouble scaling

# Computation and Bayesian Estimation

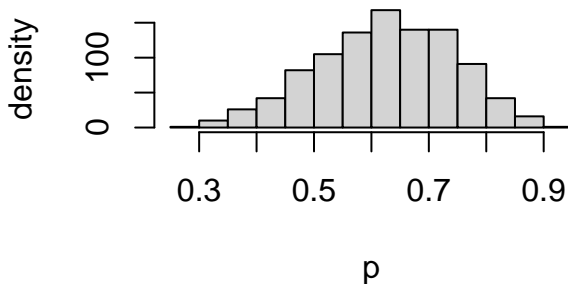
## Markov Chain Monte Carlo (MCMC)

- ▶ Use simulation to draw samples from the posterior distribution
  - ▶ A computationally intensive approach
  - ▶ Samples provide an approximation for complex spaces
  - ▶ More efficient for complex models than quadratic approximation
- ▶ MCMC has led to major advances in Bayesian methods since the 1990s (see McElreath 2.4.5).

# Computation and Bayesian Estimation

MCMC intuition: Sampling from the posterior<sup>6</sup>

## Sampling from Beta(10,6)

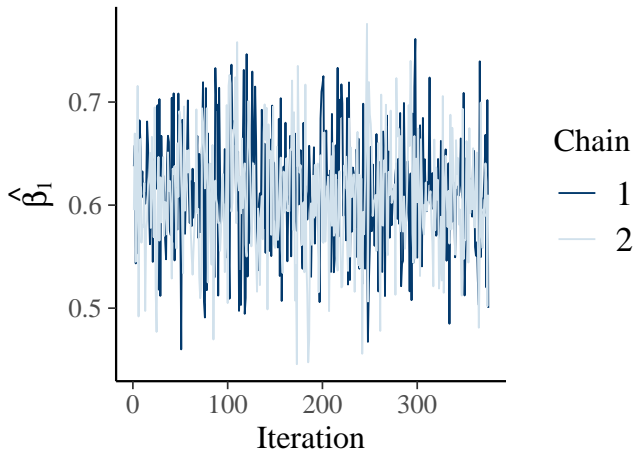


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<sup>6</sup> Note this is a trivial case since we already know the exact posterior distribution!

# Computation and Bayesian Estimation

## Samples from a Markov Chain





# Computation and Bayesian Estimation

## Stan and Hamiltonian Monte Carlo

- ▶ Stan is a programming language developed for statistical computing
- ▶ It implements **Hamiltonian Monte Carlo (HMC)** sampling
  - ▶ A variant of MCMC methods based on Hamiltonian physics
  - ▶ Approximates the posterior by “flicking” a particle and observing its movement
- ▶ HMC is highly effective at solving complex problems<sup>7</sup>
  - ▶ It provides lots of useful diagnostics making it easier to debug than early MCMC approaches
  - ▶ Greater flexibility as it not require *conjugacy*

---

<sup>7</sup> See McElreath Chapter 9 and [Betancourt 2018](#) for a more advanced conceptual overview. [Link to simulation.](#)

# Bayesian Regression in R

- ▶ We will be using `stan_glm` to estimate regression models via HMC in R
- ▶ The *posterior distributions* of the parameters are analyzed to make inferences about the relationship between  $x$  and  $y$
- ▶ We can also use the posterior to generate new data consistent with the model and to calculate new kinds of regression diagnostics

# OLS regression

```
model <- summary(lm(y ~ x, data=df))
```

## Estimating $\beta_0$ and $\beta_1$ using `stan_glm()`

We can also run the same model using Bayesian estimation.

```
library(rstanarm)
model2 <- stan_glm(y ~ x, data = df)
```

```
##
```

```
## SAMPLING FOR MODEL 'continuous' NOW (CHAIN 1).
```

```
## Chain 1:
```

```
## Chain 1: Gradient evaluation took 1.8e-05 seconds
```

```
## Chain 1: 1000 transitions using 10 leapfrog steps per transition would
```

```
## Chain 1: Adjust your expectations accordingly!
```

```
## Chain 1:
```

```
## Chain 1:
```

```
## Chain 1: Iteration:      1 / 2000 [  0%] (Warmup)
```

```
## Chain 1: Iteration:    200 / 2000 [ 10%] (Warmup)
```

```
## Chain 1: Iteration:    400 / 2000 [ 20%] (Warmup)
```

```
## Chain 1: Iteration:    600 / 2000 [ 30%] (Warmup)
```

```
## Chain 1: Iteration:    800 / 2000 [ 40%] (Warmup)
```

```
## Chain 1: Iteration:   1000 / 2000 [ 50%] (Warmup)
```

```
## Chain 1: Iteration:   1001 / 2000 [ 50%] (Sampling)
```

## Comparing `lm` and `stan_glm`

Let's compare the coefficients across the two models. We can see that they are very close. We will discuss the differences in these approaches more in lab and next week.

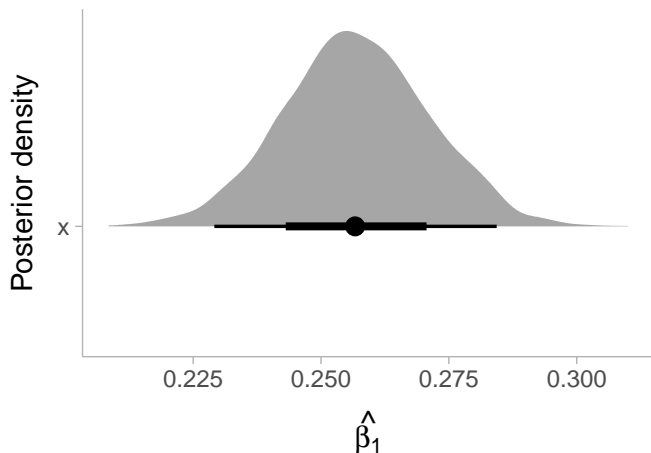
```
print(model$coefficients) # lm
```

	Estimate	Std. Error	t value	Pr(> t )
## (Intercept)	-0.02504433	0.06071423	-0.4124953	6.800650e-01
## x	0.25709671	0.01379670	18.6346586	9.737972e-67

```
print(model2$coefficients) # stan_glm
```

## (Intercept)	x
## -0.02285928	0.25667226

## The posterior distribution of $\hat{\beta}_1$



## Comparing lm and stan\_glm

We can also compare the standard deviations of the residuals,  $\sigma$ . The results are almost identical, showing that both models fit the data well.

```
model$sigma
```

```
## [1] 0.8763714
```

```
sigma(model2)
```

```
## [1] 0.876987
```

## Final remarks

**"All models are wrong, but some are useful" - George Box<sup>8</sup>**

---

<sup>8</sup>This aphorism is attributed to statistician George Box. See [Wikipedia](#) for further discussion.



# Next week

- ▶ Multivariate regression

# Lab

- ▶ Bivariate Bayesian regression using `stan_glm`