## Homework 2

Due on October 26, 2025 at 11:59 pm

Your name here

Submit your Rmarkdown (.qmd) and the compiled pdf on Gauchospace.

## 1. Trend in Same-sex Marriage

A 2017 Pew Research survey found that 10.2% of LGBT adults in the U.S. were married to a same-sex spouse. Now it's the 2020s, and Bayard guesses that  $\pi$ , the percent of LGBT adults in the U.S. who are married to a same-sex spouse, has most likely increased to about 15% but could reasonably range from 10% to 25%.

**1a.** Identify a Beta model that reflects Bayard's prior ideas about  $\pi$  by specifying the parameters of the Beta,  $\alpha$  and  $\beta$ .

```
alpha <- NULL # YOUR CODE HERE
beta <- NULL # YOUR CODE HERE
. = ottr::check("tests/q1a.R")</pre>
```

**1b.** Bayard wants to update his prior, so he randomly selects 90 US LGBT adults and 30 of them are married to a same-sex partner. What is the posterior model for  $\pi$ ?

```
posterior_alpha <- NULL # YOUR CODE HERE
posterior_beta <- NULL # YOUR CODE HERE</pre>
```

1c. Use R to compute the posterior mean and standard deviation of  $\pi$ .

```
posterior_mean <- NULL # YOUR CODE HERE
posterior_sd <- NULL # YOUR CODE HERE

print(sprintf("The posterior mean is %f", posterior_mean))
print(sprintf("The posterior sd is %f", posterior_sd))</pre>
```

1d. Does the posterior model more closely reflect the prior information or the data? Explain your reasoning. Hint: in the recorded lecture we showed a special way in which we can write the posterior mean in a Beta-Binomial model. How can this help? Check the lectures notes.

```
# YOUR CODE HERE
```

Type your answer here, replacing this text.

## 2. Cancer Research in Laboratory Mice

A laboratory is estimating the rate of tumorigenesis (the formation of tumors) in two strains of mice, A and B. They have tumor count data for 10 mice in strain A and 13 mice in strain B. Type A mice have been well studied, and information from other laboratories suggests that type A mice have tumor counts that are approximately Poisson-distributed. Tumor count rates for type B mice are unknown, but type B mice are related to type A mice. Assuming a Poisson sampling distribution for each group with rates  $\theta_A$  and  $\theta_B$ . Based on previous research you settle on the following prior distribution:

$$\theta_A \sim \text{gamma}(120, 10), \ \theta_B \sim \text{gamma}(12, 1)$$

**2a.** Before seeing any data, which group do you expect to have a higher average incidence of cancer? Which group are you more certain about a priori? You answers should be based on the priors specified above.

Type your answer here, replacing this text.

**2b.** After you the complete of the experiment, you observe the following tumor counts for the two populations:

$$\begin{aligned} y_A &= (12, 9, 12, 14, 13, 13, 15, 8, 15, 6) \\ y_B &= (11, 11, 10, 9, 9, 8, 7, 10, 6, 8, 8, 9, 7) \end{aligned}$$

Compute the posterior parameters, posterior means, posterior variances and 95% quantile-based credible intervals for  $\theta_A$  and  $\theta_B$ . Same them in the appropriate variables in the code cell below. You do not need to show your work, but you cannot get partial credit unless you do show work.

```
. = ottr::check("tests/q2b.R")
```

**2c.** Compute and plot the posterior expectation of  $\theta_B$  given  $y_B$  under the prior distribution gamma $(12 \times n_0, n_0)$  for each value of  $n_0 \in \{1, 2, ..., 50\}$ . As a reminder,  $n_0$  can be thought of as the number of prior observations (or pseudo-counts).