Problem set 5

Your name here

Due 11/1/2022 at 5pm

NOTE1: Start with the file ps5_2022.Rmd (available from the github repository at https://github.com/UCh icago-pol-methods/IntroQSS-F22/tree/main/assignments). Modify that file to include your answers. Make sure you can "knit" the file (e.g. in RStudio by clicking on the Knit button). Submit both the Rmd file and the knitted PDF via Canvas

Question 1

(1a) Write code to draw one sample of size 5 from a normal distribution with mean 3 and variance of 2.

```
# your code here
# you can make the rest of the code chunks yourself
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- (1b) Draw m=2000 such samples, compute the sample mean for each sample, and plot a histogram of the result.
- (1c) Compute the variance across your sample means. Compare it to the theoretical sampling variance of the sample mean. Explain any difference you find.
- (1d) Repeat (1b) and (1c) but set the sample size to 500.
- (1e) Show the histogram of Democratic vote share by county in the US in the 2020 presidential election. (You used this data in problem set 2. You might use the pull() function to extract a single column.) Does the distribution look symmetric?
- (1f) Obtain 2000 random samples of 5 counties (without replacement), compute the mean of Democratic vote share for each sample, and plot the distribution of sample means. Does the distribution look symmetric?
- (1g) Compute the variance across your sample means. Compare it to the theoretical variance of the sample mean. Explain any difference you find.
- (1h) Repeat (1f) and (1g) but set the sample size to 500.

Question 2

(2a) Using the plug-in sample variance (Definition 3.2.18 in Aronow & Miller) as a guide, write the formula for plug-in sample covariance.

Your math = here

(2b) Pretending for a moment that U.S. county election results in 2020 are a sample from a super-population, use the formula you wrote above to estimate the covariance between the total votes cast in a county in 2020 and the proportion of votes cast for the Democrat in 2020. Compare it to the unbiased sample covariance, which you can compute using R's cov() function.