

FOR THIS HOMEWORK, YOU WILL NEED TO PRINT YOUR CODE AND ATTACH IT TO THE BACK OF YOUR SUBMISSION.

1. t-procedures. Consider n observations X_1, X_2, \dots, X_n which are iid samples from a Log Normal distribution with population mean μ and population skew γ . The following code can be used to simulate n observations from this population.

```
generate_data <- function(n, mean, skew){
  zeta <- optim(skew^.67, function(z) ((z+2)*sqrt(z-1)-skew)^2,
               method='Brent', lower=1, upper=1e9)$par
  omega <- sqrt(log(zeta))
  theta <- log(mean) - omega^2/2
  x <- rlnorm(n, theta, omega)
}
```

For the rest of this problem, assume that $\mu = 25$.

a) Use the provided function to simulate $n = 30$ observations from the population with $\gamma = 0.5$. Make a histogram of the data, and explain why the t-procedures assumptions are (roughly) met.

b) Use t-procedures to construct a 95% confidence interval for μ . Interpret this interval.

c) Perform a *simulation study* by repeating the process above (a) and b)) $M = 10,000$ times. You can use the following code as a "skeleton". Count the number of times that the 95% CI actually succeeds in capturing the true value of μ . How close is this "coverage" to the desired value of 95%?

```
M <- 10000 #Number of simulations
n <- 30 #Sample size
skew <- .5 #Population skewness
count <- 0. #Number of times CI succeeds
#Start simulations
for(i in 1:M){
  #Simulate data
  x <- generate_data(n, 25, skew)
  #Calculate CI endpoints here (use t-procedures)

  #Check to see if CI captures the true value
  if(change_this_part <- TRUE){
    count <- count + 1
  }
}
print(100*count/M)
```

- d) Repeat your simulation study for sample sizes of $n = 100$ and $n = 500$. What happens and why?
- e) Repeat the entire simulation study (for all three values of n) after increasing the skew to $\gamma = 10$. Explain what happens and why. Repeat this again for $\gamma = 50$.

2. The Bootstrap and the Accelerated Bootstrap. The iris data set is a famous data set consisting of measurements for 150 iris flowers. The "petal length" of these 150 flowers can be obtained by typing `iris$Petal.Length` in R. Create a histogram of the data, and note that the data is "bimodal".

The Bimodality coefficient is a statistic which can be calculated from the data as

$$BC = \frac{\hat{\gamma}^2 + 1}{\hat{\kappa} + 3 \left(\frac{(n-1)^2}{(n-2)(n-3)} - 1 \right)}$$

where $\hat{\gamma}$ and $\hat{\kappa}$ are the sample *skew* and sample *kurtosis* of the data respectively

$$\hat{\gamma} = \frac{1}{n} \sum_{i=1}^n \left(\frac{x_i - \bar{x}}{s_x} \right)^3 \quad \hat{\kappa} = \frac{1}{n} \sum_{i=1}^n \left(\frac{x_i - \bar{x}}{s_x} \right)^4 .$$

- Write a function in R which calculates the bimodality coefficient from the data. What is the estimated value of BC for the petal length data?
- Roughly speaking, larger values of BC indicate bimodality, where 0.47 is often used as a cut-off (this is the value of BC for a uniform distribution). Does the estimate of BC for the petal length data seem to capture the bimodality of the data?
- Use the Bootstrap algorithm to approximate the sampling distribution of BC for the petal length data. Provide a histogram of the bootstrap distribution and also calculate a 95% Bootstrap CI for the true value of the bimodality coefficient. Do you feel confident in asserting that the true value of BC is greater than 0.47?
- Challenge:** For (even more) challenge points, construct a CI using *accelerated bootstrap* technique. (Link on course web-page).

3. Asymptotic normality of the MLE. Consider $X_1, X_2, \dots, X_n \stackrel{iid}{\sim} \text{Beta}(\theta, 1)$. The density function corresponding to this distribution is

$$f(x_i|\theta) = \theta x_i^{\theta-1}, \quad 0 < x_i < 1$$

- Find the MLE of θ . (Hint: We did this one in class).
- Find the asymptotic standard deviation of $\hat{\theta}$.

$$SD(\hat{\theta}) \approx \left(-E \left[\frac{d^2 \log f(x_i|\theta)}{d\theta^2} \right] \right)^{-1/2}$$

- Use the fact that, as $n \rightarrow \infty$,

$$Z = \frac{\hat{\theta} - \theta}{SD(\hat{\theta})/\sqrt{n}} \stackrel{approx}{\sim} N(0, 1)$$

to construct a 95% CI for θ_i

- Simulate data `rbeta(300, 4, 1)` and use your answer to c) to calculate a 95% CI for θ . Did it capture the true value?