RStudio: View PDF

Simulation Study to Understand Sampling Distribution Problem 2

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Problem 2: Simulation Study to Understand Sampling Distribution

Part A Suppose $X_1, X_2, \dots, X_n \stackrel{iid}{\sim} Gamma(\alpha, \sigma)$, with pdf as

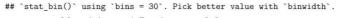
$$f(x|\alpha,\sigma) = \frac{1}{\sigma^{\alpha}\Gamma(\alpha)}e^{-x/\sigma}x^{\alpha-1}, \quad \ 0 < x < \infty,$$

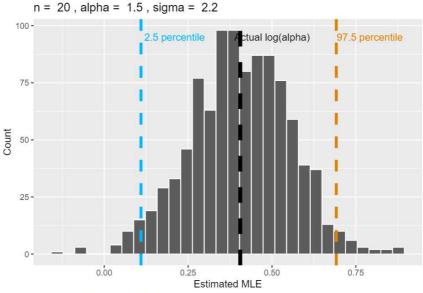
The mean and variance are $E(X) = \alpha \sigma$ and $Var(X) = \alpha \sigma^2$. Note that shape = α and scale = σ .

- 1. Write a function in R which will compute the MLE of $\theta = \log(\alpha)$ using optim function in R. You can name it MyMLE
- 2. Choose n=20, and alpha=1.5 and sigma=2.2
 - (i) Simulate $\{X_1, X_2, \dots, X_n\}$ from rgamma(n=20,shape=1.5,scale=2.2)
 - (ii) Apply the MyMLE to estimate θ and append the value in a vector
 - (iii) Repeat the step (i) and (ii) 1000 times
 - (iv) Draw histogram of the estimated MLEs of θ .
 - (v) Draw a vertical line using abline function at the true value of θ .
 - (vi) Use quantile function on estimated θ 's to find the 2.5 and 97.5-percentile points.
- 3. Choose n=40, and alpha=1.5 and repeat the (2).
- Choose n=100, and alpha=1.5 and repeat the (2).
- 5. Check if the gap between 2.5 and 97.5-percentile points are shrinking as sample size ${\tt n}$ is increasing?

Hint: Perhaps you should think of writing a single function where you will provide the values of n, sim_size, alpha and sigma; and it will return the desired output.

```
1
mle <- function(log_alpha, data, sigma) {
    1 = sum(log(dgamma(data, shape = exp(log_alpha), scale = sigma)))
    # print(paste("l is ", l))
    return(-1)
}
MyMLE <- function(data, sigma) {
    log alpha initial <- log(mean(data)^2/var(data))
    # print(paste("log alpha initial is ", log_alpha_initial))
    estimator <- optim(log_alpha_initial,
                 mle,
                 data = data,
                 sigma = sigma)
    log_alpha_hat <- estimator$par
    return(log alpha hat)
get_estimates <- function(n, alpha, sigma) {</pre>
    estimates <- c()
    for (i in 1:1000) {
        samples <- rgamma(n, shape = alpha, scale = sigma)
        # print(paste("some of the samples are ", samples[1:5]))
        estimates <- append(estimates, MyMLE(data = samples, sigma = sigma))
    return(estimates)
  2
n = 20
alpha = 1.5
sigma = 2.2
estimated mle <- tibble(get estimates(n = n, alpha = alpha, sigma = sigma))
colnames(estimated_mle) <- c("estimate")</pre>
perc_2.5 <- quantile(estimated_mle$estimate, probs = 0.025, names = FALSE)
perc 97.5 <- quantile(estimated mle$estimate, probs = 0.975, names = FALSE)
estimated_mle %>%
    ggplot(aes(estimate)) +
    geom histogram(color = "white", fill = "#5D5D5D") +
    geom_vline(xintercept = log(alpha),
               size = 2,
               linetype = "dashed") +
    annotate("text", label = "Actual log(alpha)", x = 0.5, y = 95, color = "black") +
    geom_vline(xintercept = perc_2.5,
               color = "#00B9FF", size = 1.5, linetype = "dashed") +
    annotate("text", label = "2.5 percentile", x = perc_2.5 + 0.1, y = 95, color = "#00B9FF") +
    geom_vline(xintercept = perc_97.5,
               color = "#E08304", size = 1.5, linetype = "dashed") +
    annotate("text", label = "97.5 percentile", x = perc_97.5 + 0.1, y = 95, color = "#E08304") +
    labs(title = paste("n = ", n, ", alpha = ", alpha, ", sigma = ", sigma),
        x = "Estimated MLE",
         y = "Count")
```





```
diff_20 <- perc_97.5 - perc_2.5
```

```
3.
n = 40
alpha = 1.5
sigma = 2.2
estimated_mle <- tibble(get_estimates(n = n, alpha = alpha, sigma = sigma))
colnames(estimated mle) <- c("estimate")
perc_2.5 <- quantile(estimated_mle$estimate, probs = 0.025, names = FALSE)
perc 97.5 <- quantile(estimated mle$estimate, probs = 0.975, names = FALSE)
estimated mle %>%
    ggplot(aes(estimate)) +
    geom_histogram(color = "white", fill = "#5D5D5D") +
    geom_vline(xintercept = log(alpha),
              size = 2,
               linetype = "dashed") +
    annotate("text", label = "Actual log(alpha)", x = log(alpha) + 0.1, y = 95, color = "black") +
    geom_vline(xintercept = perc_2.5,
               color = "#00B9FF", size = 1.5, linetype = "dashed") +
    annotate("text", label = "2.5 percentile", x = perc_2.5 + 0.1, y = 95, color = "#00B9FF") +
    geom_vline(xintercept = perc_97.5,
               color = "#E08304", size = 1.5, linetype = "dashed") +
    annotate("text", label = "97.5 percentile", x = perc_97.5 + 0.1, y = 95, color = "#E08304") +
    labs(title = paste("n = ", n, ", alpha = ", alpha, ", sigma = ", sigma),
        x = "Estimated MLE",
```

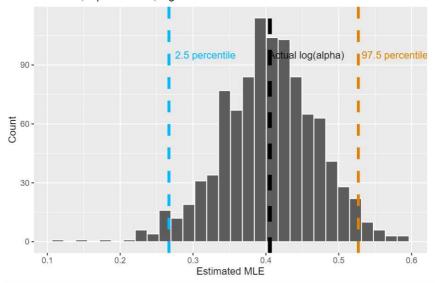
```
diff_40 <- perc_97.5 - perc_2.5
```

```
4.
n = 100
alpha = 1.5
sigma = 2.2
estimated_mle <- tibble(get_estimates(n = n, alpha = alpha, sigma = sigma))
colnames(estimated_mle) <- c("estimate")</pre>
perc_2.5 <- quantile(estimated_mle$estimate, probs = 0.025, names = FALSE)
perc_97.5 <- quantile(estimated_mle$estimate, probs = 0.975, names = FALSE)
estimated_mle %>%
    ggplot(aes(estimate)) +
    geom_histogram(color = "white", fill = "#5D5D5D") +
    geom_vline(xintercept = log(alpha),
               size = 2,
               linetype = "dashed") +
    annotate("text", label = "Actual log(alpha)", x = log(alpha) + 0.05, y = 95, color = "black") +
    geom_vline(xintercept = perc_2.5,
               color = "#00B9FF", size = 1.5, linetype = "dashed") +
    annotate("text", label = "2.5 percentile", x = perc_2.5 + 0.05, y = 95, color = "#00B9FF") +
    geom_vline(xintercept = perc_97.5,
               color = "#E08304", size = 1.5, linetype = "dashed") +
    annotate("text", label = "97.5 percentile", x = perc_97.5 + 0.05, y = 95, color = "#E08304") +
```

```
labs(title = paste("n = ", n, ", alpha = ", alpha, ", sigma = ", sigma),
    x = "Estimated MLE",
    y = "Count")
```

`stat_bin()` using `bins = 30`. Pick better value with `binwidth`.

n = 100, alpha = 1.5, sigma = 2.2



```
diff_100 <- perc_97.5 - perc_2.5
```

5. diff_20

[1] 0.5821454

diff_40

[1] 0.4001029

diff_100

[1] 0.2599853

We can see that the gap between the percentile points is decreasing as the sample size increases.