

# STAT 403 Spring 2018

## HW06

Nan Tang

1662478

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**Q1**

**Q1-1**

**Q1-2**

$$\begin{aligned} \text{Var}(\bar{X}^*) &= \text{Var}\left(\frac{1}{n} \sum_{i=1}^n X_i\right), \text{ since } \bar{X}^* = \frac{1}{n} \sum_{i=1}^n X_i \\ &= \frac{\text{Var}(\sum_{i=1}^n X_i)}{n^2}, \text{ by property of variance} \\ &= \frac{\sum_{i=1}^n \text{Var}(X_i)}{n^2} \\ &= \frac{\text{Var}(X)}{n} \end{aligned}$$

Note that  $\text{Var}(X) = \frac{1}{n} \sum_{i=1}^n (X_i - \bar{X})^2$

$$\text{Var}(\bar{X}^*) = \frac{\sum_{i=1}^n (X_i - \bar{X})^2}{n^2}$$

$$S_n^2 = \frac{1}{n-1} \sum_{i=1}^n (X_i - \bar{X})^2$$

$$(n-1)S_n^2 = \sum_{i=1}^n (X_i - \bar{X})^2$$

$$\text{Var}(\bar{X}^*) = \frac{(n-1)S_n^2}{n^2}$$

As  $n \rightarrow \infty$ ,  $\frac{n-1}{n^2} \rightarrow \frac{1}{n}$ . Therefore, when bootstrap sample size is large enough, variance of bootstrap sample mean is equal to  $\frac{S_n^2}{n}$ .

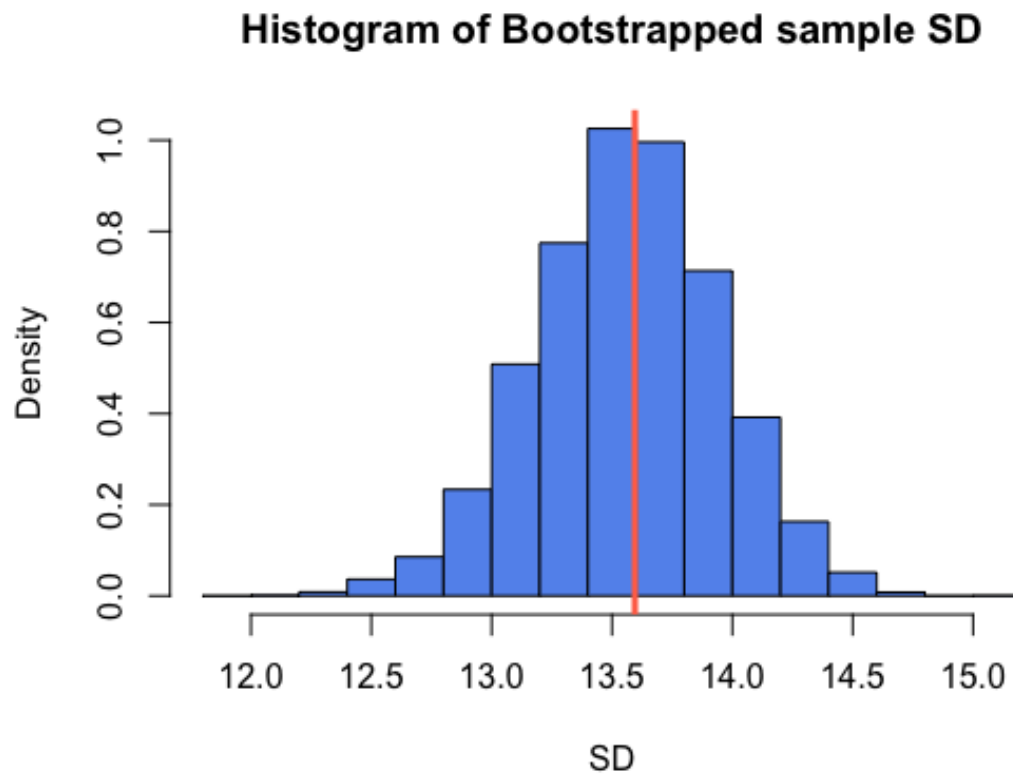
## Q2

### Q2-a

```
origin_sd <- sd(faithful$waiting)
bt_size <- 10000
sample_size <- length(faithful$waiting)
bt_result <- rep(NA, bt_size)

for (ii in 1:bt_size) {
  index <- sample(sample_size, sample_size, replace=T)
  bt_dt <- faithful$waiting[index]
  bt_result[ii] <- sd(bt_dt)
}

hist(bt_result, probability=T, col='cornflowerblue', xlab='SD',
      ylab='Density', main='Histogram of Bootstrapped sample SD')
abline(v=origin_sd, lwd=3, lty=1, col='coral1')
```



## Q2-b

```
> sd(bt_result)^2
[1] 0.1472021
> mean((bt_result - origin_sd)^2)
[1] 0.1483897
```

Variance of sample SD is 0.1472021, and MSE is 0.1483897.

## Q2-c

```
bt_sd <- sd(bt_result)
lower_bd <- origin_sd + qnorm(0.975) * bt_sd
upper_bd <- origin_sd - qnorm(0.975) * bt_sd
> lower_bd
[1] 14.34695
> upper_bd
[1] 12.843
```

```
> quantile(bt_result, c(0.025, 0.975))
      2.5%      97.5%
12.78773 14.29303
```

By theorem, the 95% confidence interval is [12.843, 14.34695]. In bootstrap samples, 95% interquantile is [12.78773, 14.29303].

## Q2-d

p-value = P(getting value more extreme than sample SD| $H_0$ )

```
bt_var <- sd(bt_result)^2
p_value <- 2 * pnorm(origin_sd, 15, bt_var)
> p_value
[1] 1.362628e-21
```

## Q2-e

The variance of bootstrapped samples is  $\frac{\sigma^2}{n}$ , indicating the value will decrease as sample size increases. Larger sample size minimizes the margin of error, of which small margin of error make the estimation more accurate.

## Q3

### Q3-a

```
admission_dt <- UCBAmissions[, , 1]

origin_or <- (admission_dt[1,1] * admission_dt[2,2]) /
  (admission_dt[2,1] * admission_dt[1,2])

## generate dataset based on given information
n_male <- admission_dt[1,1] + admission_dt[2,1]
n_female <- admission_dt[1,2] + admission_dt[2,2]
n <- n_male + n_female

data_pull <- c(rep(1, admission_dt[1,1] + admission_dt[1,2]),
               rep(0, admission_dt[2,1] + admission_dt[2,2]))

N <- 10000
bt_or <- rep(NA, N)
for (ii in 1:N) {
```

```

sp_index <- sample(n, n, replace=T)
sp_dt <- data_pull[sp_index]
sp_male <- sp_dt[1:n_male]
sp_female <- sp_dt[(n_male+1):n]
male_admit <- sum(sp_male)
male_reject <- length(sp_male) - male_admit
female_admit <- sum(sp_female)
female_reject <- length(sp_female) - female_admit
sp_or <- (male_admit * female_reject) / (male_reject * female_admit)
bt_or[iii] <- sp_or
}

bt_mse <- mean((bt_or - mean(bt_or))^2)

> bt_mse
[1] 0.04906438

```

Under bootstrap size of 10000, I get the MSE of OR is 0.04906438.

### Q3-b

```

bt_sd <- sd(bt_or)
p_value <- pnorm(origin_or, 1, bt_sd)

> p_value
[1] 0.001652304

```

### Q3-c