# Solution 10: Bootstrap

Yaqi Shi

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Applying bootstrap

The following code generates  $(X_i, Y_i)$  pairs.

```
fin_pairs <- generate_pairs(100) # Generate 100 (X,Y) pairs.
head(fin_pairs)</pre>
```

```
## X Y
## 1 2.459012 0.1854205
## 2 2.633453 0.7209364
## 3 3.102875 -0.3329833
## 4 3.765494 -2.6460032
## 5 2.643346 0.4392245
## 6 2.808591 0.5727379
```

We are interested in

$$\hat{\alpha} = \frac{\hat{\sigma}_Y^2 - \hat{\sigma}_{XY}}{\hat{\sigma}_X^2 + \hat{\sigma}_Y^2 - 2\hat{\sigma}_{XY}}$$

```
Sigmahat <- cov(fin_pairs)
Sigmahat
```

```
## X Y
## X 0.7730102 -0.1072715
## Y -0.1072715 2.0284446

sigma2hatXX <- Sigmahat[1,1]
sigma2hatYY <- Sigmahat[2,2]
sigmahatXY <- Sigmahat[1,2]</pre>
```

#### The $\hat{\alpha}$ is

```
alphahat <- (sigma2hatYY - sigmahatXY)/(sigma2hatXX + sigma2hatYY -2*sigmahatXY)
alphahat</pre>
```

#### ## [1] 0.7081292

While the true value of alpha is

```
sigma2XX <- 1
sigma2YY <- 2
sigmaXY <- -0.25
alpha_true <-(sigma2YY - sigmaXY)/(sigma2XX + sigma2YY -2*sigmaXY)
alpha_true</pre>
```

```
## [1] 0.6428571
```

Now, again, we're going to resample with replacement from our data, and compute our statistic  $\hat{\alpha}$  on each resample. The hope is that these resampled versions of the statistic will resemble the distribution of the statistic evaluated on the original data.

- 1. Create a function to compute alphabat from a given data set.
- 2. Resample the data B=200 times, evaluating  $\hat{\alpha}$  on each resample. Then, we'll use those resampled values to estimate the variance.
- 3. Create the confidence interval at the estimate.

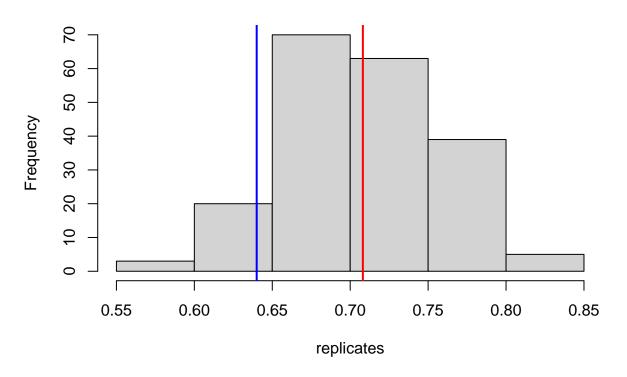
### Solution

```
compute_alphahat <- function(data) {
    # We're assuming that data is a data frame with two columns.
    Sigmahat <- cov( data )
    # Extract the variance and covariance estimates from the sample covariance
    sigma2hatXX <- Sigmahat[1,1]
    sigma2hatYY <- Sigmahat[2,2]
    sigmahatXY <- Sigmahat[1,2]
    # plug these into the definition of alpha.
    alphahat <- (sigma2hatYY - sigmahatXY)/(sigma2hatXX + sigma2hatYY -2*sigmahatXY)
    return(alphahat)
}
alphahat <- compute_alphahat(fin_pairs)</pre>
```

```
B <- 200
replicates <- rep(NA, B)
# number of observations in our data set.
n <- nrow( fin_pairs )
for( i in 1:B ) {</pre>
```

```
# To resample the data, we will sample indices, and then grab those rows.
resample_indices <- sample(1:n, n, replace=TRUE)
resampled_data <- fin_pairs[resample_indices, ]
replicates[i] <- compute_alphahat(resampled_data)
}
hist(replicates)
abline(v=alphahat, col='red', lwd=2) # alpha of true data.
abline(v=0.64, col='blue', lwd=2) # True alpha</pre>
```

## **Histogram of replicates**



```
# Estimate the variance of alphahat from our bootstrap replicates.
sd_alphahat <- sd(replicates) # estimate of the std dev of alphahat
CI <- c(alphahat - 1.96*sd_alphahat, alphahat + 1.96*sd_alphahat)
CI</pre>
```

## [1] 0.6133809 0.8028775