BoostrappingWithRobustStandardErrors.R

mbair

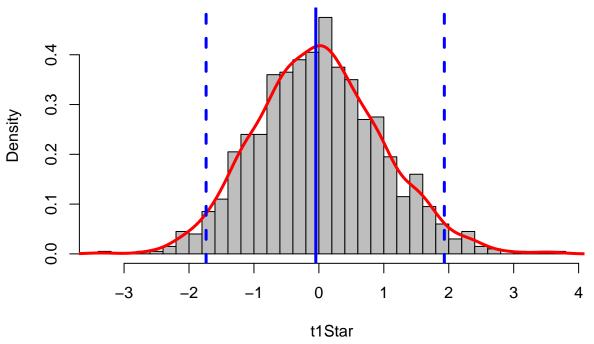
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Residual Bootstrapping

```
# Exercise
# Another bootstrap method in a linear regression model is to
# resample residuals instead of (X, Y). The former is called Residual
# Bootstrapping and the latter is called Pairwise Bootstrapping. I believe that
# boot() is based on the Pairwise Bootstrap. Estimate the standard error and
# bootstrap the 95% confidence intervals for the coeficient of horsepower with
# Residual Bootstrapping.
library(boot)
library(ISLR)
set.seed(1)
df <- Auto
dfBoot \leftarrow df[, c(1, 4, 5)]
# Sample Model ####
X1 <- dfBoot[, 2]</pre>
X2 <- dfBoot[, 3]</pre>
y <- dfBoot[, 1]
lmFit \leftarrow lm(y \sim X1 + X2)
betaOHat <- lmFit$coefficients[1]</pre>
beta1Hat <- lmFit$coefficients[2]</pre>
beta2Hat <- lmFit$coefficients[3]</pre>
# t values of sample model ####
n <- nrow(dfBoot)</pre>
num1 <- sum(((X1 - mean(X1)) ^ 2) * (lmFit$residuals^2))/n</pre>
den1 <- sum((X1-mean(X1))^2)/(n-1)
robustSE1 <- sqrt(num1)/den1
num2 \leftarrow sum(((X2 - mean(X2))^2) * (lmFit\$residuals^2))/n
den2 <- sum((X2-mean(X2))^2)/(n-1)
robustSE2 <- sqrt(num2)/den2</pre>
robT1Coef <- sqrt(n) * beta1Hat / robustSE1</pre>
robT2Coef <- sqrt(n) * beta2Hat / robustSE2</pre>
t1Coef <- summary(lmFit)$coefficients[2,3]</pre>
```

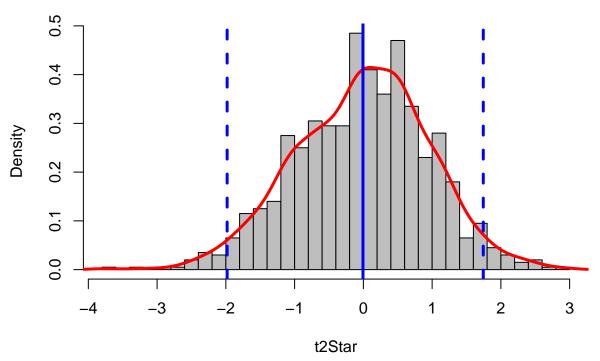
```
t2Coef <- summary(lmFit)$coefficients[3,3]
# Residual Bootstrap Model
# Initialize relevant values ####
R <- 1000
robustT1Star <- NA
robustT2Star <- NA
t1Star <- NA
t2Star <- NA
# Loop for residual bootstrapping with and without robust standard errors ####
for (r in 1:R) {
  uStar <- sample(lmFit$residual, nrow(dfBoot), replace = TRUE)
  yStar <- beta0Hat + beta1Hat * X1 + beta2Hat * X2 + uStar
  lmFitStar <- lm(yStar ~ X1 + X2)</pre>
  beta1Star <- lmFitStar$coefficients[2]</pre>
  beta2Star <- lmFitStar$coefficients[3]</pre>
  seStar1 <- summary(lmFitStar)$coefficients[2,2]</pre>
  seStar2 <- summary(lmFitStar)$coefficients[3,2]</pre>
  num1Star <- sum(((X1 - mean(X1)) ^ 2) * (lmFitStar$residuals^2))/n</pre>
  den1Star \leftarrow sum((X1-mean(X1))^2)/(n-1)
  robustSE1Star <- sqrt(num1Star)/den1Star</pre>
  num2Star <- sum(((X2 - mean(X2)) ^ 2) * (lmFitStar$residuals^2))/n</pre>
  den2Star \leftarrow sum((X2-mean(X2))^2)/(n-1)
  robustSE2Star <- sqrt(num2Star)/den2Star</pre>
  robustT1Star[r] <- sqrt(n) * (beta1Star - beta1Hat) / robustSE1Star</pre>
  robustT2Star[r] <- sqrt(n) * (beta2Star - beta2Hat) / robustSE2Star</pre>
  t1Star[r] <- (beta1Star - beta1Hat) / seStar1
  t2Star[r] <- (beta2Star - beta2Hat) / seStar2
# Sort the residual moel t values ####
robustT1Star <- sort(robustT1Star)</pre>
robustT2Star <- sort(robustT2Star)</pre>
t1Star <- sort(t1Star)</pre>
t2Star <- sort(t2Star)
# Get the 2.5% and 97.5% levels ####
critT1Star <- c(t1Star[25], t1Star[975])</pre>
critT2Star <- c(t2Star[25], t2Star[975])</pre>
critRobustT1Star <- c(robustT1Star[25], robustT1Star[975])</pre>
critRobustT2Star <- c(robustT2Star[25], robustT2Star[975])</pre>
```

Distribution of Horsepower's t* Under Residual Bootstrap



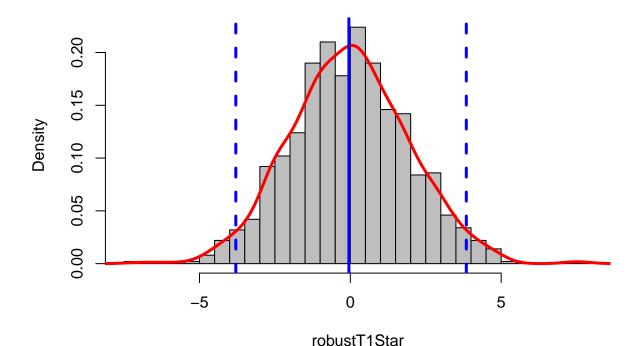
The 95% critical values are -1.737 and 1.932 .

Distribution of Weight's t* Under Residual Bootstrap



```
## The 95% critical values are -1.981 and 1.744 .
```

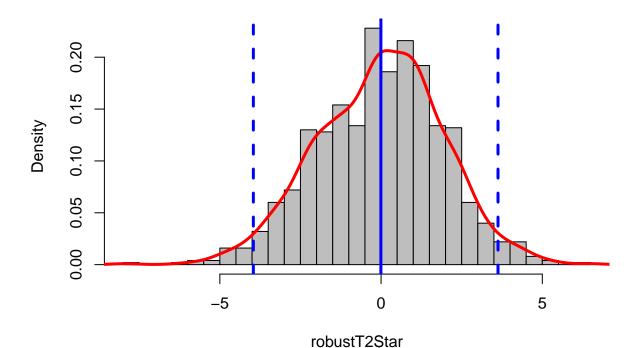
Distribution of Horsepower's robust t* Under Residual Bootstrap



```
## The 95% critical values are -3.797 and 3.844 .
```

```
hist(robustT2Star, breaks = 30, probability = TRUE, col= "grey",
    main = "Distribution of Weight's robust t* Under Residual Bootstrap")
lines(density(robustT2Star), col = "red", lwd = 3)
abline(v = c(critRobustT2Star, beta2Hat), col = c("blue"), lty = c(2, 2, 1), lwd = 3)
```

Distribution of Weight's robust t* Under Residual Bootstrap



The 95% critical values are -3.961 and 3.626 .