Problem Set 2

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```
# Load Packages/Data

library(tidyverse)
library(readr)
library(rsample)
library(broom)
library(rcfss)
library(ISLR)
library(yardstick)
library(caret)
library(randomForest)
library(pls)
library(knitr)

nes <- read_csv("./nes2008.csv")
```

1. (10 points) Estimate the MSE of the model using the traditional approach. That is, fit the linear regression model using the *entire* dataset and calculate the mean squared error for the *entire* dataset. Present and discuss your results at a simple, high level.

```
# Fit Linear Model

Model1 <- lm(nes$biden~nes$female+nes$age+nes$educ+nes$dem+nes$rep)
summary(Model1)
```

```
##
## Call:
## lm(formula = nes$biden ~ nes$female + nes$age + nes$educ + nes$dem +
##     nes$rep)
##
## Residuals:
##     Min      1Q  Median      3Q     Max
## -75.546 -11.295   1.018  12.776  53.977
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  58.81126    3.12444  18.823  < 2e-16 ***
## nes$female    4.10323    0.94823   4.327 1.59e-05 ***
## nes$age       0.04826    0.02825   1.708   0.0877 .
## nes$educ     -0.34533    0.19478  -1.773   0.0764 .
## nes$dem      15.42426    1.06803  14.442  < 2e-16 ***
## nes$rep     -15.84951    1.31136 -12.086  < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
```

```
## Residual standard error: 19.91 on 1801 degrees of freedom
## Multiple R-squared:  0.2815, Adjusted R-squared:  0.2795
## F-statistic: 141.1 on 5 and 1801 DF,  p-value: < 2.2e-16
```

```
# Calculate MSE

mse1 <- mean(Model1$residuals^2)
mse1
```

```
## [1] 395.2702
```

The mean squared error (MSE) first eliminates negative directionality by squaring the residuals, or the distance betweened the observed observation and observation expected by the regression line. It then takes arithmatic mean of those products. One easy way to intepret MSE is to convert it back to a mean error by taking the square root of the MSE. In this case, the root mean squared error (RMSE) is about 20, meaning that, on average, the expected thermometer rating was about 20 units off (either higher or lower) from the observed rating.

2. (30 points) Calculate the test MSE of the model using the simple holdout validation approach.

- (5 points) Split the sample set into a training set (50%) and a holdout set (50%).

```
# Split nes into test and train

set.seed(1)

nes_split <- initial_split(data = nes,
                           prop = 0.5)

nes_train <- training(nes_split)
nes_test <- testing(nes_split)
```

- (5 points) Fit the linear regression model using only the training observations.

```
# Fit Linear Model Using nes_train

Model2 <- lm(biden~female+age+educ+dem+rep, data = nes_train)
      mse2  <- mean(Model2$residuals^2)
summary(Model2)
```

```
##
## Call:
## lm(formula = biden ~ female + age + educ + dem + rep, data = nes_train)
##
## Residuals:
##     Min      1Q  Median      3Q     Max
## -75.875 -10.974   0.638  13.968  45.989
##
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)
## (Intercept)  61.94663    4.52928  13.677  < 2e-16 ***
## female        5.14561    1.38493   3.715 0.000215 ***
## age          -0.02402    0.04197  -0.572 0.567281
## educ         -0.46983    0.28126  -1.670 0.095179 .
```

```
## dem          16.27265    1.55652  10.454  < 2e-16 ***
## rep         -16.41671    1.96592  -8.351 2.55e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 20.74 on 898 degrees of freedom
## Multiple R-squared:  0.2799, Adjusted R-squared:  0.2759
## F-statistic:  69.8 on 5 and 898 DF,  p-value: < 2.2e-16
```

- (10 points) Calculate the MSE using only the test set observations.

```
# Calculate MSE on nse_test

mse3 <- augment(Model2, newdata = nes_test) %>%
                  mse(truth = biden, estimate = .fitted)

mse3$.estimate
```

```
## [1] 370.1792
```

- (10 points) How does this value compare to the training MSE from question 1? Present numeric comparison and discuss a bit.

The first MSE was 395.27, while the second from the split data set was lower at 370.18. We avoided an optimistically biased, overfit evaluation by estimating the model from the training data and evaluating it on the test data but since the training and test sets were generated through random processes, we can expect high variance between MSE predictions dependent on the composition of the split.

3. (30 points) Repeat the simple validation set approach from the previous question 1000 times, using 1000 different splits of the observations into a training set and a test/validation set. Visualize your results as a sampling distribution (hint: think histogram or density plots). Comment on the results obtained.

```
mc_data <- mc_cv(nes, prop = .5, times = 1000)
```

```
holdout_results <- function(splits) {

  Model3 <- lm(biden ~ female+age+educ+dem+rep, data = analysis(splits))

  res <- augment(Model3, newdata = assessment(splits)) %>%
    mse(truth = biden, estimate = .fitted)


  res
}
```

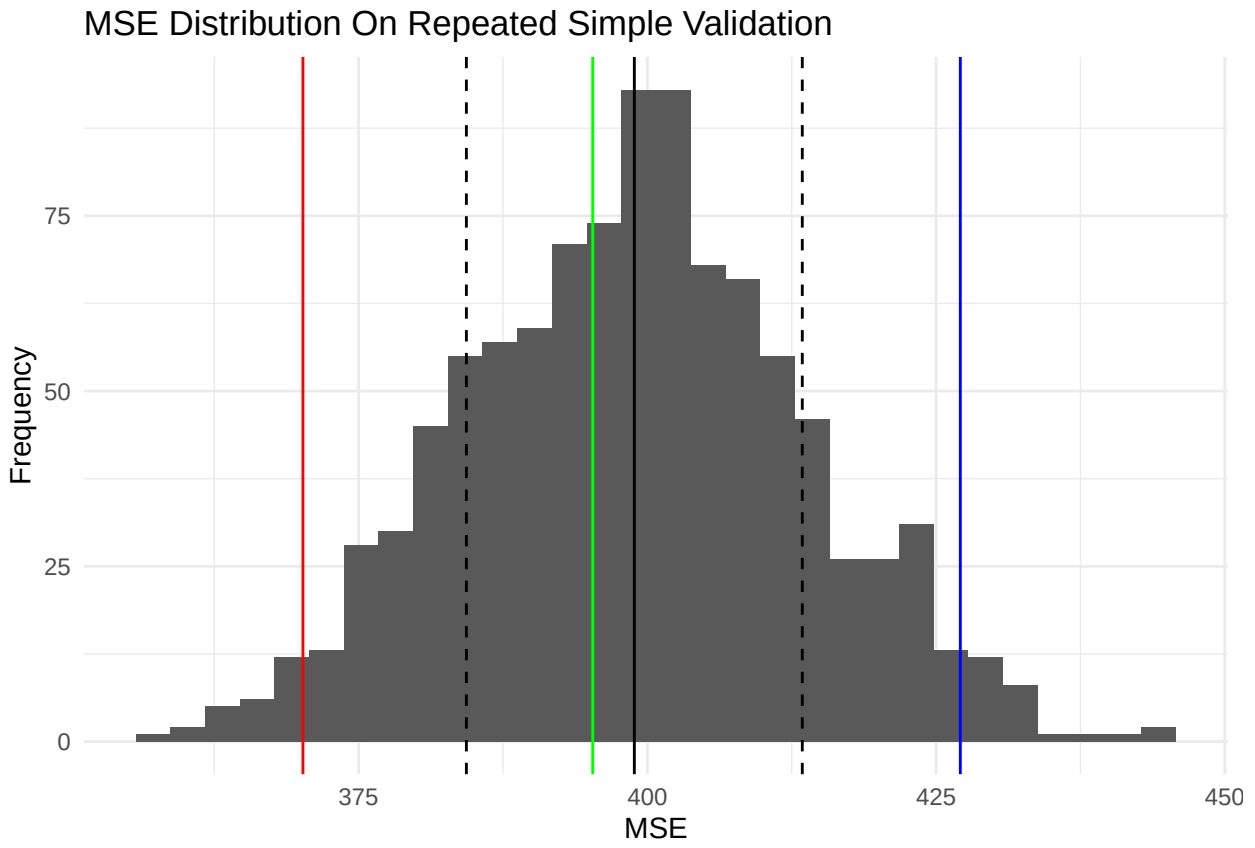
```
resample_data <- mc_data %>%
  mutate(results = map(splits, holdout_results)) %>%
  unnest(results) %>%
  spread(.metric, .estimate)

mse4 <- resample_data %>%
      summarize(mse = mean(mse))
```

```r
mse4_sd <- resample_data %>%
      summarize(mse = sd(mse))
```

```r
# Visualize distribution of RMSEs

ggplot(resample_data, aes(x = resample_data$mse)) +
      geom_histogram(bins = 30) +
      labs(title = "MSE Distribution On Repeated Simple Validation",
           x = "MSE",
           y = "Frequency") +
      theme_minimal() +
      geom_vline(xintercept =  mse1 ,color = "green") +
      geom_vline(xintercept = mse2, color = "blue") +
      geom_vline(xintercept = mse3$.estimate, color = "red") +
      geom_vline(xintercept = mse4$mse, color = "black") +
      geom_vline(xintercept = mse4$mse + mse4_sd$mse, color = "black", linetype = "dashed")+
      geom_vline(xintercept = mse4$mse - mse4_sd$mse, color = "black", linetype = "dashed")
```



The RMSE distributions suggest some instability in our model. Only two out of the three RMSE estimated by our models (the green and blue lines) fell within one standard deviation of the mean RMSE. In the histogram this is represented by the two dashed black lines about the solid black line (the mean). The RMSEs appear to be (roughly) normally distributed, so we would expect that about 68% of the RMSEs would fall within that range. The RMSE generated from our second model on the test data (the red line) fell outside of that range, however, highlighting how unrepresentative statistics can be when estimated from single, randomly generated holdouts.

4. (30 points) Compare the estimated parameters and standard errors from the original model in question 1 (the model estimated using *all of the available data*) to parameters and standard errors estimated using the bootstrap ($B = 1000$). Comparison should include, at a minimum, both numeric output as well as discussion on differences, similarities, etc. Talk also about the conceptual use and impact of bootstrapping.

```
# bootstrapped estimates of the parameter estimates and standard errors
lm_coefs <- function(splits, ...) {
  ## use `analysis` or `as.data.frame` to get the analysis data
  mod <- lm(..., data = analysis(splits))
  tidy(mod)
}
```

```
nes_female_boot <- nes %>%
            bootstraps(1000) %>%
            mutate(coef = map(splits, lm_coefs, as.formula(biden ~ poly(female, 1, raw = TRUE))))

nes_female_plot <- nes_female_boot %>%
                    unnest(coef) %>%
                    group_by(term) %>%
                    summarize(.estimate = mean(estimate),
                    .se = sd(estimate, na.rm = TRUE))
```

```
nes_age_boot <- nes %>%
            bootstraps(1000) %>%
            mutate(coef = map(splits, lm_coefs, as.formula(biden ~ poly(age, 1, raw = TRUE))))


nes_age_plot <- nes_age_boot %>%
                unnest(coef) %>%
                group_by(term) %>%
                summarize(.estimate = mean(estimate),
                  .se = sd(estimate, na.rm = TRUE))
```

```
nes_educ_boot <- nes %>%
            bootstraps(1000) %>%
            mutate(coef = map(splits, lm_coefs, as.formula(biden ~ poly(educ, 1, raw = TRUE))))

nes_educ_plot <- nes_educ_boot %>%
                 unnest(coef) %>%
                 group_by(term) %>%
                 summarize(.estimate = mean(estimate),
                  .se = sd(estimate, na.rm = TRUE))
```

```
nes_dem_boot <- nes %>%
            bootstraps(1000) %>%
            mutate(coef = map(splits, lm_coefs, as.formula(biden ~ poly(dem, 1, raw = TRUE))))

nes_dem_plot <- nes_dem_boot %>%
                 unnest(coef) %>%
                 group_by(term) %>%
                 summarize(.estimate = mean(estimate),
                  .se = sd(estimate, na.rm = TRUE))
```

```
nes_rep_boot <- nes %>%
            bootstraps(1000) %>%
            mutate(coef = map(splits, lm_coefs, as.formula(biden ~ poly(rep, 1, raw = TRUE))))

nes_rep_plot <- nes_rep_boot %>%
                unnest(coef) %>%
                group_by(term) %>%
                summarize(.estimate = mean(estimate),
                    .se = sd(estimate, na.rm = TRUE))
```

```
tidy(Model1) %>%
  kable()
```

term	estimate	std.error	statistic	p.value
(Intercept)	58.8112590	3.1244366	18.822996	0.0000000
nes$female	4.1032301	0.9482286	4.327258	0.0000159
nes$age	0.0482589	0.0282474	1.708438	0.0877274
nes$educ	-0.3453348	0.1947796	-1.772952	0.0764057
nes$dem	15.4242556	1.0680327	14.441745	0.0000000
nes$rep	-15.8495061	1.3113624	-12.086290	0.0000000

```
bind_rows(nes_female_plot[2,1:3],
      nes_age_plot[2,1:3],
      nes_educ_plot[2,1:3],
      nes_dem_plot[2,1:3],
      nes_rep_plot[2,1:3]) %>%
kable()
```

term	.estimate	.se
poly(female, 1, raw = TRUE)	6.1485008	1.0649897
poly(age, 1, raw = TRUE)	0.0612794	0.0337012
poly(educ, 1, raw = TRUE)	-0.9003602	0.2322893
poly(dem, 1, raw = TRUE)	21.8191156	0.9834975
poly(rep, 1, raw = TRUE)	-24.3992531	1.2057432

The parameter estimates from the bootstrap are similar to the ones estimated from the full dataset. The standard errors on the parameters estimated by bootstrapping are all larger than the the ones estimated from the full dataset. This finding relates to the idea that bootstrapping does not require any distributional assumptions because the standard error itself defines the distribution. We would then expect that estimates from a (correctly) assumed distribution would be more precise.

This finding implies that bootstrapping is a good method when there is no basis to make any distributional assumptions. Although the nes dataset had enough observations to generate a good estimate of the parameters, bootstrapping is also useful when working with datasets, which have few observations.