

HRP 203 Final Project

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Assignment Description

Create a 4 to 6 page Quarto reproducible document to synthesize concepts learned throughout the course and particularly from Module 3. Create the report in a repository on your GitHub account based on the cohort simulated data. Include the following features: introduction section, methods section with notation, results section with data summary table and at least 2 figures, discussion. The assignment is due at 11am Pacific time on Monday, June 10th.

Introduction

Methods

For the data analysis, I used the `cohort` simulated data set provided through the course GitHub repository. A copy of this data set can be found in the `Data` folder in the [GitHub repository](#) for this project.

	smoke	female	cardiac	age	cost
1	1	0	0	44	10566
2	0	1	0	46	9668
3	0	0	0	56	9889
4	0	0	0	35	9780
5	0	0	0	49	10200
6	0	0	0	64	10082

I included all of the variables from the data set in my analysis. Since documentation was not provided, I made assumptions about the meanings of the variables. These assumptions are outlined in Table 1.

Table 1: Definitions for five variables included in analysis

Variable Name	Description
smoke	indicator variable, equal to 1 if the patient smokes regularly (more than three times per week) and 0 if not
female	indicator variable, equal to 1 if the patient's sex is designated as female and 0 if not
cardiac	indicator variable, equal to 1 if the patient has a diagnosis of a cardiac-related problem and 0 if not
age	numeric variable, indicates the patient's age in years
cost	numeric variable, indicates the cost (in USD) of all healthcare visits for a given patient in a given year

I started the analysis with descriptive statistics for the five variables of interest.

For the next part of the analysis, I ran a regression to examine how an individual's smoking habits, sex, cardiac history, and age can be used to predict their yearly healthcare costs. I used the `lm()` function with the following equation:

$$cost = \beta_0 + \beta_1 smoke + \beta_2 female + \beta_3 cardiac + \beta_4 age$$

Finally, I generated predicted spending amounts for

Results

Data Summary Table

```
varnames <- as.matrix(names(cohort), nrow = 5, ncol = 1)
meanmat <- matrix(data = 0, nrow = 5, ncol = 1)
sdmat <- matrix(data = 0, nrow = 5, ncol = 1)
minmaxmat <- matrix(data = NA, nrow = 5, ncol = 2)
skewmat <- matrix(data = NA, nrow = 5)
for(i in 1:5){
  meanmat[i] <- round(mean(cohort[,i]), digits = 5)
  sdmat[i] <- round(sd(cohort[,i]), digits = 5)
  minmaxmat[i,1] <- round(min(cohort[,i]), digits = 5)
  minmaxmat[i,2] <- round(max(cohort[,i]), digits = 5)
  skewmat[i] <- round(skewness(cohort[,i]), digits = 5)
}
table <- cbind(varnames, minmaxmat, meanmat, sdmat, skewmat)
colnames(table) <- list("Variable", "Min", "Max", "Mean", "SD", "Skewness")
as.data.frame(table)
```

	Variable	Min	Max	Mean	SD	Skewness
1	smoke	0	1	0.1016	0.30215	2.63656
2	female	0	1	0.487	0.49988	0.052
3	cardiac	0	1	0.038	0.19122	4.83128
4	age	18	65	41.4702	13.5407	0.01173
5	cost	8478	11326	9672.2744	402.63168	0.32417

Figure 1

```
# cost regression
reg1 <- lm(cost ~ cardiac + smoke + age + female, data = cohort)
summary(reg1)
```

Call:

```
lm(formula = cost ~ cardiac + smoke + age + female, data = cohort)
```

Residuals:

	Min	1Q	Median	3Q	Max
	-700.87	-137.95	-0.95	136.99	759.92

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	8988.7981	9.5392	942.30	<2e-16 ***
cardiac	289.2236	15.2189	19.00	<2e-16 ***
smoke	592.7583	9.5149	62.30	<2e-16 ***
age	18.2124	0.2081	87.50	<2e-16 ***
female	-293.6548	5.7041	-51.48	<2e-16 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 199.2 on 4995 degrees of freedom

Multiple R-squared: 0.7555, Adjusted R-squared: 0.7553

F-statistic: 3859 on 4 and 4995 DF, p-value: < 2.2e-16

```
coefffig <- coefplot(reg1,
                      title = "Coefficients for Linear Regression",
                      color = "Maroon")
coefffig
```

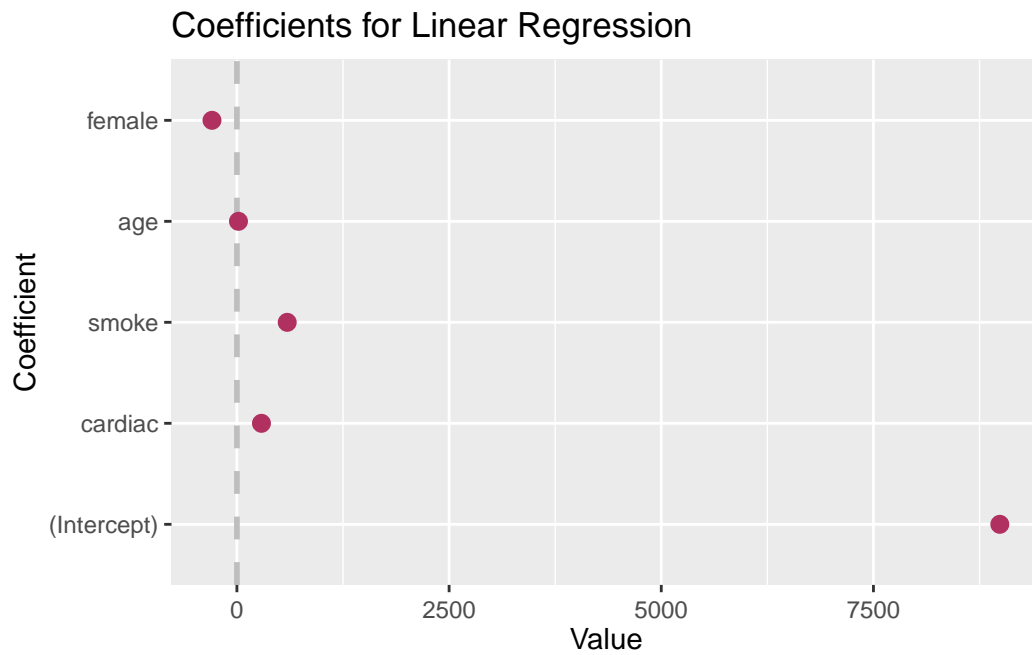
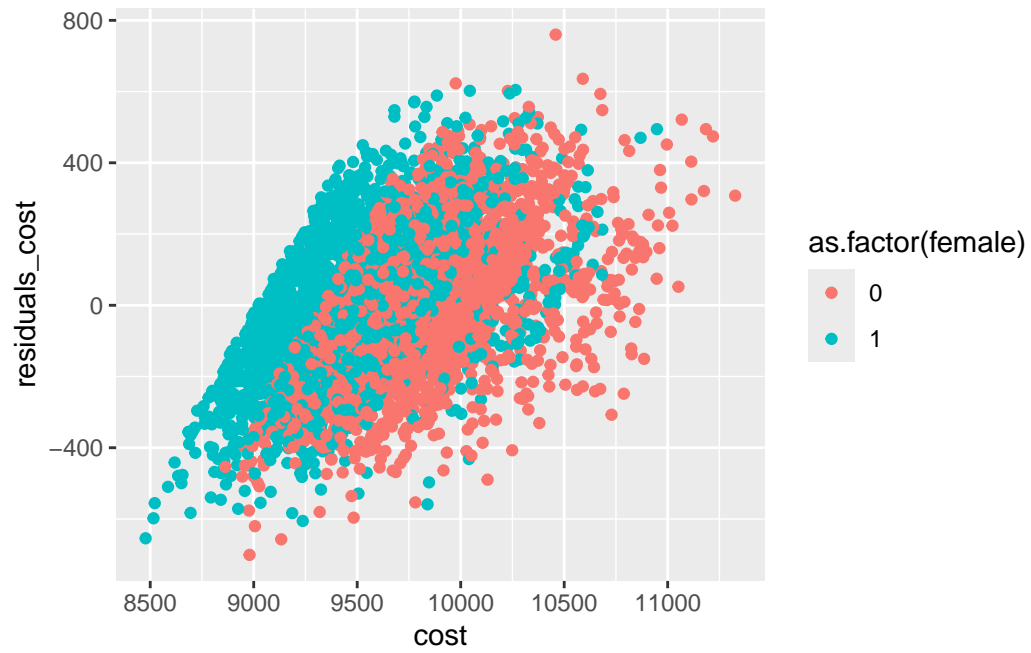


Figure 2

```
# make predictions
cohort$predict_cost <- predict(reg1, cohort)
cohort$residuals_cost <- cohort$cost - cohort$predict_cost

# plot residuals by sex
ggplot(cohort, aes(cost, residuals_cost)) +
  geom_point(aes(color = as.factor(female)))
```



Discussion

```
1 + 1
```

```
[1] 2
```