

Medical Cost Analysis Project

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R Markdown

This project analyzes the Medical Cost Personal Dataset from Kaggle, which contains 1,338 patient records with variables such as age, sex, BMI, smoking status, number of children, region, and individual medical charges.

The goal of this project is to explore how demographic and lifestyle factors influence medical costs. The response variable for the regression model is charges, representing the total medical cost billed to an individual. Predictor variables include age, BMI, number of children, and smoking status

```
install.packages("ggplot2")

## Installing package into '/cloud/lib/x86_64-pc-linux-gnu-library/4.5'
## (as 'lib' is unspecified)
install.packages("dplyr")

## Installing package into '/cloud/lib/x86_64-pc-linux-gnu-library/4.5'
## (as 'lib' is unspecified)
library(ggplot2)
library(dplyr)

##
## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
## 
##     filter, lag

## The following objects are masked from 'package:base':
## 
##     intersect, setdiff, setequal, union

Loading the Dataset

insurance <- read.csv("insurance.csv")
head(insurance)

##   age   sex   bmi children smoker   region   charges
## 1 19 female 27.900       0    yes southwest 16884.924
## 2 18 male 33.770       1     no southeast  1725.552
## 3 28 male 33.000       3     no southeast  4449.462
## 4 33 male 22.705       0     no northwest 21984.471
## 5 32 male 28.880       0     no northwest  3866.855
## 6 31 female 25.740      0     no southeast  3756.622
```

Data Exploration

```
summary(insurance)

##      age          sex          bmi        children
##  Min.   :18.00  Length:1338    Min.   :15.96  Min.   :0.000
##  1st Qu.:27.00  Class  :character  1st Qu.:26.30  1st Qu.:0.000
##  Median :39.00  Mode   :character  Median :30.40  Median :1.000
##  Mean   :39.21                  Mean   :30.66  Mean   :1.095
##  3rd Qu.:51.00                  3rd Qu.:34.69  3rd Qu.:2.000
##  Max.   :64.00                  Max.   :53.13  Max.   :5.000
##      smoker         region       charges
##  Length:1338    Length:1338    Min.   : 1122
##  Class  :character  Class  :character  1st Qu.: 4740
##  Mode   :character  Mode   :character  Median : 9382
##                           Mean   :13270
##                           3rd Qu.:16640
##                           Max.   :63770
```

```
sd(insurance$age)
```

```
## [1] 14.04996
```

```
sd(insurance$bmi)
```

```
## [1] 6.098187
```

```
sd(insurance$charges)
```

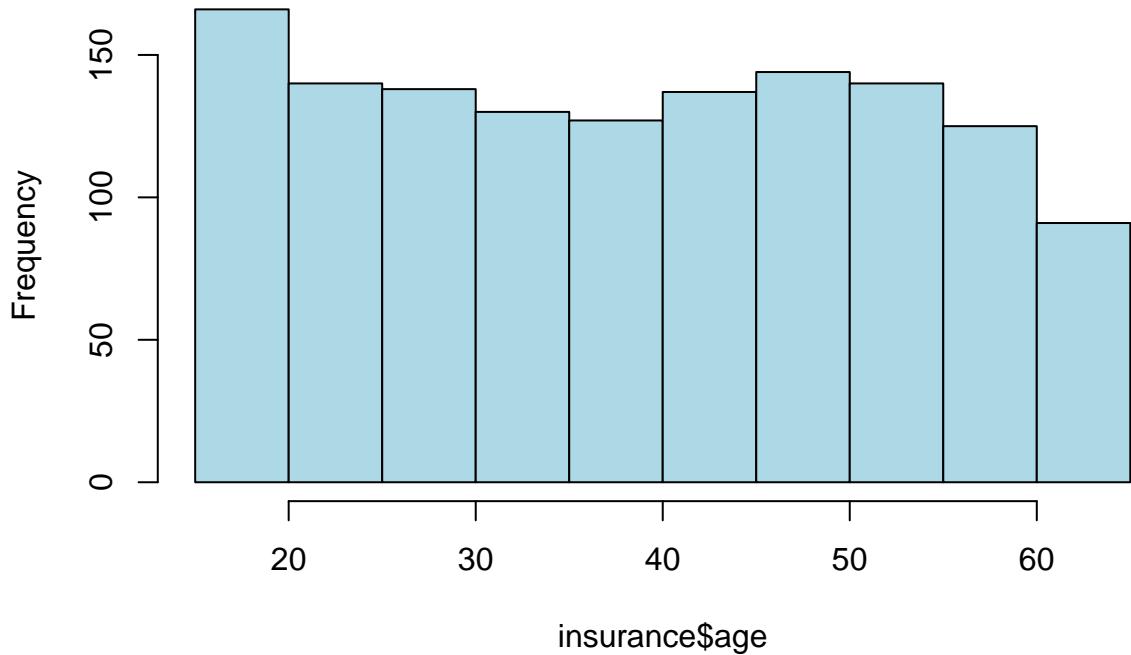
```
## [1] 12110.01
```

Creating Data Visualisations

Histograms

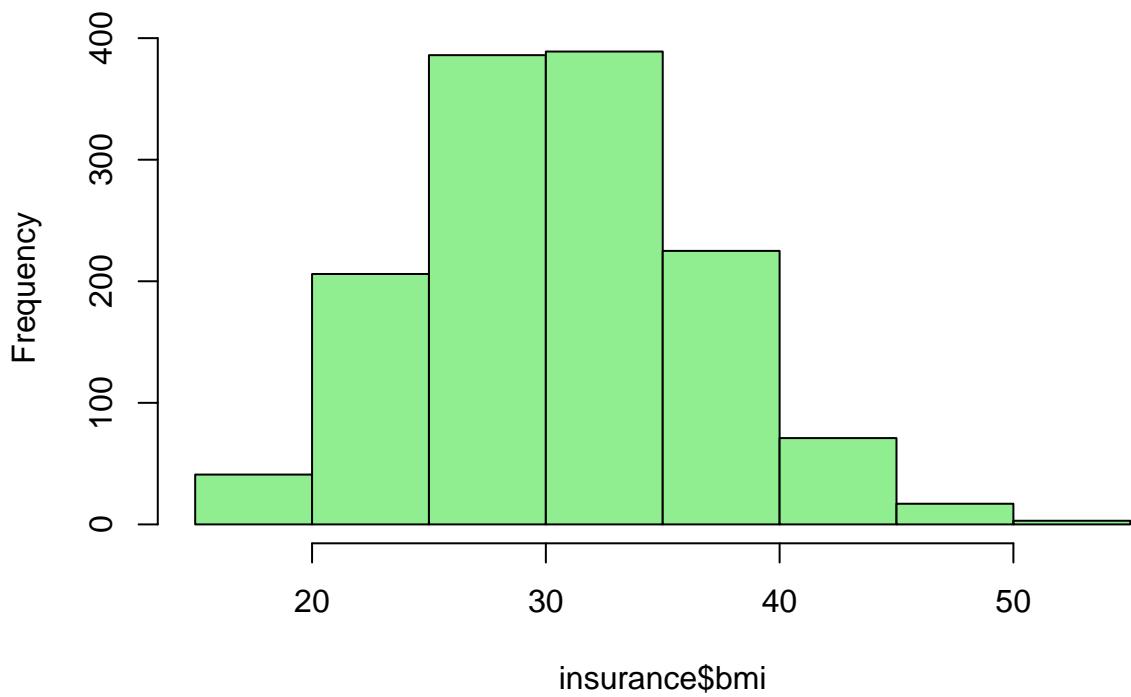
```
hist(insurance$age, main="Age Distribution", col="lightblue")
```

Age Distribution



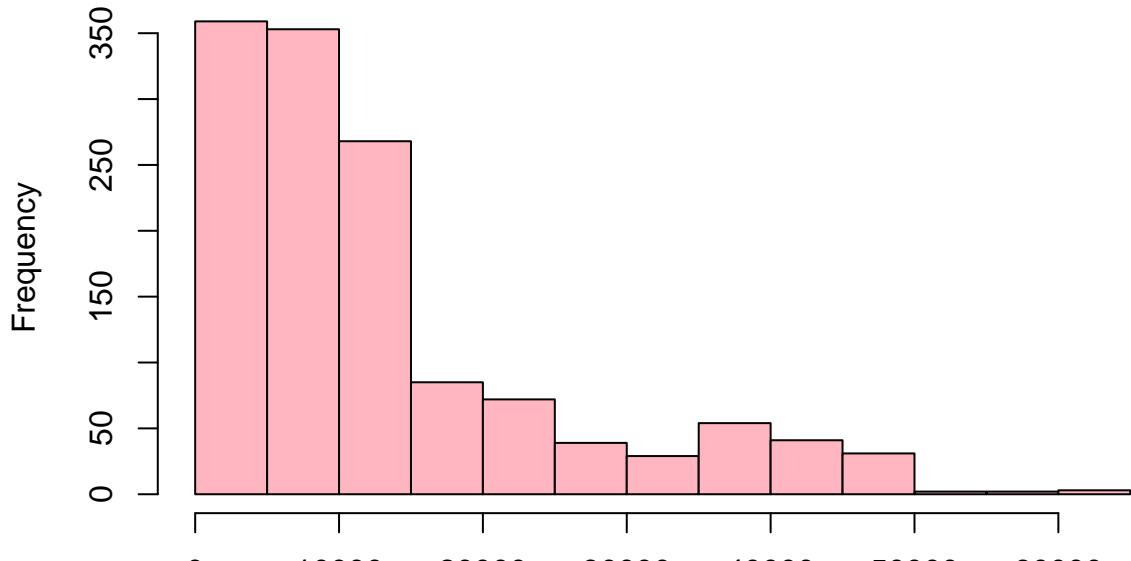
```
hist(insurance$bmi, main="BMI Distribution", col="lightgreen")
```

BMI Distribution



```
hist(insurance$charges, main="Medical Charges Distribution", col="lightpink")
```

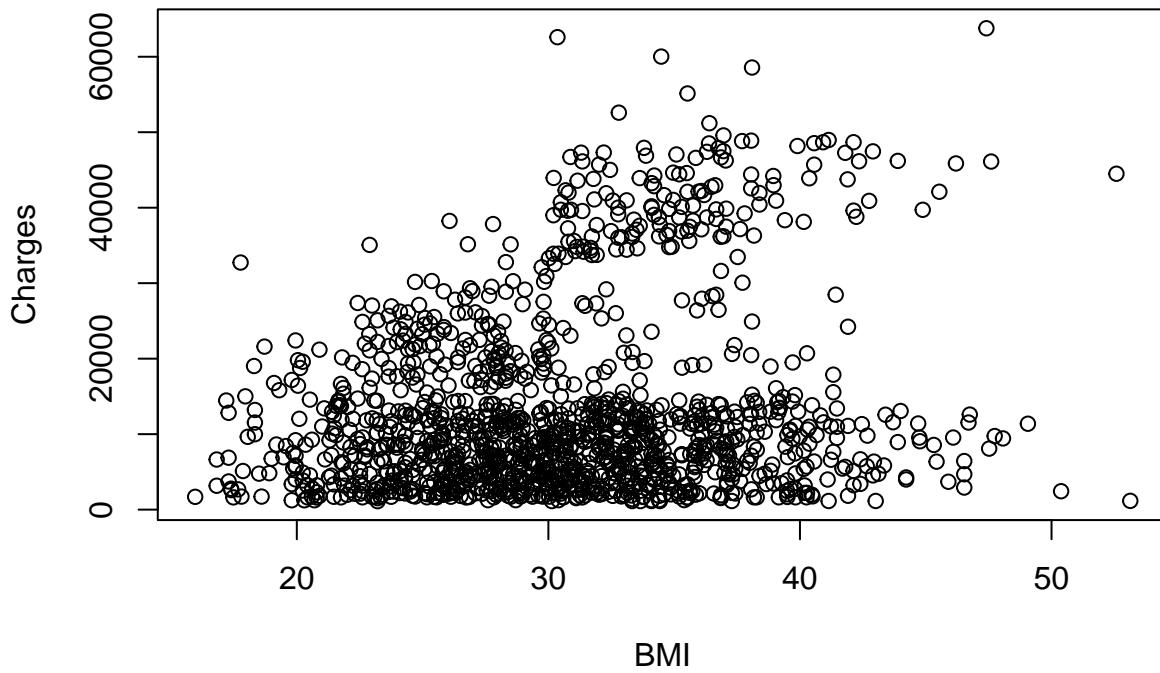
Medical Charges Distribution



insurance\$charges
Scatter-
plots

```
plot(insurance$bmi, insurance$charges,  
     main="BMI vs Charges", xlab="BMI", ylab="Charges")
```

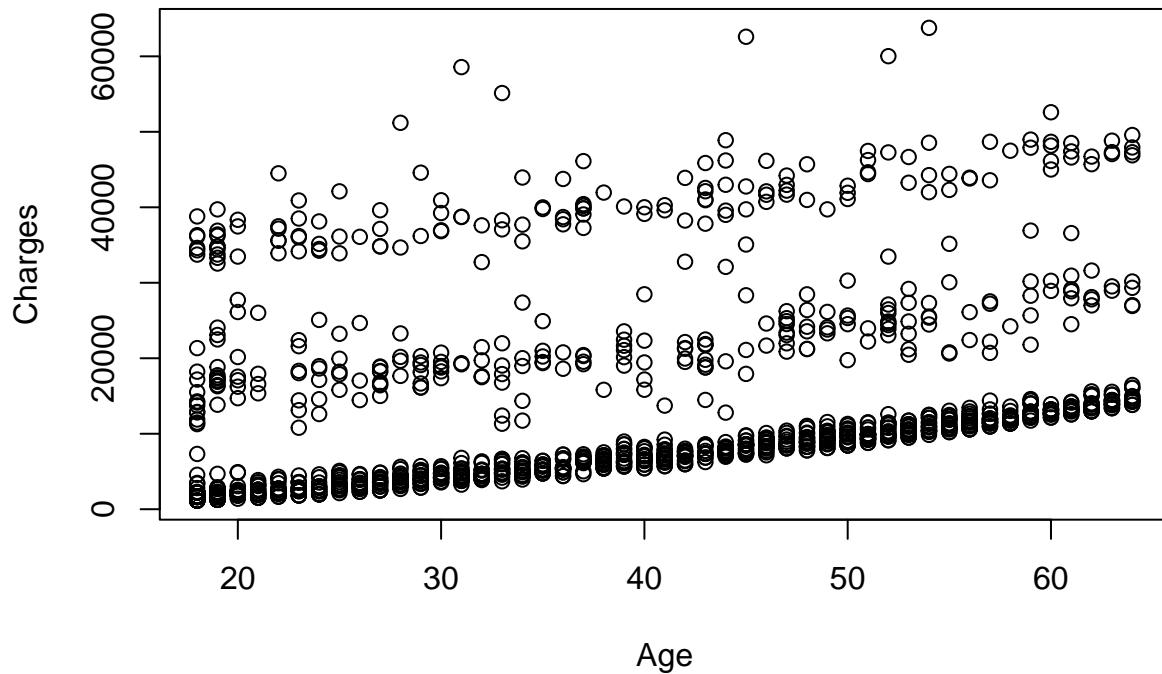
BMI vs Charges



```
plot(insurance$age, insurance$charges,
```

```
main="Age vs Charges", xlab="Age", ylab="Charges")
```

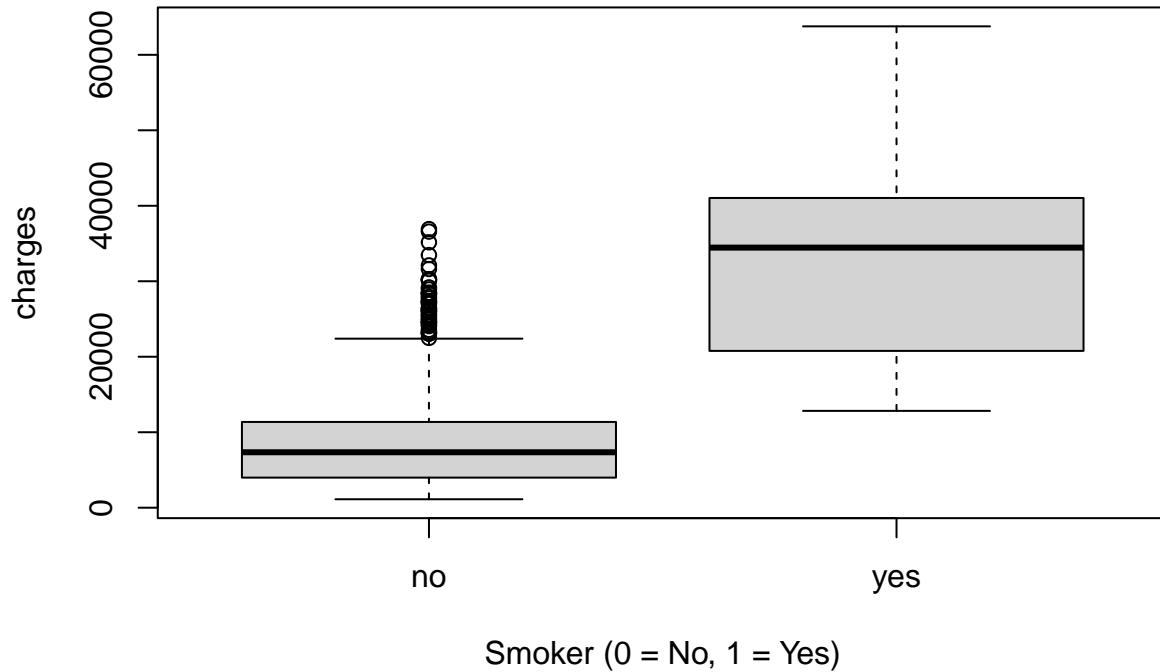
Age vs Charges



Boxplot(smoker status)

```
boxplot(charges ~ smoker, data=insurance,
        main="Charges by Smoking Status",
        xlab="Smoker (0 = No, 1 = Yes)")
```

Charges by Smoking Status



Data Preprocessing

```
insurance$smoker <- ifelse(insurance$smoker == "yes", 1, 0)
insurance$sex <- ifelse(insurance$sex == "male", 1, 0)
```

Linear Regression Model

```
model <- lm(charges ~ age + bmi + children + smoker, data=insurance)
summary(model)

##
## Call:
## lm(formula = charges ~ age + bmi + children + smoker, data = insurance)
##
## Residuals:
##      Min       1Q   Median       3Q      Max 
## -11897.9  -2920.8  -986.6  1392.2 29509.6 
##
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)    
## (Intercept) -12102.77    941.98 -12.848 < 2e-16 ***
## age          257.85     11.90   21.675 < 2e-16 ***
## bmi          321.85     27.38   11.756 < 2e-16 ***
## children     473.50     137.79   3.436 0.000608 ***
## smoker       23811.40    411.22   57.904 < 2e-16 ***
## ---        
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 6068 on 1333 degrees of freedom
## Multiple R-squared:  0.7497, Adjusted R-squared:  0.7489 
## F-statistic: 998.1 on 4 and 1333 DF,  p-value: < 2.2e-16
```

Residual Plot

```
plot(model$residuals,
  main="Residual Plot",
  xlab="Index",
  ylab="Residuals")
abline(h=0, col="red")
```

