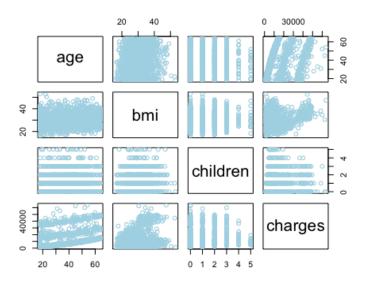
Medical Cost Analysis

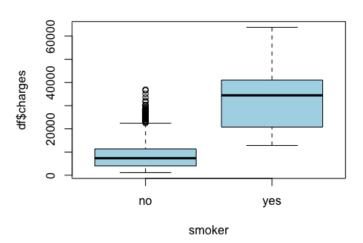
Mark Arzola

```
knitr::opts chunk$set(echo = TRUE)
install.packages("ggplot2", repos = "https://cran.r-project.org")
##
## The downloaded binary packages are in
    /var/folders/3w/rdvgs5053xz_4sgwf7mp_p300000gn/T//Rtmp55VEqi/downloaded_packages
library(ggplot2)
setwd('/Users/markarzola/Desktop/portfolio projects/Insurance Analysis Project')
df <- read.csv('insurance.csv', header=TRUE)</pre>
head(df)
##
                   bmi children smoker
                                           region
     age
            sex
                                                    charges
     19 female 27.900
                                   yes southwest 16884.924
                              0
           male 33.770
                              1
                                    no southeast 1725.552
## 2
      18
## 3
           male 33.000
      28
                              3
                                    no southeast 4449.462
           male 22.705
                                    no northwest 21984.471
## 4 33
                              0
           male 28.880
                              0
                                    no northwest 3866.855
## 5
      32
## 6
      31 female 25.740
                                    no southeast 3756.622
numericdata \leftarrow df[ , c(1,3,4,7)]
plot(numericdata, col = "lightblue")
```

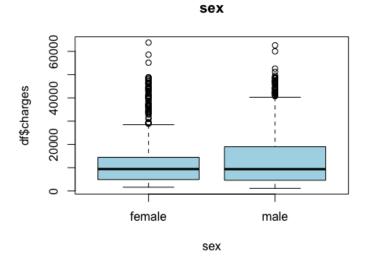


```
round(cor(numericdata),2)
##
             age bmi children charges
## age
            1.00 0.11
                          0.04
                                   0.30
                                   0.20
## bmi
            0.11 1.00
                          0.01
                                   0.07
## children 0.04 0.01
                          1.00
## charges 0.30 0.20
                          0.07
                                   1.00
smoker = as.factor(df$smoker)
sex = as.factor(df$sex)
region = as.factor(df$region)
boxplot(df$charges ~ smoker, main = 'smoker', col = "lightblue")
```

smoker



boxplot(df\$charges ~ sex, main = 'sex', col = "lightblue")



```
boxplot(df$charges ~ region, main = 'region', col = "lightblue")
```

northeast northwest southeast

region

region

```
model1 = lm(charges ~., data = df)
summary(model1)
##
## Call:
## lm(formula = charges ~ ., data = df)
##
## Residuals:
                       Median
##
        Min
                  10
                                    3Q
                                            Max
            -2848.1
                       -982.1
##
  -11304.9
                                1393.9
                                        29992.8
##
## Coefficients:
##
                   Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                   -11938.5
                                 987.8 -12.086 < 2e-16 ***
## age
                      256.9
                                  11.9 21.587 < 2e-16 ***
                                 332.9 -0.394 0.693348
## sexmale
                     -131.3
## bmi
                      339.2
                                  28.6 11.860 < 2e-16 ***
## children
                      475.5
                                 137.8
                                        3.451 0.000577 ***
                                 413.1 57.723 < 2e-16 ***
## smokeryes
                    23848.5
## regionnorthwest
                                 476.3 -0.741 0.458769
                     -353.0
## regionsoutheast
                   -1035.0
                                 478.7 -2.162 0.030782 *
## regionsouthwest
                                 477.9 -2.009 0.044765 *
                     -960.0
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 6062 on 1329 degrees of freedom
## Multiple R-squared: 0.7509, Adjusted R-squared: 0.7494
## F-statistic: 500.8 on 8 and 1329 DF, p-value: < 2.2e-16
```