Assignment 2

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Exercise 1

a) To investigate whether tree type influences total wood volume, we can perform a one-way ANOVA.

```
tree_df$type <- as.factor(tree_df$type)</pre>
tree_type_lm <- lm(volume~type, data=tree_df)</pre>
anova(tree_type_lm)
## Analysis of Variance Table
##
## Response: volume
             Df Sum Sq Mean Sq F value Pr(>F)
## type
                   380
                           380
                                    1.9
                                          0.17
## Residuals 57 11395
                           200
summary(tree_type_lm)
##
## Call:
## lm(formula = volume ~ type, data = tree_df)
##
## Residuals:
      Min
              1Q Median
                            3Q
                                   Max
## -19.97 -9.96 -2.77
                          5.94 46.83
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
##
                  30.17
                              2.54
## (Intercept)
                                      11.88
                                              <2e-16 ***
                   5.08
                              3.69
                                       1.38
                                                0.17
## typeoak
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## Residual standard error: 14.1 on 57 degrees of freedom
## Multiple R-squared: 0.0322, Adjusted R-squared: 0.0153
## F-statistic: 1.9 on 1 and 57 DF, p-value: 0.174
```

With p > 0.05, we can conclude that type does not have a significant effect on volume. Because the factor type has two levels, we can apply a two sample t-test.

```
mask <- tree_df$type == "beech"
t.test(tree_df$volume[mask], tree_df$volume[!mask])</pre>
```

This supports the result from the ANOVA test. The estimated volume is 30.2 for Beech trees and 35.2 for Oak trees.

b) To investigate this claim, we create two models, each including all three explanatory variables (type, diameter and height). In the first model, we also include the pairwise interaction between type and diameter.

```
tree_type_d_lm <- lm(volume~height+type*diameter, data=tree_df)
anova(tree_type_d_lm)</pre>
```

```
## Analysis of Variance Table
##
## Response: volume
##
                 Df Sum Sq Mean Sq F value Pr(>F)
## height
                  1
                      2188
                              2188 206.21 < 2e-16 ***
                  1
                                      40.65 4.2e-08 ***
## type
                       431
                               431
## diameter
                  1
                      8577
                              8577
                                    808.49 < 2e-16 ***
## type:diameter
                                  6
                  1
                         6
                                       0.52
                                               0.47
## Residuals
                                 11
                 54
                       573
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
summary(tree_type_d_lm)
```

```
##
## Call:
## lm(formula = volume ~ height + type * diameter, data = tree_df)
##
```

```
## Residuals:
##
     Min
              1Q Median
                            3Q
                                  Max
## -7.350 -2.194 -0.141 1.701 8.176
##
## Coefficients:
                    Estimate Std. Error t value Pr(>|t|)
##
## (Intercept)
                     -63.873
                                  5.539 -11.53 3.5e-16 ***
## height
                       0.434
                                  0.079
                                           5.49 1.1e-06 ***
## typeoak
                      -4.963
                                  5.149
                                          -0.96
                                                    0.34
## diameter
                       4.608
                                  0.207
                                          22.26 < 2e-16 ***
## typeoak:diameter
                                           0.72
                                                    0.47
                       0.259
                                  0.359
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
##
## Residual standard error: 3.26 on 54 degrees of freedom
## Multiple R-squared: 0.951, Adjusted R-squared: 0.948
## F-statistic: 264 on 4 and 54 DF, p-value: <2e-16
tree_type_h_lm <- lm(volume~diameter+type*height, data=tree_df)</pre>
anova(tree_type_h_lm)
## Analysis of Variance Table
## Response: volume
               Df Sum Sq Mean Sq F value Pr(>F)
                1 10827
                           10827 1045.97 < 2e-16 ***
## diameter
                              45
                                    4.37
                                           0.041 *
## type
                1
                      45
## height
                     324
                             324
                                   31.32 7.5e-07 ***
                1
## type:height 1
                      19
                              19
                                    1.88
                                           0.176
## Residuals
             54
                     559
                              10
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
summary(tree_type_h_lm)
##
## Call:
## lm(formula = volume ~ diameter + type * height, data = tree_df)
##
## Residuals:
##
      Min
              1Q Median
                            3Q
                                  Max
## -6.230 -2.113 -0.161 1.801 8.165
##
## Coefficients:
##
                  Estimate Std. Error t value Pr(>|t|)
                                        -8.09
                                                 7e-11 ***
## (Intercept)
                   -57.551
                                7.111
```

```
## diameter
                     4.779
                                0.173
                                        27.55
                                                <2e-16 ***
## typeoak
                   -17.471
                               11.826
                                        -1.48
                                                0.1454
## height
                     0.321
                                0.102
                                         3.14
                                                0.0027 **
## typeoak:height
                     0.212
                                         1.37
                                0.154
                                                0.1761
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
## Residual standard error: 3.22 on 54 degrees of freedom
## Multiple R-squared: 0.953, Adjusted R-squared: 0.949
## F-statistic: 271 on 4 and 54 DF, p-value: <2e-16
```

We see that both pairwise interactions are not significant. Therefore, we can conclude that both height and diameter have the same influence regardless of type. Both models suggest that all three explanatory variables have a significant effect individually.

c)

In (b), we saw that the interactions of *height* and *diameter* with *type* were not significant, and so we will investigate a purely additive model (assuming no interactions).

```
tree_add_all_lm <- lm(volume~diameter+height+type, data=tree_df)
drop1(tree_add_all_lm, test= "F")</pre>
```

```
## Single term deletions
##
## Model:
## volume ~ diameter + height + type
##
           Df Sum of Sq RSS AIC F value Pr(>F)
## <none>
                          578 143
## diameter 1
                    8577 9155 304 815.61 < 2e-16 ***
## height
             1
                                    30.82 8.4e-07 ***
                     324 903 167
## type
                      23
                         602 143
                                     2.21
                                             0.14
## ---
## Signif. codes:
                   0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
```

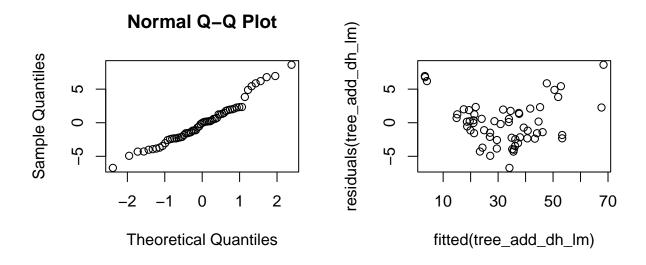
We see that the effect of type is not significant in the additive model. Therefore we will investigate an additive model that excludes type.

```
tree_add_dh_lm <- lm(volume~diameter+height, data=tree_df)
anova(tree_add_dh_lm)</pre>
```

```
## Residuals 56
                   602
                             11
## ---
                   0 '*** 0.001 '** 0.01 '* 0.05 '. ' 0.1 ' 1
## Signif. codes:
summary(tree_add_dh_lm)
##
## Call:
## lm(formula = volume ~ diameter + height, data = tree_df)
##
  Residuals:
##
      Min
              1Q Median
                             3Q
                                   Max
  -6.724 -2.278 -0.034
                         1.820
                                8.629
##
##
##
  Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
  (Intercept) -64.3697
                             5.5577
                                     -11.58
                                             < 2e-16 ***
  diameter
                 4.6325
                             0.1602
                                      28.92
                                             < 2e-16 ***
## height
                 0.4289
                             0.0755
                                       5.68
                                             5.1e-07 ***
##
                   0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
## Signif. codes:
## Residual standard error: 3.28 on 56 degrees of freedom
## Multiple R-squared: 0.949, Adjusted R-squared:
```

F-statistic: 520 on 2 and 56 DF, p-value: <2e-16

This model has almost the same R-squared value as before, while using fewer variables. Since simpler models are generally preferred, this is our model of choice to make predictions. As a final test, we need to check this model's assumptions to ensure that the conclusions we draw from it are valid:



While these plots are not perfect, we believe the model assumptions to be valid.

Therefore, the effects of type, diameter and height can be summarized as follows:

- The tree *type* does not affect volume significantly.
- Looking at the coefficients, we see that increasing both height and diameter result in an increase in volume, with diameter having a bigger impact (with a gradient of 4.63 compared to *height's* 0.43). This makes sense given that we know volume is proportional to the the square of the diameter.

To predict the volume for a tree with the overall average diameter and height, we can use the following linear regression model:

```
volume = -64.37 + 4.63 * diameter + 0.43 * height
```

```
mean_d <- mean(tree_df$diameter)
mean_h <- mean(tree_df$height)
means <- data.frame(diameter=c(mean_d), height=c(mean_h))
predict(tree_add_dh_lm, means, se.fit = TRUE)</pre>
```

```
## $fit
## 1
## 32.6
##
## $se.fit
## [1] 0.427
##
## $df
## [1] 56
##
## $residual.scale
## [1] 3.28
```

Therefore we expect the volume for such a tree to be 32.6.

d) Assuming that a tree is roughly cylindrical, we expect that *volume* would be proportional to the *height* multiplied by the square of *diameter*. We perform this transformation and add it as a new column in the data frame. We could apply the true transformation, $V = h \times \pi (d/2)^2$, but this would just add unnecessary constants which would already be captured in the regression coefficients.

```
tree_df$math_volume <- tree_df$height * tree_df$diameter^2
math_volume_lm <- lm(volume~math_volume, data=tree_df)
anova(math_volume_lm)</pre>
```

```
## Analysis of Variance Table
##
## Response: volume
```

```
## Df Sum Sq Mean Sq F value Pr(>F)

## math_volume 1 11477 11477 2201 <2e-16 ***

## Residuals 57 297 5

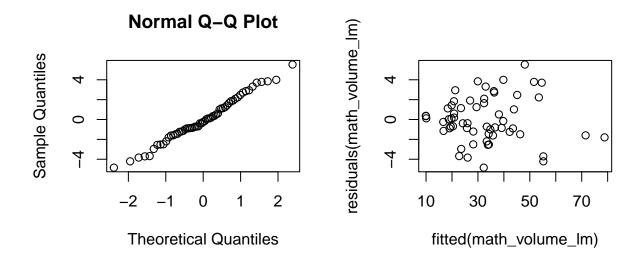
## ---

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

summary(math_volume_lm)
```

```
##
## Call:
  lm(formula = volume ~ math_volume, data = tree_df)
##
## Residuals:
##
      Min
              1Q Median
                            3Q
                                  Max
  -4.846 -1.343 -0.245
                         1.533
                                5.532
##
##
## Coefficients:
##
                Estimate Std. Error t value Pr(>|t|)
## (Intercept) -3.79e-01
                           7.63e-01
                                        -0.5
                                                 0.62
## math_volume 2.14e-03
                           4.57e-05
                                        46.9
                                               <2e-16 ***
                   0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
## Signif. codes:
##
## Residual standard error: 2.28 on 57 degrees of freedom
## Multiple R-squared: 0.975, Adjusted R-squared:
## F-statistic: 2.2e+03 on 1 and 57 DF, p-value: <2e-16
```

We see that this transformation does indeed produce an explanatory value with a significant effect. We also see that the R-squared value of 0.975 is higher than that of the previous models, indicating that it better explains the data. Finally, we check the assumptions of this model.

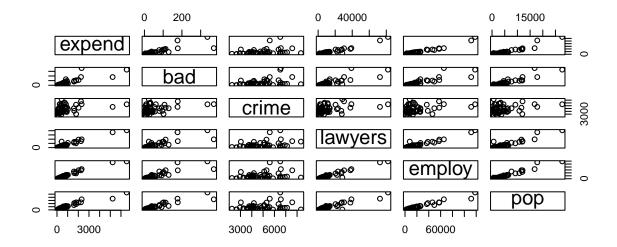


These plots are acceptable, meaning we can accept the model assumptions.

Exercise 2

a) ««« INCLUDE OTHER GRAPHICAL SUMMARIES??? »»»»

To investigate the interactions between all the variables of interest, we can plot the pairwise scatter plots for all their combinations:



We see that *expend*, our response variable, appears to have a positive correlation with all the explanatory variables except for *crime*. There appear to be several outliers at the high end of the data which could skew the model. We can also see that collinearity exists between the explanatory variables *bad*, *lawyers*, *employ* and *pop*. This is a problem since the redundant information will make the regression coefficients difficult to estimate.

We can use Cook's distance to find the influence points (a distance greater than 1 indicates an outlier)

```
crime_lm <- lm(expend~bad+crime+lawyers+employ+pop, data=crime_df)
cooks.distance(crime_lm)[cooks.distance(crime_lm) > 1]
```

```
## 5 8 35 44
## 4.91 3.51 1.09 2.70
```

We can see that indices of 5, 8, 35 and 44 are outliers, which we can remove:

```
crime_df_upd <- crime_df[-c(5,8,35,44),]</pre>
```

To further investigate collinearity, we can examine the correlations between all the explanatory variables, which confirms strong correlations between bad, lawyers, employ and pop.

```
round(cor(crime_df[, c(exp_vars)]), 2)
```

```
bad crime lawyers employ pop
##
           1.00 0.37
                         0.83
                                0.87 0.92
## bad
           0.37 1.00
## crime
                         0.38
                                0.31 0.28
## lawyers 0.83 0.38
                         1.00
                                0.97 0.93
## employ 0.87 0.31
                         0.97 1.00 0.97
## pop
           0.92 0.28
                         0.93 0.97 1.00
«««« IS VIF NECCESSARY? »»»»
To resolve the problem of collinearity, we can iteratively remove variables based on their VIF-values
as follows:
Full model:
vif(lm(expend~bad+crime+lawyers+employ+pop, data=crime_df))
##
       bad
             crime lawyers employ
                                        pop
##
      8.36
              1.49
                     16.97
                             33.59
                                      32.94
Remove employ:
vif(lm(expend~bad+crime+lawyers+pop, data=crime_df_upd))
##
       bad
             crime lawyers
                               pop
##
      7.16
              1.34
                     12.40
                             20.72
Remove pop:
vif_lm = lm(expend~bad+crime+lawyers, data=crime_df_upd)
vif(vif_lm)
##
       bad
             crime lawyers
##
      3.99
                      3.78
              1.13
summary(vif_lm)
##
## Call:
## lm(formula = expend ~ bad + crime + lawyers, data = crime_df_upd)
##
## Residuals:
              1Q Median
##
      Min
                            3Q
                                   Max
## -328.4 -42.4 -14.2 33.8 355.3
##
## Coefficients:
```

Estimate Std. Error t value Pr(>|t|)

##

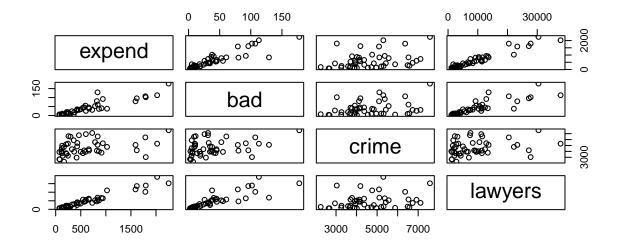
```
## (Intercept) -113.5051
                            66.5587
                                      -1.71
                                             0.09535 .
                             0.8845
## bad
                  3.7457
                                       4.23
                                             0.00012 ***
                  0.0333
                             0.0145
                                       2.30
                                             0.02655 *
## crime
                             0.0039
## lawyers
                  0.0456
                                      11.68
                                            6.3e-15 ***
## ---
                  0 '*** 0.001 '** 0.01 '* 0.05 '. ' 0.1 ' 1
## Signif. codes:
##
## Residual standard error: 118 on 43 degrees of freedom
## Multiple R-squared: 0.957, Adjusted R-squared: 0.954
## F-statistic: 322 on 3 and 43 DF, p-value: <2e-16
```

In the resulting model, all the explanatory variables are significant.

«««« SHOULD WE SHOW PLOT AGAIN?? »»»»

Therefore, after removing the influence points and collinear explanatory variables, the adjusted scatter plot appears as follows. We will work with this adjusted data for the remainder of this question.

```
pairs(crime_df_upd[, c(response, "bad", "crime", "lawyers")])
```



b)

The step-up process was carried out. The variables added in order were *employ*, *crime* and *pop*, after which no further added variables had significant p-values. Hence the final model is as follows:

```
step_up_lm <- lm(expend~employ+crime+pop, data=crime_df_upd)
summary(step_up_lm)</pre>
```

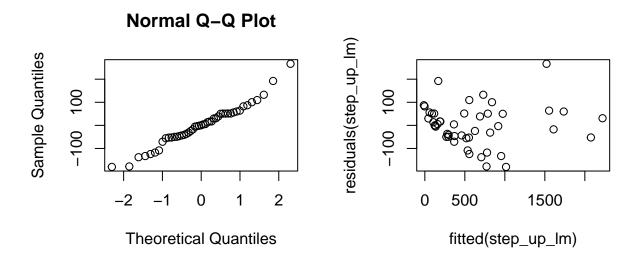
```
##
## Call:
## lm(formula = expend ~ employ + crime + pop, data = crime_df_upd)
```

```
##
## Residuals:
##
       Min
                1Q
                    Median
                                 3Q
                                        Max
## -179.99
           -49.64
                      0.48
                              51.19
                                     266.63
##
##
  Coefficients:
##
                Estimate Std. Error t value Pr(>|t|)
## (Intercept) -2.47e+02
                           5.47e+01
                                       -4.52
                                              4.8e-05 ***
                2.09e-02
                            3.95e-03
                                        5.30
## employ
                                              3.7e-06 ***
## crime
                5.43e-02
                            1.13e-02
                                        4.82
                                              1.8e-05 ***
                            1.79e-02
## pop
                7.14e-02
                                        4.00
                                              0.00025 ***
##
                   0 '*** 0.001 '** 0.01 '* 0.05 '. ' 0.1 ' ' 1
## Signif. codes:
##
## Residual standard error: 91.4 on 43 degrees of freedom
## Multiple R-squared: 0.974, Adjusted R-squared:
## F-statistic: 547 on 3 and 43 DF, p-value: <2e-16
```

Final model: expend = -247 + 0.0209*employ + 0.0543*crime + 0.0714*pop \pm error, with $R^2 = 0.974$.

We see that the step-up method naturally removes collinearity and produced a better model than was arrived upon using VIF in (a), which had an R-squared value of 0.957.

Finally, we check the model assumptions, which can be accepted based on the following plots:



Using the step-up model found in (b), the 95% prediction interval for *expend* is given by:

c)

new_data <- data.frame(bad=50, crime=5000, lawyers=5000, employ=5000, pop=5000)
predict(step_up_lm, new_data, interval="prediction", level=0.95)</pre>

```
## fit lwr upr
## 1 486 258 713
```

We can improve this interval (make it more narrow) by considering the **confidence interval**, which does not take into account the error.

```
predict(step_up_lm, new_data, interval="confidence", level=0.95)

## fit lwr upr
## 1 486 352 619
d)
```

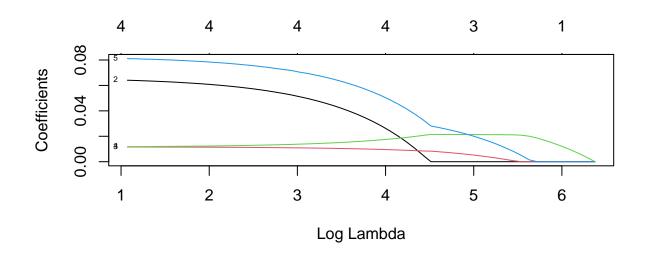
We can apply the lasso method as follows:

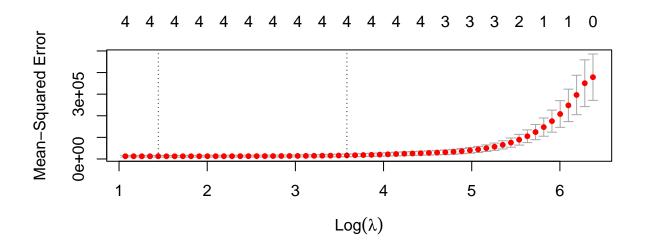
```
x <- as.matrix(crime_df_upd[, exp_vars])
y <- as.matrix(crime_df_upd[, c(response)])

# train-test splitting
train <- (sample(1:nrow(x), 0.67*nrow(x))) # train by using 2/3 of the data
x.train <- x[train,]; y.train <- y[train]
x.test <- x[-train,]; y.test <- y[-train]

# fitting the model
lasso.mod <- glmnet(x.train, y.train, alpha=1)
cv.lasso <- cv.glmnet(x.train,y.train,alpha=1,type.measure='mse')</pre>
```

plot(lasso.mod, label=T, xvar="lambda") # have a look at the lasso path





```
(lambda.min <- cv.lasso$lambda.min)</pre>
```

```
## [1] 4.24
```

```
(lambda.1se <- cv.lasso$lambda.1se)
```

[1] 36

 $\begin{tabular}{ll} \# \ https://glmnet.stanford.edu/articles/glmnet.html\#assessing-models-on-test-data-1 \\ assess.glmnet(lasso.mod, newx = x.test, newy = y.test, s=cv.lasso$lambda.1se) \\ \end{tabular}$

```
## $mse
## $1
## 4767
## attr(,"measure")
## [1] "Mean-Squared Error"
##
## $mae
## $1
## 46.8
## attr(,"measure")
## [1] "Mean Absolute Error"
```

Looking at lambda min

```
coef(lasso.mod, s=cv.lasso$lambda.min) # beta's for the best lambda
## 6 x 1 sparse Matrix of class "dgCMatrix"
##
## (Intercept) -296.8297
## bad
## crime
                  0.0631
## lawyers
                  0.0115
## employ
                  0.0120
## pop
                  0.0803
y.pred <- predict(lasso.mod, s=lambda.min, newx=x.test) # predict for test
mse.lasso <- mean((y.test - y.pred)^2); mse.lasso # mse for the predicted test rows
## [1] 6924
Looking at lambda 1se (the one we should use I think?)
coef(lasso.mod, s=cv.lasso$lambda.1se) # beta's for lambda.1se
## 6 x 1 sparse Matrix of class "dgCMatrix"
##
## (Intercept) -154.2127
## bad
## crime
                  0.0399
## lawyers
                  0.0103
## employ
                  0.0155
## pop
                  0.0615
y.pred <- predict(lasso.mod, s=lambda.1se, newx=x.test) # predict for test</pre>
mse.lasso <- mean((y.test - y.pred)^2); mse.lasso # mse for the predicted test rows</pre>
## [1] 4767
Compare to step-up model in (b).
new_data <- data.frame(x.test)</pre>
y.pred <- predict(step_up_lm, new_data, interval="confidence", level=0.95)</pre>
mse.step_up <- mean((y.test - y.pred)^2); mse.step_up # mse for the predicted test rows
## [1] 7599
```

Step-up model is better? Is this the right way to compare?

cv.lasso

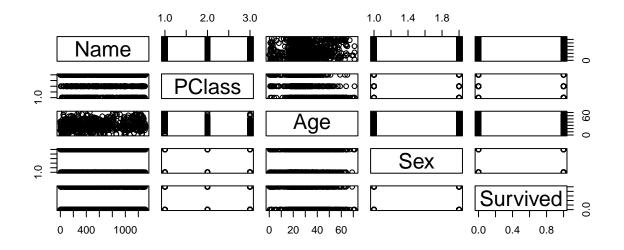
```
##
## Call: cv.glmnet(x = x.train, y = y.train, type.measure = "mse", alpha = 1)
## Measure: Mean-Squared Error
##
##
       Lambda Index Measure
                               SE Nonzero
## min
          4.2
                 54
                      12516 4093
## 1se
         36.0
                 31
                      16560 6146
                                        4
```

Exercise 3

head(titanic_df)

```
##
                                               Name PClass
                                                                     Sex Survived
                                                              Age
## 1
                      Allen, Miss Elisabeth Walton
                                                       1st 29.00 female
## 2
                       Allison, Miss Helen Loraine
                                                       1st 2.00 female
                                                                                0
               Allison, Mr Hudson Joshua Creighton
                                                       1st 30.00
                                                                    male
                                                                                0
## 4 Allison, Mrs Hudson JC (Bessie Waldo Daniels)
                                                       1st 25.00 female
                                                                                0
## 5
                     Allison, Master Hudson Trevor
                                                       1st 0.92
                                                                    male
                                                                                1
## 6
                                 Anderson, Mr Harry
                                                       1st 47.00
                                                                                1
                                                                    male
```

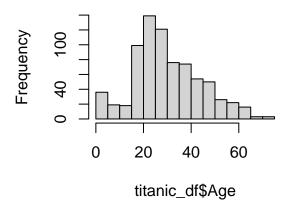
plot(titanic_df)

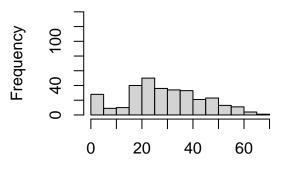


a)

```
titanic_df$PClass <- as.factor(titanic_df$PClass)</pre>
titanic_df$Sex <- as.factor(titanic_df$Sex)</pre>
summary(titanic_df)
##
        Name
                       PClass
                                                   Sex
                                                                Survived
                                      Age
## Length:1313
                       1st:322
                                 Min. : 0
                                               female:462
                                                                    :0.000
                                                            Min.
## Class :character
                       2nd:280
                                 1st Qu.:21
                                               male :851
                                                            1st Qu.:0.000
## Mode :character
                       3rd:711
                                 Median:28
                                                            Median :0.000
##
                                 Mean
                                        :30
                                                            Mean
                                                                  :0.343
                                 3rd Qu.:39
                                                             3rd Qu.:1.000
##
##
                                 Max.
                                        :71
                                                             Max. :1.000
##
                                 NA's
                                        :557
tot_comb <- xtabs(~PClass+Sex, data=titanic_df)</pre>
tot_comb
         Sex
##
## PClass female male
             143 179
##
      1st
##
      2nd
             107 173
##
      3rd
             212 499
tot_comb.surv <- xtabs(Survived~PClass+Sex, data=titanic_df)</pre>
round(tot_comb.surv/tot_comb, 2)
##
         Sex
## PClass female male
##
      1st 0.94 0.33
##
      2nd
            0.88 0.14
##
           0.38 0.12
      3rd
par(mfrow=c(1, 2))
hist(titanic_df$Age)
hist(titanic_df$Age[titanic_df$Survived == 1], ylim =c(0, 140))
```

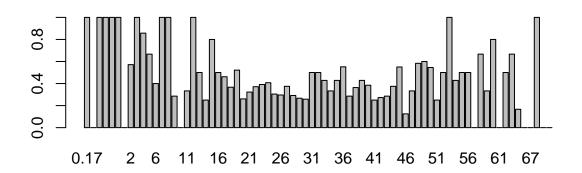
Histogram of titanic_df\$Age 'am of titanic_df\$Age[titanic_df\$Sul



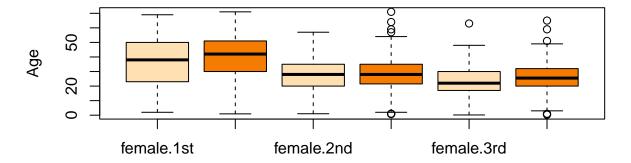


titanic_df\$Age[titanic_df\$Survived == 1

```
tot_age <- xtabs(~Age, data=titanic_df)
barplot(xtabs(Survived~Age, data=titanic_df)/tot_age)</pre>
```



boxplot(Age ~ Sex + PClass, data=titanic_df, col = c("#FFE0B2", "#F57C00"))



Sex: PClass

```
titanic_df_upd <- na.omit(titanic_df)</pre>
titanic_df_upd$PClass <- as.factor(titanic_df_upd$PClass)</pre>
titanic_df_upd$Sex <- as.factor(titanic_df_upd$Sex)</pre>
# head(titanic_df_upd)
base_lm <- glm(Survived ~ Age+PClass+Sex, data = titanic_df_upd, family = binomial)</pre>
summary(base_lm)
##
## Call:
## glm(formula = Survived ~ Age + PClass + Sex, family = binomial,
##
       data = titanic_df_upd)
##
## Deviance Residuals:
##
      Min
               1Q Median
                                3Q
                                       Max
## -2.723 -0.707 -0.392
                                     2.529
                            0.649
##
## Coefficients:
               Estimate Std. Error z value Pr(>|z|)
                                      9.46 < 2e-16 ***
                           0.39757
## (Intercept) 3.75966
## Age
               -0.03918
                           0.00762
                                      -5.14 2.7e-07 ***
## PClass2nd
               -1.29196
                           0.26008
                                      -4.97 6.8e-07 ***
## PClass3rd
               -2.52142
                           0.27666
                                      -9.11 < 2e-16 ***
## Sexmale
               -2.63136
                           0.20151 -13.06 < 2e-16 ***
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
```

```
Null deviance: 1025.57 on 755 degrees of freedom
## Residual deviance: 695.14 on 751
                                       degrees of freedom
## AIC: 705.1
##
## Number of Fisher Scoring iterations: 5
exp(coef(base_lm))
## (Intercept)
                             PClass2nd
                                         PClass3rd
                                                        Sexmale
                       Age
##
       42.9339
                    0.9616
                                0.2747
                                             0.0803
                                                         0.0720
TODO: add discussion of odds from the paper
b)
anova(glm(Survived ~ Age*PClass, data = titanic_df_upd, family = binomial), test="Chisq")
## Analysis of Deviance Table
##
## Model: binomial, link: logit
## Response: Survived
##
## Terms added sequentially (first to last)
##
##
              Df Deviance Resid. Df Resid. Dev Pr(>Chi)
##
## NULL
                                755
                                           1026
               1
                                754
                      2.8
                                           1023
                                                   0.091 .
## Age
## PClass
               2
                    112.8
                                752
                                            910
                                                  <2e-16 ***
                                750
## Age:PClass 2
                      1.2
                                            909
                                                   0.558
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
anova(glm(Survived ~ Age*Sex, data = titanic_df_upd, family = binomial), test="Chisq")
## Analysis of Deviance Table
##
## Model: binomial, link: logit
##
## Response: Survived
##
## Terms added sequentially (first to last)
##
##
##
           Df Deviance Resid. Df Resid. Dev Pr(>Chi)
```

```
## NULL
                               755
                                         1026
                              754
                                         1023
## Age
            1
                    2.8
                                                  0.091 .
## Sex
            1
                  227.1
                              753
                                          796
                                                < 2e-16 ***
            1
                   25.0
                              752
                                          771
                                               5.6e-07 ***
## Age:Sex
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
Therefore we decided to keep following model as Age: Sex intersection is significant.
final_lm <- glm(Survived ~ PClass+Age*Sex, data = titanic_df_upd, family = binomial)</pre>
anova(final_lm, test="Chisq")
## Analysis of Deviance Table
##
## Model: binomial, link: logit
## Response: Survived
##
## Terms added sequentially (first to last)
##
##
           Df Deviance Resid. Df Resid. Dev Pr(>Chi)
##
## NULL
                              755
                                         1026
## PClass
            2
                   78.0
                              753
                                          948
                                               < 2e-16 ***
            1
                   37.6
                              752
                                          910
                                               8.6e-10 ***
## Age
## Sex
            1
                  214.8
                              751
                                          695
                                                < 2e-16 ***
                              750
## Age:Sex
                   28.1
                                          667
                                               1.2e-07 ***
            1
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
newdata <- data.frame(Age=c(55, 55, 55, 55, 55, 55), PClass=c("1st", "1st", "2nd", "2nd", "3rd
predict(final_lm, newdata, type="response")
##
                                      5
                                              6
        1
                       3
## 0.9474 0.1450 0.7937 0.0350 0.5590 0.0118
For "female" all the probs > 0.5 and for the "male" probs are < 0.5.
c) Use confusion matrix, log likelihood as quality measures
d)
table(titanic_df_upd$PClass, titanic_df_upd$Sex)
##
##
         female male
##
     1st
            101
                 125
##
     2nd
             85
                 127
```

102 216

##

3rd

```
chisq.test(x=titanic_df_upd$Survived, y=titanic_df_upd$Sex)
##
   Pearson's Chi-squared test with Yates' continuity correction
## data: titanic_df_upd$Survived and titanic_df_upd$Sex
## X-squared = 219, df = 1, p-value <2e-16
For 2x2 tables we can obtain exact p-value using the Fisher test.
fisher.test(x=titanic_df_upd$Survived, y=titanic_df_upd$Sex)
##
   Fisher's Exact Test for Count Data
##
##
## data: titanic_df_upd$Survived and titanic_df_upd$Sex
## p-value <2e-16
## alternative hypothesis: true odds ratio is not equal to 1
## 95 percent confidence interval:
## 0.0586 0.1215
## sample estimates:
## odds ratio
##
       0.0848
chisq.test(x=titanic_df_upd$Survived, y=titanic_df_upd$PClass)
##
##
   Pearson's Chi-squared test
##
## data: titanic_df_upd$Survived and titanic_df_upd$PClass
## X-squared = 76, df = 2, p-value <2e-16
We reject null hypothesis, meaning that rows and cols are actually dependent: Sex and PClass
have influence on Survived variable.
e)
TODO: comparison btw c) and d)?
```

Exercise 4

characteristics

contingency table tells us only about the presence of effect and doesn't provide some quantitative

```
head(coups_df)
##
             miltcoup oligarchy pollib parties pctvote
                                                          popn size numelec
## Benin
                              7
                                      1
                                             34
                                                   45.7 4.600 113
## Burkina
                    6
                             13
                                      2
                                             62
                                                   17.5 8.800 274
                                                                           5
## Burundi
                    2
                             13
                                      2
                                             10
                                                   34.4 5.300
                                                                           3
                                                                 28
## Cameroon
                    0
                              0
                                      2
                                             34
                                                   30.3 11.600 475
                                                                          14
                                                                           2
## Capeverde
                    1
                              0
                                      2
                                              5
                                                   30.5 0.361
                                                                  4
                                      2
## CAR
                             14
                                             14
                                                   16.2 3.000 623
                                                                           6
##
             numregim
## Benin
## Burkina
                    3
## Burundi
                    3
                    3
## Cameroon
## Capeverde
                    1
## CAR
# coups_df$pollib <- as.factor(coups_df$pollib)</pre>
# coups_df$numregim <- as.factor(coups_df$numregim)</pre>
a)
poison_glm <- glm(miltcoup ~ oligarchy + pollib + parties + pctvote + popn + size + numelec + :
summary(poison_glm)
##
## Call:
## glm(formula = miltcoup ~ oligarchy + pollib + parties + pctvote +
       popn + size + numelec + numregim, family = poisson, data = coups_df)
##
## Deviance Residuals:
##
     \mathtt{Min}
               1Q Median
                               ЗQ
                                      Max
## -1.344 -0.954 -0.259
                                     1.695
                            0.391
##
## Coefficients:
                Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) -0.510269
                           0.905330
                                       -0.56
                                               0.5730
## oligarchy
                0.073081
                           0.034596
                                       2.11
                                               0.0346 *
## pollib
               -0.712978
                           0.272563
                                      -2.62 0.0089 **
## parties
               0.030774
                           0.011187
                                       2.75
                                               0.0059 **
## pctvote
                           0.009753
                                        1.42
                                               0.1549
                0.013872
```

1.42

-0.76

-0.25

0.84

0.1566

0.4445

0.8060

0.4030

popn

size

numelec

numregim

0.009343

-0.000190

-0.016078

0.191735

0.006595

0.000248

0.065484

0.229289

```
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for poisson family taken to be 1)
##
## Null deviance: 65.945 on 35 degrees of freedom
## Residual deviance: 28.668 on 27 degrees of freedom
## AIC: 111.5
##
## Number of Fisher Scoring iterations: 6
```

Through summary(drop1)we can find the variables that are significant in predicting number of successful military coups: oligarchy, pollib, parties.

b)

```
summary(glm(miltcoup ~ oligarchy + pollib + parties + pctvote + popn + size + numelec, data = -
##
## Call:
## glm(formula = miltcoup ~ oligarchy + pollib + parties + pctvote +
       popn + size + numelec, family = poisson, data = coups_df)
##
## Deviance Residuals:
      Min
               1Q Median
                               3Q
                                      Max
## -1.353 -0.965 -0.195
                            0.483
                                    1.617
## Coefficients:
##
                Estimate Std. Error z value Pr(>|z|)
## (Intercept) -0.114181
                           0.756715
                                      -0.15
                                              0.8801
## oligarchy
                                              0.0052 **
                0.086007
                           0.030774
                                       2.79
## pollib
               -0.689010
                           0.270392
                                      -2.55
                                              0.0108 *
## parties
                0.029183
                           0.011006
                                       2.65
                                              0.0080 **
## pctvote
                0.014150
                           0.009753
                                       1.45
                                              0.1468
## popn
                0.006272
                           0.005440
                                       1.15
                                              0.2490
               -0.000195
## size
                           0.000247
                                      -0.79
                                              0.4297
## numelec
                0.000168
                                       0.00
                                              0.9978
                           0.062185
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for poisson family taken to be 1)
##
       Null deviance: 65.945
                              on 35
                                     degrees of freedom
## Residual deviance: 29.363 on 28 degrees of freedom
## AIC: 110.2
##
## Number of Fisher Scoring iterations: 5
```

```
summary(glm(miltcoup ~ oligarchy + pollib + parties + pctvote + popn + size, data = coups_df, :
##
## Call:
## glm(formula = miltcoup ~ oligarchy + pollib + parties + pctvote +
      popn + size, family = poisson, data = coups_df)
##
## Deviance Residuals:
     Min
              1Q Median
##
                             3Q
                                    Max
## -1.352 -0.965 -0.195 0.483
                                  1.618
##
## Coefficients:
               Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) -0.112687  0.516303  -0.22  0.82723
## oligarchy
             0.085962 0.025910 3.32 0.00091 ***
## pollib
             0.029194 0.010195 2.86 0.00419 **
## parties
## pctvote
              0.014159 0.009198 1.54 0.12372
## popn
              0.006274 0.005399
                                    1.16 0.24527
## size
              -0.000195 0.000242 -0.80 0.42138
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for poisson family taken to be 1)
##
##
      Null deviance: 65.945 on 35 degrees of freedom
## Residual deviance: 29.363 on 29 degrees of freedom
## AIC: 108.2
##
## Number of Fisher Scoring iterations: 5
summary(glm(miltcoup ~ oligarchy + pollib + parties + pctvote + popn, data = coups_df), test="0"
##
## glm(formula = miltcoup ~ oligarchy + pollib + parties + pctvote +
      popn, data = coups_df)
##
##
## Deviance Residuals:
      Min
                1Q
                    Median
                                 3Q
                                         Max
## -2.0424 -0.8843 -0.0798
                                      2.0044
                             1.1548
##
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                1.0174
                          0.7904
                                    1.29
                                          0.2079
## oligarchy
                          0.0393
                                    3.25
                                          0.0028 **
                0.1277
```

```
## pollib
                           0.3811
                                  -2.65
                                           0.0127 *
               -1.0098
## parties
                0.0504
                          0.0193
                                     2.62 0.0138 *
## pctvote
                0.0152
                           0.0126
                                     1.20 0.2393
## popn
                0.0178
                                     1.55
                           0.0115
                                           0.1325
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for gaussian family taken to be 1.59)
##
##
      Null deviance: 108.750 on 35 degrees of freedom
## Residual deviance: 47.787 on 30 degrees of freedom
## AIC: 126.4
##
## Number of Fisher Scoring iterations: 2
summary(glm(miltcoup ~ oligarchy + pollib + parties + popn, data = coups_df), test="Chisq")
##
## Call:
## glm(formula = miltcoup ~ oligarchy + pollib + parties + popn,
##
      data = coups_df)
##
## Deviance Residuals:
     Min
              10 Median
                              30
                                     Max
## -1.928 -0.800 -0.220
                           0.987
                                   2.208
##
## Coefficients:
##
              Estimate Std. Error t value Pr(>|t|)
                       0.7209 1.97 0.0578.
## (Intercept)
                1.4201
## oligarchy
                0.1343
                           0.0392
                                     3.43 0.0017 **
## pollib
                                  -2.43
               -0.9083
                          0.3743
                                            0.0212 *
                                    2.40
## parties
                0.0454
                         0.0190
                                            0.0228 *
## popn
                0.0150
                          0.0113
                                  1.33 0.1947
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 1.62)
##
      Null deviance: 108.750 on 35 degrees of freedom
## Residual deviance: 50.083 on 31 degrees of freedom
## AIC: 126
##
## Number of Fisher Scoring iterations: 2
final_plm <- glm(miltcoup ~ oligarchy + parties + pollib, data = coups_df, family = poisson)</pre>
summary(final_plm, test="Chisq")
```

```
## glm(formula = miltcoup ~ oligarchy + parties + pollib, family = poisson,
       data = coups_df)
##
## Deviance Residuals:
      Min
                1Q Median
                                 3Q
                                        Max
## -1.358 -1.042 -0.286
                              0.628
                                      1.752
##
## Coefficients:
##
                Estimate Std. Error z value Pr(>|z|)
                                        0.67
## (Intercept)
                0.25138
                            0.37269
                                                 0.500
## oligarchy
                 0.09262
                             0.02178
                                        4.25
                                               2.1e-05 ***
## parties
                 0.02206
                             0.00896
                                        2.46
                                                 0.014 *
## pollib
                -0.57410
                            0.20438
                                       -2.81
                                                 0.005 **
## ---
## Signif. codes:
                    0 '*** 0.001 '** 0.01 '* 0.05 '. ' 0.1 ' ' 1
##
## (Dispersion parameter for poisson family taken to be 1)
##
##
       Null deviance: 65.945
                               on 35
                                       degrees of freedom
## Residual deviance: 32.856 on 32 degrees of freedom
## AIC: 105.7
##
## Number of Fisher Scoring iterations: 5
After step-down approach there are oligarchy, parties and pollib left. (For treating vars as factors
only oligarchy left) In comparison with a) all the same factors are significant.
\mathbf{c}
coups_df$pollib <- as.factor(coups_df$pollib)</pre>
coups_df$numregim <- as.factor(coups_df$numregim)</pre>
mean(coups_df$oligarchy); mean(coups_df$parties)
## [1] 5.22
## [1] 17.1
newdata <- data.frame(pollib=c(0, 1, 2), oligarchy=c(5.22, 5.22, 5.22), parties=c(17.1, 17.1,
predict(final_plm, newdata, type="response")
##
       1
## 3.041 1.713 0.965
```

##

Call:

Our model is predicting there will be roughly 3 successful coups for pollib=0, roughly 2 successful coups for pollib=1 and 1 successful coup for pollib=2.