

ECON 187: Project 1

Austin Pham (905318112), Shannan Liu (305172952), Christina Zhang(605325840), Zachary Wrubel (20

For this project you will need to use two different datasets of your choice. One will be used for classification and the other for regularization. For the classification dataset make sure you have more than two classes. For your regularization dataset, since the methods focus on variable selection, please make sure to have as many predictors as possible (e.g., 10s or 100s).

Classification

```
body <- read_csv("bodyPerformance.csv")
body <- body %>% mutate(gender = factor(ifelse(gender == "M", 1, 0)))
sum(is.na(body))
```

```
## [1] 0
```

```
head(body)
```

```
## # A tibble: 6 x 12
##   age gender height_cm weight_kg 'body fat_%' diastolic systolic gripForce
##   <dbl> <fct>      <dbl>      <dbl>      <dbl>      <dbl>      <dbl>      <dbl>
## 1    27 1          172.        75.2        21.3         80        130        54.9
## 2    25 1          165        55.8        15.7         77        126        36.4
## 3    31 1          180.         78         20.1         92        152        44.8
## 4    32 1          174.        71.1        18.4         76        147        41.4
## 5    28 1          174.        67.7        17.1         70        127        43.5
## 6    36 0          165.        55.4         22         64        119        23.8
## # ... with 4 more variables: 'sit and bend forward_cm' <dbl>,
## #   'sit-ups counts' <dbl>, 'broad jump_cm' <dbl>, class <chr>
```

Below are the graphs that shows the possible correlation between all the predictors with Class A,B,C,and D. From the grpahs, we can see that all the correlations are significant, so we will be using all the predictors for classification analysis.

```
plot1 <- body %>% ggpairs(columns = c(1:4,12), ggplot2::aes(col = class, fill = class))
plot2 <- body %>% ggpairs(columns = c(5:8,12), ggplot2::aes(col = class, fill = class))
plot3 <- body %>% ggpairs(columns = c(9:12), ggplot2::aes(col = class, fill = class))
plot1
plot2
plot2
```

Logistic Regression

```
set.seed(42)
fit.control.cv <- trainControl(method = "repeatedcv", number = 10, repeats = 3)
fit.control.boot <- trainControl(method = "boot")
mn.fit.cv <- train(class ~ ., data = body, method = "multinom",
                  trControl = fit.control.cv, trace = FALSE,
                  preProcess = c("center", "scale"))
mn.fit.boot <- train(class ~ ., data = body, method = "multinom",
                    trControl = fit.control.boot, trace = FALSE,
                    preProcess = c("center", "scale"))
```

```
mn.fit.cv$finalModel
```

```
## Call:
## nnet::multinom(formula = .outcome ~ ., data = dat, decay = param$decay,
##   trace = FALSE)
##
## Coefficients:
##   (Intercept)      age  gender1  height_cm weight_kg '\body fat_%'\
## B   1.6436941 -1.031072  1.228708 -0.07014251 0.8745149      0.029745697
## C   1.8396621 -1.891377  1.977937  0.11469195 1.2557692      0.005482127
## D   0.4281742 -2.824718  2.598717 -0.30909023 2.4963687      0.459483497
##   diastolic    systolic gripForce '\sit and bend forward_cm'\
## B 0.05764353 -0.01869305 -0.920936      -1.331405
## C 0.13202220 -0.07684313 -1.519526      -2.285391
## D 0.25219705 -0.14232365 -2.101412      -3.597352
##   '\sit-ups counts'\ '\broad jump_cm'\
## B           -1.423940      -0.6803174
## C           -2.603735      -1.1445756
## D           -4.161081      -1.2372507
##
## Residual Deviance: 23128.6
## AIC: 23200.6
```

```
mn.fit.boot$finalModel
```

```
## Call:
## nnet::multinom(formula = .outcome ~ ., data = dat, decay = param$decay,
##   trace = FALSE)
##
## Coefficients:
##   (Intercept)      age  gender1  height_cm weight_kg '\body fat_%'\
## B   1.6436941 -1.031072  1.228708 -0.07014251 0.8745149      0.029745697
## C   1.8396621 -1.891377  1.977937  0.11469195 1.2557692      0.005482127
## D   0.4281742 -2.824718  2.598717 -0.30909023 2.4963687      0.459483497
##   diastolic    systolic gripForce '\sit and bend forward_cm'\
## B 0.05764353 -0.01869305 -0.920936      -1.331405
## C 0.13202220 -0.07684313 -1.519526      -2.285391
## D 0.25219705 -0.14232365 -2.101412      -3.597352
##   '\sit-ups counts'\ '\broad jump_cm'\
## B           -1.423940      -0.6803174
```

```
## C          -2.603735          -1.1445756
## D          -4.161081          -1.2372507
##
## Residual Deviance: 23128.6
## AIC: 23200.6
```

```
confusionMatrix(mn.fit.cv)
```

```
## Cross-Validated (10 fold, repeated 3 times) Confusion Matrix
##
## (entries are percentual average cell counts across resamples)
##
##           Reference
## Prediction  A    B    C    D
##           A 18.4  5.9  2.0  0.3
##           B  6.1 11.1  5.4  1.3
##           C  0.5  7.1 12.9  4.1
##           D  0.0  0.8  4.7 19.2
##
## Accuracy (average) : 0.6173
```

```
confusionMatrix(mn.fit.boot)
```

```
## Bootstrapped (25 reps) Confusion Matrix
##
## (entries are percentual average cell counts across resamples)
##
##           Reference
## Prediction  A    B    C    D
##           A 18.4  6.0  2.0  0.3
##           B  6.1 11.1  5.4  1.3
##           C  0.5  7.0 13.0  4.1
##           D  0.0  0.8  4.6 19.2
##
## Accuracy (average) : 0.6177
```

```
mn.fit.cv$results
```

```
##   decay Accuracy      Kappa AccuracySD      KappaSD
## 1 0e+00 0.6172122 0.4896154 0.009252803 0.01233862
## 2 1e-04 0.6172122 0.4896154 0.009252803 0.01233862
## 3 1e-01 0.6173367 0.4897813 0.009319778 0.01242802
```

```
mn.fit.boot$results
```

```
##   decay Accuracy      Kappa AccuracySD      KappaSD
## 1 0e+00 0.6177344 0.4903212 0.005948138 0.007938624
## 2 1e-04 0.6177344 0.4903212 0.005948138 0.007938624
## 3 1e-01 0.6177418 0.4903313 0.005897619 0.007870582
```

The logistic regression method shows an overall accuracy of 0.6177.

LDA

```
lda.fit.cv <- train(class ~ ., data = body, method = "lda",
                    trControl = fit.control.cv, trace = FALSE,
                    preProcess = c("center", "scale"))
lda.fit.boot <- train(class ~ ., data = body, method = "lda",
                      trControl = fit.control.boot, trace = FALSE,
                      preProcess = c("center", "scale"))
```

```
lda.fit.cv$finalModel
```

```
## Call:
## lda(x, grouping = y, trace = FALSE)
##
## Prior probabilities of groups:
##      A      B      C      D
## 0.2499813 0.2499067 0.2500560 0.2500560
##
## Group means:
##      age      gender1      height_cm      weight_kg      'body fat_%'      diastolic
## A -0.110591989 -0.15645294 -0.081738935 -0.25349141  -0.3722011 -0.08317758
## B  0.021943176  0.02852539  0.002433177 -0.06986770  -0.1653865 -0.01306765
## C -0.005540626  0.07416582  0.071483362 -0.05749133  -0.0820744 -0.02291650
## D  0.094169521  0.05373205  0.007799443  0.38073302   0.6194521  0.11912909
##      systolic      gripForce      'sit and bend forward_cm'      'sit-ups counts'
## A -0.06421067  0.15546979                0.73108017          0.56575012
## B  0.02749309  0.08908903                0.26704551          0.20088507
## C -0.02103213 -0.03587942                -0.09680201          -0.07361708
## D  0.05774696 -0.20857978                -0.90094590          -0.69272922
##      'broad jump_cm'
## A      0.31640410
## B      0.13050414
## C     -0.03766627
## D     -0.40906956
##
## Coefficients of linear discriminants:
##                                LD1          LD2          LD3
## age                -0.64678307  0.22309394  0.47011181
## gender1              0.69584600 -1.28409193  1.15159573
## height_cm           -0.05207736 -0.93671269 -0.35916540
## weight_kg            0.53802507  1.12001910  0.15508149
## 'body fat_%'         0.16993470  0.16486574  0.50244600
## diastolic            0.06595464  0.07418465 -0.27129251
## systolic            -0.04539913 -0.07760683  0.30887408
## gripForce           -0.48748656  0.34068971 -0.19140457
## 'sit and bend forward_cm' -0.74403894 -0.28310233  0.43563764
## 'sit-ups counts'     -1.04324820  0.33643160  0.13732175
## 'broad jump_cm'      -0.26279633  0.68788193  0.04374994
##
## Proportion of trace:
##      LD1      LD2      LD3
## 0.9785 0.0195 0.0019
```

```
lda.fit.boot$finalModel
```

```
## Call:
## lda(x, grouping = y, trace = FALSE)
##
## Prior probabilities of groups:
##      A      B      C      D
## 0.2499813 0.2499067 0.2500560 0.2500560
##
## Group means:
##      age      gender1      height_cm      weight_kg      'body fat_%'      diastolic
## A -0.110591989 -0.15645294 -0.081738935 -0.25349141 -0.3722011 -0.08317758
## B  0.021943176  0.02852539  0.002433177 -0.06986770 -0.1653865 -0.01306765
## C -0.005540626  0.07416582  0.071483362 -0.05749133 -0.0820744 -0.02291650
## D  0.094169521  0.05373205  0.007799443  0.38073302  0.6194521  0.11912909
##      systolic      gripForce      'sit and bend forward_cm'      'sit-ups counts'
## A -0.06421067  0.15546979      0.73108017      0.56575012
## B  0.02749309  0.08908903      0.26704551      0.20088507
## C -0.02103213 -0.03587942      -0.09680201      -0.07361708
## D  0.05774696 -0.20857978      -0.90094590      -0.69272922
##      'broad jump_cm'
## A      0.31640410
## B      0.13050414
## C     -0.03766627
## D     -0.40906956
##
## Coefficients of linear discriminants:
##      LD1      LD2      LD3
## age      -0.64678307  0.22309394  0.47011181
## gender1      0.69584600 -1.28409193  1.15159573
## height_cm     -0.05207736 -0.93671269 -0.35916540
## weight_kg      0.53802507  1.12001910  0.15508149
## 'body fat_%'      0.16993470  0.16486574  0.50244600
## diastolic      0.06595464  0.07418465 -0.27129251
## systolic     -0.04539913 -0.07760683  0.30887408
## gripForce     -0.48748656  0.34068971 -0.19140457
## 'sit and bend forward_cm' -0.74403894 -0.28310233  0.43563764
## 'sit-ups counts'     -1.04324820  0.33643160  0.13732175
## 'broad jump_cm'     -0.26279633  0.68788193  0.04374994
##
## Proportion of trace:
##      LD1      LD2      LD3
## 0.9785 0.0195 0.0019
```

```
confusionMatrix(lda.fit.cv)
```

```
## Cross-Validated (10 fold, repeated 3 times) Confusion Matrix
##
## (entries are percentual average cell counts across resamples)
##
##      Reference
## Prediction  A   B   C   D
##      A 18.2  6.1  2.1  0.3
```

```
##           B  6.2 11.0  5.4  1.4
##           C  0.6  7.4 14.1  5.0
##           D  0.0  0.5  3.4 18.3
##
## Accuracy (average) : 0.6156
```

```
confusionMatrix(lda.fit.boot)
```

```
## Bootstrapped (25 reps) Confusion Matrix
##
## (entries are percentual average cell counts across resamples)
##
##           Reference
## Prediction   A    B    C    D
##           A 18.2  6.2  2.1  0.3
##           B  6.4 10.9  5.5  1.6
##           C  0.6  7.3 13.9  4.9
##           D  0.0  0.5  3.4 18.3
##
## Accuracy (average) : 0.6126
```

```
lda.fit.cv$results
```

```
## parameter Accuracy      Kappa AccuracySD      KappaSD
## 1      none 0.615571 0.4874285 0.01318156 0.0175775
```

```
lda.fit.boot$results
```

```
## parameter Accuracy      Kappa AccuracySD      KappaSD
## 1      none 0.6125766 0.4834807 0.004671325 0.006265792
```

LDA shows an overall accuracy of 0.6126.

QDA

```
qda.fit.cv <- train(class ~ ., data = body, method = "qda",
                    trControl = fit.control.cv, trace = FALSE,
                    preProcess = c("center", "scale"))
qda.fit.boot <- train(class ~ ., data = body, method = "qda",
                     trControl = fit.control.boot, trace = FALSE,
                     preProcess = c("center", "scale"))
```

```
qda.fit.cv$finalModel
```

```
## Call:
## qda(x, grouping = y, trace = FALSE)
##
## Prior probabilities of groups:
##           A           B           C           D
```

```
## 0.2499813 0.2499067 0.2500560 0.2500560
##
## Group means:
##      age      gender1      height_cm      weight_kg      'body fat_%'      diastolic
## A -0.110591989 -0.15645294 -0.081738935 -0.25349141 -0.3722011 -0.08317758
## B  0.021943176  0.02852539  0.002433177 -0.06986770 -0.1653865 -0.01306765
## C -0.005540626  0.07416582  0.071483362 -0.05749133 -0.0820744 -0.02291650
## D  0.094169521  0.05373205  0.007799443  0.38073302  0.6194521  0.11912909
##      systolic      gripForce      'sit and bend forward_cm'      'sit-ups counts'
## A -0.06421067  0.15546979      0.73108017      0.56575012
## B  0.02749309  0.08908903      0.26704551      0.20088507
## C -0.02103213 -0.03587942      -0.09680201      -0.07361708
## D  0.05774696 -0.20857978      -0.90094590      -0.69272922
##      'broad jump_cm'
## A      0.31640410
## B      0.13050414
## C     -0.03766627
## D     -0.40906956
```

```
qda.fit.boot$finalModel
```

```
## Call:
## qda(x, grouping = y, trace = FALSE)
##
## Prior probabilities of groups:
##      A      B      C      D
## 0.2499813 0.2499067 0.2500560 0.2500560
##
## Group means:
##      age      gender1      height_cm      weight_kg      'body fat_%'      diastolic
## A -0.110591989 -0.15645294 -0.081738935 -0.25349141 -0.3722011 -0.08317758
## B  0.021943176  0.02852539  0.002433177 -0.06986770 -0.1653865 -0.01306765
## C -0.005540626  0.07416582  0.071483362 -0.05749133 -0.0820744 -0.02291650
## D  0.094169521  0.05373205  0.007799443  0.38073302  0.6194521  0.11912909
##      systolic      gripForce      'sit and bend forward_cm'      'sit-ups counts'
## A -0.06421067  0.15546979      0.73108017      0.56575012
## B  0.02749309  0.08908903      0.26704551      0.20088507
## C -0.02103213 -0.03587942      -0.09680201      -0.07361708
## D  0.05774696 -0.20857978      -0.90094590      -0.69272922
##      'broad jump_cm'
## A      0.31640410
## B      0.13050414
## C     -0.03766627
## D     -0.40906956
```

```
confusionMatrix(qda.fit.cv)
```

```
## Cross-Validated (10 fold, repeated 3 times) Confusion Matrix
##
## (entries are percentual average cell counts across resamples)
##
##      Reference
## Prediction  A      B      C      D
```

```
##          A 19.0  6.0  1.9  0.3
##          B  5.5 12.2  5.7  1.6
##          C  0.4  6.2 15.8  4.4
##          D  0.1  0.6  1.6 18.6
##
## Accuracy (average) : 0.6572
```

```
confusionMatrix(qda.fit.boot)
```

```
## Bootstrapped (25 reps) Confusion Matrix
##
## (entries are percentual average cell counts across resamples)
##
##          Reference
## Prediction   A    B    C    D
##          A 18.8  6.1  1.9  0.4
##          B  5.7 11.9  5.5  1.7
##          C  0.5  6.3 15.7  4.4
##          D  0.1  0.6  1.7 18.6
##
## Accuracy (average) : 0.6506
```

```
qda.fit.cv$results
```

```
##   parameter Accuracy      Kappa AccuracySD      KappaSD
## 1      none 0.6571846 0.5429133 0.01127552 0.01503324
```

```
qda.fit.boot$results
```

```
##   parameter Accuracy      Kappa AccuracySD      KappaSD
## 1      none 0.6505498 0.534103 0.01300762 0.01732938
```

QDA shows an overall accuracy of 0.6549.

kNN

```
knn.fit.cv <- train(class ~ ., data = body, method = "knn",
                    trControl = fit.control.cv, preProcess = c("center", "scale"))
knn.fit.boot <- train(class ~ ., data = body, method = "knn",
                      trControl = fit.control.boot, preProcess = c("center", "scale"))
```

```
knn.fit.cv$finalModel
```

```
## 9-nearest neighbor model
## Training set outcome distribution:
##
##      A      B      C      D
## 3348 3347 3349 3349
```



```
knn.fit.boot$finalModel
```

```
## 9-nearest neighbor model
## Training set outcome distribution:
##
##      A      B      C      D
## 3348 3347 3349 3349
```

```
confusionMatrix(knn.fit.cv)
```

```
## Cross-Validated (10 fold, repeated 3 times) Confusion Matrix
##
## (entries are percentual average cell counts across resamples)
##
##           Reference
## Prediction      A      B      C      D
##           A 19.9  8.2  2.8  0.6
##           B  4.5 11.7  7.4  2.0
##           C  0.6  4.4 13.6  5.7
##           D  0.1  0.6  1.2 16.6
##
## Accuracy (average) : 0.6179
```

```
confusionMatrix(knn.fit.boot)
```

```
## Bootstrapped (25 reps) Confusion Matrix
##
## (entries are percentual average cell counts across resamples)
##
##           Reference
## Prediction      A      B      C      D
##           A 18.5  8.2  2.9  0.7
##           B  5.1 11.0  7.6  2.2
##           C  1.1  5.0 12.5  5.9
##           D  0.2  0.9  1.9 16.4
##
## Accuracy (average) : 0.584
```

```
knn.fit.cv$results
```

```
##      k Accuracy      Kappa AccuracySD      KappaSD
## 1 5 0.6045941 0.4727952 0.01216431 0.01622369
## 2 7 0.6142261 0.4856382 0.01132453 0.01510491
## 3 9 0.6178839 0.4905149 0.01147146 0.01529822
```

```
knn.fit.boot$results
```

```
##      k Accuracy      Kappa AccuracySD      KappaSD
## 1 5 0.5599129 0.4132660 0.006783944 0.009123717
## 2 7 0.5754512 0.4339888 0.006548934 0.008788444
## 3 9 0.5840070 0.4454149 0.006918906 0.009270053
```

KNN yields an accuracy of 0.5849.

k-Means

```
#k means

bp_ <- body[,c(-2,-12)]
bp_ <- na.omit(bp_)
bp_ <-scale(bp_)

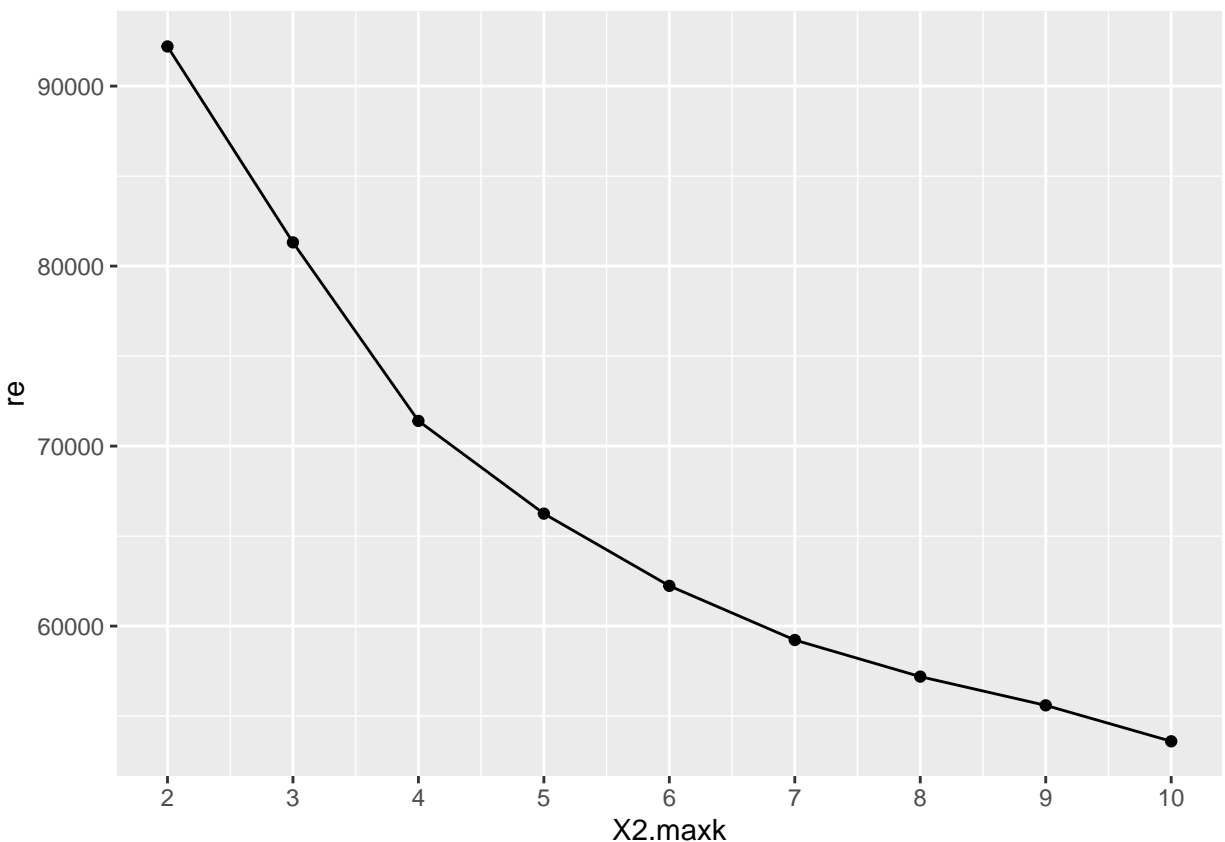
#create the function that runs the k-mean algorithm and get the clusters sum of squares

kmean_withinss <- function(k) {
  cluster <- kmeans(bp_, k)
  return (cluster$tot.withinss)
}

maxk <-10
re <- sapply(2:maxk, kmean_withinss)

re. <-data.frame(2:maxk, re) #create a data frame to store all values

ggplot(re., aes(x = X2.maxk, y = re)) +
  geom_point() +
  geom_line() +
  scale_x_continuous(breaks = seq(1, 10, by = 1))
```



From the graph, we can see the optimal k is 6, where the curve starts to diminish the return.

```
set.seed(240) # Setting seed
kmeans.re <- kmeans(bp_, centers=6, nstart = 20)
```

```
# Confusion Matrix
table(body$class, kmeans.re$cluster)
```

```
##
##      1      2      3      4      5      6
## A  404      2 1169   751   225   797
## B  580     36   794   816   361   760
## C  552    127   661   844   480   685
## D  394  1144   370   361   828   252
```

```
body_kmeans <- cbind(body, cluster = kmeans.re$cluster)
head(body_kmeans)
```

```
##   age gender height_cm weight_kg body fat_% diastolic systolic gripForce
## 1  27      1    172.3    75.24    21.3      80      130     54.9
## 2  25      1    165.0    55.80    15.7      77      126     36.4
## 3  31      1    179.6    78.00    20.1      92      152     44.8
## 4  32      1    174.5    71.10    18.4      76      147     41.4
## 5  28      1    173.8    67.70    17.1      70      127     43.5
## 6  36      0    165.4    55.40    22.0      64      119     23.8
##   sit and bend forward_cm sit-ups counts broad jump_cm class cluster
## 1              18.4           60      217      C      6
## 2              16.3           53      229      A      4
## 3              12.0           49      181      C      6
## 4              15.2           53      219      B      6
## 5              27.1           45      217      B      4
## 6              21.0           27      153      B      3
```

```
kmeans.re$centers #matrix of cluster centers
```

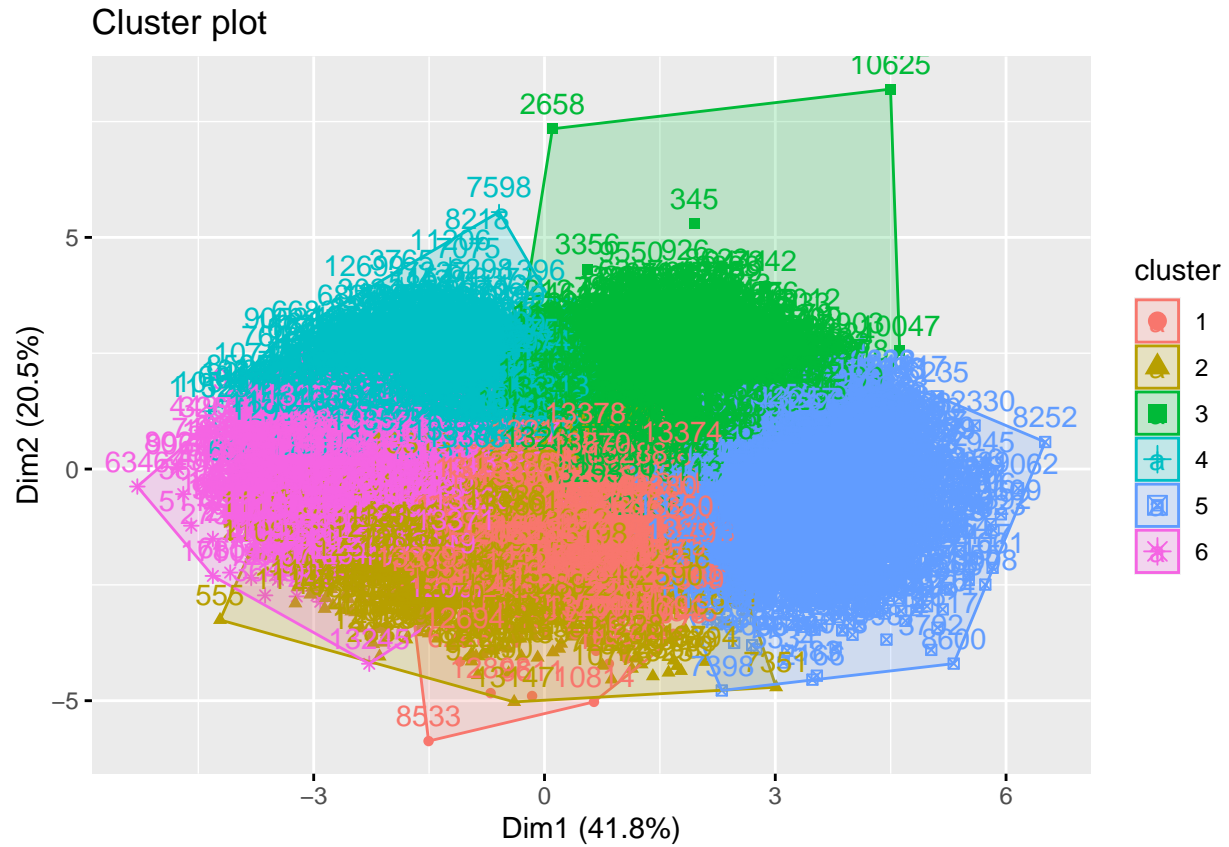
```
##      age      height_cm      weight_kg      body fat_%      diastolic      systolic
## 1  1.3093285  0.05239229  0.1066542 -0.1490933  0.54992415  0.64165957
## 2 -0.2044529  0.75095325  1.2803784  0.5784224  0.48411057  0.37676075
## 3 -0.5843503 -0.75152887 -0.9759586  0.3929658 -0.52772620 -0.76164170
## 4 -0.5628393  0.61639755  0.2828347 -0.8674661 -0.67966223 -0.50721974
## 5  1.1668925 -1.33496525 -0.7806493  1.2249225 -0.05727179 -0.01854236
## 6 -0.4650096  0.79620397  0.6955469 -0.6260350  0.75278844  0.79787684
##   gripForce sit and bend forward_cm sit-ups counts broad jump_cm
## 1  0.2046815      -0.22392263      -0.37074296      -0.11269288
## 2  0.4189320      -1.44220187      -0.28534222       0.07211187
## 3 -0.9398905       0.60664814      -0.09882091      -0.56221858
## 4  0.6526715      -0.05376574       0.77779309       0.84769071
## 5 -1.2337413       0.07878567      -1.50290672      -1.51799134
## 6  0.9615539       0.20189534       0.83214898       0.93490866
```

```
kmeans.re$size #number of points in each clusters
```

```
## [1] 1930 1309 2994 2772 1894 2494
```

The cluster centers shows that cluster 5 have the highest body fat % and age average among all clusters, and cluster 3 has the highest average weight among all.

```
fviz_cluster(kmeans.re, data = bp_)
```



The plot displays that each group have similar dimensions but there are still some overlaps.

```
fviz_cluster(kmeans(bp_, centers=4, nstart = 20), data = bp_)
```

Based on my fits, a non-linear is more appropriate since the accuracy generated from QDA has the highest accuracy, though the accuracy are pretty close to each other in all classification methods.

For this portion of the project, we're going to use a data set containing information on the Australian housing market and apply regularization techniques to make predictions on house prices. The data set will have 81 columns of data.

Categorical pipeline: - In each vector of data, we will fill missing values with the most commonly observed sample - After that, we will transform each vector to numeric representation using label encoding or by making them dummy variables, where appropriate

For the LASSO, Ridge, and Elastic Net regressions, we have first remove multicollinearity before applying them. To address this issue, we will examine each variable's variance inflation factor. As a rule of thumb, a $VIF > 5$ implies that the variable is a potential cause of collinearity, so we will remove it from our dataset.

```
rm(list = ls())
# importing data
df = read.csv('house_data.csv')
cat("Number of variables:",length(df))
```

```
## Number of variables: 81
```

```
head(df) # check
```

```
##      Id MSSubClass MSZoning LotFrontage LotArea Street Alley LotShape LandContour
## 1 1      60      RL      65      8450   Pave  <NA>      Reg      Lvl
## 2 2      20      RL      80      9600   Pave  <NA>      Reg      Lvl
## 3 3      60      RL      68     11250   Pave  <NA>      IR1      Lvl
## 4 4      70      RL      60     9550   Pave  <NA>      IR1      Lvl
## 5 5      60      RL      84     14260   Pave  <NA>      IR1      Lvl
## 6 6      50      RL      85     14115   Pave  <NA>      IR1      Lvl
##      Utilities LotConfig LandSlope Neighborhood Condition1 Condition2 BldgType
## 1 AllPub      Inside      Gtl      CollgCr      Norm      Norm      1Fam
## 2 AllPub      FR2      Gtl      Veenker      Feedr      Norm      1Fam
## 3 AllPub      Inside      Gtl      CollgCr      Norm      Norm      1Fam
## 4 AllPub      Corner      Gtl      Crawfor      Norm      Norm      1Fam
## 5 AllPub      FR2      Gtl      NoRidge      Norm      Norm      1Fam
## 6 AllPub      Inside      Gtl      Mitchel      Norm      Norm      1Fam
##      HouseStyle OverallQual OverallCond YearBuilt YearRemodAdd RoofStyle RoofMatl
## 1 2Story      7      5      2003      2003      Gable      CompShg
## 2 1Story      6      8      1976      1976      Gable      CompShg
## 3 2Story      7      5      2001      2002      Gable      CompShg
## 4 2Story      7      5      1915      1970      Gable      CompShg
## 5 2Story      8      5      2000      2000      Gable      CompShg
## 6 1.5Fin      5      5      1993      1995      Gable      CompShg
##      Exterior1st Exterior2nd MasVnrType MasVnrArea ExterQual ExterCond Foundation
## 1 VinylSd      VinylSd      BrkFace      196      Gd      TA      PConc
## 2 MetalSd      MetalSd      None      0      TA      TA      CBlocc
## 3 VinylSd      VinylSd      BrkFace      162      Gd      TA      PConc
## 4 Wd Sdng      Wd Shng      None      0      TA      TA      BrkTil
## 5 VinylSd      VinylSd      BrkFace      350      Gd      TA      PConc
## 6 VinylSd      VinylSd      None      0      TA      TA      Wood
##      BsmtQual BsmtCond BsmtExposure BsmtFinType1 BsmtFinSF1 BsmtFinType2
## 1 Gd      TA      No      GLQ      706      Unf
## 2 Gd      TA      Gd      ALQ      978      Unf
## 3 Gd      TA      Mn      GLQ      486      Unf
## 4 TA      Gd      No      ALQ      216      Unf
## 5 Gd      TA      Av      GLQ      655      Unf
## 6 Gd      TA      No      GLQ      732      Unf
##      BsmtFinSF2 BsmtUnfSF TotalBsmtSF Heating HeatingQC CentralAir Electrical
## 1 0      150      856      GasA      Ex      Y      SBrkr
## 2 0      284      1262      GasA      Ex      Y      SBrkr
## 3 0      434      920      GasA      Ex      Y      SBrkr
## 4 0      540      756      GasA      Gd      Y      SBrkr
## 5 0      490      1145      GasA      Ex      Y      SBrkr
## 6 0      64      796      GasA      Ex      Y      SBrkr
##      X1stFlrSF X2ndFlrSF LowQualFinSF GrLivArea BsmtFullBath BsmtHalfBath FullBath
## 1 856      854      0      1710      1      0      2
```

```
## 2      1262      0      0      1262      0      1      2
## 3      920     866      0     1786      1      0      2
## 4      961     756      0     1717      1      0      1
## 5     1145    1053      0     2198      1      0      2
## 6      796     566      0     1362      1      0      1
##      HalfBath BedroomAbvGr KitchenAbvGr KitchenQual TotRmsAbvGrd Functional
## 1      1      3      1      Gd      8      Typ
## 2      0      3      1      TA      6      Typ
## 3      1      3      1      Gd      6      Typ
## 4      0      3      1      Gd      7      Typ
## 5      1      4      1      Gd      9      Typ
## 6      1      1      1      TA      5      Typ
##      Fireplaces FireplaceQu GarageType GarageYrBlt GarageFinish GarageCars
## 1      0      <NA>      Attchd      2003      RFn      2
## 2      1      TA      Attchd      1976      RFn      2
## 3      1      TA      Attchd      2001      RFn      2
## 4      1      Gd      Detchd      1998      Unf      3
## 5      1      TA      Attchd      2000      RFn      3
## 6      0      <NA>      Attchd      1993      Unf      2
##      GarageArea GarageQual GarageCond PavedDrive WoodDeckSF OpenPorchSF
## 1      548      TA      TA      Y      0      61
## 2      460      TA      TA      Y      298      0
## 3      608      TA      TA      Y      0      42
## 4      642      TA      TA      Y      0      35
## 5      836      TA      TA      Y      192      84
## 6      480      TA      TA      Y      40      30
##      EnclosedPorch X3SsnPorch ScreenPorch PoolArea PoolQC Fence MiscFeature
## 1      0      0      0      0      <NA> <NA> <NA>
## 2      0      0      0      0      <NA> <NA> <NA>
## 3      0      0      0      0      <NA> <NA> <NA>
## 4      272      0      0      0      <NA> <NA> <NA>
## 5      0      0      0      0      <NA> <NA> <NA>
## 6      0      320      0      0      <NA> MnPrv      Shed
##      MiscVal MoSold YrSold SaleType SaleCondition SalePrice
## 1      0      2      2008      WD      Normal      208500
## 2      0      5      2007      WD      Normal      181500
## 3      0      9      2008      WD      Normal      223500
## 4      0      2      2006      WD      Abnorml      140000
## 5      0      12     2008      WD      Normal      250000
## 6     700     10     2009      WD      Normal      143000
```

```
# convert ID column into row names
# and drop it
rownames(df) <- df$Id
drops <- c('Id')
df <- df[ , !(names(df) %in% drops)]

# check length of our data
cat("Length of data:",length(df$MSSubClass))
```

```
## Length of data: 1460
```

```
# check if our target variable has any missing values
cat("Missing values in our target var:",sum(is.na(df$SalePrice)))
```

```
## Missing values in our target var: 0
```

```
# great, now we can train-test-split
set.seed(42)
train_index = createDataPartition(df$SalePrice, p = .7, list = FALSE)
train <- df[train_index,]
test  <- df[-train_index,]

# drop columns with high number of NA values
# define "high" as >20% null values
drops <- c()

# find the columns with a high number of NAs
for (col in names(train)){
  if (sum(is.na(df[col])) > 0.2*nrow(df[col])){
    drops <- c(drops,col)
  }
}

# drop those columns from our data
train <- train[ , !(names(train) %in% drops)]
test <- test[ , !(names(test) %in% drops)]

# find if there are any variables that don't
# give any information i.e. 0 variance
# this would be the case if a column only
# has 1 unique value
cat("In the training set these variables have 0 variance:",names(sapply(lapply(train, unique), length))
```

```
## In the training set these variables have 0 variance:
```

```
cat("In the test set these variables have 0 variance:",names(sapply(lapply(test, unique), length))[sapply(
  '\n')
```

```
## In the test set these variables have 0 variance: Utilities
```

```
# we will drop the utilities variable
drops <- c('Utilities')
train<- train[ , !(names(train) %in% drops)]
test <- test[ , !(names(test) %in% drops)]

# split into train-test sets
drops <- c('SalePrice')
X_train <-train[ , !(names(train) %in% drops)]
y_train <- train$SalePrice
X_test <- test[ , !(names(test) %in% drops)]
y_test <- test$SalePrice
```


Pipeline

```
# handling numeric data
# (1) impute > median
# (2) scale
X_train_scaled <- X_train %>% mutate_if(is.numeric,function(x) ifelse(is.na(x),median(x,na.rm=T),x)) %>%

X_test_scaled <- X_test %>% mutate_if(is.numeric,function(x) ifelse(is.na(x),median(x,na.rm=T),x)) %>%

# handling categorical data
# (1) impute with mode
X_train_scaled <- X_train_scaled %>% mutate_if(is.character,function(x) ifelse(is.na(x),mode(x),x))
X_test_scaled <- X_test_scaled %>% mutate_if(is.character,function(x) ifelse(is.na(x),mode(x),x))

# (2) encode data
X_train_scaled <- X_train_scaled %>% mutate_if(is.character,function(x) as.integer(factor(x)))
X_test_scaled <- X_test_scaled %>% mutate_if(is.character,function(x) as.integer(factor(x)))
```

PCR

```
library(pls)
```

```
##
## Attaching package: 'pls'

## The following object is masked from 'package:caret':
##
##      R2

## The following object is masked from 'package:stats':
##
##      loadings
```

```
set.seed(42)

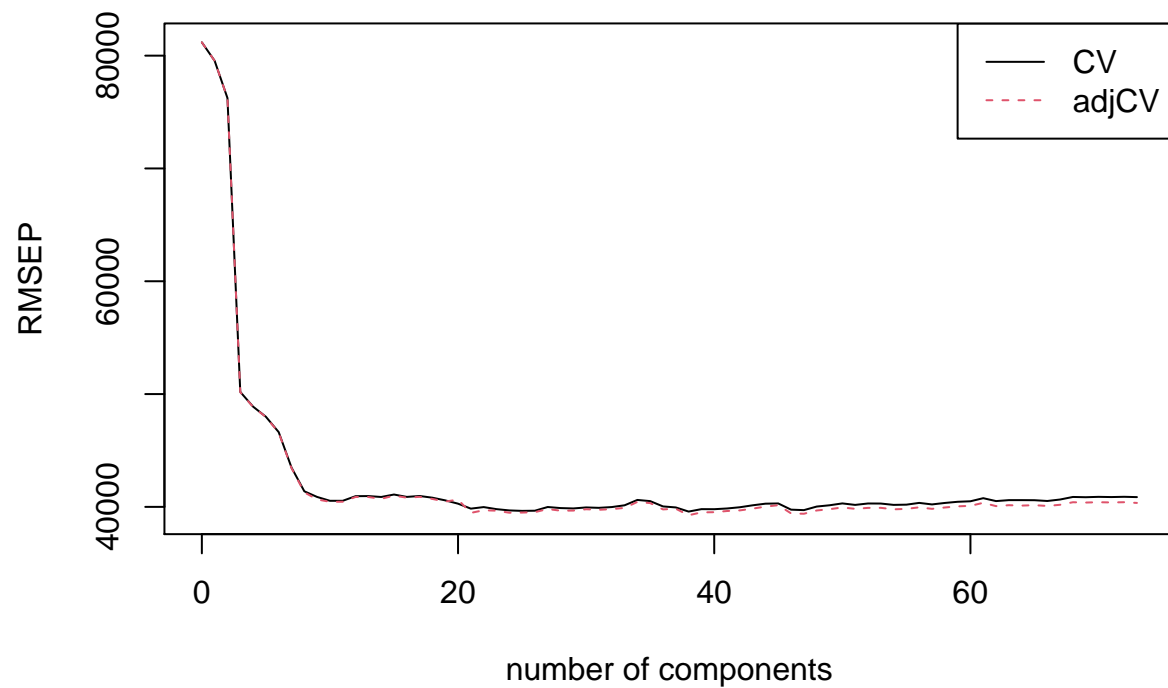
# combine y_train and X_train_scaled
train_scaled <- X_train_scaled
train_scaled['SalePrice'] <- y_train

# combine y_test and X_test_scaled
test_scaled <- X_test_scaled
test_scaled['SalePrice'] <- y_test

# fit principal component analysis regression
pcr.fit <- pcr(SalePrice ~ .,data = train_scaled,validation = "CV")

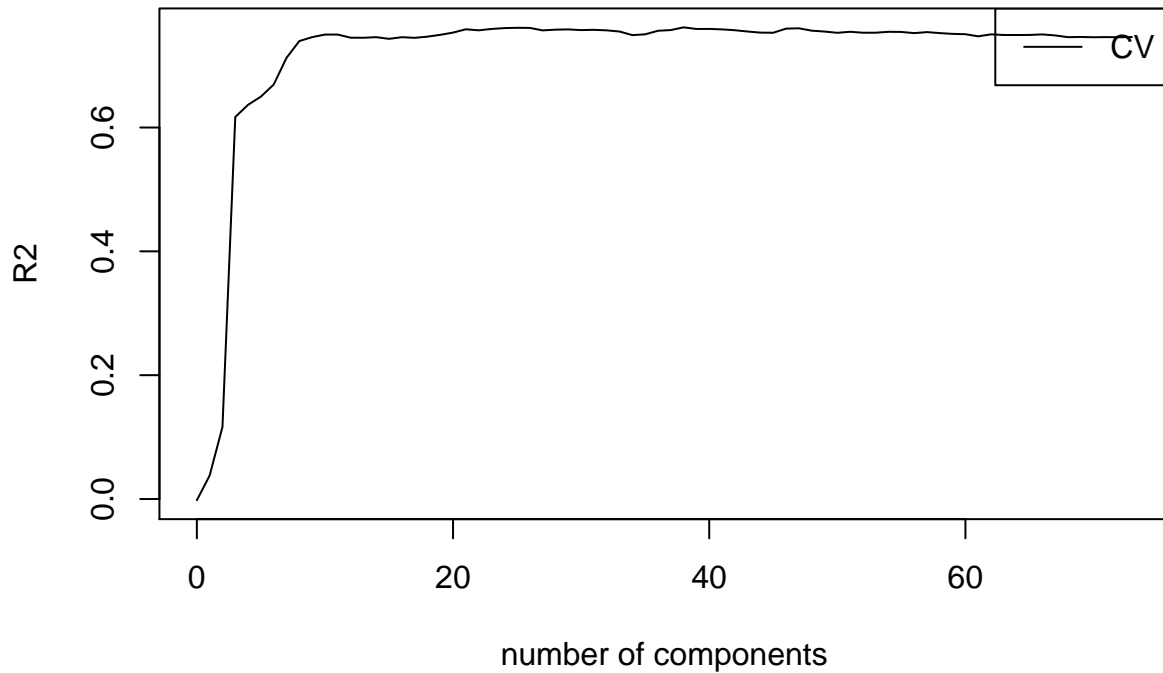
# plot RMSE vs number of components
validationplot(pcr.fit, val.type = "RMSEP",
               legendpos='topright',
               main = 'Number of Principal Components needed to minimise RMSE to
Predict Sale Price')
```

Number of Principal Components needed to minimise RMSE to Predict Sale Price



```
# plot of Rsquared vs number of components
validationplot(pcr.fit, val.type = "R2",
               legendpos='topright',
               main = 'Principal Components needed to maximise R-squared to
Predict Sale Price')
```

Principal Components needed to maximise R-squared to Predict Sale Price



From the validation plot, we observe that the number of components with the lowest cross-validation error is in the range of 21 to 40 components. Therefore, we will test `ncomps` in [21,40] to see which principal component regression model performs best on our test set.

```
ncomps = seq(1:20) + 20
ncomp_score <- c()
for (n in ncomps){
  pcr.pred <- predict(pcr.fit, X_test_scaled, ncomp = n)
  ncomp_score <- c(ncomp_score, sqrt(mean((pcr.pred - y_test)^2)))
}
```

```
# table of ncomps and respective test scores
data.frame(ncomps, ncomp_score)
```

```
##      ncomps ncomp_score
## 1         21    28476.00
## 2         22    28580.16
## 3         23    28582.13
## 4         24    28344.71
## 5         25    28349.25
## 6         26    28379.93
## 7         27    28271.16
## 8         28    28265.73
## 9         29    28308.92
## 10        30    28243.94
## 11        31    28141.38
```

```
## 12      32      28215.65
## 13      33      28715.08
## 14      34      28714.30
## 15      35      28804.00
## 16      36      28346.10
## 17      37      28302.48
## 18      38      28882.26
## 19      39      29028.45
## 20      40      29050.29
```

This table shows that $\text{ncomps} = 31$ performs the best with an RMSE of 28,141.38. This performance will later be compared against the rest of the regressions.

Ridge

Removing linearly dependent variables

Before moving onto any of the other regressions, we're going to eliminate the linearly dependent variables from our data. We didn't confront this problem earlier because principal component analysis naturally eliminates multicollinearity.

```
library(car)

## Loading required package: carData

##
## Attaching package: 'car'

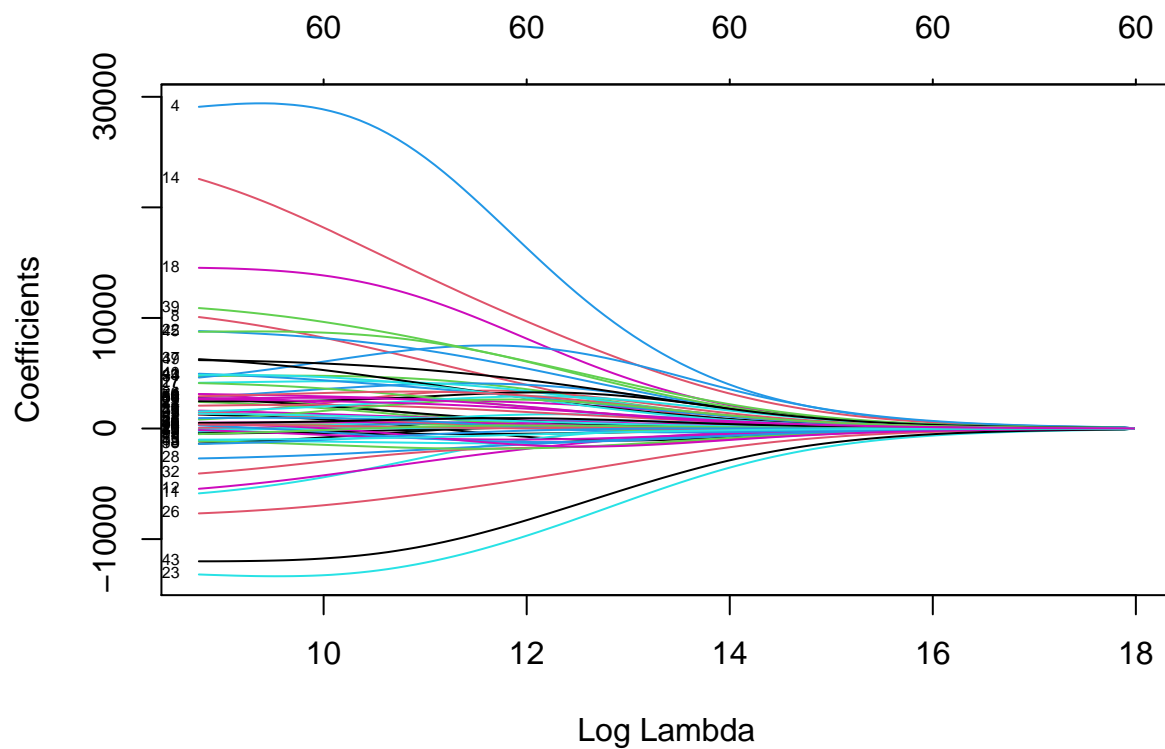
## The following object is masked from 'package:gtools':
##
##      logit

## The following object is masked from 'package:dplyr':
##
##      recode

## The following object is masked from 'package:purrr':
##
##      some

# find linearly dependent variables
fit <- lm(SalePrice~.,data = train_scaled)
ld_vars <- attributes(alias(fit)$Complete)$dimnames[[1]]
cat('Linearly dependent variables:',ld_vars)

## Linearly dependent variables: TotalBsmtSF GrLivArea
```

This plot shows how increasing the size of lambda affects the coefficients of different variables. We observe that our 4th, 14th, 45th, 23rd, 36th, 37th, and 26th variables are highly important, while the other variables are less so.

Now let's take a closer look at our important variables by creating a variable importance plot.

```
library(vip)
```

```
##
```

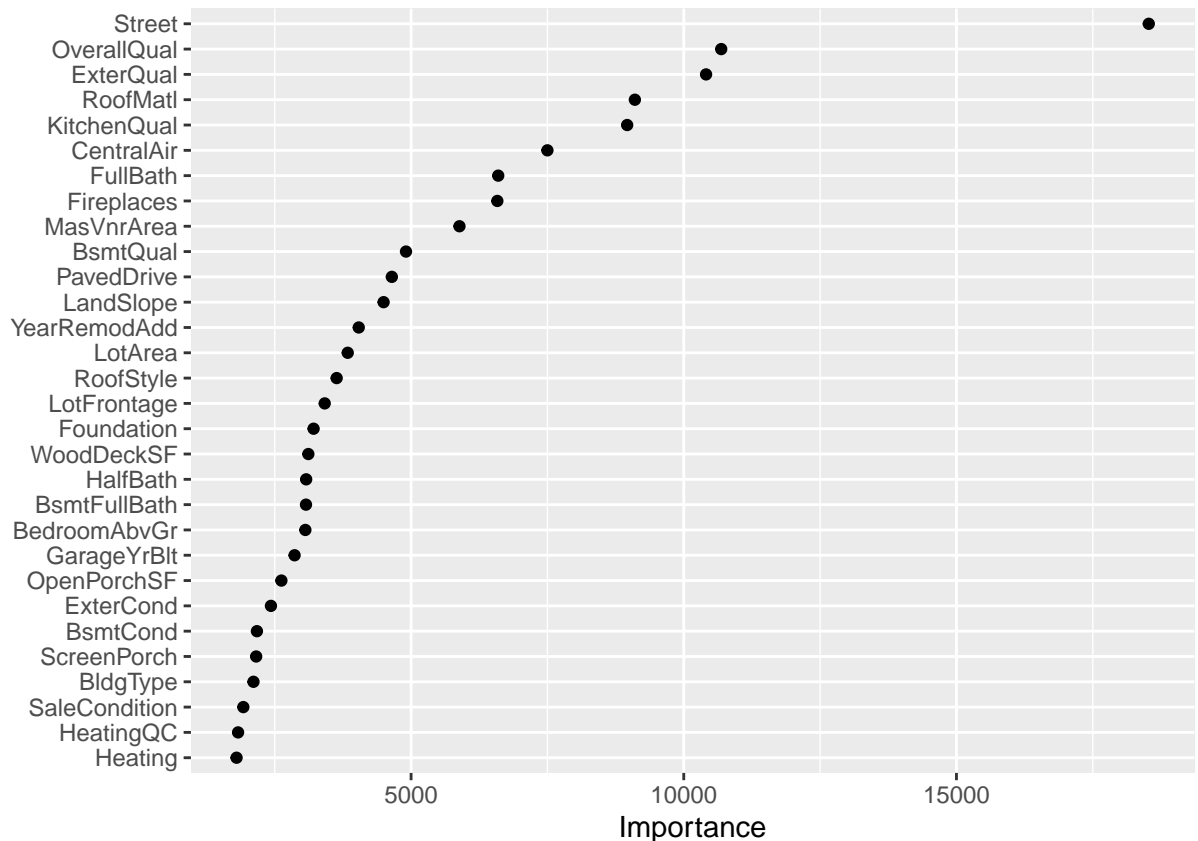
```
## Attaching package: 'vip'
```

```
## The following object is masked from 'package:utils':
```

```
##
```

```
## vi
```

```
vip(model_ride, num_features = 30, geom = "point")
```



This shows us that our street variable (which is the type of street the property is on) is the most important variable by far, then exterior quality, overall quality, and kitchen quality are the next most important variables. On a qualitative level, this makes sense because most people evaluate real-estate from the outside first, then the inside. Kitchen quality is unsurprising because many people spend a lot of time in their kitchens, either eating food or preparing food. Finally, on the less important side of the 30 variables, we see heating, basement condition, and so forth, which is also expected – at least in Australia, where it’s generally quite warm, and because the basement aesthetics are less noticeable.

```
# we can utilize cross validation to train our ridge model
# and find the best lambda
train_control <- trainControl(method = "repeatedcv",
                              number = 5,
                              repeats = 1,
                              search = "random",
                              verboseIter = FALSE)

ridge_model <- train(SalePrice ~ .,
                    data = train_scaled_reg,
                    metrics = 'RMSE',
                    method = "ridge",
                    tuneLength = 25,
                    trControl = train_control)

# Predict using the testing data
ridge_preds = predict(ridge_model, newdata = X_test_scaled)
```

```
# Evaluate performance
postResample(pred = ridge_preds, obs = y_test)
```

##	RMSE	Rsquared	MAE
##	3.314731e+04	8.070257e-01	2.371836e+04

This shows us that our ridge model's best cross-validated performance is 33,211.91, which is worse than our PCR model (by roughly 4,000). The Rsquared is 0.8071006.

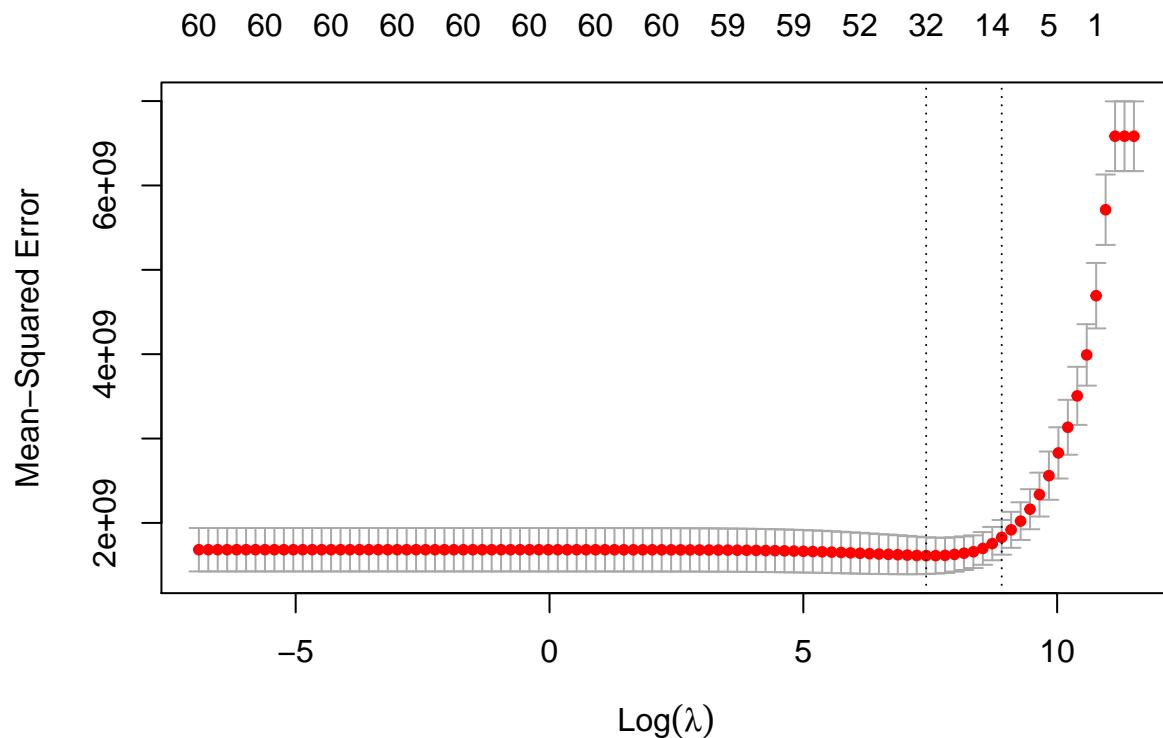
LASSO

We will now try a LASSO model to regularize our data. The LASSO model works similarly to Ridge, except by using a different penalty term that takes the absolute value of β , which results in the coefficients of less important variables can be reduced to exactly 0 (not just near-zero as in our Ridge Model). Thus, by using the penalty term $\lambda \sum_{i=1}^p |\beta_i|$, our LASSO model will have high predictive power and be simple to interpret.

```
# Perform 10-fold cross-validation to select lambda
lambdas_to_try <- 10^seq(-3, 5, length.out = 100)

# Setting alpha = 1 implements lasso regression
lasso_cv <- cv.glmnet(x = as.matrix(train_scaled_reg[, -which(names(train_scaled_reg) %in% c("SalePrice",
                                                    "standardize = TRUE, nfolds = 10)

# Plot cross-validation results
plot(lasso_cv)
```



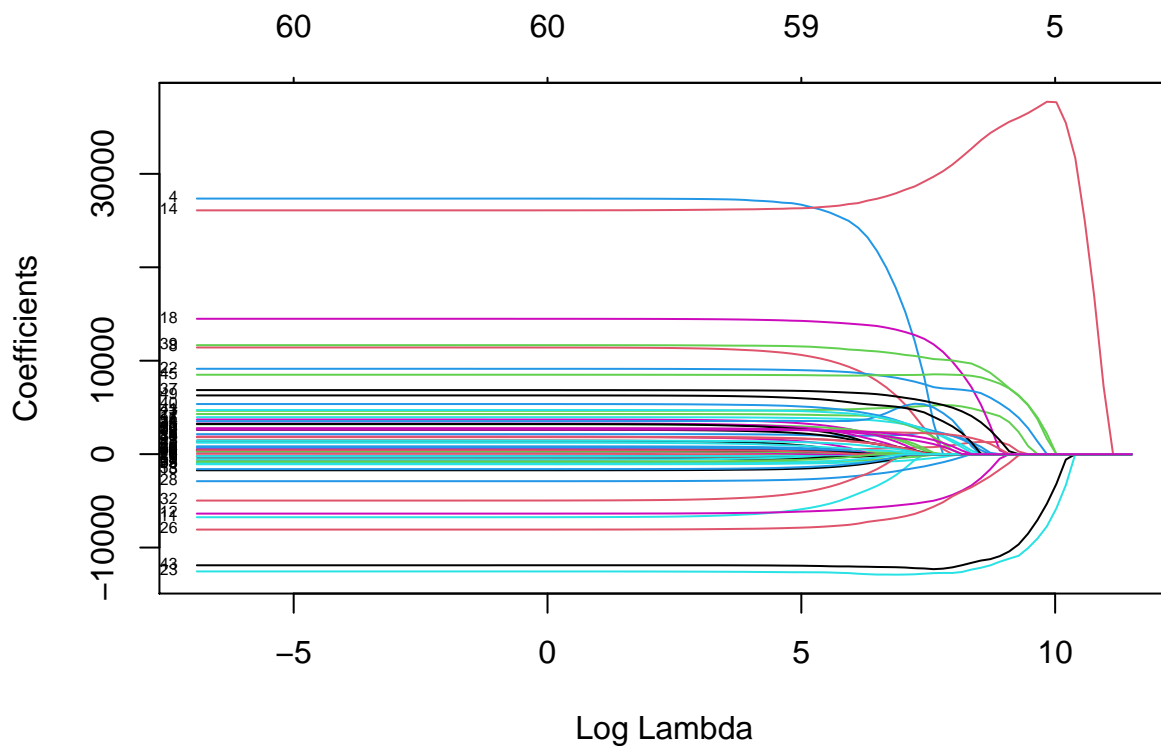

```

# Best cross-validated lambda
lambda_cv <- lasso_cv$lambda.min

# Fit final model
model_lasso <- glmnet(x = as.matrix(train_scaled_reg[, -which(names(train_scaled_reg) %in% c("SalePrice"

# Compare variables across lambdas
plot(lasso_cv$glmnet.fit, "lambda", label=TRUE)

```



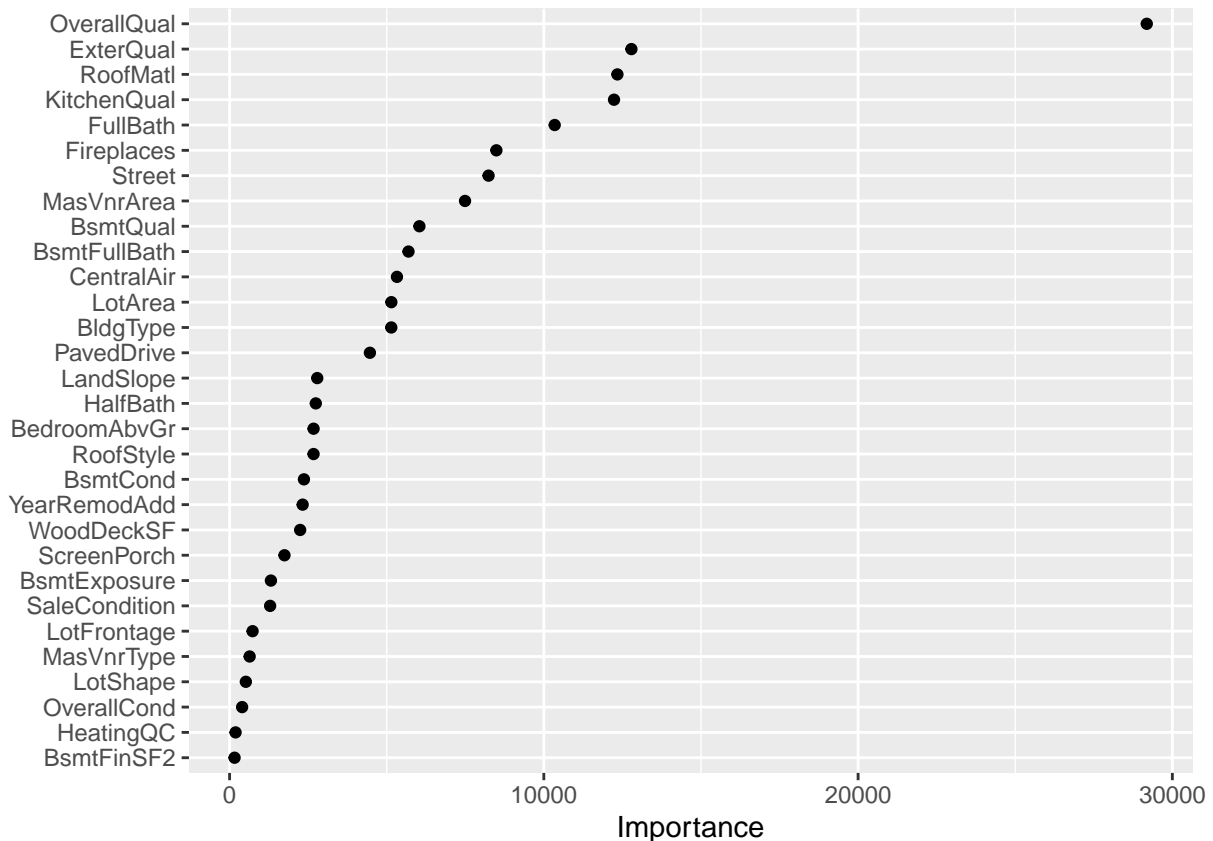
As with the Ridge Model, this plot shows how the coefficients of different variables react to increasing lambda. From this plot we find the 14th, 23rd, 43rd, 30th, 45th, and 22nd variables are highly important, while other variables are less so.

We will now create a variable importance plot to closer examine these important variables.

```

library(vip)
vip(model_lasso, num_features = 30, geom = "point")

```



This shows us that Overall Quality is the most important variable and Street is the second most important and both of these variables are far more important than the rest. While our Ridge Model had Street as the most important variable and Overall Quality at second, both our Ridge Model and Lasso Model agree on the top two most important variables. Our Lasso Model then shows that Roof Material, External Quality, Kitchen Quality, and Full Bathrooms are the next most important variables. Our Ridge Model resulted in the same aside from Central Air replacing Full Bathrooms. On the other end, both our Lasso and our Ridge Models agree that heating is not that important of a variable.

We will now train our Lasso Model and predict our test set to measure RMSE:

```
# we can utilize cross validation to train our lasso model
# and find the best lambda
train_control <- trainControl(method = "repeatedcv",
                              number = 5,
                              repeats = 5,
                              search = "random",
                              verboseIter = FALSE)

lasso_model <- train(SalePrice ~ .,
                    data = train_scaled_reg,
                    metrics = 'RMSE',
                    method = "glmnet",
                    tuneGrid = expand.grid(alpha = 1,
                                           lambda = 1),
                    tuneLength = 25,
                    trControl = train_control)
```

```
# Predict using the testing data
lasso_preds = predict(lasso_model, newdata = X_test_scaled)

# Evaluate performance
postResample(pred = lasso_preds, obs = y_test)
```

```
##          RMSE      Rsquared      MAE
## 33128.049508    0.806975 23602.019008
```

This shows us that our LASSO model's best cross-validated performance is 33,128.05, which is worse than our PCR model (by roughly 4,000). The Rsquared is 0.806975.

Elastic Net

We will now try an Elastic Net model to regularize our data. Elastic Net improves upon Ridge and Lasso by combining the penalties from each model resulting in a shrinkage model less dependent on the data. The new penalty term is $\frac{1-\alpha}{2} \sum_{i=1}^p \beta_i^2 + \alpha \sum_{i=1}^p |\beta_i|$ where $\alpha = 0$ (Ridge) % $\alpha = 1$ (Lasso).

```
#use cross validation to train our elastic net model
train_control <- trainControl(method = "repeatedcv",
                              number = 5,
                              repeats = 5,
                              search = "random",
                              verboseIter = TRUE)

# Train the model
elastic_net_model <- train(SalePrice ~ .,
                           data = train_scaled_reg,
                           metrics = 'RMSE',
                           method = "glmnet",
                           preProcess = c("center", "scale"),
                           tuneLength = 25,
                           trControl = train_control)
```

```
## + Fold1.Rep1: alpha=0.63608, lambda=0.0017708
## - Fold1.Rep1: alpha=0.63608, lambda=0.0017708
## + Fold1.Rep1: alpha=0.97864, lambda=0.1779302
## - Fold1.Rep1: alpha=0.97864, lambda=0.1779302
## + Fold1.Rep1: alpha=0.42631, lambda=0.0633234
## - Fold1.Rep1: alpha=0.42631, lambda=0.0633234
## + Fold1.Rep1: alpha=0.37574, lambda=0.0015142
## - Fold1.Rep1: alpha=0.37574, lambda=0.0015142
## + Fold1.Rep1: alpha=0.80531, lambda=0.0023148
## - Fold1.Rep1: alpha=0.80531, lambda=0.0023148
## + Fold1.Rep1: alpha=0.06683, lambda=0.1639706
## - Fold1.Rep1: alpha=0.06683, lambda=0.1639706
## + Fold1.Rep1: alpha=0.01230, lambda=0.3197005
## - Fold1.Rep1: alpha=0.01230, lambda=0.3197005
## + Fold1.Rep1: alpha=0.75340, lambda=0.1553943
## - Fold1.Rep1: alpha=0.75340, lambda=0.1553943
## + Fold1.Rep1: alpha=0.65786, lambda=0.0962132
## - Fold1.Rep1: alpha=0.65786, lambda=0.0962132
```

```

## + Fold1.Rep1: alpha=0.42268, lambda=4.0716342
## - Fold1.Rep1: alpha=0.42268, lambda=4.0716342
## + Fold1.Rep1: alpha=0.87538, lambda=0.0525596
## - Fold1.Rep1: alpha=0.87538, lambda=0.0525596
## + Fold1.Rep1: alpha=0.49128, lambda=6.5121646
## - Fold1.Rep1: alpha=0.49128, lambda=6.5121646
## + Fold1.Rep1: alpha=0.59242, lambda=0.0997138
## - Fold1.Rep1: alpha=0.59242, lambda=0.0997138
## + Fold1.Rep1: alpha=0.29222, lambda=0.0009879
## - Fold1.Rep1: alpha=0.29222, lambda=0.0009879
## + Fold1.Rep1: alpha=0.29635, lambda=1.0847852
## - Fold1.Rep1: alpha=0.29635, lambda=1.0847852
## + Fold1.Rep1: alpha=0.89740, lambda=0.0104120
## - Fold1.Rep1: alpha=0.89740, lambda=0.0104120
## + Fold1.Rep1: alpha=0.38484, lambda=0.0798488
## - Fold1.Rep1: alpha=0.38484, lambda=0.0798488
## + Fold1.Rep1: alpha=0.50231, lambda=0.0405177
## - Fold1.Rep1: alpha=0.50231, lambda=0.0405177
## + Fold1.Rep1: alpha=0.87686, lambda=2.7659135
## - Fold1.Rep1: alpha=0.87686, lambda=2.7659135
## + Fold1.Rep1: alpha=0.78346, lambda=0.3752296
## - Fold1.Rep1: alpha=0.78346, lambda=0.3752296
## + Fold1.Rep1: alpha=0.81532, lambda=0.0358768
## - Fold1.Rep1: alpha=0.81532, lambda=0.0358768
## + Fold1.Rep1: alpha=0.93018, lambda=0.0013239
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## Aggregating results
## Selecting tuning parameters
## Fitting alpha = 0.0668, lambda = 0.164 on full training set

```

Our cross validation results in the tuning parameters $\alpha = 0.001489799$ and $\lambda = 0.007510219$ for our Elastic Net Model. We will now predict the testing data in order to measure the RMSE of our model.

We will now look at the variables our E-Net model finds the most important:

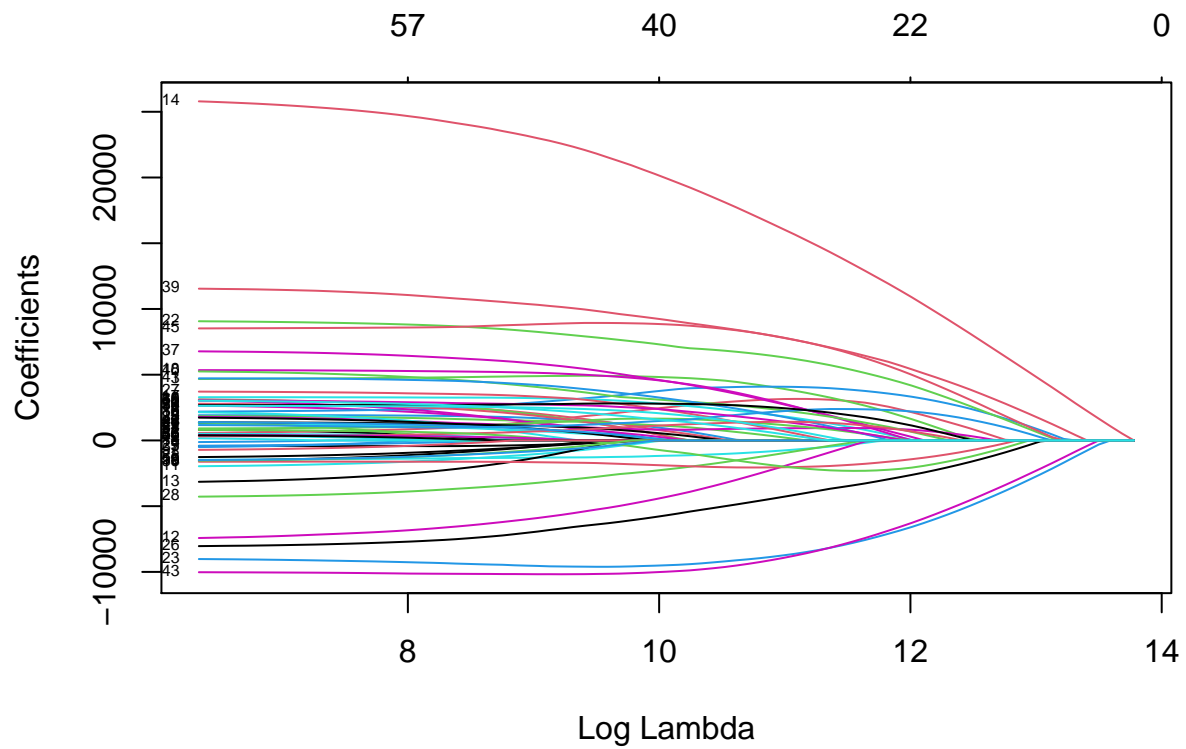
```
print(elastic_net_model)
```

```
## glmnet
##
## 1024 samples
## 60 predictor
##
## Pre-processing: centered (60), scaled (60)
## Resampling: Cross-Validated (5 fold, repeated 5 times)
## Summary of sample sizes: 820, 818, 820, 819, 819, 819, ...
## Resampling results across tuning parameters:
##
##  alpha      lambda      RMSE      Rsquared    MAE
##  0.01229978 0.3197005277 40745.39 0.7545629 25312.08
##  0.06683004 0.1639706105 40737.38 0.7547006 25306.71
##  0.29222196 0.0009879343 40753.04 0.7546061 25324.78
##  0.29634765 1.0847852177 40752.62 0.7546095 25324.72
##  0.31997283 0.0066008589 40753.54 0.7546021 25325.25
##  0.37573718 0.0015141704 40756.00 0.7545802 25327.66
##  0.38483874 0.0798487895 40756.26 0.7545791 25327.66
##  0.42268111 4.0716341923 40757.09 0.7545720 25328.96
##  0.42630757 0.0633233870 40757.22 0.7545708 25329.08
##  0.49128262 6.5121645952 40758.09 0.7545660 25329.57
##  0.50231095 0.0405176509 40758.35 0.7545634 25329.82
##  0.59242344 0.0997138392 40759.41 0.7545555 25331.33
##  0.63608417 0.0017707770 40760.68 0.7545379 25332.49
##  0.65786011 0.0962131861 40760.70 0.7545378 25332.62
##  0.74423815 0.0070055641 40761.24 0.7545346 25333.33
##  0.75339509 0.1553943124 40760.86 0.7545406 25333.29
##  0.78346385 0.3752296113 40760.74 0.7545452 25333.03
##  0.80531494 0.0023148114 40760.99 0.7545428 25333.27
##  0.81532431 0.0358768471 40760.21 0.7545502 25332.06
##  0.87537882 0.0525596350 40761.54 0.7545354 25333.37
##  0.87685746 2.7659135237 40761.55 0.7545353 25333.38
##  0.89739502 0.0104119842 40761.19 0.7545399 25333.04
##  0.91107522 4.6314855213 40761.31 0.7545388 25333.15
##  0.93017705 0.0013239303 40761.42 0.7545377 25333.26
##  0.97864070 0.1779302308 40761.77 0.7545346 25333.58
##
## RMSE was used to select the optimal model using the smallest value.
## The final values used for the model were alpha = 0.06683004 and lambda
## = 0.1639706.
```

```
elastic_net_model$bestTune
```

```
##      alpha      lambda
## 2 0.06683004 0.1639706
```

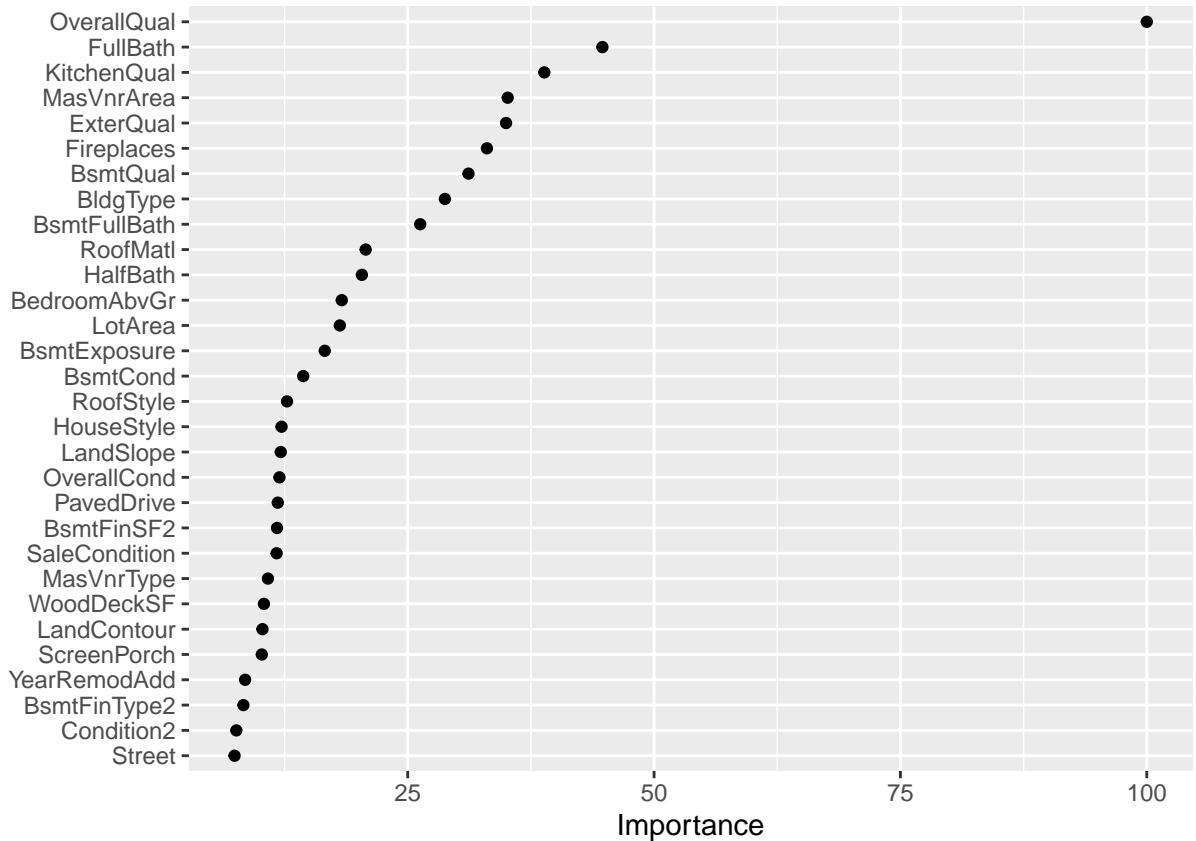
```
plot(elastic_net_model$finalModel, "lambda", label=TRUE)
```



This plot again shows how the coefficients of different variables react to increasing lambda. This plot demonstrates that the 14th, 39th, 43rd, 23rd, 22nd, 46th, and 26th are highly important, while other variables are less so.

We will now create a variable importance plot to closer examine these important variables.

```
vip(elastic_net_model, num_features = 30, geom = "point")
```



Our Elastic Net Model shows that Overall Quality is the most important variable by far, then followed by Full Bathrooms, Kitchen Quality, and External Quality - which is in agreement with our Ridge and Lasso models aside from the Street variable.

We will now use our elastic net model fit on the full training set to predict the test set in order to measure RMSE and R-squared.

```
# Predict using the testing data
enet_preds = predict(elastic_net_model, newdata = X_test_scaled)

# Evaluate performance
postResample(enet_preds, y_test)
```

```
##          RMSE      Rsquared        MAE
## 3.311439e+04 8.070347e-01 2.354398e+04
```

This shows us that our Elastic Net Model's best cross-validated performance is 33,115.49 which is almost 4,000 higher than our Ridge model's performance.

Model Comparison

Comparing the RMSEs from predicting the test set Sale Price using our four regularization models, we find our principal component regression model performed best with the lowest RMSE of 28,141.38.

RMSE Comparison: PCR: 28,141.38 Ridge: 33,211.91 Lasso: 33,128.05 E-Net: 33,115.49

This PCR model found that the number of components with the lowest cross-validation error was `ncomps = 31`. Thus our final regularization model is a PCR model with `ncomps = 31`:

```
#Final PCR fit
pcr.fit.final <- pcr(SalePrice ~., data = train_scaled, ncomp = 31)
summary(pcr.fit.final)
```

```
## Data:      X dimension: 1024 73
## Y dimension: 1024 1
## Fit method: svdpc
## Number of components considered: 31
## TRAINING: % variance explained
##           1 comps  2 comps  3 comps  4 comps  5 comps  6 comps  7 comps
## X           26.031   41.39   50.09   54.59   58.06   61.12   63.75
## SalePrice    4.143   12.34   62.22   64.24   65.48   67.62   71.83
##           8 comps  9 comps 10 comps 11 comps 12 comps 13 comps 14 comps
## X           65.99   68.02   70.02   71.83   73.35   74.84   76.07
## SalePrice    74.57   75.20   75.74   75.77   75.78   75.82   76.24
##          15 comps 16 comps 17 comps 18 comps 19 comps 20 comps 21 comps
## X           77.27   78.39   79.50   80.51   81.51   82.41   83.29
## SalePrice    76.29   76.49   76.58   76.85   77.29   77.29   79.01
##          22 comps 23 comps 24 comps 25 comps 26 comps 27 comps 28 comps
## X           84.12   84.93   85.72   86.49   87.22   87.94   88.60
## SalePrice    79.10   79.10   79.29   79.30   79.31   79.43   79.82
##          29 comps 30 comps 31 comps
## X           89.24   89.87   90.49
## SalePrice    79.84   79.85   79.94
```

```
#Predicting Sale Price
pcr.pred.final <- predict(pcr.fit.final, X_test_scaled, ncomp = 31)
```

This PCR model reduced dimensionality from 73 variables down to 31 components and outperformed Ridge, Lasso, and Elastic Net regularization methods making it our model of choice.

Sources

<https://remiller1450.github.io/s230f19/caret3.html> <https://dataaspirant.com/knn-implementation-r-using-caret-package/> <https://www.rdocumentation.org/packages/caret/versions/4.47/topics/train>