

Statistics, Modeling, and Machine Learning

Reproducibility

Steps of data analysis: -Define the question

-Define the ideal data set

-Determine what data you can access

-Obtain the data

-Clean the Data

-Exploratory data analysis

-Statistical prediction/Modeling

-Interpret results

-Challenge results

-Synthesize/Write up results

-Create reproducible code

Probability

$P(A \cup B) = P(A) + P(B)$ - The probability of the union between scenario A and B is simply the sum of each. Random variable = is the numerical outcome of an experiment (discrete or continuous) - e.g. the flip of a coin (discrete random variable) - e.g. the roll of a dice (also discrete random variable because it's limited to 1-6) - e.g. BMI of a person is a continuous random variable

For random variables, the probabilities must add up to 1, and components are larger or equal to 0.

Probability mass function - for discrete random variables

Probability density function - for continuous random variables

in R, probability of some instance occurring within a continuous distribution can be calculated

Conditional probability- think lightning strike on a storming vs. sunny day.

$P(A|B) = P(A \cap B)/P(B)$ is the conditional probability. Written as the probability of A given

mean = center of a distribution variance and standard deviations = how spread out the distribution is

```
## Beta distribution
pbeta(0.75, 2, 1) ##here 0.75 is 75% probability of some density.
```

```
## [1] 0.5625
```

```
pbeta(c(.4, .5, .6), 2, 1) ##40-60% probability example of the same density.
```

```
## [1] 0.16 0.25 0.36
```

```
pnorm(70, mean = 80, sd = 10) ## probability to have value 70 given a set mean and standard dev. Data i
```

```
## [1] 0.1586553
```

```
qnorm(0.95, mean=1100, sd = 75) ## will return a value of a normal distribution given a specified perc
```

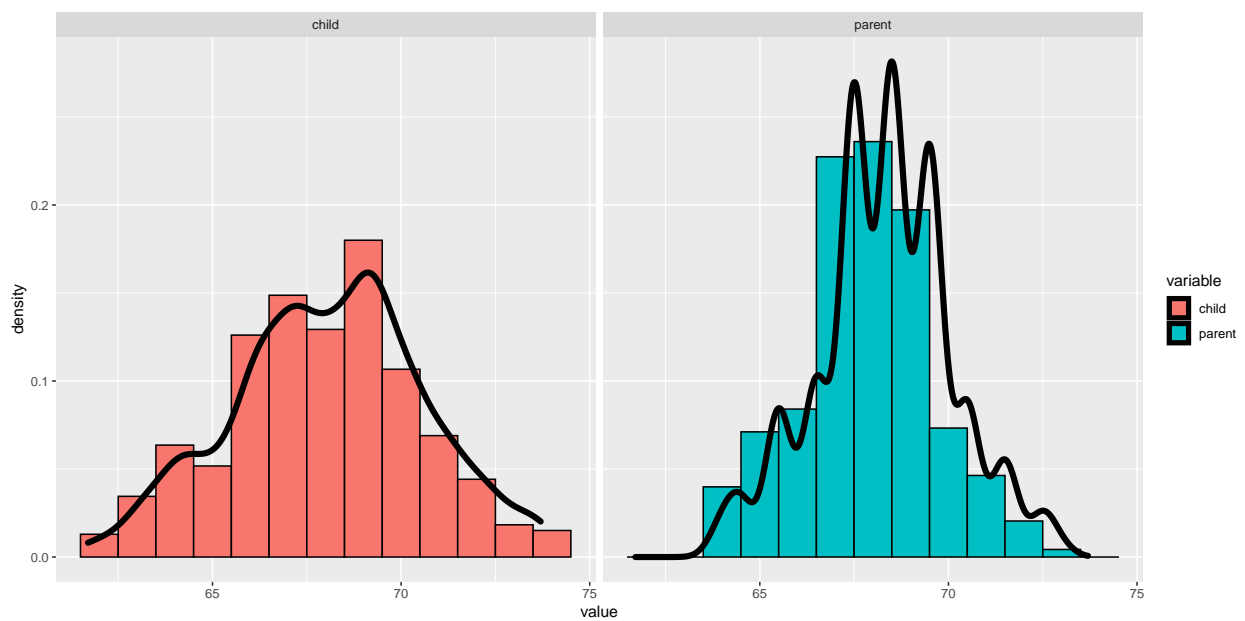
```
## [1] 1223.364
```

```
pnorm(16, mean = 15, sd = 1) - pnorm(14, mean = 15, sd = 1) ##probability that a an output is between tw
```

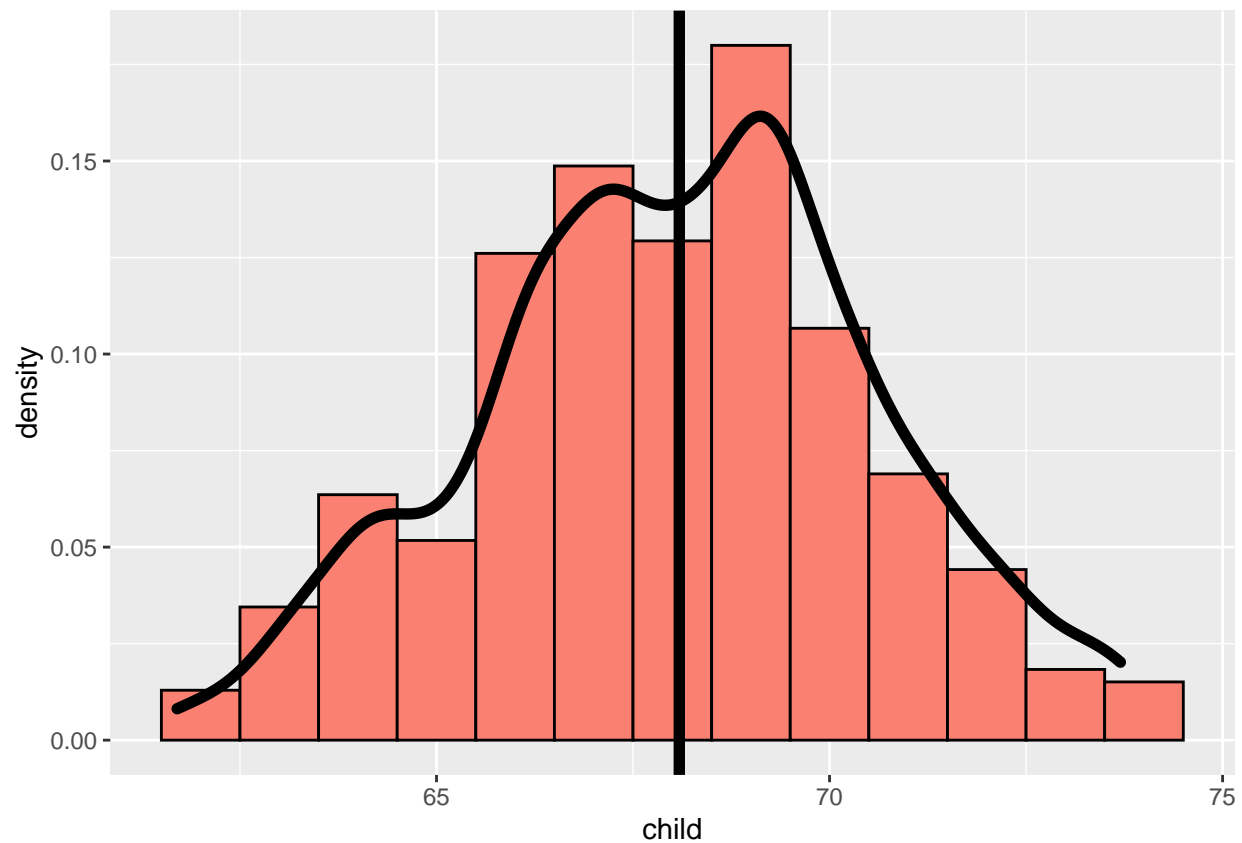
```
## [1] 0.6826895
```

```
ppois(10, lambda = 15) ##poisson distributed data. Probability of seeing 10 or less of an instance, whe
```

```
## [1] 0.1184644
```



```
g <- ggplot(galton, aes(x = child))
g <- g + geom_histogram(fill = "salmon",
  binwidth=1, aes(y = ..density..), colour = "black")
g <- g + geom_density(size = 2)
g <- g + geom_vline(xintercept = mean(galton$child), size = 2)
g
```



Variance = Total sum of squares

```
lambda <- 0.2
nsim <- 1:1000 # Number of Simulations/rows
n <- 40

Ematrix <- data.frame(x = sapply(nsim, function(x) {mean(rexp(n, lambda))})) ##creating a matrix of values
head(Ematrix)
```

```
##           x
## 1 3.349103
## 2 5.582835
## 3 4.038226
## 4 5.064299
## 5 5.987589
## 6 5.290295
```

```
Smean <- apply(Ematrix, 2, mean) ## calculating the simulated mean from above matrix.
Smean
```

```
##           x
## 4.971585
```

```
Tmean <- 1/lambda ##calculating the theoretical mean given lambda
Tmean
```

```
## [1] 5
```

```
SSD <- sd(Ematrix$x) ##The simulated standard deviation
SSD
```

```
## [1] 0.7920842
```

```
SVar <- var(Ematrix$x) ##The simulated variance from the above matrix.
SVar
```

```
## [1] 0.6273973
```

```
TSD <- (1/lambda)/sqrt(n) ##The theoretical standard deviation.
TSD
```

```
## [1] 0.7905694
```

```
TVar <- TSD^2 ##The theoretical calculation for the Variance
TVar
```

```
## [1] 0.625
```

t-test

$(\bar{X} - X^*)/(s/\sqrt{n})$ in other words, sample mean minus the hypothesized/test mean or value divided by standard error of the mean divided by square root of sample size, n. This follows a t-distribution with n-1 degrees of freedom. You can calculate the T-distribution using qt(.95, 15), where .95 is the percentile or an alpha of 0.05, and 15 is n-1 in this example.

```
library(UsingR); data(father.son)
t.test(father.son$sheight - father.son$fheight) ##testing whether there are significant differences be
```

```
##
## One Sample t-test
##
## data: father.son$sheight - father.son$fheight
## t = 11.789, df = 1077, p-value < 2.2e-16
## alternative hypothesis: true mean is not equal to 0
## 95 percent confidence interval:
## 0.8310296 1.1629160
## sample estimates:
## mean of x
## 0.9969728
```

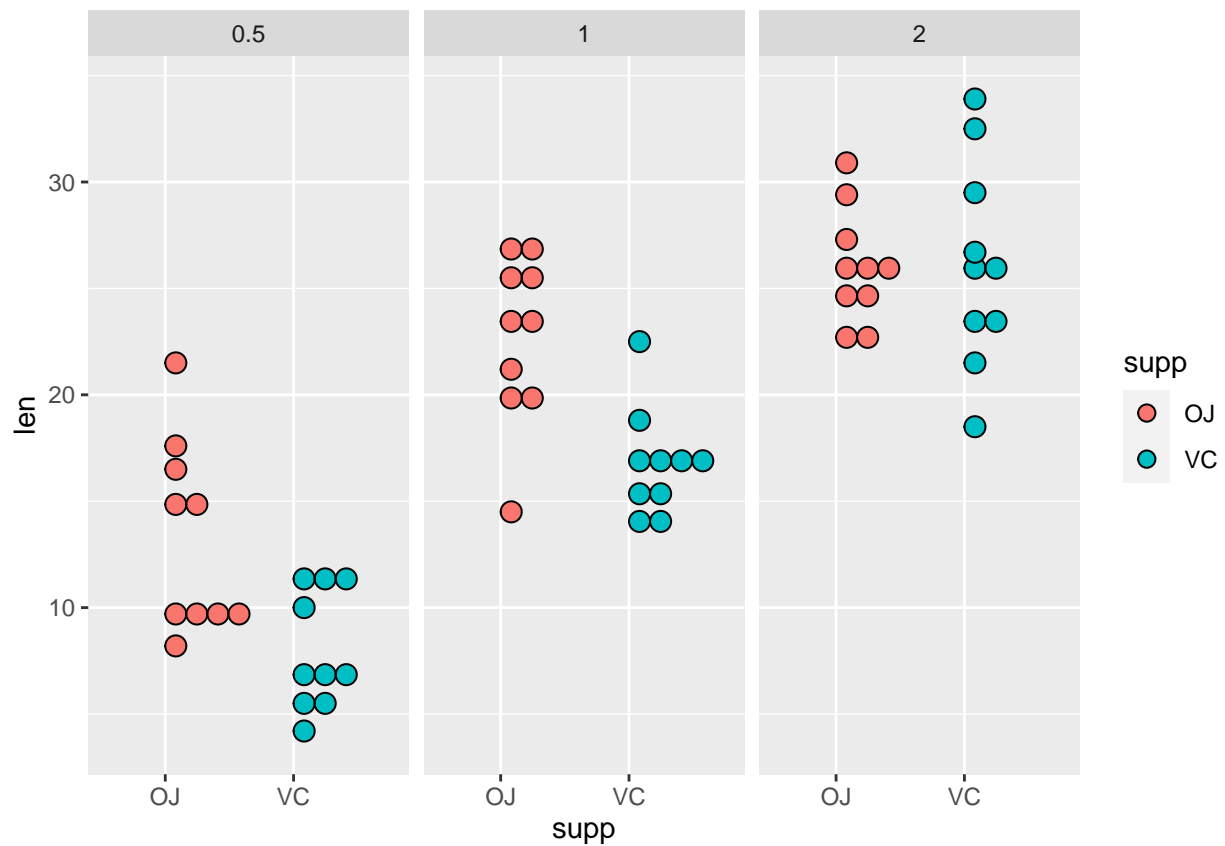
```
library(datasets)
data(ToothGrowth)
head(ToothGrowth)
```

```
##      len supp dose
## 1  4.2   VC  0.5
## 2 11.5   VC  0.5
## 3  7.3   VC  0.5
## 4  5.8   VC  0.5
## 5  6.4   VC  0.5
## 6 10.0   VC  0.5
```

```
library(ggplot2)

plot0 <- ggplot(ToothGrowth, aes(supp, len, fill = supp))
plot1 <- plot0 + geom_dotplot(binaxis = "y") +
  facet_grid(.~ dose)
plot1
```

Bin width defaults to 1/30 of the range of the data. Pick better value with 'binwidth'.



```
t.test(len ~ supp, data = ToothGrowth) ##testing sig difference between supp types and length
```

```
##
## Welch Two Sample t-test
##
## data: len by supp
## t = 1.9153, df = 55.309, p-value = 0.06063
## alternative hypothesis: true difference in means between group OJ and group VC is not equal to 0
## 95 percent confidence interval:
## -0.1710156 7.5710156
## sample estimates:
## mean in group OJ mean in group VC
## 20.66333 16.96333
```

```
test1 <- subset(ToothGrowth, dose %in% c(.5,1)) ##subsetting doses to make pairwise comparisons
t.test(len ~ dose, data = test1) ##testing sig differences between discrete doses and length
```

```
##
## Welch Two Sample t-test
##
## data: len by dose
## t = -6.4766, df = 37.986, p-value = 1.268e-07
## alternative hypothesis: true difference in means between group 0.5 and group 1 is not equal to 0
## 95 percent confidence interval:
## -11.983781 -6.276219
## sample estimates:
## mean in group 0.5 mean in group 1
## 10.605 19.735
```

```
test2 <- subset(ToothGrowth, dose %in% c(1,2)) ##subsetting doses to make pairwise comparisons
t.test(len ~ dose, data = test2)
```

```
##
## Welch Two Sample t-test
##
## data: len by dose
## t = -4.9005, df = 37.101, p-value = 1.906e-05
## alternative hypothesis: true difference in means between group 1 and group 2 is not equal to 0
## 95 percent confidence interval:
## -8.996481 -3.733519
## sample estimates:
## mean in group 1 mean in group 2
## 19.735 26.100
```

t-test Confidence Interval

```
## Prompt for below: a sample of 9 men yielded a sample average brain volume of 1,100cc and a standard deviation of 100cc
mn = 1100
s = 30
n = 9
round(1100 + c(-1,1)*qt(.975, df = 8)*s/sqrt(n))
```

```
## [1] 1077 1123
```

```
## diet pill is given to 9 subjects over six weeks. The average difference in weight (follow up - baseline)
n = 9
mn_dif = 2
t = .95
(y_d <- round(mn_dif*sqrt(n) / qt(.975, df = 8), 2))
```

```
## [1] 2.6
```

Two sample t-test

```
##Running a two-sample t-test
data(ChickWeight); library(reshape2)
wideCW <- dcast(ChickWeight, Diet + Chick ~ Time, value.var = "weight") ##dcasting the long-form of the data to wide
names(wideCW)[- (1:2)] <- paste( "time", names(wideCW)[- (1:2)], sep= "") ##renames columns after the first two
library(dplyr)
```

```
##
```

```
## Attaching package: 'dplyr'
```

```
## The following objects are masked from 'package:Hmisc':
```

```
##
```

```
##      src, summarize
```

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

```
##
```

```
##      select
```

```
## The following objects are masked from 'package:stats':
```

```
##
```

```
##      filter, lag
```

```
## The following objects are masked from 'package:base':
```

```
##
```

```
##      intersect, setdiff, setequal, union
```

```
wideCW <- mutate(wideCW, gain = time21-time0) ##adds a column titled gain, where it subtracts or takes the difference between time21 and time0
```

```
wideCW_14 <- subset(wideCW, Diet %in% c(1,4))
```

```
t.test(gain~Diet, paired = FALSE, var.equal = TRUE, data = wideCW_14)
```

```
##
```

```
## Two Sample t-test
```

```
##
```

```
## data: gain by Diet
```

```
## t = -2.7252, df = 23, p-value = 0.01207
```

```
## alternative hypothesis: true difference in means between group 1 and group 4 is not equal to 0
```

```
## 95 percent confidence interval:
```

```
## -108.14679 -14.81154
```

```
## sample estimates:
```

```
## mean in group 1 mean in group 4
```

```
##      136.1875      197.6667
```

P-Values

```
pt(2.5, 15, lower.tail = FALSE) ##the probability of getting a T-statistic of 2.5 (first element of th

## [1] 0.0122529
```

Least Squares (Regression)

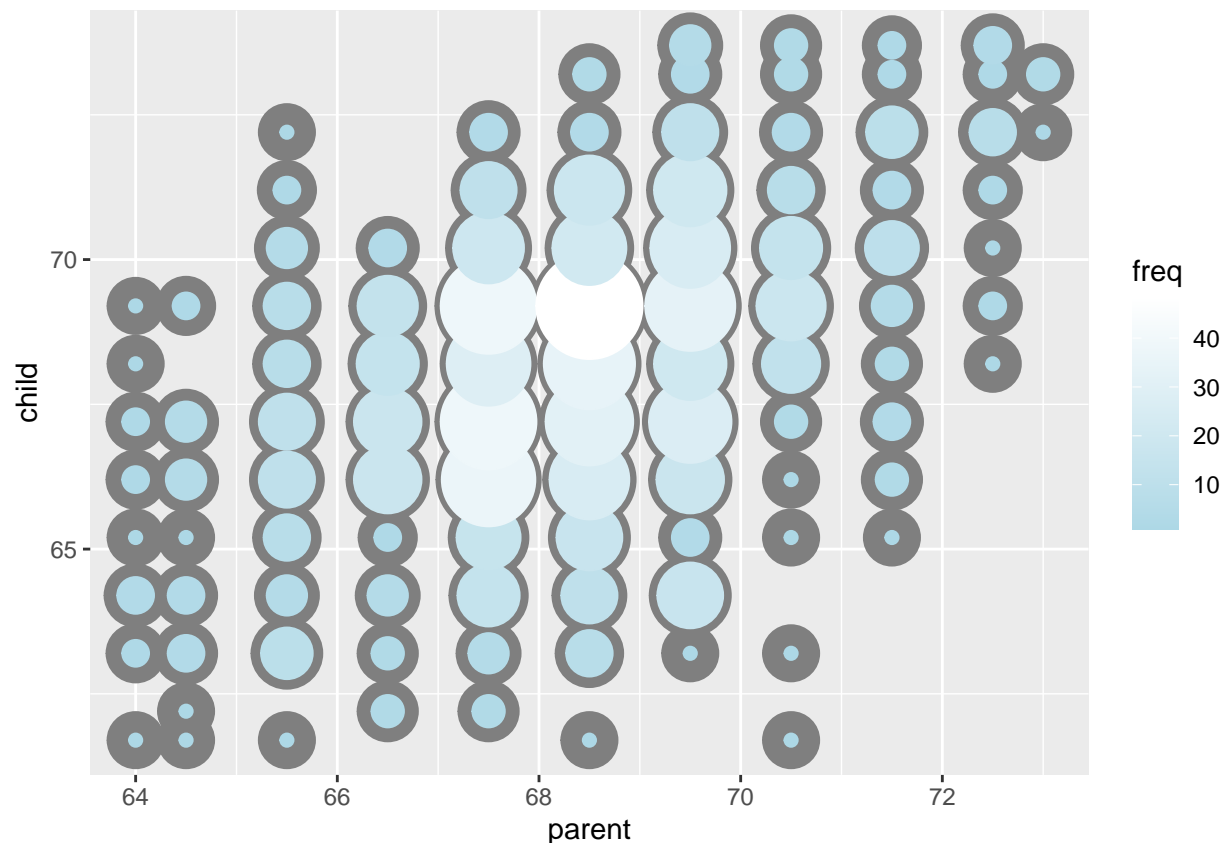
Sum[i=1 to n] of $[Y_i - (B_0 + B_1 X_i)]^2$ Least squares best fit is the sum of i to the n of Y_i which is the i'th sample data y-coordinate minus the intercept plus the slope times the i'th sample data x-coordinate, squared. B_0 = intercept of y-axis B_1 = slope $Y_i = y$ of sample i $X_i = x$ of sample i Think of it as subtracting the distance between sample's y-coordinate from the best fit, which is $y - (ax + b)$ which is a linear regression of the best fit, or $Y_i - Y_{fit}$. Then square it.

```
library(UsingR)
library(ggplot2)

freqData <- as.data.frame(table(galton$child, galton$parent))
names(freqData) <- c("child", "parent", "freq")
freqData$child <- as.numeric(as.character(freqData$child))
freqData$parent <- as.numeric(as.character(freqData$parent))
g <- ggplot(filter(freqData, freq > 0), aes(x = parent, y = child))
g <- g + scale_size(range = c(2, 20), guide = "none" )
g <- g + geom_point(colour="grey50", aes(size = freq+5, show_guide = FALSE))
```

```
## Warning: Ignoring unknown aesthetics: show_guide
```

```
g <- g + geom_point(aes(colour=freq, size = freq-5))
g <- g + scale_colour_gradient(low = "lightblue", high="white")
g
```

```
###-----Linear Least Squares calculated
y <- galton$child
x <- galton$parent
beta1 <- cor(y, x) * sd(y) / sd(x) ##The slope, which is the correlation between y and x times the st.
beta0 <- mean(y) - beta1 * mean(x) ##calculates the y-intercept of the best fit.
rbind(c(beta0, beta1), coef(lm(y ~ x))) ##lm stands for linear model. coef takes the output and gives u

##      (Intercept)          x
## [1,]    23.94153  0.6462906
## [2,]    23.94153  0.6462906

##This example gives you the same output, because one is the manual calculation, while the other is done

beta1 <- cor(y, x) * sd(x) / sd(y) ##if we swapped the predictor
beta0 <- mean(x) - beta1 * mean(y)
rbind(c(beta0, beta1), coef(lm(x ~ y))) ##shows us the slope is the same

##      (Intercept)          y
## [1,]    46.13535  0.3256475
## [2,]    46.13535  0.3256475

yc <- y - mean(y)
xc <- x - mean(x)
beta1 <- sum(yc * xc) / sum(xc ^ 2)
c(beta1, coef(lm(y ~ x))[2]) ##returns the same slope
```

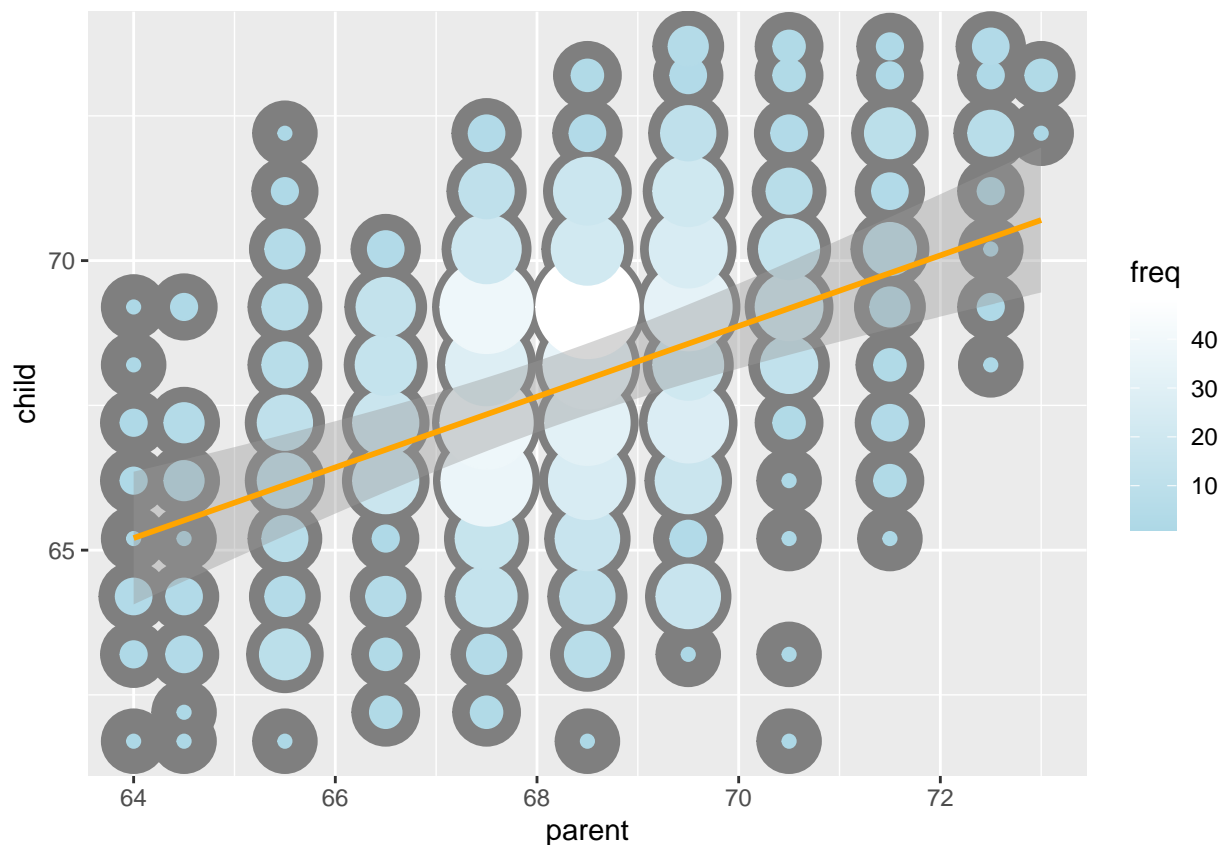
```
##                                x
## 0.6462906 0.6462906
```

```
###simplified
```

```
g <- ggplot(filter(freqData, freq > 0), aes(x = parent, y = child))
g <- g + scale_size(range = c(2, 20), guide = "none" )
g <- g + geom_point(colour="grey50", aes(size = freq+5, show_guide = FALSE))
```

```
## Warning: Ignoring unknown aesthetics: show_guide
```

```
g <- g + geom_point(aes(colour=freq, size = freq-10))
g <- g + scale_colour_gradient(low = "lightblue", high="white")
g <- g + geom_smooth(method="lm", col ="orange", formula=y~x) ##for fitting the linear best fit to a pl
g
```



Linear Regression for Prediction

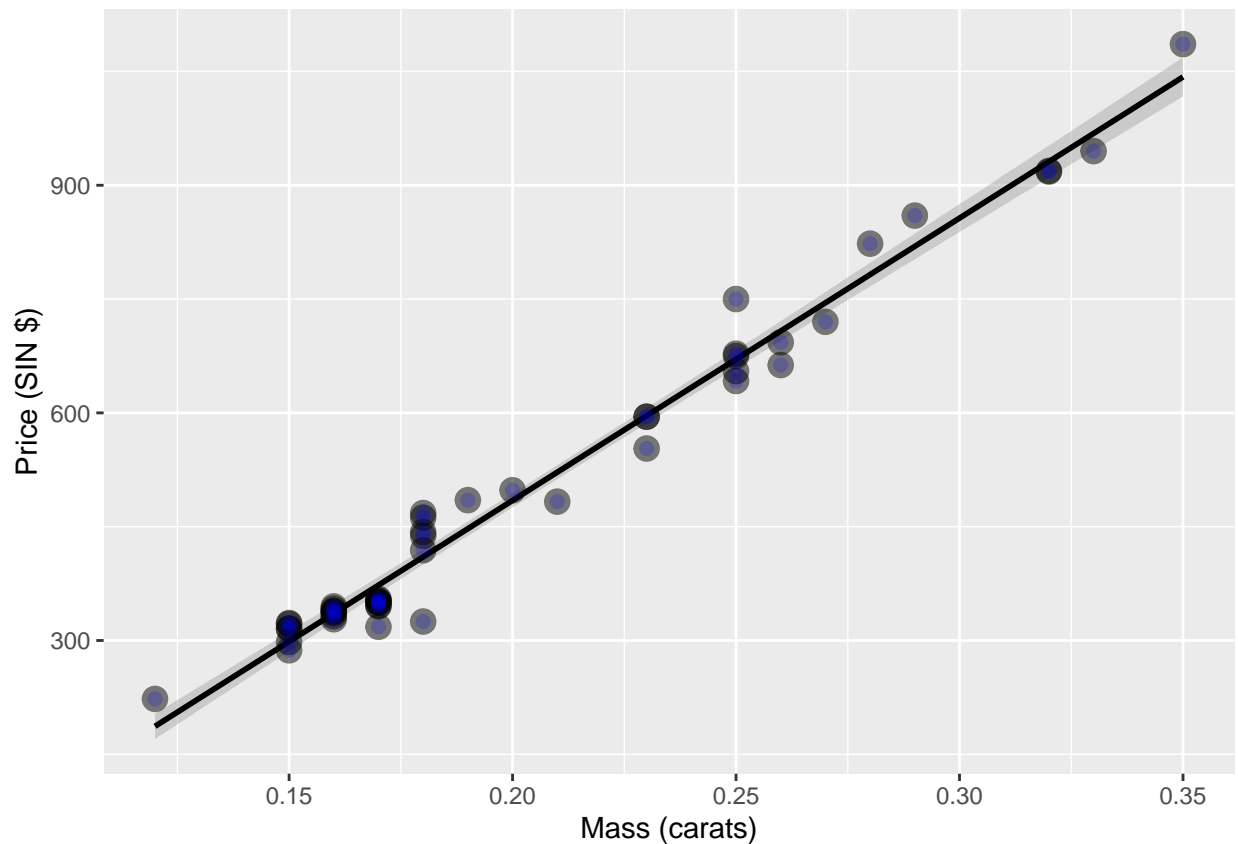
```
library(UsingR)
data(diamond)
library(ggplot2)
g = ggplot(diamond, aes(x = carat, y = price))
g = g + xlab("Mass (carats)")
```

```

g = g + ylab("Price (SIN $)")
g = g + geom_smooth(method = "lm", colour = "black")
g = g + geom_point(size = 4, colour = "black", alpha=0.5)
g = g + geom_point(size = 2, colour = "blue", alpha=0.2)
g

```

```
## 'geom_smooth()' using formula 'y ~ x'
```



```

fit <- lm(price~ carat, data = diamond)  ## prints coefficient Bo and B1, slope and intercept
coef(fit)

```

```

## (Intercept)      carat
##  -259.6259    3721.0249

```

```
summary(fit) ##gives you the complete statistic
```

```

##
## Call:
## lm(formula = price ~ carat, data = diamond)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -85.159  -21.448   -0.869   18.972   79.370

```

```
##
## Coefficients:
##           Estimate Std. Error t value Pr(>|t|)
## (Intercept)  -259.63      17.32  -14.99  <2e-16 ***
## carat        3721.02      81.79   45.50  <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 31.84 on 46 degrees of freedom
## Multiple R-squared:  0.9783, Adjusted R-squared:  0.9778
## F-statistic: 2070 on 1 and 46 DF,  p-value: < 2.2e-16
```

```
fit2 <- lm(price ~ I(carat - mean(carat)), data=diamond) ##how you mean center your predictor, carat. I
coef(fit2)
```

```
##           (Intercept) I(carat - mean(carat))
##           500.0833      3721.0249
```

```
fit3 <- lm(price ~ I(carat * 10), data=diamond) ##shortcut where fit3 produces the price change for 1/10
coef(fit3)
```

```
##      (Intercept) I(carat * 10)
##      -259.6259      372.1025
```

```
newx <- c(0.16, 0.27, 0.34) ## an example of a list of carat sizes for diamonds, and we want to predict
predict(fit, newdata = data.frame(carat = newx)) ##taking original fit data, and calling to the newx c
```

```
##           1           2           3
## 335.7381 745.0508 1005.5225
```

Residuals

```
data(diamond)
y <- diamond$price; x <- diamond$carat; n <- length(y)
fit <- lm(y ~ x)
e <- resid(fit) ##easiest way to calculate residuals
yhat <- predict(fit)
max(abs(e - (y - yhat)))
```

```
## [1] 9.485746e-13
```

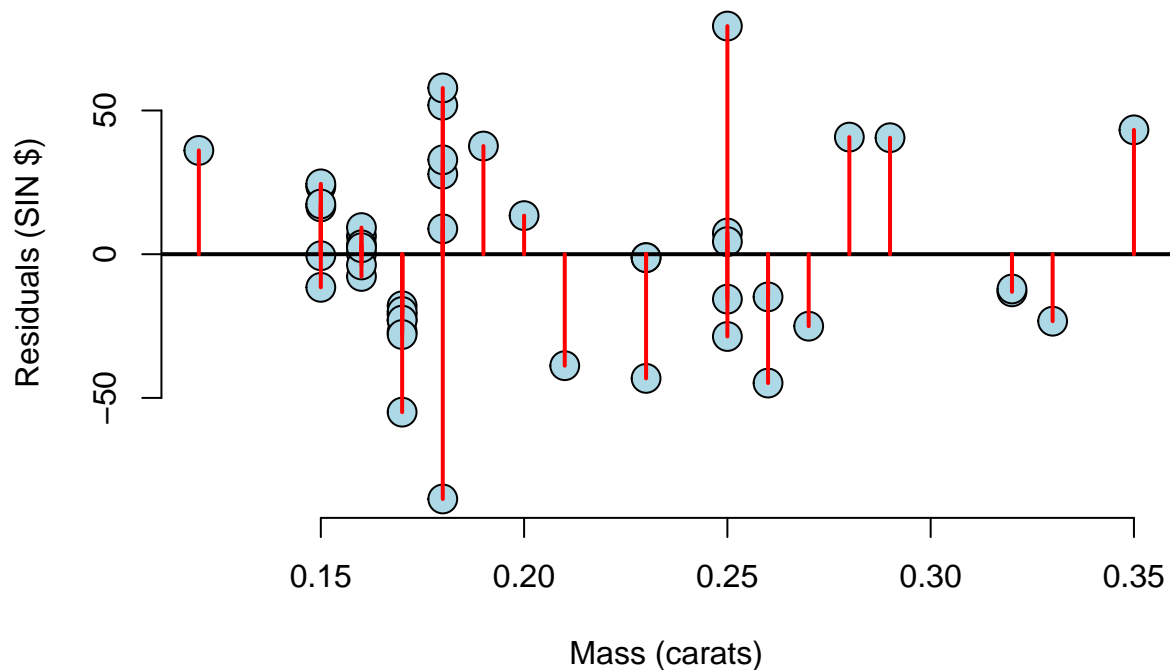
```
max(abs(e - (y - coef(fit)[1] - coef(fit)[2] * x))) ##will return the same as resid()
```

```
## [1] 9.485746e-13
```

```

plot(x, e,
     xlab = "Mass (carats)",
     ylab = "Residuals (SIN $)",
     bg = "lightblue",
     col = "black", cex = 2, pch = 21, frame = FALSE)
abline(h = 0, lwd = 2)
for (i in 1 : n)
  lines(c(x[i], x[i]), c(e[i], 0), col = "red" , lwd = 2)

```



```

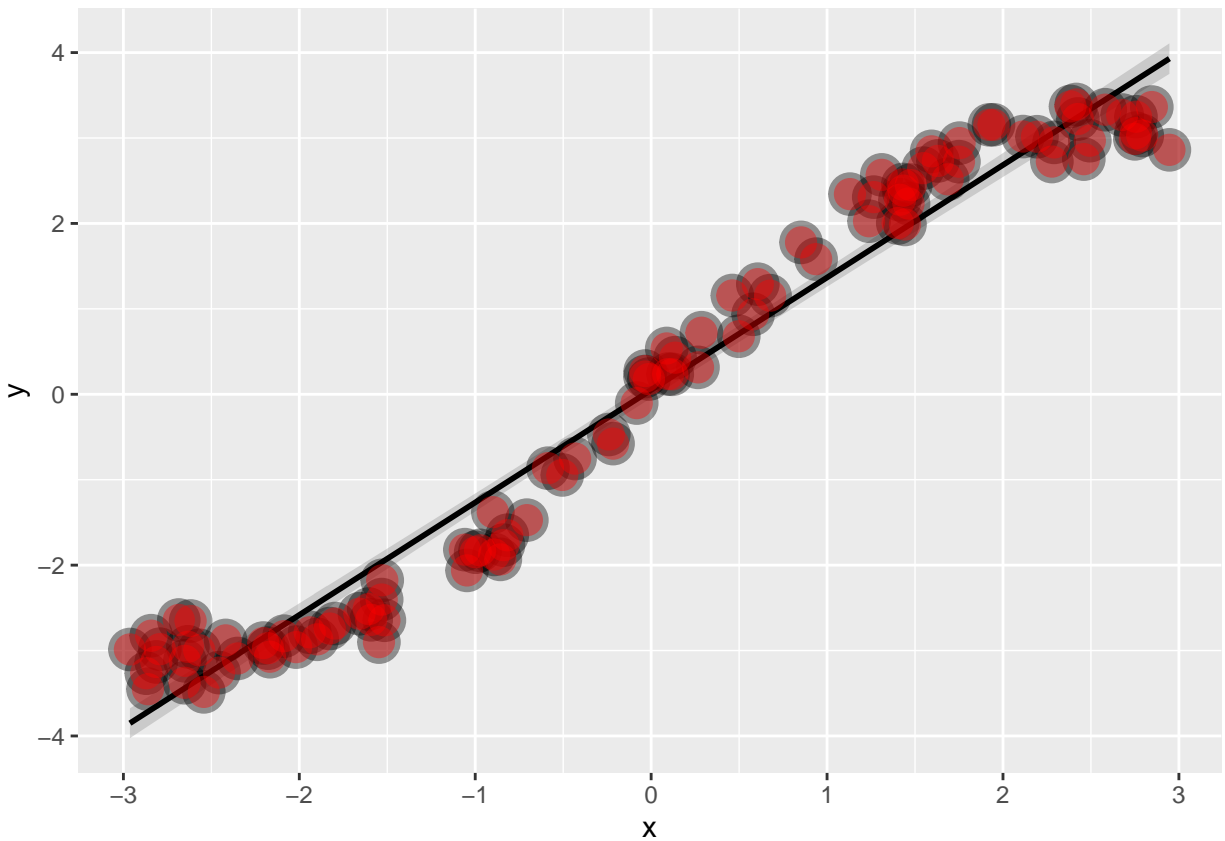
## ----- Example of non-linear data and residuals
x = runif(100, -3, 3); y = x + sin(x) + rnorm(100, sd = .2);
library(ggplot2)
library(RColorBrewer)
g = ggplot(data.frame(x = x, y = y), aes(x = x, y = y))
g = g + geom_smooth(method = "lm", colour = "black")
g = g + geom_point(size = 7, colour = "black", alpha = 0.4)
g = g + geom_point(size = 5, colour = "red", alpha = 0.4)
g

```

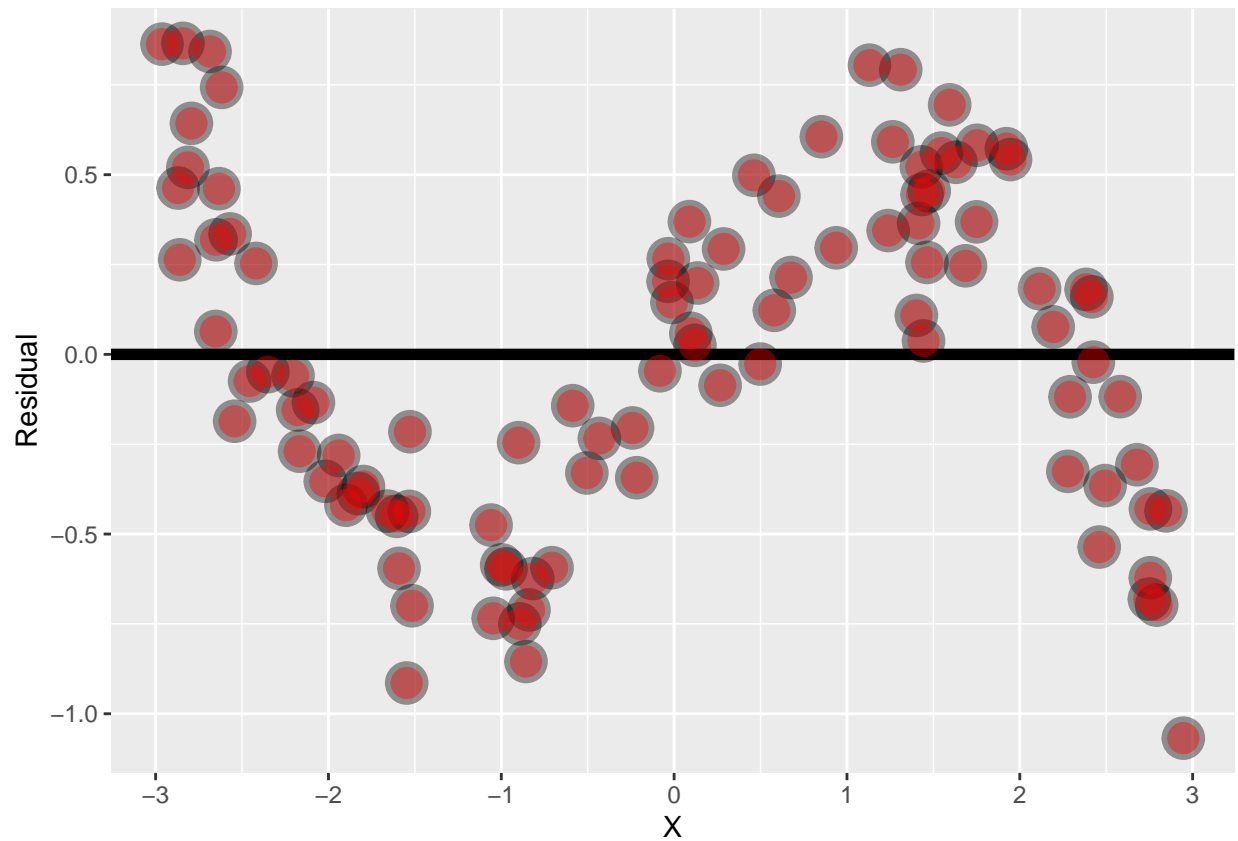
```

## 'geom_smooth()' using formula 'y ~ x'

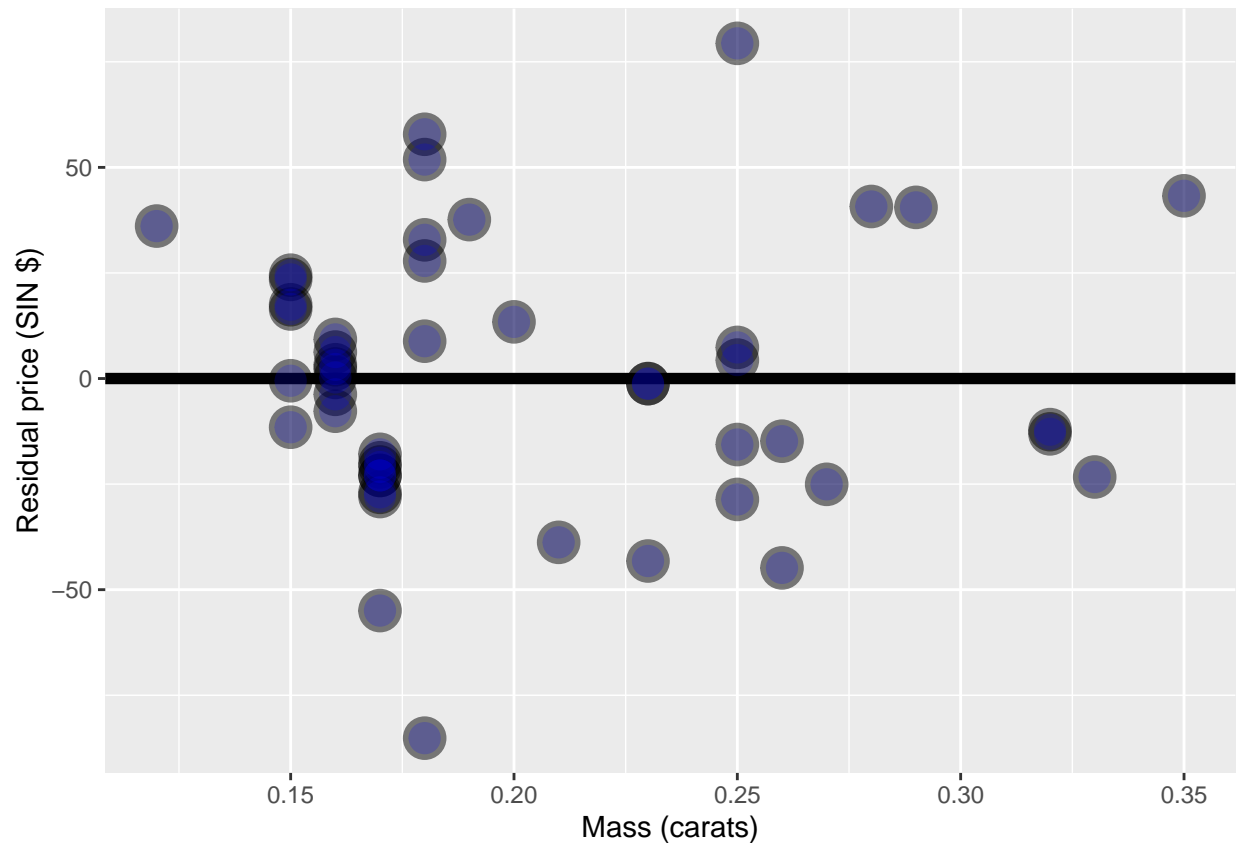
```



```
## -----
g = ggplot(data.frame(x = x, y = resid(lm(y ~ x))), ##How to plot residuals
           aes(x = x, y = y))
g = g + geom_hline(yintercept = 0, size = 2);
g = g + geom_point(size = 7, colour = "black", alpha = 0.4)
g = g + geom_point(size = 5, colour = "red", alpha = 0.4)
g = g + xlab("X") + ylab("Residual")
g
```



```
##----- Calculating the residual and plotting for diamond data
diamond$e <- resid(lm(price ~ carat, data = diamond)) ##adding a new column into data.frame for residu
g = ggplot(diamond, aes(x = carat, y = e))
g = g + xlab("Mass (carats)")
g = g + ylab("Residual price (SIN $)")
g = g + geom_hline(yintercept = 0, size = 2)
g = g + geom_point(size = 7, colour = "black", alpha=0.5)
g = g + geom_point(size = 5, colour = "blue", alpha=0.2)
g
```



Predictions

```
fit <- lm(mpg~wt, data= mtcars) ## creating a linear model of mtcars dataset, where weight is the predi
predict(fit, wt=3000) ##using the model above, and predicting the mpg for all cars weighing 3000lbs.
```

| | | | | |
|----|-------------------|------------------|--------------------|---------------------|
| ## | Mazda RX4 | Mazda RX4 Wag | Datsun 710 | Hornet 4 Drive |
| ## | 23.282611 | 21.919770 | 24.885952 | 20.102650 |
| ## | Hornet Sportabout | Valiant | Duster 360 | Merc 240D |
| ## | 18.900144 | 18.793255 | 18.205363 | 20.236262 |
| ## | Merc 230 | Merc 280 | Merc 280C | Merc 450SE |
| ## | 20.450041 | 18.900144 | 18.900144 | 15.533127 |
| ## | Merc 450SL | Merc 450SLC | Cadillac Fleetwood | Lincoln Continental |
| ## | 17.350247 | 17.083024 | 9.226650 | 8.296712 |
| ## | Chrysler Imperial | Fiat 128 | Honda Civic | Toyota Corolla |
| ## | 8.718926 | 25.527289 | 28.653805 | 27.478021 |
| ## | Toyota Corona | Dodge Challenger | AMC Javelin | Camaro Z28 |
| ## | 24.111004 | 18.472586 | 18.926866 | 16.762355 |
| ## | Pontiac Firebird | Fiat X1-9 | Porsche 914-2 | Lotus Europa |
| ## | 16.735633 | 26.943574 | 25.847957 | 29.198941 |
| ## | Ford Pantera L | Ferrari Dino | Maserati Bora | Volvo 142E |
| ## | 20.343151 | 22.480940 | 18.205363 | 22.427495 |

Multivariate Regressions

Shapiro Wilks test for Normal distribution

```
data <- rnorm(100)

#perform Shapiro-Wilk test for normality
shapiro.test(data) ## where we reject normality when p < 0.05, otherwise we accept that the test data .

##
## Shapiro-Wilk normality test
##
## data: data
## W = 0.9854, p-value = 0.3395
```

Machine Learning

question - input data - features - algorithm - parameters - evaluation

Prediction has accuracy tradeoffs

In-sample error: the error rate you get on the same data set you used to build your predictor. Sometimes called re-substitution error. Out of sample error: the error rate you get on a new data set, sometimes called generalization error. Out of sample error is what we should care about In sample error is always less than out of sample error The reason is over-fitting. True positive vs. false positive, true negative vs. false negative Sensitivity - probability that the test positively predicted, and that the prediction was right Specificity- probability that the test was negative, and so too was the outcome Receiver Operating Characteristic (ROC) - Area under the curve is the measure of goodness of fit (model/prediction to data/outcome). AUC of 0.5 is random guessing. AUC > 0.8 is a good model. Cross-validation. 1.) use the training set 2.) split the training set into a test set and a smaller training set 3.) Build a model on the training set 4.) Evaluate the model on the test set 5.) repeat and average the estimated errors k-fold cross-validation is popular technique.

```
library(kernlab)
```

```
##
## Attaching package: 'kernlab'

## The following object is masked from 'package:ggplot2':
##
## alpha
```

```
data(spam)
head(spam)
```

```
## make address all num3d our over remove internet order mail receive will
## 1 0.00 0.64 0.64 0 0.32 0.00 0.00 0.00 0.00 0.00 0.00 0.64
## 2 0.21 0.28 0.50 0 0.14 0.28 0.21 0.07 0.00 0.94 0.21 0.79
## 3 0.06 0.00 0.71 0 1.23 0.19 0.19 0.12 0.64 0.25 0.38 0.45
## 4 0.00 0.00 0.00 0 0.63 0.00 0.31 0.63 0.31 0.63 0.31 0.31
## 5 0.00 0.00 0.00 0 0.63 0.00 0.31 0.63 0.31 0.63 0.31 0.31
```

```

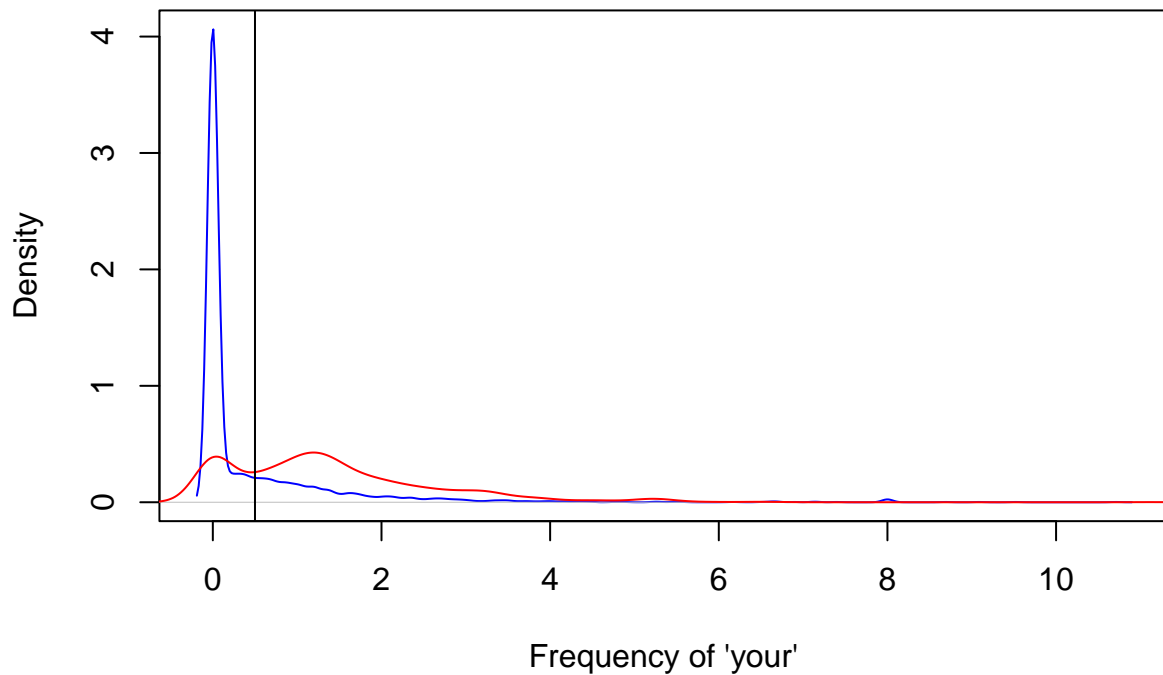
## 6 0.00    0.00 0.00    0 1.85 0.00    0.00    1.85 0.00 0.00    0.00 0.00
##   people report addresses free business email  you credit your font num000
## 1  0.00    0.00    0.00 0.32    0.00 1.29 1.93    0.00 0.96    0  0.00
## 2  0.65    0.21    0.14 0.14    0.07 0.28 3.47    0.00 1.59    0  0.43
## 3  0.12    0.00    1.75 0.06    0.06 1.03 1.36    0.32 0.51    0  1.16
## 4  0.31    0.00    0.00 0.31    0.00 0.00 3.18    0.00 0.31    0  0.00
## 5  0.31    0.00    0.00 0.31    0.00 0.00 3.18    0.00 0.31    0  0.00
## 6  0.00    0.00    0.00 0.00    0.00 0.00 0.00    0.00 0.00    0  0.00
##   money hp hpl george num650 lab labs telnet num857 data num415 num85
## 1  0.00 0  0    0    0  0  0  0    0    0    0    0    0
## 2  0.43 0  0    0    0  0  0  0    0    0    0    0    0
## 3  0.06 0  0    0    0  0  0  0    0    0    0    0    0
## 4  0.00 0  0    0    0  0  0  0    0    0    0    0    0
## 5  0.00 0  0    0    0  0  0  0    0    0    0    0    0
## 6  0.00 0  0    0    0  0  0  0    0    0    0    0    0
##   technology num1999 parts pm direct cs meeting original project re edu
## 1      0    0.00    0  0  0.00 0    0    0.00    0 0.00 0.00
## 2      0    0.07    0  0  0.00 0    0    0.00    0 0.00 0.00
## 3      0    0.00    0  0  0.06 0    0    0.12    0 0.06 0.06
## 4      0    0.00    0  0  0.00 0    0    0.00    0 0.00 0.00
## 5      0    0.00    0  0  0.00 0    0    0.00    0 0.00 0.00
## 6      0    0.00    0  0  0.00 0    0    0.00    0 0.00 0.00
##   table conference charSemicolon charRoundbracket charSquarebracket
## 1      0          0          0.00          0.000          0
## 2      0          0          0.00          0.132          0
## 3      0          0          0.01          0.143          0
## 4      0          0          0.00          0.137          0
## 5      0          0          0.00          0.135          0
## 6      0          0          0.00          0.223          0
##   charExclamation charDollar charHash capitalAve capitalLong capitalTotal type
## 1      0.778      0.000      0.000      3.756        61        278 spam
## 2      0.372      0.180      0.048      5.114       101       1028 spam
## 3      0.276      0.184      0.010      9.821       485       2259 spam
## 4      0.137      0.000      0.000      3.537        40        191 spam
## 5      0.135      0.000      0.000      3.537        40        191 spam
## 6      0.000      0.000      0.000      3.000        15         54 spam

```

```

plot(density(spam$your[spam$type=="nonspam"]),
     col="blue",main="",xlab="Frequency of 'your'")
lines(density(spam$your[spam$type=="spam"]),col="red")
abline(v=0.5,col="black")

```



##We want to find a value C, which is the cutoff point between spam frequencies, and non-spam frequencies

```
prediction <- ifelse(spam$your > 0.5,"spam","nonspam")
table(prediction,spam$type)/length(spam$type)
```

```
##
## prediction  nonspam      spam
## nonspam  0.4590306 0.1017170
## spam     0.1469246 0.2923278
```

###Caret for machine learning package

obj Class Package Predict Function Syntax lda MASS Predict(obj) (no options needed) glm stats predict(obj, type = "response") bgm gbm predict(obj, type = "response", n.trees) mda mda predict(obj, type = "posterior") rpart rpart predict(obj, type = "prob") Weka RWeka predict(obj, type = "probability") LogitBoost caTools predict(obj, type = "raw", nIter)

** Different types of predictors have different object class syntax requirements.

```
library(caret)
```

```
##
## Attaching package: 'caret'

## The following object is masked from 'package:survival':
##
## cluster
```

```

library(kernlab) ## to get spam dataset

data(spam)
inTrain <- createDataPartition(y=spam$type, p= 0.75, list = FALSE) ## data is partitioned by spam type,
training <- spam[inTrain,] ##subsetting the output of the partition function by spam
testing <- spam[-inTrain,] ##subsetting the output of the partition function by not spam
dim(training) ##shows the dimensions of the dataframe

## [1] 3451 58

set.seed(11111)
modelFit <- train(type ~., data= training, method = "glm") ##training a model, specifically a glm model.

## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
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## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred

```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
modelFit
```

```
## Generalized Linear Model
##
## 3451 samples
## 57 predictor
## 2 classes: 'nonspam', 'spam'
##
## No pre-processing
## Resampling: Bootstrapped (25 reps)
## Summary of sample sizes: 3451, 3451, 3451, 3451, 3451, ...
## Resampling results:
##
## Accuracy Kappa
## 0.919168 0.8302427
```

```
modelFit$finalModel ##will return the fitted values for all of the other columns of data. The higher t
```

```
##
## Call: NULL
##
## Coefficients:
## (Intercept) make address all
## -1.671e+00 -2.453e-01 -1.538e-01 8.485e-02
## num3d our over remove
## 1.661e+00 5.668e-01 6.307e-01 2.443e+00
## internet order mail receive
## 5.826e-01 2.844e-01 5.552e-02 -5.869e-01
## will people report addresses
## -1.568e-01 -1.696e-01 9.178e-02 9.142e-01
## free business email you
## 9.522e-01 9.009e-01 6.647e-02 1.119e-01
## credit your font num000
## 9.564e-01 2.165e-01 1.969e-01 2.149e+00
## money hp hpl george
## 7.201e-01 -1.595e+00 -1.439e+00 -4.871e+00
## num650 lab labs telnet
## 5.969e-01 -4.523e+00 -9.483e-03 -6.249e+00
## num857 data num415 num85
## 2.468e+00 -7.011e-01 -7.162e+00 -2.197e+00
```

```
##      technology          num1999          parts          pm
##      7.988e-01          4.972e-02          -6.271e-01          -9.098e-01
##      direct          cs          meeting          original
##      4.704e-01          -3.570e+02          -2.181e+00          -2.297e+00
##      project          re          edu          table
##      -1.505e+00          -8.806e-01          -1.683e+00          -2.575e+00
##      conference          charSemicolon          charRoundbracket          charSquarebracket
##      -3.942e+00          -1.278e+00          -1.816e-01          -8.319e-01
##      charExclamation          charDollar          charHash          capitalAve
##      2.443e-01          6.219e+00          1.576e+00          8.079e-02
##      capitalLong          capitalTotal
##      1.132e-02          5.696e-04
##
## Degrees of Freedom: 3450 Total (i.e. Null); 3393 Residual
## Null Deviance: 4628
## Residual Deviance: 1369 AIC: 1485
```

```
predictions <- predict(modelFit, newdata = testing) ##how we can predict on new samples, here using the
predictions
```

```
##      [1] spam      spam      spam      spam      spam      spam      spam      nonspam nonspam nonspam
##      [10] spam      nonspam spam      spam      spam      spam      spam      spam      spam      spam
##      [19] spam      spam      spam      spam      spam      spam      nonspam spam      spam      spam
##      [28] spam      spam      nonspam spam      spam      spam      nonspam spam      spam      nonspam
##      [37] spam      spam      spam      spam      spam      spam      spam      spam      spam      spam
##      [46] spam      nonspam spam      spam      spam      spam      spam      nonspam spam      spam
##      [55] spam      spam      nonspam spam      spam      spam      spam      spam      spam      spam
##      [64] spam      spam      spam      spam      spam      spam      nonspam spam      spam      spam
##      [73] spam      nonspam spam      spam      spam      spam      spam      spam      spam      spam
##      [82] nonspam spam      nonspam nonspam spam      spam      spam      spam      spam      spam
##      [91] spam      spam      spam      spam      spam      spam      spam      spam      spam      spam
##      [100] nonspam spam      nonspam spam      spam      spam      spam      spam      spam      spam
##      [109] spam      nonspam spam      spam      spam      spam      spam      spam      spam      nonspam
##      [118] spam      spam      spam      spam      spam      spam      spam      spam      spam      spam
##      [127] spam      spam      spam      spam      spam      spam      spam      spam      spam      spam
##      [136] spam      nonspam spam      spam      spam      spam      spam      nonspam spam      spam
##      [145] spam      spam      spam      spam      spam      spam      spam      spam      spam      spam
##      [154] nonspam spam      spam      spam      nonspam spam      spam      spam      spam      spam
##      [163] spam      spam      spam      spam      spam      spam      spam      spam      spam      spam
##      [172] spam      spam      spam      nonspam spam      spam      spam      nonspam spam      spam
##      [181] spam      spam      spam      spam      spam      spam      spam      spam      nonspam spam
##      [190] spam      spam      spam      spam      spam      spam      spam      spam      spam      spam
##      [199] spam      spam      spam      spam      spam      spam      spam      spam      spam      spam
##      [208] spam      spam      nonspam spam      spam      spam      spam      spam      spam      spam
##      [217] spam      spam      spam      spam      spam      spam      spam      spam      spam      spam
##      [226] spam      spam      spam      spam      spam      spam      spam      spam      spam      spam
##      [235] spam      spam      spam      spam      spam      spam      spam      spam      spam      spam
##      [244] spam      nonspam spam      spam      spam      spam      spam      spam      spam      spam
##      [253] spam      spam      spam      spam      spam      spam      nonspam spam      spam      spam
##      [262] spam      spam      spam      spam      nonspam spam      spam      spam      spam      spam
##      [271] spam      spam      spam      spam      spam      spam      spam      spam      spam      spam
##      [280] spam      spam      spam      spam      spam      spam      spam      spam      spam      spam
##      [289] spam      spam      spam      spam      spam      nonspam spam      spam      spam      spam
##      [298] spam      spam      spam      spam      spam      nonspam spam      spam      spam      spam
```

[illegible]

```

## [793] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [802] nonspam spam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [811] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [820] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [829] nonspam spam spam nonspam spam nonspam nonspam nonspam nonspam nonspam
## [838] nonspam nonspam nonspam spam nonspam nonspam nonspam nonspam nonspam nonspam
## [847] nonspam nonspam spam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [856] nonspam spam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [865] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam spam
## [874] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [883] nonspam nonspam spam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [892] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [901] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [910] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [919] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [928] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam spam nonspam
## [937] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam spam
## [946] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [955] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [964] nonspam nonspam spam spam nonspam nonspam nonspam nonspam nonspam nonspam
## [973] nonspam spam nonspam nonspam nonspam nonspam nonspam nonspam spam nonspam
## [982] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [991] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [1000] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [1009] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [1018] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [1027] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [1036] nonspam nonspam spam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [1045] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [1054] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [1063] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [1072] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam spam nonspam
## [1081] nonspam nonspam nonspam nonspam spam nonspam nonspam spam nonspam nonspam
## [1090] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [1099] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [1108] nonspam nonspam spam nonspam spam nonspam nonspam nonspam nonspam nonspam
## [1117] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [1126] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [1135] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## [1144] nonspam nonspam nonspam nonspam nonspam nonspam nonspam nonspam
## Levels: nonspam spam

```

```

confusionMatrix(predictions, testing$type) ##using the confusingMatrix argument to determine how well

```

```

## Confusion Matrix and Statistics
##
##           Reference
## Prediction nonspam spam
##   nonspam    664    52
##   spam       33   401
##
##           Accuracy : 0.9261
##           95% CI : (0.9094, 0.9405)
##   No Information Rate : 0.6061

```



```
##      P-Value [Acc > NIR] : < 2e-16
##
##              Kappa : 0.8441
##
## Mcnemar's Test P-Value : 0.05089
##
##      Sensitivity : 0.9527
##      Specificity : 0.8852
##      Pos Pred Value : 0.9274
##      Neg Pred Value : 0.9240
##      Prevalence : 0.6061
##      Detection Rate : 0.5774
##      Detection Prevalence : 0.6226
##      Balanced Accuracy : 0.9189
##
##      'Positive' Class : nonspam
##
```

Data Slicing with caret

```
library(caret)
library(kernlab) ## to get spam dataset

data(spam)
inTrain <- createDataPartition(y=spam$type, p= 0.75, list = FALSE) ## data is partitioned by spam type,
training <- spam[inTrain,] ##subsetting the output of the partition function by spam
testing <- spam[-inTrain,] ##subsetting the output of the partition function by not spam
dim(training)
```

```
## [1] 3451  58
```

```
set.seed(11111)
folds <- createFolds(y= spam$type, k=10, list= TRUE, returnTrain=TRUE) ##k is the number of folds, sma
sapply(folds, length)
```

```
## Fold01 Fold02 Fold03 Fold04 Fold05 Fold06 Fold07 Fold08 Fold09 Fold10
##   4141   4140   4141   4142   4142   4141   4140   4140   4141   4141
```

Plotting Predictors + Hmisc for cutting data.frames

```
library(ISLR)
library(ggplot2)
library(caret)
```

```
data(wage)
```

```
## Warning in data(wage): data set 'wage' not found
```

```
summary(Wage)
```

```
##          year          age          maritl          race
## Min.      :2003    Min.      :18.00    1. Never Married: 648    1. White:2480
## 1st Qu.:2004    1st Qu.:33.75    2. Married      :2074    2. Black: 293
## Median :2006    Median :42.00    3. Widowed      : 19    3. Asian: 190
## Mean      :2006    Mean      :42.41    4. Divorced     : 204    4. Other:  37
## 3rd Qu.:2008    3rd Qu.:51.00    5. Separated    :  55
## Max.      :2009    Max.      :80.00
##
##          education          region          jobclass
## 1. < HS Grad      :268    2. Middle Atlantic :3000    1. Industrial :1544
## 2. HS Grad        :971    1. New England   :  0    2. Information:1456
## 3. Some College   :650    3. East North Central:  0
## 4. College Grad   :685    4. West North Central:  0
## 5. Advanced Degree:426    5. South Atlantic    :  0
##                      6. East South Central:  0
##                      (Other)              :  0
##
##          health    health_ins    logwage          wage
## 1. <=Good      : 858    1. Yes:2083    Min.      :3.000    Min.      : 20.09
## 2. >=Very Good:2142    2. No : 917    1st Qu.:4.447    1st Qu.: 85.38
##                      Median :4.653    Median :104.92
##                      Mean      :4.654    Mean      :111.70
##                      3rd Qu.:4.857    3rd Qu.:128.68
##                      Max.      :5.763    Max.      :318.34
##
```

```
head(Wage, n=20)
```

```
##          year age          maritl          race          education          region
## 231655 2006  18 1. Never Married 1. White      1. < HS Grad 2. Middle Atlantic
## 86582  2004  24 1. Never Married 1. White      4. College Grad 2. Middle Atlantic
## 161300 2003  45      2. Married 1. White      3. Some College 2. Middle Atlantic
## 155159 2003  43      2. Married 3. Asian      4. College Grad 2. Middle Atlantic
## 11443  2005  50      4. Divorced 1. White      2. HS Grad 2. Middle Atlantic
## 376662 2008  54      2. Married 1. White      4. College Grad 2. Middle Atlantic
## 450601 2009  44      2. Married 4. Other      3. Some College 2. Middle Atlantic
## 377954 2008  30 1. Never Married 3. Asian      3. Some College 2. Middle Atlantic
## 228963 2006  41 1. Never Married 2. Black      3. Some College 2. Middle Atlantic
## 81404  2004  52      2. Married 1. White      2. HS Grad 2. Middle Atlantic
## 302778 2007  45      4. Divorced 1. White      3. Some College 2. Middle Atlantic
## 305706 2007  34      2. Married 1. White      2. HS Grad 2. Middle Atlantic
## 8690   2005  35 1. Never Married 1. White      2. HS Grad 2. Middle Atlantic
## 153561 2003  39      2. Married 1. White      4. College Grad 2. Middle Atlantic
## 449654 2009  54      2. Married 1. White      2. HS Grad 2. Middle Atlantic
## 447660 2009  51      2. Married 1. White      3. Some College 2. Middle Atlantic
## 160191 2003  37 1. Never Married 3. Asian      4. College Grad 2. Middle Atlantic
## 230312 2006  50      2. Married 1. White 5. Advanced Degree 2. Middle Atlantic
## 301585 2007  56      2. Married 1. White      4. College Grad 2. Middle Atlantic
## 153682 2003  37 1. Never Married 1. White      3. Some College 2. Middle Atlantic
##
##          jobclass    health health_ins    logwage          wage
## 231655 1. Industrial    1. <=Good    2. No 4.318063 75.04315
```

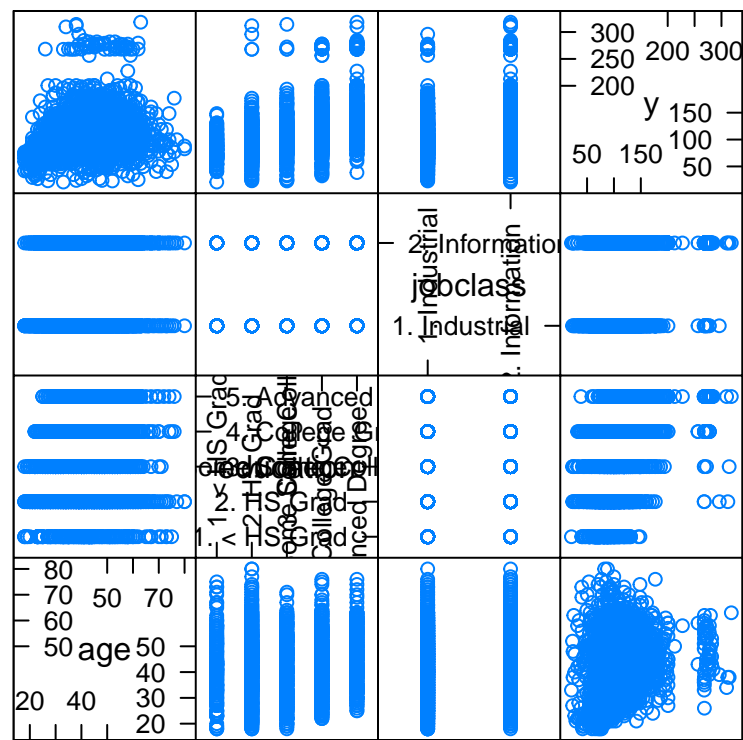
```
## 86582 2. Information 2. >=Very Good      2. No 4.255273  70.47602
## 161300 1. Industrial      1. <=Good      1. Yes 4.875061 130.98218
## 155159 2. Information 2. >=Very Good      1. Yes 5.041393 154.68529
## 11443 2. Information      1. <=Good      1. Yes 4.318063  75.04315
## 376662 2. Information 2. >=Very Good      1. Yes 4.845098 127.11574
## 450601 1. Industrial 2. >=Very Good      1. Yes 5.133021 169.52854
## 377954 2. Information      1. <=Good      1. Yes 4.716003 111.72085
## 228963 2. Information 2. >=Very Good      1. Yes 4.778151 118.88436
## 81404 2. Information 2. >=Very Good      1. Yes 4.857332 128.68049
## 302778 2. Information      1. <=Good      1. Yes 4.763428 117.14682
## 305706 1. Industrial 2. >=Very Good      2. No 4.397940  81.28325
## 8690 2. Information 2. >=Very Good      1. Yes 4.494155  89.49248
## 153561 1. Industrial 2. >=Very Good      1. Yes 4.903090 134.70538
## 449654 2. Information 2. >=Very Good      1. Yes 4.903090 134.70538
## 447660 1. Industrial 2. >=Very Good      1. Yes 4.505150  90.48191
## 160191 1. Industrial 2. >=Very Good      2. No 4.414973  82.67964
## 230312 2. Information 2. >=Very Good      2. No 5.360552 212.84235
## 301585 1. Industrial      1. <=Good      1. Yes 4.861026 129.15669
## 153682 1. Industrial 2. >=Very Good      1. Yes 4.591065  98.59934
```

```
inTrain <- createDataPartition(y=Wage$wage, p=0.7, list=FALSE)
training <- Wage[inTrain,]
testing <- Wage[-inTrain,]
dim(training); dim(testing) ##returns the number of rows and columns
```

```
## [1] 2102  11
```

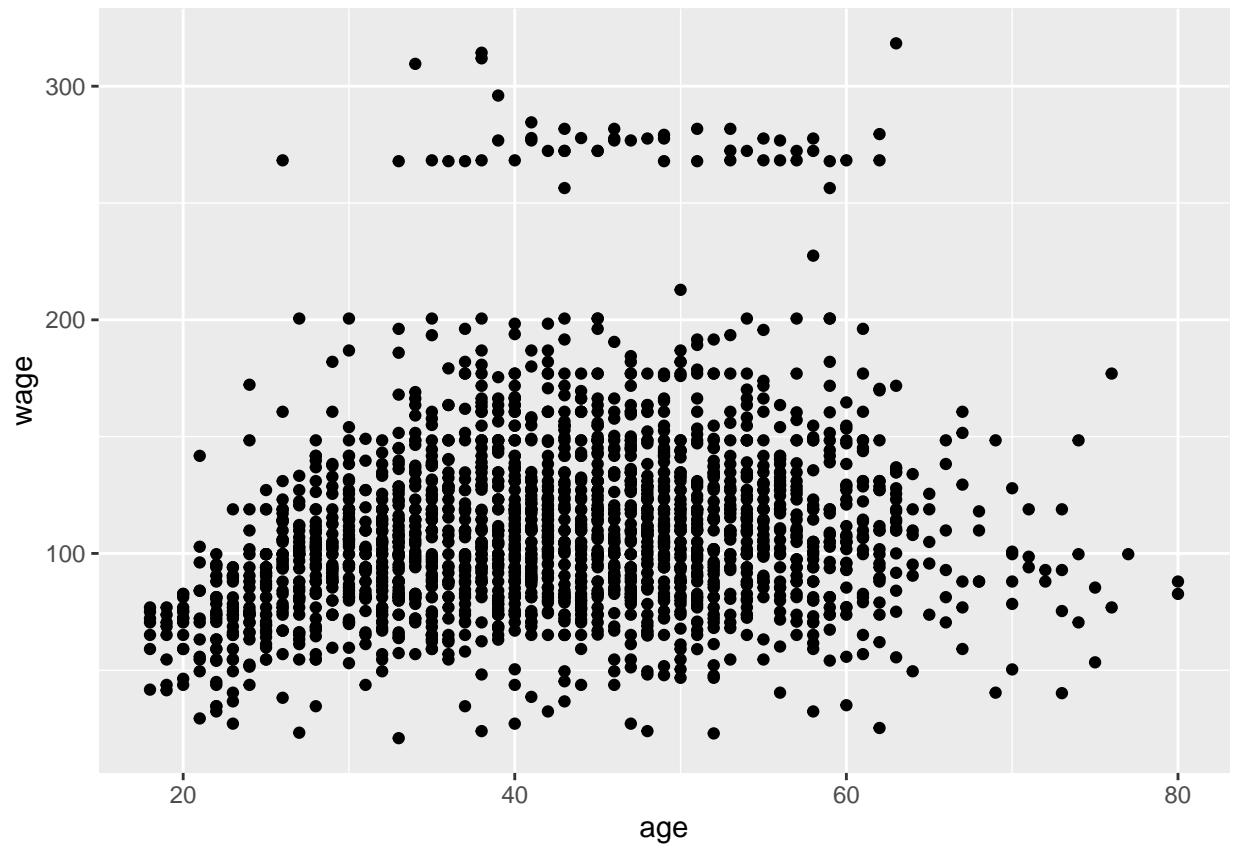
```
## [1] 898  11
```

```
featurePlot(x=training[, c("age", "education", "jobclass")], ##a plotting function that comes with care
            y= training$wage, ##this is your output of interest
            plot="pairs")
```

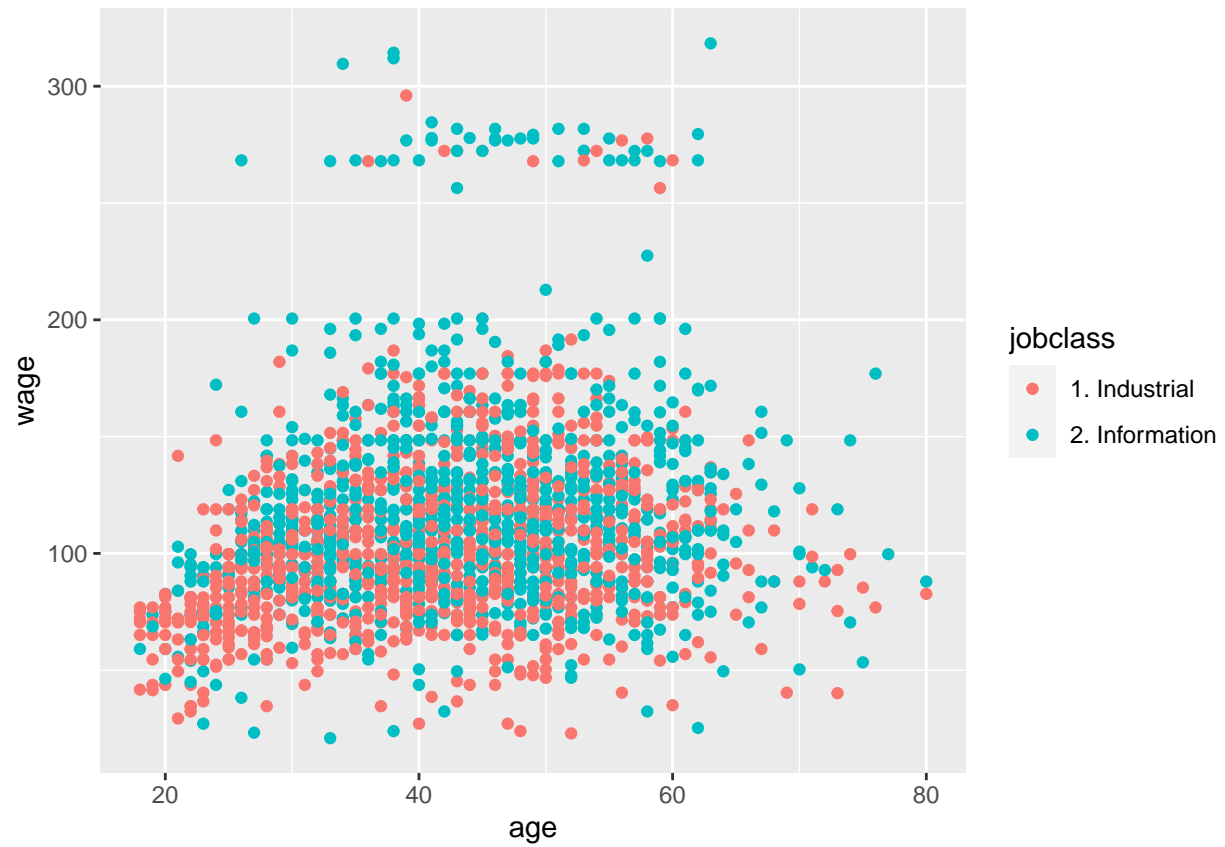


Scatter Plot Matrix

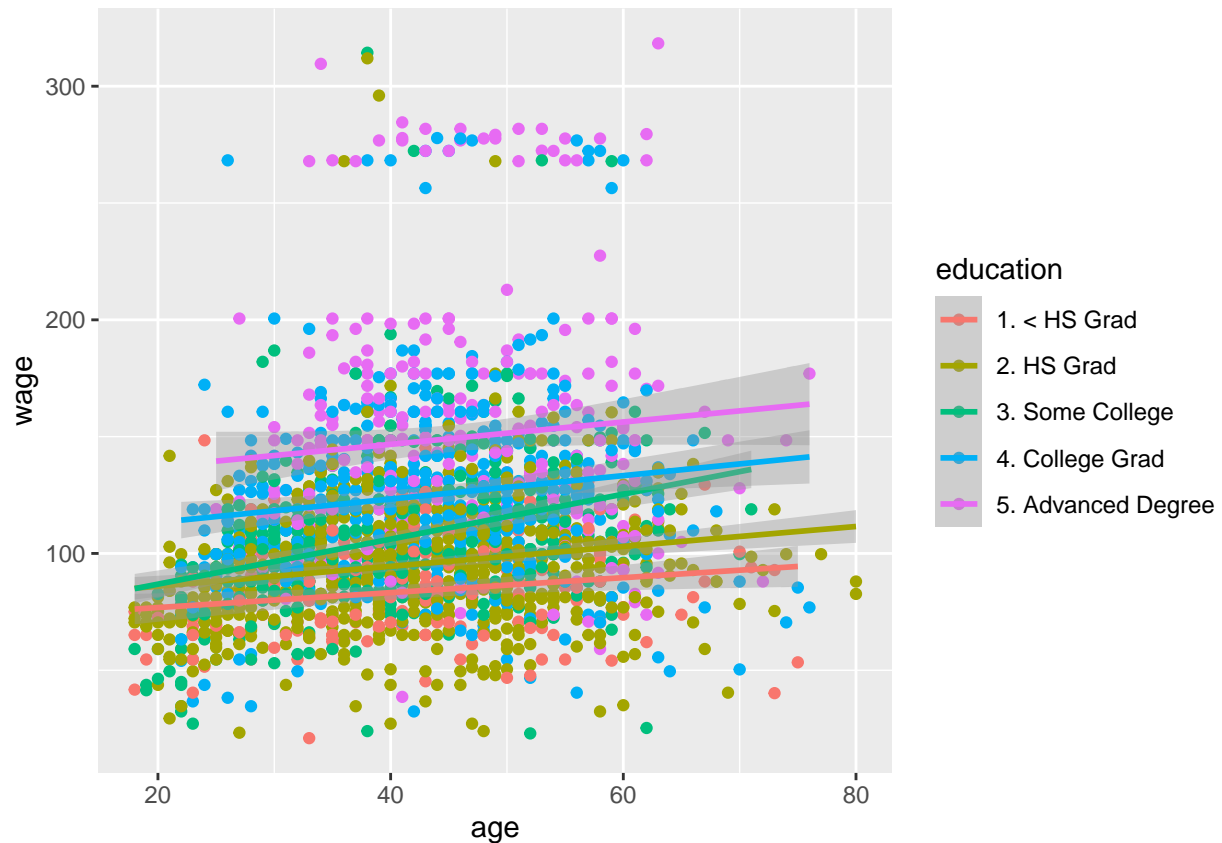
```
qplot(age, wage, data=training) ##here we see two distinct groups
```



```
qplot(age, wage, colour= jobclass, data= training)
```



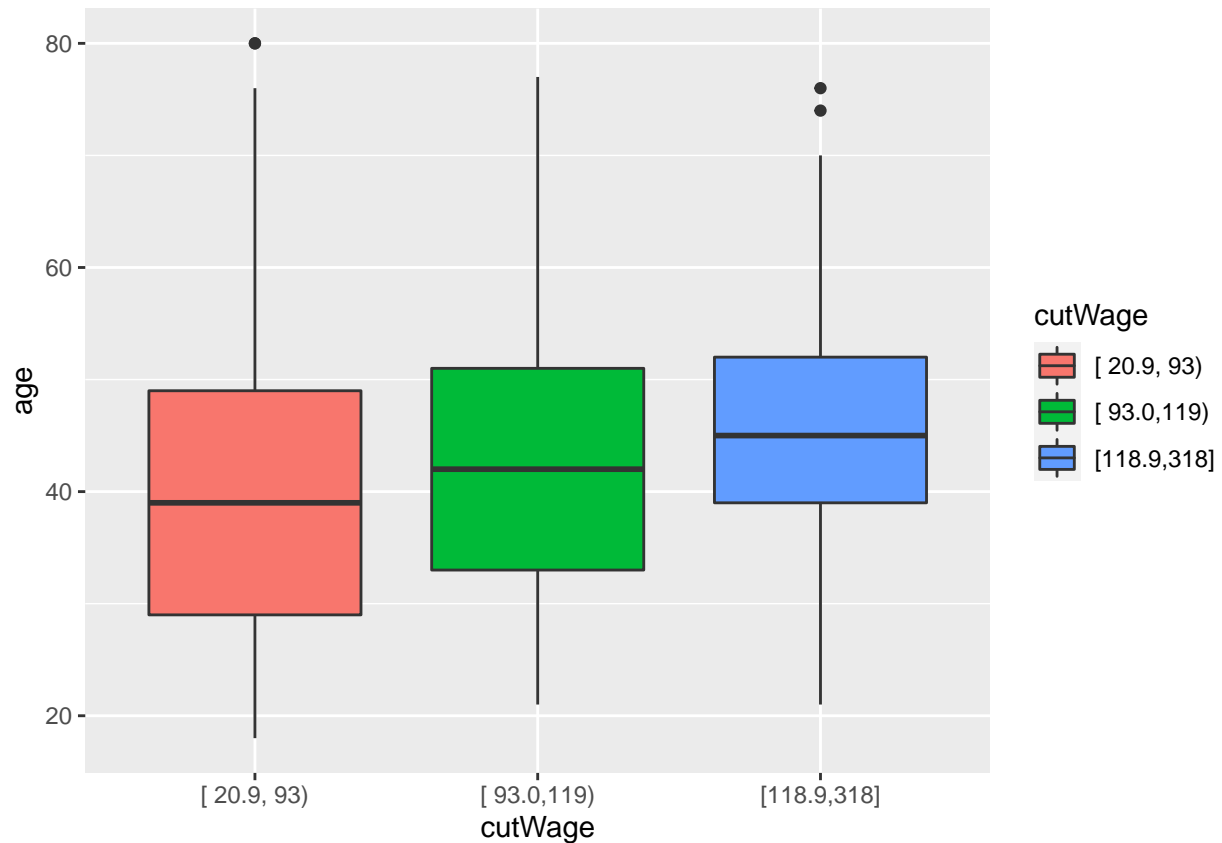
```
qp <- qplot(age, wage, color= education, data=training)
qp + geom_smooth(method = 'lm', formula= y~x)
```



```
library(Hmisc) ##good library for cutting data.frames into pieces.
cutWage <- cut2(training$wage, g=3) ##cut2 argument performs the cut, by $wage column, into g=3 pieces.
table(cutWage)
```

```
## cutWage
## [ 20.9, 93) [ 93.0,119) [118.9,318]
##          707          718          677
```

```
p1 <- qplot(cutWage, age, data=training, fill=cutWage, geom=c("boxplot"))
p1
```



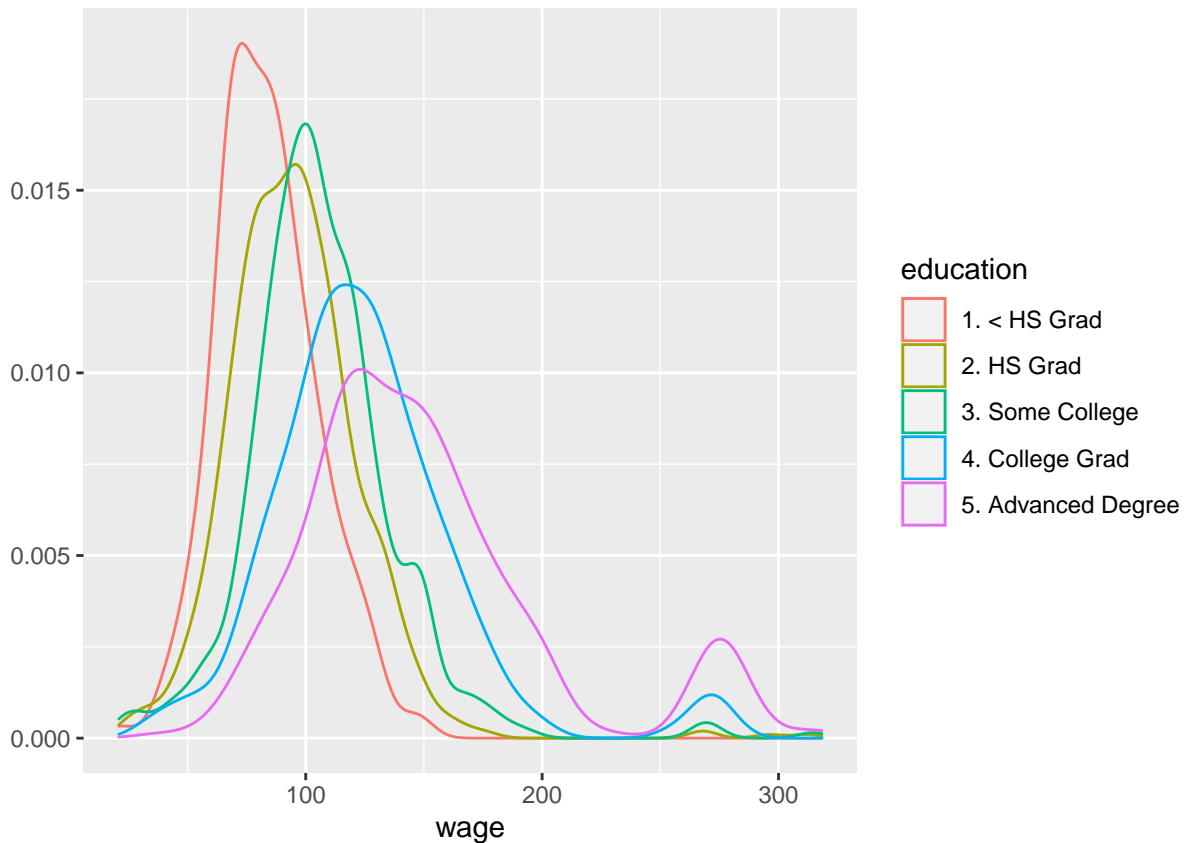
```
t1 <- table(cutWage, training$jobclass) ##great way to make quick tables
t1
```

```
##
## cutWage      1. Industrial 2. Information
## [ 20.9, 93)      455        252
## [ 93.0, 119)     362        356
## [118.9, 318]     276        401
```

```
prop.table(t1,1) ##proportion table, where the 1 represents proportion in each row. a 2 would be each
```

```
##
## cutWage      1. Industrial 2. Information
## [ 20.9, 93)    0.6435644    0.3564356
## [ 93.0, 119)    0.5041783    0.4958217
## [118.9, 318]    0.4076809    0.5923191
```

```
qplot(wage, color=education, data= training, geom="density")
```

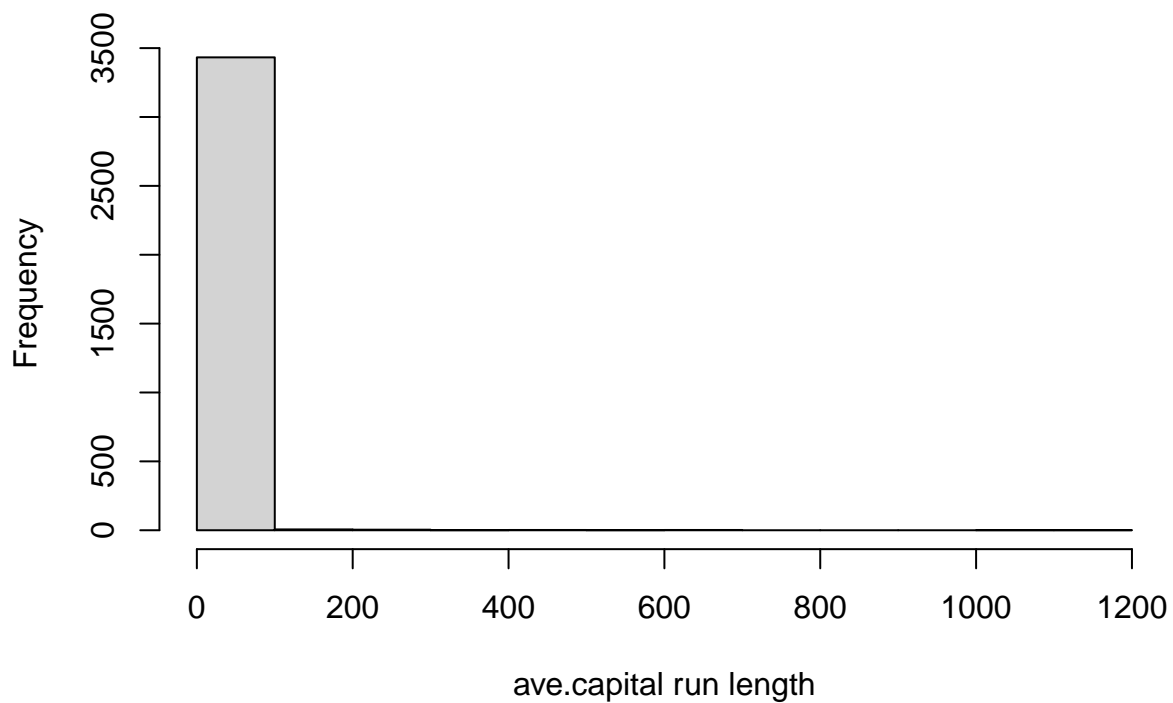
Pre-processing predictor variables

```
library(caret)
library(kernlab) ## to get spam dataset
library(RANN)

data(spam)
inTrain <- createDataPartition(y=spam$type, p= 0.75, list = FALSE) ## data is partitioned by spam type,
training <- spam[inTrain,] ##subsetting the output of the partition function by spam
testing <- spam[-inTrain,] ##subsetting the output of the partition function by not spam
dim(training)

## [1] 3451 58

hist(training$capitalAve, main="", xlab="ave.capital run length")
```



```
mean(training$capitalAve)
```

```
## [1] 5.577148
```

```
sd(training$capitalAve) ## what we see here is that the standard deviation is significantly higher than
```

```
## [1] 35.92989
```

```
trainCapAve <- training$capitalAve  
trainCapAveS <- (trainCapAve - mean(trainCapAve))/sd(trainCapAve) ## A way of standardizing the data  
mean(trainCapAveS)
```

```
## [1] -1.162129e-17
```

```
sd(trainCapAveS)
```

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

```
##If we want to then standardize the test set, we must use the mean from the training set, and the st.d.  
testCapAve <- testing$capitalAve  
testCapAveS <- (testCapAve - mean(trainCapAve))/sd(trainCapAve)  
mean(testCapAveS)
```

```
## [1] -0.04294102
```

```
sd(testCapAveS)
```

```
## [1] 0.3437084
```

```
##Alternative is passing preProcess into the train() argument
```

```
modelFit <- train(type~., data= training, preProcess= c("center", "scale"), method="glm") ##This process
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Warning: glm.fit: algorithm did not converge
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Warning: glm.fit: algorithm did not converge
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
## Warning: glm.fit: algorithm did not converge

## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred

## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred

## Warning: glm.fit: algorithm did not converge

## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred

## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred

## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
```

```
modelFit
```

```
## Generalized Linear Model
##
## 3451 samples
## 57 predictor
## 2 classes: 'nonspam', 'spam'
##
## Pre-processing: centered (57), scaled (57)
## Resampling: Bootstrapped (25 reps)
## Summary of sample sizes: 3451, 3451, 3451, 3451, 3451, 3451, ...
## Resampling results:
##
## Accuracy Kappa
## 0.9167869 0.8257754
```

```
## Prediction models tend to fail with NA. We'll use K nearest imputation statistical function.
# Make some values NA
training$capAve <- training$capitalAve
selectNA <- rbinom(dim(training)[1],size=1,prob=0.05)==1 ## we are adding NA's into the dataset here.
training$capAve[selectNA] <- NA
# Impute and standardize
preObj <- preProcess(training[, -58],method="knnImpute") ##removing the 58th column here
capAve <- predict(preObj,training[, -58])$capAve
# Standardize true values
capAveTruth <- training$capitalAve
capAveTruth <- (capAveTruth-mean(capAveTruth))/sd(capAveTruth)
```

Covariate (predictor) creation

```
##think about creating or calculating variables/features to summarize the sample data, in this instance

library(ISLR); library(caret); data(Wage);
inTrain <- createDataPartition(y=Wage$wage,
                                p=0.7, list=FALSE)
training <- Wage[inTrain,]
```

```
testing <- Wage[-inTrain,]
```

```
##turning qualitative variables into quantitative ones
table(training$jobclass)
```

```
##
## 1. Industrial 2. Information
##      1067      1035
```

```
dummies <- dummyVars(wage ~ jobclass,data=training) ##using the dummyVars argument to turn categorical
head(predict(dummies,newdata=training))
```

```
##      jobclass.1. Industrial jobclass.2. Information
## 161300      1      0
## 155159      0      1
## 376662      0      1
## 450601      1      0
## 377954      0      1
## 228963      0      1
```

```
##-----
##A way to throw out less-meaningful predictors, i.e. near-zero variations in the data.
nsv <- nearZeroVar(training,saveMetrics=TRUE)
nsv ##in this example we can throw out sex and region
```

```
##      freqRatio percentUnique zeroVar  nzv
## year      1.008721    0.33301618  FALSE FALSE
## age      1.013699    2.85442436  FALSE FALSE
## maritl    3.152505    0.23786870  FALSE FALSE
## race      8.964103    0.19029496  FALSE FALSE
## education 1.432150    0.23786870  FALSE FALSE
## region    0.000000    0.04757374   TRUE  TRUE
## jobclass  1.030918    0.09514748  FALSE FALSE
## health    2.434641    0.09514748  FALSE FALSE
## health_ins 2.269051    0.09514748  FALSE FALSE
## logwage   1.000000    19.36251189  FALSE FALSE
## wage      1.000000    19.36251189  FALSE FALSE
```

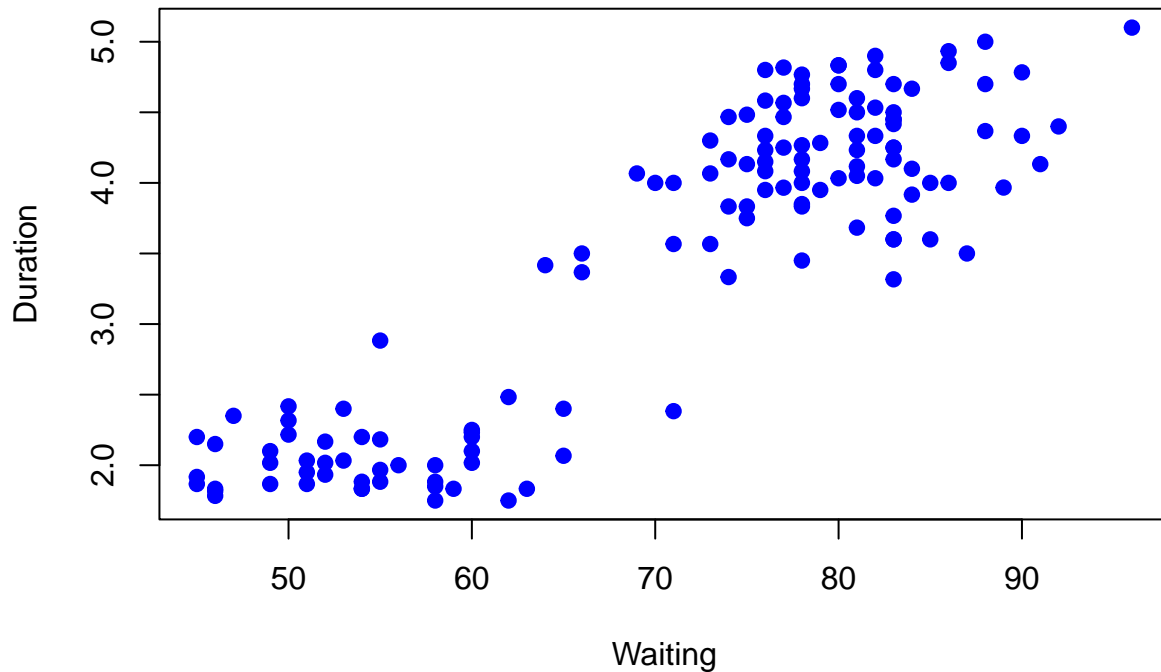
Predicting with Regression

```
##Using the old faithful dataset
library(caret);data(faithful); set.seed(333)
inTrain <- createDataPartition(y=faithful$waiting,
                                p=0.5, list=FALSE)
trainFaith <- faithful[inTrain,]; testFaith <- faithful[-inTrain,]
head(trainFaith)
```

```
##      eruptions waiting
## 3      3.333      74
```

```
## 6      2.883      55
## 7      4.700      88
## 8      3.600      85
## 9      1.950      51
## 11     1.833      54
```

```
plot(trainFaith$waiting,trainFaith$eruptions,pch=19,col="blue",xlab="Waiting",ylab="Duration") ##plotti
```



```
lm1 <- lm(eruptions ~ waiting, data=trainFaith)
summary(lm1)
```

```
##
## Call:
## lm(formula = eruptions ~ waiting, data = trainFaith)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -1.13375 -0.36778  0.06064  0.36578  0.96057
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -1.648629   0.226603  -7.275 2.55e-11 ***
## waiting      0.072211   0.003136  23.026 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
##
## Residual standard error: 0.4941 on 135 degrees of freedom
## Multiple R-squared:  0.7971, Adjusted R-squared:  0.7956
## F-statistic: 530.2 on 1 and 135 DF,  p-value: < 2.2e-16

##output is y(eruption duration) = 0.073 * (waiting time) - 1.792

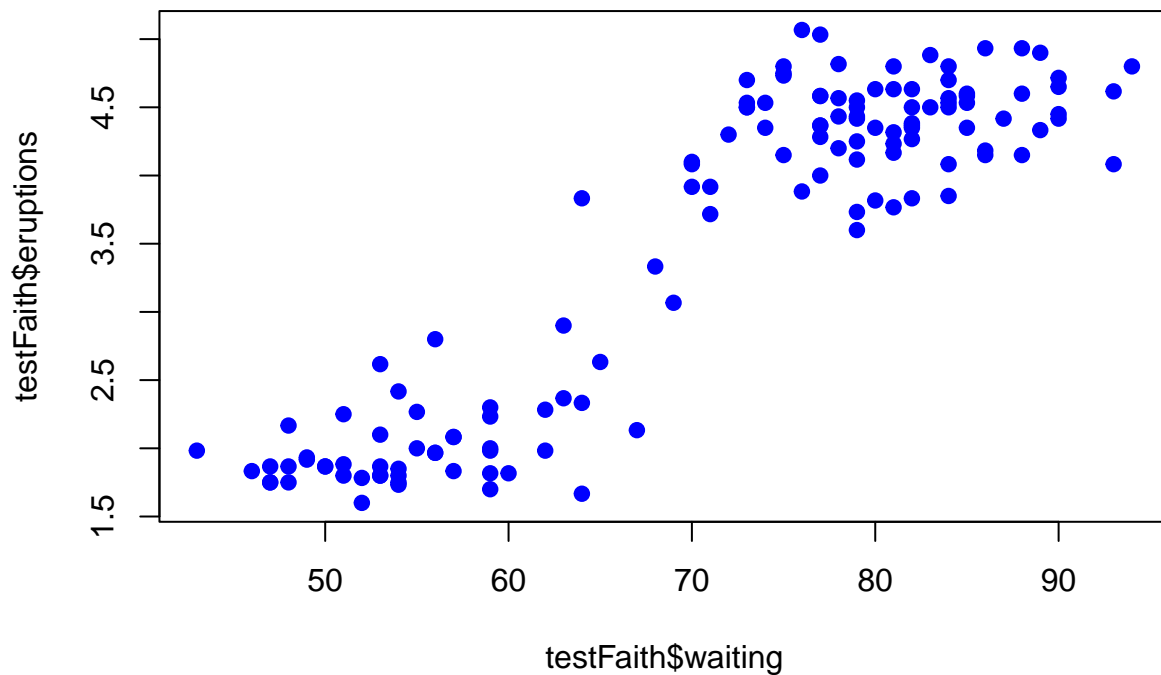
## To predict a new value, we can automate this by:
coef(lm1)[1] + coef(lm1)[2]*80 ## coef(lm1)[1] returns the intercept, and [2] returns the slope. We are

## (Intercept)
##      4.128276

newdata <- data.frame(waiting=80)
predict(lm1,newdata) ## a shortcut so we don't have to continuously calculate the coefficients.

##      1
## 4.128276

##We can not use the predictions from the training set on the TEST set
pred1 <- predict(lm1,newdata=testFaith,interval="prediction")
ord <- order(testFaith$waiting)
plot(testFaith$waiting,testFaith$eruptions,pch=19,col="blue")
```



```
##-----
##We can use CARET to do the same, and much faster
modFit <- train(eruptions ~ waiting,data=trainFaith,method="lm") ##eruptions is outcome, waiting is pr
summary(modFit$finalModel) ##How we get final model output

##
## Call:
## lm(formula = .outcome ~ ., data = dat)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -1.13375 -0.36778  0.06064  0.36578  0.96057
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -1.648629   0.226603  -7.275 2.55e-11 ***
## waiting      0.072211   0.003136  23.026 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.4941 on 135 degrees of freedom
## Multiple R-squared:  0.7971, Adjusted R-squared:  0.7956
## F-statistic: 530.2 on 1 and 135 DF,  p-value: < 2.2e-16
```

Predicting Multiple Covariate Regression

```
library(ISLR); library(ggplot2); library(caret);
data(Wage); Wage <- subset(Wage,select=-c(logwage)) ##here we are subsetting and removing the variable
summary(Wage)
```

```
##      year      age      maritl      race
## Min.   :2003   Min.   :18.00   1. Never Married: 648   1. White:2480
## 1st Qu.:2004   1st Qu.:33.75   2. Married   :2074   2. Black: 293
## Median :2006   Median :42.00   3. Widowed   : 19    3. Asian: 190
## Mean   :2006   Mean   :42.41   4. Divorced  : 204    4. Other:  37
## 3rd Qu.:2008   3rd Qu.:51.00   5. Separated :  55
## Max.   :2009   Max.   :80.00
##
##      education      region      jobclass
## 1. < HS Grad      :268   2. Middle Atlantic :3000   1. Industrial :1544
## 2. HS Grad        :971   1. New England   :  0   2. Information:1456
## 3. Some College   :650   3. East North Central:  0
## 4. College Grad   :685   4. West North Central:  0
## 5. Advanced Degree:426   5. South Atlantic    :  0
##                      6. East South Central:  0
##                      (Other)              :  0
##      health      health_ins      wage
## 1. <=Good      : 858   1. Yes:2083   Min.   : 20.09
## 2. >=Very Good:2142   2. No : 917   1st Qu.: 85.38
##                      Median :104.92
##                      Mean   :111.70
```



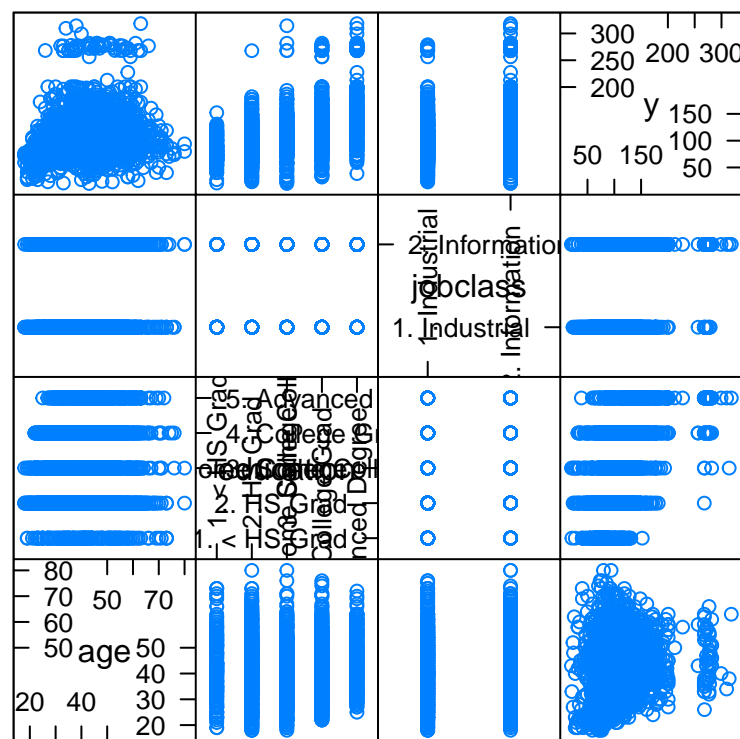
```
##                               3rd Qu.:128.68
##                               Max.    :318.34
##
```

```
inTrain <- createDataPartition(y=Wage$wage,##subsetting into test and train
                                p=0.7, list=FALSE)
training <- Wage[inTrain,]
testing  <- Wage[-inTrain,]
dim(training); dim(testing)
```

```
## [1] 2102  10
```

```
## [1] 898  10
```

```
featurePlot(x=training[,c("age","education","jobclass")],
            y = training$wage,
            plot="pairs")
```

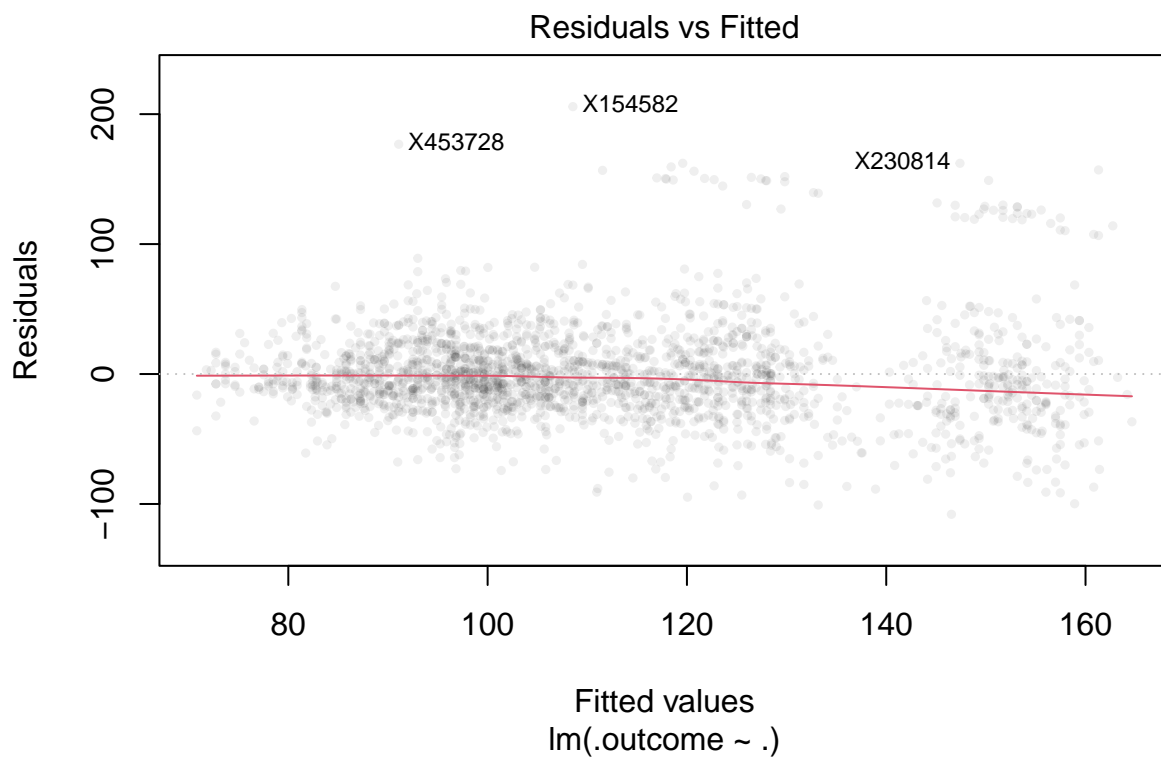


Scatter Plot Matrix

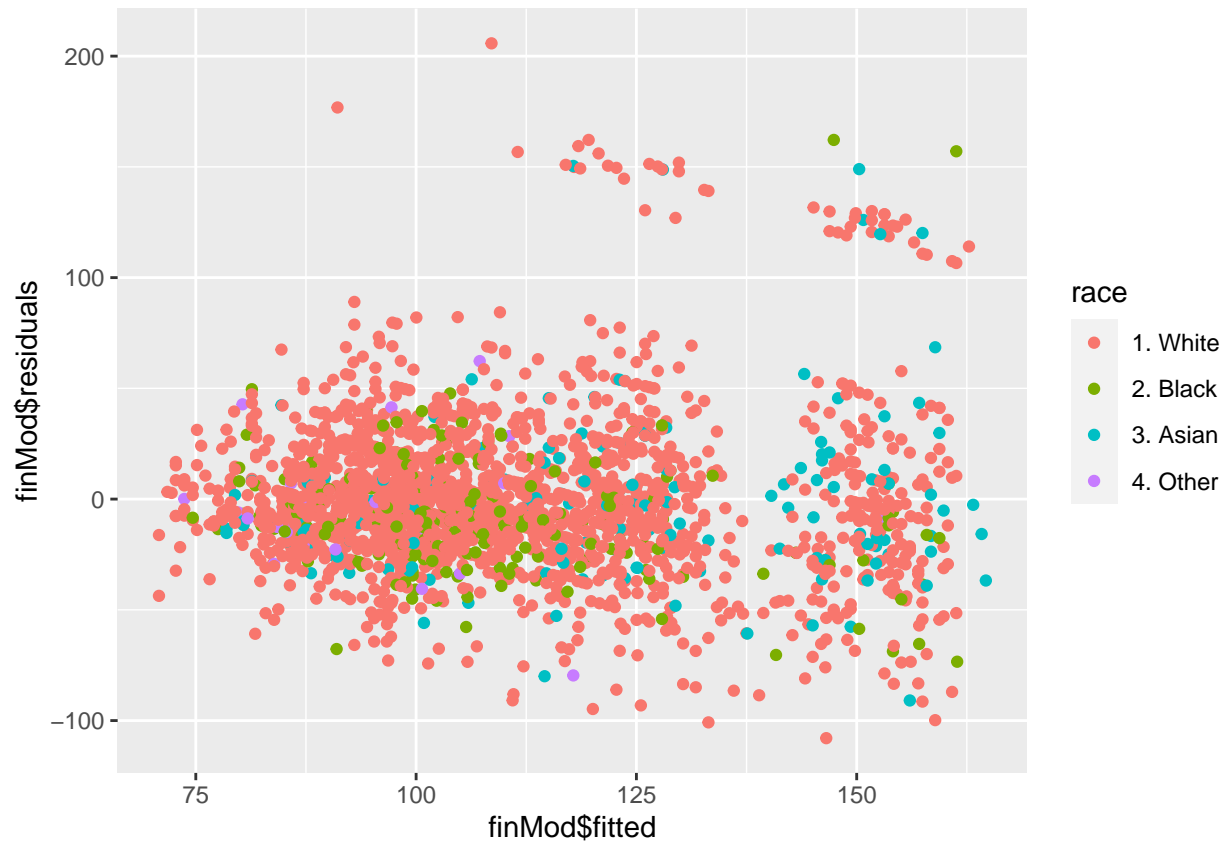
```
## -----
modFit<- train(wage ~ age + jobclass + education,
              method = "lm",data=training)
finMod <- modFit$finalModel
print(modFit)
```

```
## Linear Regression
##
## 2102 samples
##    3 predictor
##
## No pre-processing
## Resampling: Bootstrapped (25 reps)
## Summary of sample sizes: 2102, 2102, 2102, 2102, 2102, 2102, ...
## Resampling results:
##
##    RMSE      Rsquared   MAE
##  35.56759  0.2589245  24.87554
##
## Tuning parameter 'intercept' was held constant at a value of TRUE
```

```
plot(finMod,1,pch=19,cex=0.5,col="#00000010") ##residuals vs. fitted. We want to see a straight line
```



```
qplot(finMod$fitted,finMod$residuals,colour=race,data=training) ##also fitted vs. residuals. Trying to
```



Predicting with Trees

Better performance in non-linear settings Can cause over-fitting, so be careful Considered non-linear models, so they use interactions between variables

1.) start with all variables in one group 2.) Find the variable/split that best separates the outcomes 3.) Divide the data into two groups “leaves” on that node 4.) Within each split, find the best variable/split that separates the outcomes 5.) Continue until the groups are too small or sufficiently “pure”

```
data(iris); library(ggplot2)
names(iris)
```

```
## [1] "Sepal.Length" "Sepal.Width" "Petal.Length" "Petal.Width" "Species"
```

```
table(iris$Species)
```

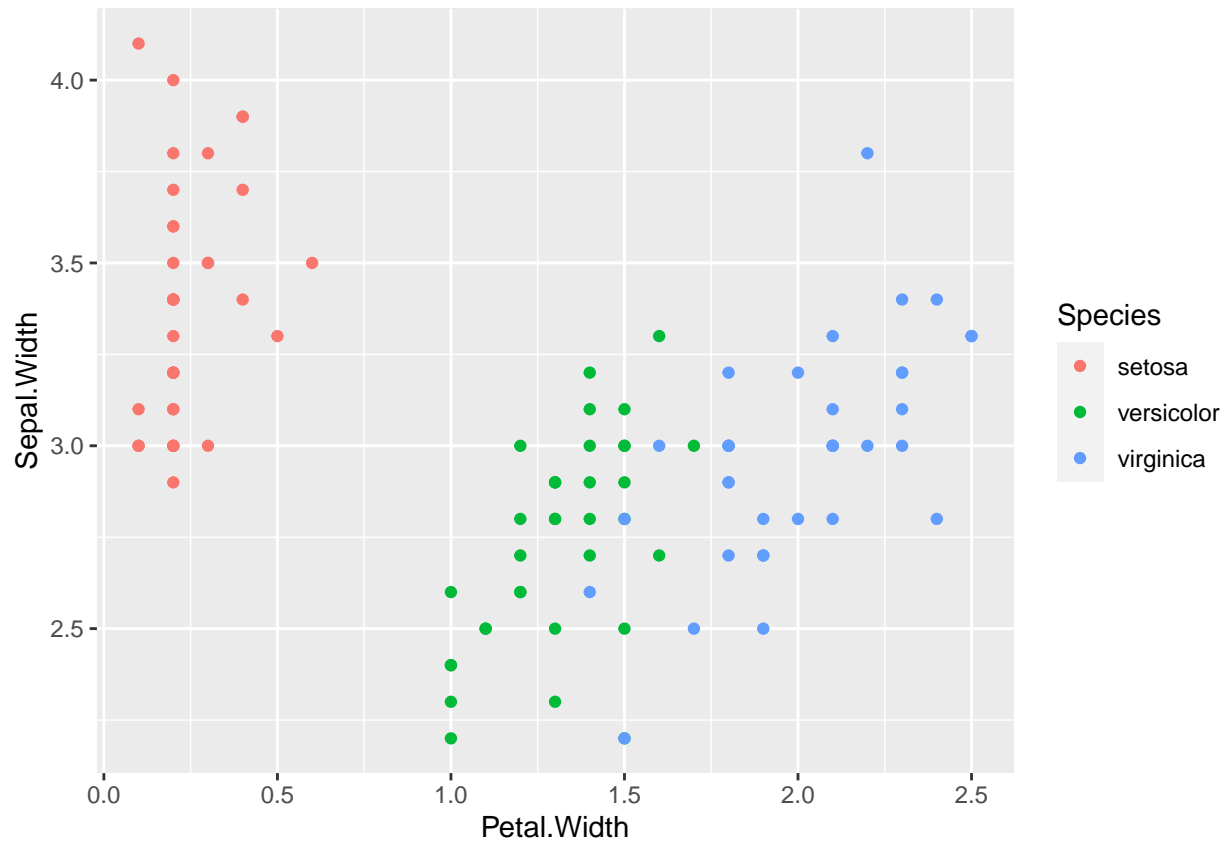
```
##
##      setosa versicolor  virginica
##         50         50         50
```

```
inTrain <- createDataPartition(y=iris$Species,
                               p=0.7, list=FALSE)
training <- iris[inTrain,]
testing  <- iris[-inTrain,]
dim(training); dim(testing)
```

```
## [1] 105 5
```

```
## [1] 45 5
```

```
qplot(Petal.Width,Sepal.Width,colour=Species,data=training)
```

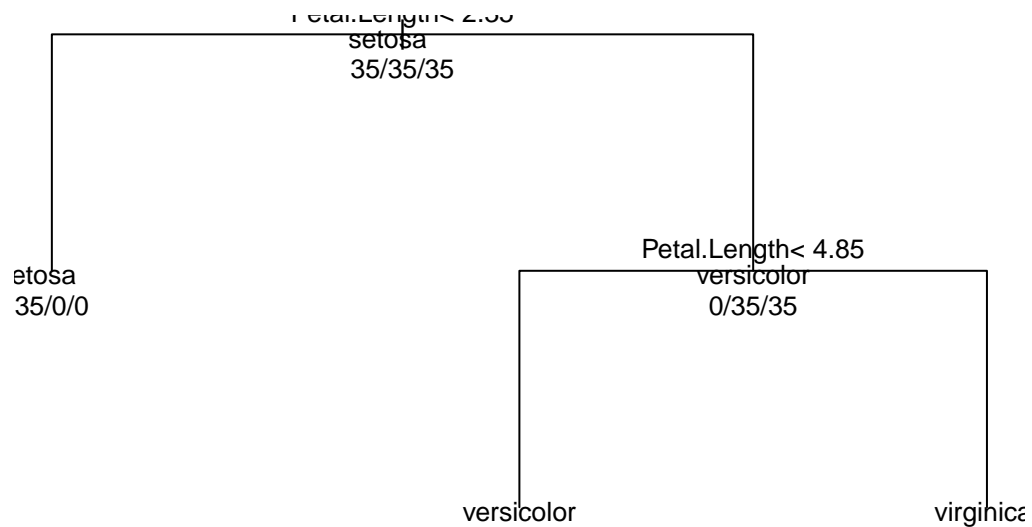


```
library(caret)
modFit <- train(Species ~ .,method="rpart",data=training) ##train function fits the model here. looking
print(modFit$finalModel)
```

```
## n= 105
##
## node), split, n, loss, yval, (yprob)
##      * denotes terminal node
##
## 1) root 105 70 setosa (0.33333333 0.33333333 0.33333333)
## 2) Petal.Length< 2.35 35 0 setosa (1.00000000 0.00000000 0.00000000) *
## 3) Petal.Length>=2.35 70 35 versicolor (0.00000000 0.50000000 0.50000000)
## 6) Petal.Length< 4.85 33 1 versicolor (0.00000000 0.96969697 0.03030303) *
## 7) Petal.Length>=4.85 37 3 virginica (0.00000000 0.08108108 0.91891892) *
```

```
plot(modFit$finalModel, uniform=TRUE,
      main="Classification Tree")
text(modFit$finalModel, use.n=TRUE, all=TRUE, cex=.8) ##plotting the dendogram tree
```

Classification Tree



```
library(rattle)
```

```
## Loading required package: tibble
```

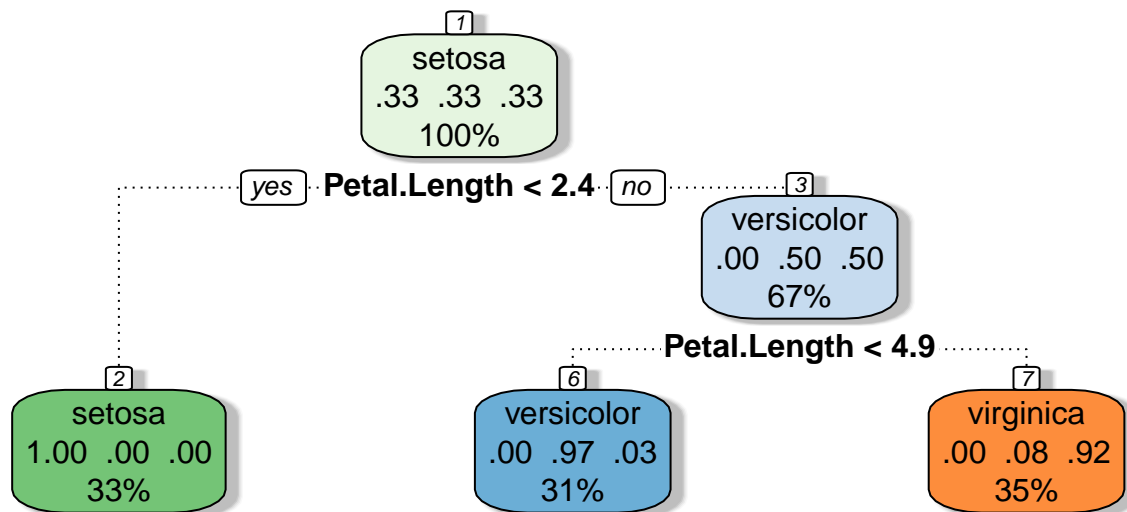
```
## Loading required package: bitops
```

```
## Rattle: A free graphical interface for data science with R.
```

```
## Version 5.5.1 Copyright (c) 2006-2021 Togaware Pty Ltd.
```

```
## Type 'rattle()' to shake, rattle, and roll your data.
```

```
fancyRpartPlot(modFit$finalModel) ##makes a fancier version of the same dendrogram
```



Rattle 2022–Nov–07 14:54:57 payashome

Bootstrap-aggregating “Bagging”

Re-sample the data, and recalculate predictions, then average or majority vote. Useful for non-linear functions. When using bagging in caret, consider using “bagEarth”, “treebag”, “bagFDA” inside of the train function.

Below is the syntax/code, but ozone dataset not available

```

predictors = data.frame(ozone=ozone, temperature =
  ozonetemperature)
treebag <- bag(predictors, temperature, B = 10, bagControl = bagControl(fit =
  ctreeBagfit, predict = ctreeBagpred, aggregate = ctreeBag$aggregate))

```

Random Forests

1.) Bootstrap samples 2.) at each split, bootstrap the variables again 3.) Grow multiple trees and vote
Cons- prone to overfitting, but it is very accurate

```
data(iris); library(ggplot2); library(randomForest); library(caret)
```

```
## randomForest 4.7-1.1
```

```
## Type rfNews() to see new features/changes/bug fixes.
```

```
##
```

```
## Attaching package: 'randomForest'
```

```
## The following object is masked from 'package:rattle':
##
## importance
```

```
## The following object is masked from 'package:dplyr':
##
## combine
```

```
## The following object is masked from 'package:ggplot2':
##
## margin
```

```
inTrain <- createDataPartition(y=iris$Species,
                               p=0.7, list=FALSE)
training <- iris[inTrain,]
testing <- iris[-inTrain,]
```

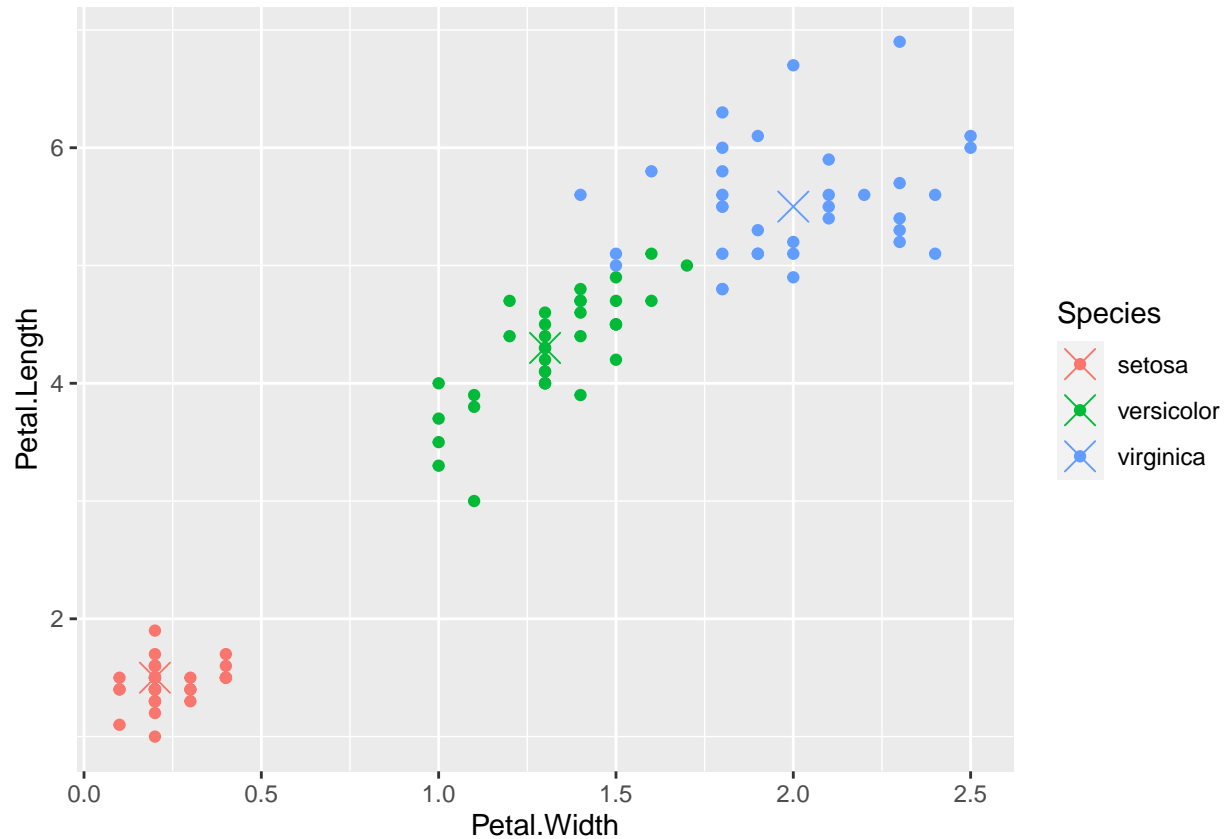
```
modFit <- train(Species~ .,data=training,method="rf",prox=TRUE) ##using train, here method is "rf", wh
modFit
```

```
## Random Forest
##
## 105 samples
## 4 predictor
## 3 classes: 'setosa', 'versicolor', 'virginica'
##
## No pre-processing
## Resampling: Bootstrapped (25 reps)
## Summary of sample sizes: 105, 105, 105, 105, 105, 105, ...
## Resampling results across tuning parameters:
##
## mtry Accuracy Kappa
## 2 0.9497883 0.9240399
## 3 0.9508409 0.9256164
## 4 0.9508994 0.9256779
##
## Accuracy was used to select the optimal model using the largest value.
## The final value used for the model was mtry = 4.
```

```
getTree(modFit$finalModel,k=2) ##to view the individual trees, you can use this function, which is look
```

```
## left daughter right daughter split var split point status prediction
## 1 2 3 3 2.70 1 0
## 2 0 0 0 0.00 -1 1
## 3 4 5 4 1.75 1 0
## 4 6 7 3 5.35 1 0
## 5 0 0 0 0.00 -1 3
## 6 0 0 0 0.00 -1 2
## 7 0 0 0 0.00 -1 3
```

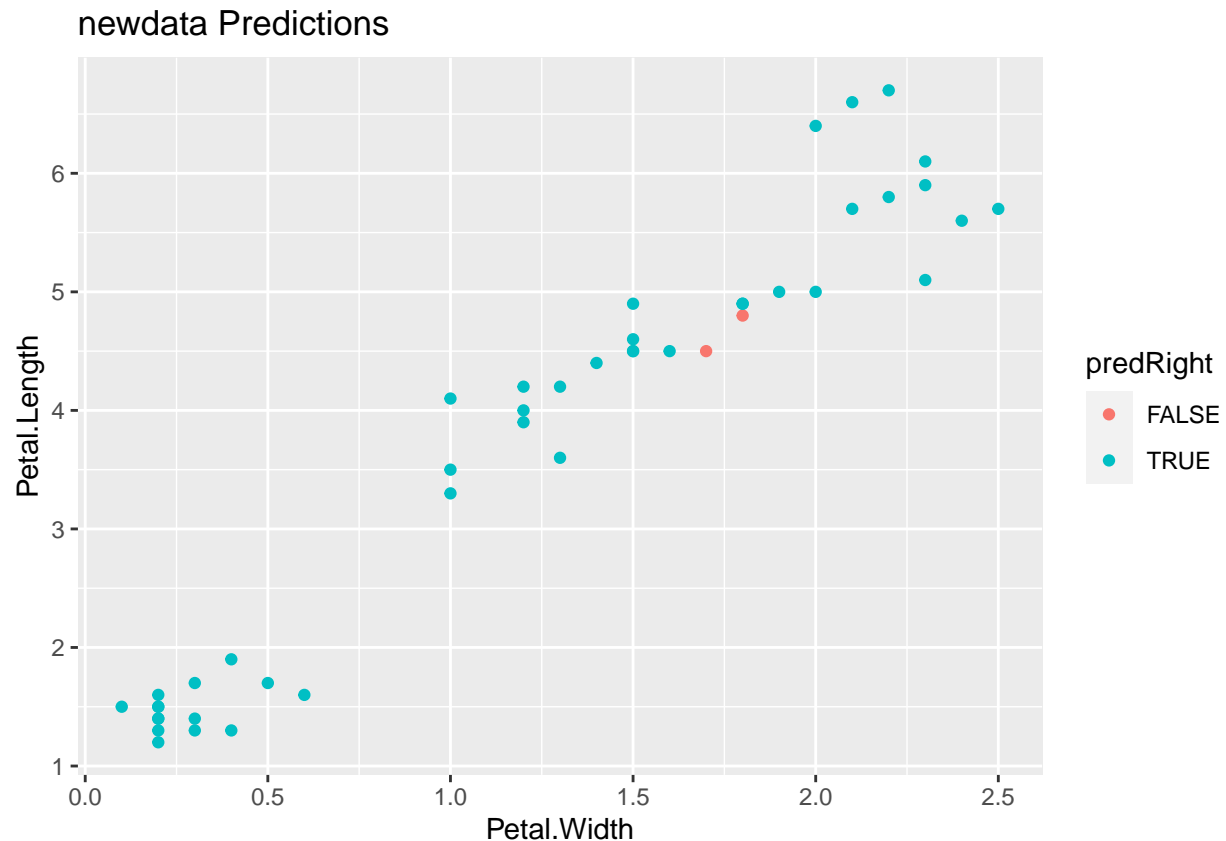
```
irisP <- classCenter(training[,c(3,4)], training$Species, modFit$finalModel$prox) ##This centers your
irisP <- as.data.frame(irisP); irisP$Species <- rownames(irisP)
p <- qplot(Petal.Width, Petal.Length, col=Species, data=training)
p + geom_point(aes(x=Petal.Width, y=Petal.Length, col=Species), size=5, shape=4, data=irisP)
```



```
pred <- predict(modFit,testing); testing$predRight <- pred==testing$Species
table(pred,testing$Species)
```

```
##
## pred      setosa versicolor virginica
## setosa    15      0          0
## versicolor 0      14         1
## virginica  0      1         14
```

```
qplot(Petal.Width,Petal.Length,colour=predRight,data=testing,main="newdata Predictions")
```

Boosting

1.) Takes a lot of weak predictors 2.) Weight them and add them up 3.) Get a stronger predictor

```
library(ISLR); data(Wage); library(ggplot2); library(caret);
Wage <- subset(Wage,select=-c(logwage)) ##removes the variable were interested in
inTrain <- createDataPartition(y=Wage$wage,
                               p=0.7, list=FALSE)
training <- Wage[inTrain,]; testing <- Wage[-inTrain,]

modFit <- train(wage ~ ., method="gbm",data=training,verbose=FALSE) ## "gbm" is boosting with trees us

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 14: region2. Middle Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 15: region3. East North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 16: region4. West North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 17: region5. South Atlantic has no variation.
```

```

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 18: region6. East South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 19: region7. West South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 20: region8. Mountain has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 21: region9. Pacific has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 14: region2. Middle Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 15: region3. East North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 16: region4. West North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 17: region5. South Atlantic has no variation.

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## "bernoulli", : variable 18: region6. East South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 19: region7. West South Central has no variation.

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## "bernoulli", : variable 16: region4. West North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 17: region5. South Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 18: region6. East South Central has no variation.

```

```

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 19: region7. West South Central has no variation.

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## "bernoulli", : variable 20: region8. Mountain has no variation.

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## "bernoulli", : variable 19: region7. West South Central has no variation.

```

```

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## "bernoulli", : variable 20: region8. Mountain has no variation.

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## "bernoulli", : variable 20: region8. Mountain has no variation.

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```

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```



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## "bernoulli", : variable 14: region2. Middle Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 15: region3. East North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 16: region4. West North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 17: region5. South Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 18: region6. East South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 19: region7. West South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 20: region8. Mountain has no variation.

```

```

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 21: region9. Pacific has no variation.

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## "bernoulli", : variable 14: region2. Middle Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 15: region3. East North Central has no variation.

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## "bernoulli", : variable 16: region4. West North Central has no variation.

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```

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## "bernoulli", : variable 15: region3. East North Central has no variation.

```

```

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```



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## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 21: region9. Pacific has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 14: region2. Middle Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 15: region3. East North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 16: region4. West North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 17: region5. South Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 18: region6. East South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 19: region7. West South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 20: region8. Mountain has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 21: region9. Pacific has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 14: region2. Middle Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 15: region3. East North Central has no variation.

```

```

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 16: region4. West North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 17: region5. South Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 18: region6. East South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 19: region7. West South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 20: region8. Mountain has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 21: region9. Pacific has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 14: region2. Middle Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 15: region3. East North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 16: region4. West North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 17: region5. South Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 18: region6. East South Central has no variation.

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## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 14: region2. Middle Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 15: region3. East North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 16: region4. West North Central has no variation.

```



```

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 17: region5. South Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 18: region6. East South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
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## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
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## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
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## "bernoulli", : variable 16: region4. West North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 17: region5. South Atlantic has no variation.

```

```

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
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## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
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## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 18: region6. East South Central has no variation.

```

```

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
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## "bernoulli", : variable 19: region7. West South Central has no variation.

```

```

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## "bernoulli", : variable 20: region8. Mountain has no variation.

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## "bernoulli", : variable 20: region8. Mountain has no variation.

```

```

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## "bernoulli", : variable 21: region9. Pacific has no variation.

```

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## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
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## "bernoulli", : variable 14: region2. Middle Atlantic has no variation.

```

```

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
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## "bernoulli", : variable 15: region3. East North Central has no variation.

```

```

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```



```

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
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## "bernoulli", : variable 18: region6. East South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 19: region7. West South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 20: region8. Mountain has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 21: region9. Pacific has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 14: region2. Middle Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 15: region3. East North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 16: region4. West North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 17: region5. South Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 18: region6. East South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 19: region7. West South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 20: region8. Mountain has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 21: region9. Pacific has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 14: region2. Middle Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 15: region3. East North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 16: region4. West North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 17: region5. South Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 18: region6. East South Central has no variation.

```

```

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 19: region7. West South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 20: region8. Mountain has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 21: region9. Pacific has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 14: region2. Middle Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 15: region3. East North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 16: region4. West North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 17: region5. South Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 18: region6. East South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 19: region7. West South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 20: region8. Mountain has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 21: region9. Pacific has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 14: region2. Middle Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 15: region3. East North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 16: region4. West North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 17: region5. South Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 18: region6. East South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 19: region7. West South Central has no variation.

```

```

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 20: region8. Mountain has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 21: region9. Pacific has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 14: region2. Middle Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 15: region3. East North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 16: region4. West North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 17: region5. South Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 18: region6. East South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 19: region7. West South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 20: region8. Mountain has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 21: region9. Pacific has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 14: region2. Middle Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 15: region3. East North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 16: region4. West North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 17: region5. South Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 18: region6. East South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 19: region7. West South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 20: region8. Mountain has no variation.

```

```
## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 21: region9. Pacific has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 14: region2. Middle Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 15: region3. East North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 16: region4. West North Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 17: region5. South Atlantic has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 18: region6. East South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 19: region7. West South Central has no variation.

## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 20: region8. Mountain has no variation.

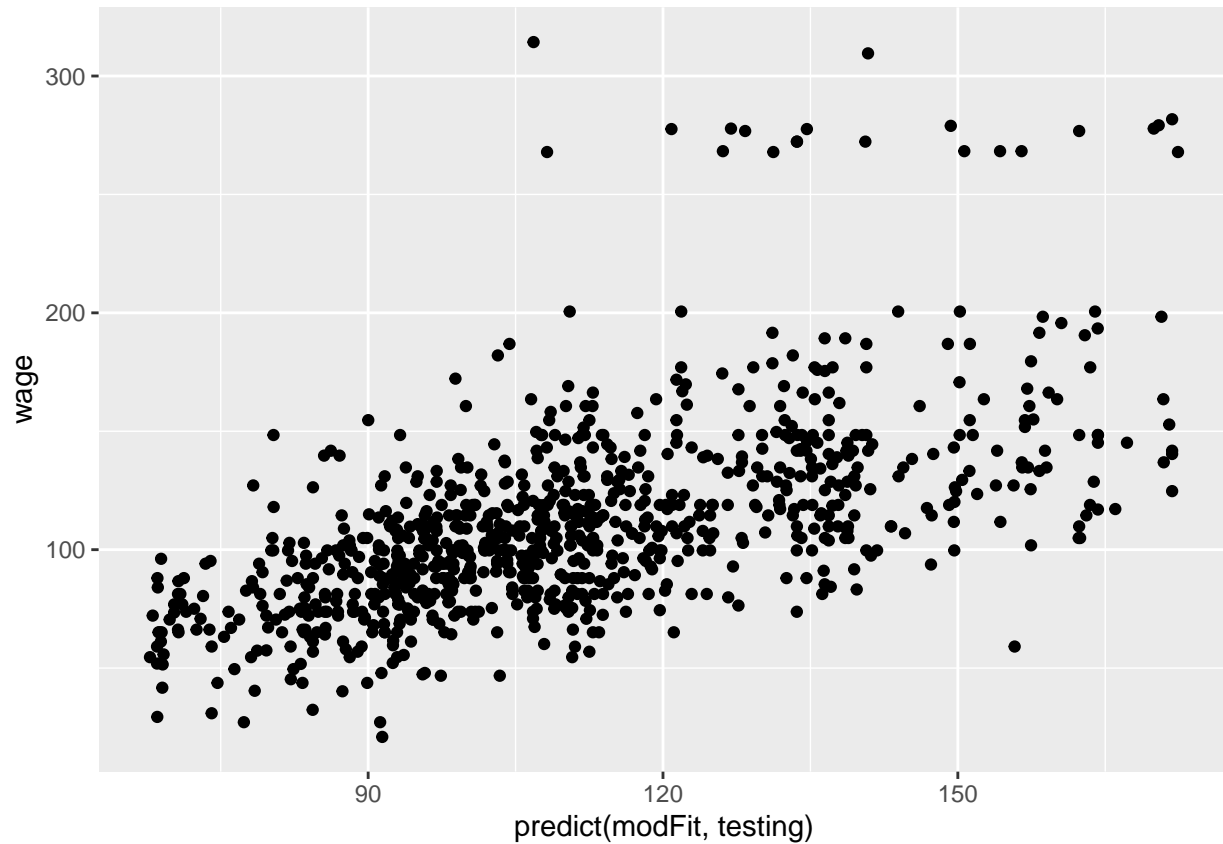
## Warning in (function (x, y, offset = NULL, misc = NULL, distribution =
## "bernoulli", : variable 21: region9. Pacific has no variation.
```

```
print(modFit)
```

```
## Stochastic Gradient Boosting
##
## 2102 samples
## 9 predictor
##
## No pre-processing
## Resampling: Bootstrapped (25 reps)
## Summary of sample sizes: 2102, 2102, 2102, 2102, 2102, 2102, ...
## Resampling results across tuning parameters:
##
## interaction.depth n.trees RMSE Rsquared MAE
## 1 50 35.65470 0.3067370 24.18107
## 1 100 35.08875 0.3164737 23.77081
## 1 150 35.00625 0.3181686 23.78886
## 2 50 34.99228 0.3216033 23.70188
## 2 100 34.86318 0.3238252 23.70463
## 2 150 34.91481 0.3222133 23.80882
## 3 50 34.84892 0.3253748 23.62091
## 3 100 34.95760 0.3212939 23.80812
## 3 150 35.11354 0.3165658 23.99817
##
## Tuning parameter 'shrinkage' was held constant at a value of 0.1
```

```
##
## Tuning parameter 'n.minobsinnode' was held constant at a value of 10
## RMSE was used to select the optimal model using the smallest value.
## The final values used for the model were n.trees = 50, interaction.depth =
## 3, shrinkage = 0.1 and n.minobsinnode = 10.
```

```
qplot(predict(modFit,testing),wage,data=testing)
```



Practical Example 1

1. Subset the data to a training set and testing set based on the Case variable in the data set.
2. Set the seed to 125 and fit a caret model with the rpart method using all predictor variables and default caret settings.
3. In the final model what would be the final model prediction for cases with the following variable values:
 - a. TotalIntench2 = 23,000; FiberWidthCh1 = 10; PerimStatusCh1 = 2.
 - b. TotalIntench2 = 50,000; FiberWidthCh1 = 10; VarIntenCh4 = 100.
 - c. TotalIntench2 = 57,000; FiberWidthCh1 = 8; VarIntenCh4 = 100.

d. FiberWidthCh1 = 8; VarIntenCh4 = 100; PerimStatusCh1 = 2.

```
library(rattle)
library(AppliedPredictiveModeling)
data(segmentationOriginal)
seg1 <- segmentationOriginal
head(seg1, n=20)
```

| ## | Cell | Case | Class | AngleCh1 | AngleStatusCh1 | AreaCh1 | AreaStatusCh1 |
|-------|-------------------|-------------------|-------------------|------------------------|-------------------|---------|---------------|
| ## 1 | 207827637 | Test | PS | 143.247705 | 1 | 185 | 0 |
| ## 2 | 207932307 | Train | PS | 133.752037 | 0 | 819 | 1 |
| ## 3 | 207932463 | Train | WS | 106.646387 | 0 | 431 | 0 |
| ## 4 | 207932470 | Train | PS | 69.150325 | 0 | 298 | 0 |
| ## 5 | 207932455 | Test | PS | 2.887837 | 2 | 285 | 0 |
| ## 6 | 207827656 | Test | WS | 40.748298 | 2 | 172 | 0 |
| ## 7 | 207827659 | Test | WS | 173.957833 | 1 | 177 | 0 |
| ## 8 | 207827661 | Test | PS | 179.800467 | 1 | 251 | 0 |
| ## 9 | 207932479 | Test | WS | 18.936420 | 2 | 495 | 0 |
| ## 10 | 207932480 | Test | WS | 153.252427 | 1 | 384 | 0 |
| ## 11 | 207827711 | Test | WS | 13.666671 | 2 | 424 | 0 |
| ## 12 | 207932484 | Train | WS | 109.416426 | 0 | 256 | 0 |
| ## 13 | 207827674 | Test | PS | 52.205447 | 2 | 236 | 0 |
| ## 14 | 207932302 | Test | PS | 92.901464 | 0 | 187 | 0 |
| ## 15 | 207932459 | Train | PS | 104.278654 | 0 | 258 | 0 |
| ## 16 | 207827779 | Train | PS | 77.991943 | 0 | 358 | 0 |
| ## 17 | 207827784 | Train | PS | 13.659972 | 2 | 158 | 0 |
| ## 18 | 207827790 | Test | WS | 23.071066 | 2 | 313 | 0 |
| ## 19 | 207827645 | Train | WS | 106.844369 | 0 | 315 | 0 |
| ## 20 | 207932306 | Test | PS | 93.591140 | 0 | 762 | 1 |
| ## | AvgIntenCh1 | AvgIntenCh2 | AvgIntenCh3 | AvgIntenCh4 | AvgIntenStatusCh1 | | |
| ## 1 | 15.71186 | 3.954802 | 9.548023 | 2.214689 | | 0 | |
| ## 2 | 31.92327 | 205.878517 | 69.916880 | 164.153453 | | 0 | |
| ## 3 | 28.03883 | 115.315534 | 63.941748 | 106.696602 | | 0 | |
| ## 4 | 19.45614 | 101.294737 | 28.217544 | 31.028070 | | 0 | |
| ## 5 | 24.27574 | 111.415441 | 20.474265 | 40.577206 | | 0 | |
| ## 6 | 325.93333 | 652.606061 | 128.690909 | 346.854545 | | 1 | |
| ## 7 | 260.05917 | 595.420118 | 123.840237 | 273.295858 | | 1 | |
| ## 8 | 18.33750 | 4.729167 | 17.233333 | 1.554167 | | 0 | |
| ## 9 | 16.05285 | 88.482030 | 13.716702 | 51.427061 | | 0 | |
| ## 10 | 17.74387 | 88.931880 | 20.444142 | 63.111717 | | 0 | |
| ## 11 | 174.35556 | 388.427160 | 38.762963 | 126.728395 | | 1 | |
| ## 12 | 18.82857 | 125.938776 | 13.600000 | 46.800000 | | 0 | |
| ## 13 | 18.19027 | 5.327434 | 17.088496 | 1.911504 | | 0 | |
| ## 14 | 40.24022 | 213.189944 | 44.195531 | 136.251397 | | 0 | |
| ## 15 | 17.57085 | 124.368421 | 22.461538 | 71.206478 | | 0 | |
| ## 16 | 42.28363 | 217.131579 | 42.321637 | 67.540936 | | 0 | |
| ## 17 | 31.41060 | 102.211921 | 41.490066 | 116.119205 | | 0 | |
| ## 18 | 215.41137 | 568.913043 | 150.652174 | 378.612040 | | 1 | |
| ## 19 | 294.76744 | 491.734219 | 193.435216 | 400.451827 | | 1 | |
| ## 20 | 35.64099 | 163.719395 | 77.522696 | 213.110041 | | 0 | |
| ## | AvgIntenStatusCh2 | AvgIntenStatusCh3 | AvgIntenStatusCh4 | ConvexHullAreaRatioCh1 | | | |
| ## 1 | 2 | 2 | 2 | 1.124509 | | | |
| ## 2 | 0 | 0 | 0 | 1.263158 | | | |
| ## 3 | 0 | 0 | 0 | 1.053310 | | | |

| | | | | |
|-------|---|---|---|----------|
| ## 4 | 0 | 0 | 2 | 1.202625 |
| ## 5 | 0 | 0 | 0 | 1.109333 |
| ## 6 | 1 | 1 | 1 | 1.005831 |
| ## 7 | 1 | 1 | 0 | 1.014164 |
| ## 8 | 2 | 0 | 2 | 1.200389 |
| ## 9 | 0 | 0 | 0 | 1.192308 |
| ## 10 | 0 | 0 | 0 | 1.160365 |
| ## 11 | 1 | 0 | 0 | 1.038462 |
| ## 12 | 0 | 0 | 0 | 1.081871 |
| ## 13 | 2 | 0 | 2 | 1.292658 |
| ## 14 | 0 | 0 | 0 | 1.035230 |
| ## 15 | 0 | 0 | 0 | 1.075728 |
| ## 16 | 0 | 0 | 0 | 1.037830 |
| ## 17 | 0 | 0 | 0 | 1.379310 |
| ## 18 | 1 | 1 | 1 | 1.040258 |
| ## 19 | 1 | 1 | 1 | 1.014516 |
| ## 20 | 0 | 0 | 0 | 1.202662 |

| ## | ConvexHullAreaRatioStatusCh1 | ConvexHullPerimRatioCh1 |
|----|------------------------------|-------------------------|
|----|------------------------------|-------------------------|

| | | |
|-------|---|-----------|
| ## 1 | 0 | 0.9196827 |
| ## 2 | 1 | 0.7970801 |
| ## 3 | 0 | 0.9354750 |
| ## 4 | 0 | 0.8658291 |
| ## 5 | 0 | 0.9568124 |
| ## 6 | 0 | 0.9926363 |
| ## 7 | 0 | 0.9840211 |
| ## 8 | 0 | 0.8310554 |
| ## 9 | 0 | 0.8219601 |
| ## 10 | 0 | 0.8649792 |
| ## 11 | 0 | 0.9725156 |
| ## 12 | 0 | 0.9204595 |
| ## 13 | 1 | 0.8013622 |
| ## 14 | 0 | 0.9796029 |
| ## 15 | 0 | 0.9313002 |
| ## 16 | 0 | 0.9612698 |
| ## 17 | 1 | 0.8960387 |
| ## 18 | 0 | 0.9576618 |
| ## 19 | 0 | 0.9775437 |
| ## 20 | 0 | 0.8487132 |

| ## | ConvexHullPerimRatioStatusCh1 | DiffIntenDensityCh1 | DiffIntenDensityCh3 |
|----|-------------------------------|---------------------|---------------------|
|----|-------------------------------|---------------------|---------------------|

| | | | |
|-------|---|----------|----------|
| ## 1 | 0 | 29.51923 | 13.77564 |
| ## 2 | 2 | 31.87500 | 43.12228 |
| ## 3 | 0 | 32.48771 | 35.98577 |
| ## 4 | 2 | 26.73228 | 22.91732 |
| ## 5 | 0 | 31.58065 | 21.70968 |
| ## 6 | 1 | 92.56291 | 61.92715 |
| ## 7 | 1 | 89.22654 | 47.81553 |
| ## 8 | 2 | 28.61792 | 19.22170 |
| ## 9 | 2 | 26.98404 | 15.06613 |
| ## 10 | 2 | 25.76036 | 18.23077 |
| ## 11 | 0 | 67.25131 | 27.18848 |
| ## 12 | 0 | 28.02004 | 14.87082 |
| ## 13 | 2 | 28.52041 | 16.91327 |
| ## 14 | 0 | 32.11585 | 25.60976 |
| ## 15 | 0 | 27.87611 | 16.11947 |

| | | | | |
|-------|---------------------------|---------------------------|---------------------------|-----------------|
| ## 16 | | 0 | 43.30938 | 43.36563 |
| ## 17 | | 0 | 38.45038 | 37.64885 |
| ## 18 | | 0 | 73.87410 | 78.32014 |
| ## 19 | | 0 | 154.57801 | 100.31560 |
| ## 20 | | 2 | 35.00584 | 54.41065 |
| ## | DiffIntenDensityCh4 | DiffIntenDensityStatusCh1 | DiffIntenDensityStatusCh3 | |
| ## 1 | 6.826923 | 2 | 2 | |
| ## 2 | 79.308424 | 0 | 0 | |
| ## 3 | 51.357050 | 0 | 0 | |
| ## 4 | 26.393701 | 2 | 0 | |
| ## 5 | 25.032258 | 0 | 0 | |
| ## 6 | 145.708609 | 1 | 1 | |
| ## 7 | 139.291262 | 1 | 0 | |
| ## 8 | 5.382075 | 2 | 0 | |
| ## 9 | 36.117446 | 2 | 2 | |
| ## 10 | 38.059172 | 2 | 0 | |
| ## 11 | 84.141361 | 0 | 0 | |
| ## 12 | 32.697105 | 2 | 2 | |
| ## 13 | 5.918367 | 2 | 0 | |
| ## 14 | 90.274390 | 0 | 0 | |
| ## 15 | 36.185841 | 2 | 0 | |
| ## 16 | 56.043750 | 0 | 0 | |
| ## 17 | 74.923664 | 0 | 0 | |
| ## 18 | 191.553957 | 1 | 1 | |
| ## 19 | 237.407801 | 1 | 1 | |
| ## 20 | 121.671043 | 0 | 0 | |
| ## | DiffIntenDensityStatusCh4 | EntropyIntenCh1 | EntropyIntenCh3 | EntropyIntenCh4 |
| ## 1 | 2 | 4.969781 | 4.371017 | 2.718884 |
| ## 2 | 0 | 6.087592 | 6.642761 | 7.880155 |
| ## 3 | 0 | 5.883557 | 6.683000 | 7.144601 |
| ## 4 | 2 | 5.420065 | 5.436732 | 5.778329 |
| ## 5 | 2 | 5.658248 | 5.285671 | 5.235857 |
| ## 6 | 1 | 6.998111 | 6.805538 | 7.119323 |
| ## 7 | 1 | 6.896239 | 6.660887 | 7.032449 |
| ## 8 | 2 | 5.318070 | 5.085643 | 2.295804 |
| ## 9 | 0 | 5.189203 | 4.493851 | 6.281189 |
| ## 10 | 0 | 5.334973 | 5.320425 | 6.740386 |
| ## 11 | 0 | 7.383121 | 5.247167 | 5.997840 |
| ## 12 | 2 | 5.383272 | 4.151183 | 6.194035 |
| ## 13 | 2 | 5.293391 | 4.993778 | 2.423698 |
| ## 14 | 0 | 5.929336 | 6.055303 | 6.821641 |
| ## 15 | 0 | 5.184278 | 5.486225 | 6.624639 |
| ## 16 | 0 | 6.136583 | 5.870027 | 5.995699 |
| ## 17 | 0 | 5.756261 | 5.814347 | 6.121377 |
| ## 18 | 1 | 7.468859 | 6.813520 | 7.540135 |
| ## 19 | 1 | 7.402635 | 6.761025 | 7.560942 |
| ## 20 | 0 | 6.249204 | 7.027566 | 7.240787 |
| ## | EntropyIntenStatusCh1 | EntropyIntenStatusCh3 | EntropyIntenStatusCh4 | |
| ## 1 | 2 | 0 | 2 | |
| ## 2 | 0 | 1 | 1 | |
| ## 3 | 0 | 1 | 0 | |
| ## 4 | 2 | 0 | 0 | |
| ## 5 | 2 | 0 | 0 | |
| ## 6 | 0 | 1 | 0 | |

| | | | | |
|-------|-----------------------|-----------------------------|------------------------|-----------------------|
| ## 7 | 0 | 1 | 0 | |
| ## 8 | 2 | 0 | 2 | |
| ## 9 | 2 | 0 | 0 | |
| ## 10 | 2 | 0 | 0 | |
| ## 11 | 1 | 0 | 0 | |
| ## 12 | 2 | 0 | 0 | |
| ## 13 | 2 | 0 | 2 | |
| ## 14 | 0 | 0 | 0 | |
| ## 15 | 2 | 0 | 0 | |
| ## 16 | 0 | 0 | 0 | |
| ## 17 | 2 | 0 | 0 | |
| ## 18 | 1 | 1 | 1 | |
| ## 19 | 1 | 1 | 1 | |
| ## 20 | 0 | 1 | 0 | |
| ## | EqCircDiamCh1 | EqCircDiamStatusCh1 | EqEllipseLWRCh1 | EqEllipseLWRStatusCh1 |
| ## 1 | 15.36954 | 0 | 3.060676 | 1 |
| ## 2 | 32.30558 | 1 | 1.558394 | 0 |
| ## 3 | 23.44892 | 0 | 1.375386 | 0 |
| ## 4 | 19.50279 | 0 | 3.391220 | 1 |
| ## 5 | 19.05279 | 0 | 2.741380 | 0 |
| ## 6 | 14.83940 | 2 | 1.039726 | 0 |
| ## 7 | 15.01819 | 2 | 1.088583 | 0 |
| ## 8 | 17.89698 | 0 | 2.866415 | 0 |
| ## 9 | 25.12493 | 0 | 1.506684 | 0 |
| ## 10 | 22.13132 | 0 | 1.755000 | 0 |
| ## 11 | 23.24886 | 0 | 1.465172 | 0 |
| ## 12 | 18.08245 | 0 | 1.378748 | 0 |
| ## 13 | 17.36715 | 0 | 3.378319 | 1 |
| ## 14 | 15.45613 | 0 | 1.535489 | 0 |
| ## 15 | 18.15611 | 0 | 1.616053 | 0 |
| ## 16 | 21.36423 | 0 | 1.468210 | 0 |
| ## 17 | 14.19589 | 2 | 1.477694 | 0 |
| ## 18 | 19.97606 | 0 | 1.762458 | 0 |
| ## 19 | 20.04276 | 0 | 1.016647 | 0 |
| ## 20 | 31.14881 | 1 | 1.871580 | 0 |
| ## | EqEllipseOblateVolCh1 | EqEllipseOblateVolStatusCh1 | EqEllipseProlateVolCh1 | |
| ## 1 | 336.9691 | 0 | 110.0963 | |
| ## 2 | 2232.9055 | 1 | 1432.8246 | |
| ## 3 | 802.1945 | 0 | 583.2504 | |
| ## 4 | 724.7143 | 0 | 213.7031 | |
| ## 5 | 607.5185 | 0 | 221.6105 | |
| ## 6 | 176.7693 | 0 | 170.0153 | |
| ## 7 | 187.4918 | 0 | 172.2348 | |
| ## 8 | 514.8823 | 0 | 179.6259 | |
| ## 9 | 1032.8193 | 0 | 685.4917 | |
| ## 10 | 761.8320 | 0 | 434.0923 | |
| ## 11 | 806.9533 | 0 | 550.7567 | |
| ## 12 | 368.3097 | 0 | 267.1334 | |
| ## 13 | 510.7808 | 0 | 151.1938 | |
| ## 14 | 242.7306 | 0 | 158.0803 | |
| ## 15 | 403.6405 | 0 | 249.7694 | |
| ## 16 | 626.8373 | 0 | 426.9397 | |
| ## 17 | 184.4927 | 0 | 124.8518 | |
| ## 18 | 561.4198 | 0 | 318.5436 | |

| | | | |
|-------|------------------------------|----------------------|-----------------------|
| ## 19 | 430.6817 | 0 | 423.6295 |
| ## 20 | 2193.4483 | 1 | 1171.9766 |
| ## | EqEllipseProlateVolStatusCh1 | EqSphereAreaCh1 | EqSphereAreaStatusCh1 |
| ## 1 | 0 | 742.1156 | 0 |
| ## 2 | 1 | 3278.7256 | 1 |
| ## 3 | 0 | 1727.4104 | 0 |
| ## 4 | 0 | 1194.9320 | 0 |
| ## 5 | 0 | 1140.4263 | 0 |
| ## 6 | 0 | 691.8027 | 0 |
| ## 7 | 0 | 708.5737 | 0 |
| ## 8 | 0 | 1006.2585 | 0 |
| ## 9 | 0 | 1983.1678 | 0 |
| ## 10 | 0 | 1538.7370 | 0 |
| ## 11 | 0 | 1698.0612 | 0 |
| ## 12 | 0 | 1027.2222 | 0 |
| ## 13 | 0 | 947.5601 | 0 |
| ## 14 | 0 | 750.5011 | 0 |
| ## 15 | 0 | 1035.6077 | 0 |
| ## 16 | 0 | 1433.9184 | 0 |
| ## 17 | 0 | 633.1043 | 0 |
| ## 18 | 0 | 1253.6304 | 0 |
| ## 19 | 0 | 1262.0159 | 0 |
| ## 20 | 1 | 3048.1247 | 1 |
| ## | EqSphereVolCh1 | EqSphereVolStatusCh1 | FiberAlign2Ch3 |
| ## 1 | 1900.996 | 0 | 0.0000000 |
| ## 2 | 17653.525 | 1 | 0.4879354 |
| ## 3 | 6750.985 | 0 | 0.3005220 |
| ## 4 | 3884.084 | 0 | 0.2204239 |
| ## 5 | 3621.385 | 0 | 0.4913869 |
| ## 6 | 1710.989 | 0 | 0.3095039 |
| ## 7 | 1773.582 | 0 | 0.2449735 |
| ## 8 | 3001.499 | 0 | 0.0000000 |
| ## 9 | 8304.491 | 0 | 0.3552463 |
| ## 10 | 5675.713 | 0 | 0.7624199 |
| ## 11 | 6579.666 | 0 | 0.5247012 |
| ## 12 | 3095.782 | 0 | 0.3642360 |
| ## 13 | 2742.736 | 0 | 0.0000000 |
| ## 14 | 1933.307 | 0 | 0.6259856 |
| ## 15 | 3133.767 | 0 | 0.3594184 |
| ## 16 | 5105.761 | 0 | 0.4789638 |
| ## 17 | 1497.913 | 0 | 0.7245099 |
| ## 18 | 4173.766 | 0 | 0.4876284 |
| ## 19 | 4215.713 | 0 | 0.2735832 |
| ## 20 | 15824.242 | 1 | 0.4549632 |
| ## | FiberAlign2StatusCh3 | FiberAlign2StatusCh4 | FiberLengthCh1 |
| ## 1 | 2 | 2 | 26.98132 |
| ## 2 | 0 | 0 | 64.28230 |
| ## 3 | 0 | 0 | 21.14115 |
| ## 4 | 0 | 1 | 43.14112 |
| ## 5 | 0 | 0 | 34.74977 |
| ## 6 | 0 | 0 | 12.38420 |
| ## 7 | 0 | 0 | 12.59624 |
| ## 8 | 2 | 2 | 37.84863 |
| ## 9 | 0 | 0 | 44.14773 |

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|-------|----------------------|-----------------------|----------------------------|-----------------|
| ## 10 | 1 | 1 | 37.24447 | |
| ## 11 | 0 | 0 | 20.24908 | |
| ## 12 | 0 | 0 | 22.29743 | |
| ## 13 | 2 | 2 | 43.64276 | |
| ## 14 | 1 | 1 | 13.70788 | |
| ## 15 | 0 | 0 | 26.52856 | |
| ## 16 | 0 | 0 | 25.12696 | |
| ## 17 | 1 | 1 | 27.63249 | |
| ## 18 | 0 | 0 | 18.68085 | |
| ## 19 | 0 | 2 | 16.75358 | |
| ## 20 | 0 | 0 | 55.03983 | |
| ## | FiberLengthStatusCh1 | FiberWidthCh1 | FiberWidthStatusCh1 | IntenCoocASMCh3 |
| ## 1 | 0 | 7.410365 | 2 | 0.011183899 |
| ## 2 | 1 | 13.167079 | 0 | 0.028051061 |
| ## 3 | 0 | 21.141150 | 1 | 0.006862315 |
| ## 4 | 0 | 7.404412 | 2 | 0.030962071 |
| ## 5 | 0 | 8.483574 | 2 | 0.022768411 |
| ## 6 | 0 | 12.384199 | 0 | 0.008914083 |
| ## 7 | 0 | 12.596237 | 0 | 0.008166438 |
| ## 8 | 0 | 7.117398 | 2 | 0.006116300 |
| ## 9 | 0 | 11.728889 | 0 | 0.089600535 |
| ## 10 | 0 | 10.792994 | 0 | 0.012277917 |
| ## 11 | 0 | 20.249084 | 1 | 0.095391688 |
| ## 12 | 0 | 12.057876 | 0 | 0.108161298 |
| ## 13 | 0 | 5.806209 | 2 | 0.005475966 |
| ## 14 | 0 | 13.707875 | 0 | 0.009545651 |
| ## 15 | 0 | 10.174239 | 0 | 0.013034881 |
| ## 16 | 0 | 14.610886 | 0 | 0.024982639 |
| ## 17 | 0 | 6.050330 | 2 | 0.012319717 |
| ## 18 | 0 | 17.422216 | 1 | 0.022057270 |
| ## 19 | 0 | 16.753579 | 1 | 0.033156657 |
| ## 20 | 1 | 14.306907 | 0 | 0.013880529 |
| ## | IntenCoocASMCh4 | IntenCoocASMStatusCh3 | IntenCoocASMStatusCh4 | |
| ## 1 | 0.050448005 | 0 | 0 | |
| ## 2 | 0.012594975 | 0 | 0 | |
| ## 3 | 0.006141691 | 0 | 0 | |
| ## 4 | 0.011033195 | 0 | 0 | |
| ## 5 | 0.079690842 | 0 | 0 | |
| ## 6 | 0.009155300 | 0 | 0 | |
| ## 7 | 0.008311984 | 0 | 0 | |
| ## 8 | 0.127326686 | 0 | 0 | |
| ## 9 | 0.019221940 | 0 | 0 | |
| ## 10 | 0.008028741 | 0 | 0 | |
| ## 11 | 0.093600421 | 0 | 0 | |
| ## 12 | 0.009951830 | 0 | 0 | |
| ## 13 | 0.103488064 | 0 | 0 | |
| ## 14 | 0.008644145 | 0 | 0 | |
| ## 15 | 0.008962990 | 0 | 0 | |
| ## 16 | 0.044225099 | 0 | 0 | |
| ## 17 | 0.020887482 | 0 | 0 | |
| ## 18 | 0.010350158 | 0 | 0 | |
| ## 19 | 0.009490846 | 0 | 0 | |
| ## 20 | 0.038698122 | 0 | 0 | |
| ## | IntenCoocContrastCh3 | IntenCoocContrastCh4 | IntenCoocContrastStatusCh3 | |

| | | | |
|-------|----------------------------|---------------------------|---------------------|
| ## 1 | 40.751777 | 13.895439 | 1 |
| ## 2 | 8.227953 | 6.984046 | 0 |
| ## 3 | 14.446074 | 16.700843 | 0 |
| ## 4 | 7.299457 | 13.390884 | 0 |
| ## 5 | 15.854691 | 3.538895 | 0 |
| ## 6 | 8.248344 | 9.867550 | 0 |
| ## 7 | 12.881127 | 17.409342 | 0 |
| ## 8 | 32.702178 | 8.833734 | 1 |
| ## 9 | 9.943647 | 9.568663 | 0 |
| ## 10 | 18.139723 | 12.330637 | 1 |
| ## 11 | 4.302937 | 3.941435 | 0 |
| ## 12 | 6.160585 | 10.590685 | 0 |
| ## 13 | 38.771563 | 11.704375 | 1 |
| ## 14 | 16.421096 | 16.470915 | 1 |
| ## 15 | 9.402056 | 10.301528 | 0 |
| ## 16 | 11.841077 | 8.084776 | 0 |
| ## 17 | 11.913578 | 6.154895 | 0 |
| ## 18 | 4.818421 | 5.870657 | 0 |
| ## 19 | 5.696809 | 8.225177 | 0 |
| ## 20 | 7.776731 | 3.789387 | 0 |
| ## | IntenCoocContrastStatusCh4 | IntenCoocEntropyCh3 | IntenCoocEntropyCh4 |
| ## 1 | 1 | 7.199458 | 5.249744 |
| ## 2 | 0 | 6.822138 | 7.098988 |
| ## 3 | 1 | 7.580100 | 7.671478 |
| ## 4 | 1 | 6.312641 | 7.197026 |
| ## 5 | 2 | 6.778093 | 5.502166 |
| ## 6 | 0 | 7.016333 | 6.967390 |
| ## 7 | 1 | 7.172120 | 7.142348 |
| ## 8 | 0 | 7.557944 | 4.177876 |
| ## 9 | 0 | 5.711782 | 6.960814 |
| ## 10 | 1 | 7.309512 | 7.406901 |
| ## 11 | 2 | 5.518710 | 5.550419 |
| ## 12 | 0 | 5.044067 | 7.126900 |
| ## 13 | 1 | 7.739265 | 4.512367 |
| ## 14 | 1 | 7.236606 | 7.142452 |
| ## 15 | 0 | 6.957487 | 7.144784 |
| ## 16 | 0 | 6.794864 | 6.361635 |
| ## 17 | 0 | 6.796920 | 6.485465 |
| ## 18 | 0 | 6.546452 | 7.087970 |
| ## 19 | 0 | 6.390147 | 7.161842 |
| ## 20 | 2 | 7.096212 | 6.233320 |
| ## | IntenCoocEntropyStatusCh3 | IntenCoocEntropyStatusCh4 | IntenCoocMaxCh3 |
| ## 1 | 0 | 0 | 0.07741935 |
| ## 2 | 0 | 0 | 0.15321477 |
| ## 3 | 1 | 1 | 0.02835052 |
| ## 4 | 0 | 0 | 0.16279070 |
| ## 5 | 0 | 0 | 0.12741313 |
| ## 6 | 0 | 0 | 0.01986755 |
| ## 7 | 0 | 0 | 0.01948052 |
| ## 8 | 1 | 2 | 0.01643192 |
| ## 9 | 0 | 0 | 0.29519451 |
| ## 10 | 1 | 1 | 0.08630952 |
| ## 11 | 0 | 0 | 0.30079156 |
| ## 12 | 0 | 0 | 0.31531532 |

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|-------|--------------------|-----------------------|-----------------------|----------------------------|--------------|
| ## 13 | | 1 | | 2 | 0.02083333 |
| ## 14 | | 0 | | 0 | 0.06586826 |
| ## 15 | | 0 | | 0 | 0.07391304 |
| ## 16 | | 0 | | 0 | 0.14012739 |
| ## 17 | | 0 | | 0 | 0.06818182 |
| ## 18 | | 0 | | 0 | 0.12727273 |
| ## 19 | | 0 | | 0 | 0.16312057 |
| ## 20 | | 0 | | 0 | 0.08333333 |
| ## | IntenCoocMaxCh4 | IntenCoocMaxStatusCh3 | IntenCoocMaxStatusCh4 | KurtIntenCh1 | |
| ## 1 | 0.17197452 | 0 | | 0 | -0.656744087 |
| ## 2 | 0.07387141 | 0 | | 0 | -0.248769067 |
| ## 3 | 0.02319588 | 2 | | 2 | -0.293484630 |
| ## 4 | 0.07751938 | 0 | | 0 | 0.625856124 |
| ## 5 | 0.27848101 | 0 | | 0 | 0.042106843 |
| ## 6 | 0.01986755 | 2 | | 2 | -0.196841369 |
| ## 7 | 0.01948052 | 2 | | 2 | 0.250897694 |
| ## 8 | 0.34123223 | 2 | | 1 | 0.112969507 |
| ## 9 | 0.11212815 | 0 | | 0 | -0.219208563 |
| ## 10 | 0.04705882 | 0 | | 0 | 0.144554913 |
| ## 11 | 0.30079156 | 0 | | 0 | -0.625392634 |
| ## 12 | 0.05855856 | 0 | | 0 | -0.364691438 |
| ## 13 | 0.29500000 | 2 | | 0 | -0.001236013 |
| ## 14 | 0.03592814 | 0 | | 0 | -0.930684432 |
| ## 15 | 0.03478261 | 0 | | 0 | -0.556261231 |
| ## 16 | 0.19938650 | 0 | | 0 | -0.261787171 |
| ## 17 | 0.11363636 | 0 | | 0 | -0.365908061 |
| ## 18 | 0.05454545 | 0 | | 0 | -0.370863990 |
| ## 19 | 0.04609929 | 0 | | 0 | 38.974284980 |
| ## 20 | 0.17903930 | 0 | | 0 | 0.230696287 |
| ## | KurtIntenCh3 | KurtIntenCh4 | KurtIntenStatusCh1 | KurtIntenStatusCh3 | |
| ## 1 | -0.60805827 | 0.7258145 | 0 | 0 | |
| ## 2 | -0.33078390 | -0.2652638 | 0 | 0 | |
| ## 3 | 1.05128134 | 0.1506140 | 0 | 0 | |
| ## 4 | 0.12774061 | -0.3472936 | 0 | 0 | |
| ## 5 | 0.95229003 | -0.1954188 | 0 | 0 | |
| ## 6 | -0.86079937 | -0.4050669 | 0 | 0 | |
| ## 7 | -0.59831464 | -0.8256696 | 0 | 0 | |
| ## 8 | -0.74863969 | 2.0335593 | 0 | 0 | |
| ## 9 | 1.88482515 | -0.7610117 | 0 | 0 | |
| ## 10 | -0.56811998 | 0.1918431 | 0 | 0 | |
| ## 11 | -0.35835018 | -0.5315540 | 0 | 0 | |
| ## 12 | 1.08333286 | -0.6262704 | 0 | 0 | |
| ## 13 | -0.09820689 | 0.2876581 | 0 | 0 | |
| ## 14 | -0.58857201 | 6.7741213 | 0 | 0 | |
| ## 15 | -0.51231861 | -0.6471575 | 0 | 0 | |
| ## 16 | 0.80950143 | 0.7286793 | 0 | 0 | |
| ## 17 | -0.27882467 | -0.7073985 | 0 | 0 | |
| ## 18 | -0.42094795 | -0.4443304 | 0 | 0 | |
| ## 19 | -1.19024918 | -1.3328752 | 1 | 0 | |
| ## 20 | 5.19308674 | 2.7868190 | 0 | 1 | |
| ## | KurtIntenStatusCh4 | LengthCh1 | LengthStatusCh1 | MemberAvgAvgIntenStatusCh2 | |
| ## 1 | 0 | 26.20779 | 0 | 0 | |
| ## 2 | 0 | 47.21855 | 1 | 0 | |
| ## 3 | 0 | 28.14303 | 0 | 0 | |

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|-------|------------------------------|--------------------------|--------------------------|---|
| ## 4 | 0 | 37.85957 | 0 | 0 |
| ## 5 | 0 | 35.99415 | 0 | 0 |
| ## 6 | 0 | 16.20494 | 2 | 0 |
| ## 7 | 0 | 15.91843 | 2 | 0 |
| ## 8 | 0 | 29.70812 | 0 | 0 |
| ## 9 | 0 | 32.37446 | 0 | 0 |
| ## 10 | 0 | 31.13589 | 0 | 0 |
| ## 11 | 0 | 28.28436 | 0 | 0 |
| ## 12 | 0 | 23.05547 | 0 | 0 |
| ## 13 | 0 | 34.75543 | 0 | 0 |
| ## 14 | 1 | 18.71589 | 0 | 0 |
| ## 15 | 0 | 26.31960 | 0 | 0 |
| ## 16 | 0 | 27.80421 | 0 | 0 |
| ## 17 | 0 | 20.56691 | 0 | 0 |
| ## 18 | 0 | 26.61822 | 0 | 0 |
| ## 19 | 0 | 21.28784 | 0 | 0 |
| ## 20 | 0 | 43.93737 | 1 | 0 |
| ## | MemberAvgTotalIntenStatusCh2 | NeighborAvgDistCh1 | NeighborAvgDistStatusCh1 | |
| ## 1 | 0 | 370.4543 | 1 | |
| ## 2 | 0 | 174.4442 | 2 | |
| ## 3 | 0 | 158.4774 | 2 | |
| ## 4 | 0 | 206.3344 | 0 | |
| ## 5 | 0 | 204.9213 | 0 | |
| ## 6 | 0 | 231.9933 | 0 | |
| ## 7 | 0 | 230.2935 | 0 | |
| ## 8 | 0 | 218.2644 | 0 | |
| ## 9 | 0 | 235.0062 | 0 | |
| ## 10 | 0 | 220.1542 | 0 | |
| ## 11 | 0 | 334.7284 | 1 | |
| ## 12 | 0 | 263.6345 | 0 | |
| ## 13 | 0 | 204.2607 | 0 | |
| ## 14 | 0 | 190.9553 | 2 | |
| ## 15 | 0 | 231.1529 | 0 | |
| ## 16 | 0 | 213.9487 | 0 | |
| ## 17 | 0 | 228.9701 | 0 | |
| ## 18 | 0 | 314.1311 | 1 | |
| ## 19 | 0 | 288.1973 | 0 | |
| ## 20 | 0 | 213.4731 | 0 | |
| ## | NeighborMinDistCh1 | NeighborMinDistStatusCh1 | NeighborVarDistCh1 | |
| ## 1 | 99.10349 | 1 | 127.96080 | |
| ## 2 | 30.11114 | 0 | 81.38063 | |
| ## 3 | 34.94477 | 0 | 90.43768 | |
| ## 4 | 33.08030 | 0 | 116.89276 | |
| ## 5 | 27.02935 | 0 | 110.96662 | |
| ## 6 | 15.92674 | 0 | 95.96952 | |
| ## 7 | 15.92674 | 0 | 94.81307 | |
| ## 8 | 29.31744 | 0 | 105.76881 | |
| ## 9 | 38.43038 | 0 | 79.58161 | |
| ## 10 | 36.18265 | 0 | 73.85064 | |
| ## 11 | 28.66667 | 0 | 130.46044 | |
| ## 12 | 38.43038 | 0 | 88.50411 | |
| ## 13 | 29.31744 | 0 | 78.31229 | |
| ## 14 | 23.90107 | 0 | 75.44312 | |
| ## 15 | 29.84892 | 0 | 103.45812 | |

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|-------|--------------------------|--------------|-------------------|--------------|--------------------|
| ## 16 | 11.44654 | 2 | 105.15059 | | |
| ## 17 | 17.40476 | 0 | 112.56123 | | |
| ## 18 | 22.73226 | 0 | 126.52233 | | |
| ## 19 | 49.92581 | 1 | 119.63999 | | |
| ## 20 | 46.97267 | 1 | 127.92813 | | |
| ## | NeighborVarDistStatusCh1 | PerimCh1 | PerimStatusCh1 | ShapeBFRCh1 | |
| ## 1 | 0 | 68.78338 | 0 | 0.6651480 | |
| ## 2 | 2 | 154.89876 | 1 | 0.5397584 | |
| ## 3 | 2 | 84.56460 | 0 | 0.7243116 | |
| ## 4 | 0 | 101.09107 | 0 | 0.5891625 | |
| ## 5 | 0 | 86.46670 | 0 | 0.6001347 | |
| ## 6 | 0 | 49.53679 | 0 | 0.6869375 | |
| ## 7 | 0 | 50.38495 | 0 | 0.7468435 | |
| ## 8 | 0 | 89.93206 | 0 | 0.6858659 | |
| ## 9 | 2 | 111.75324 | 0 | 0.6318210 | |
| ## 10 | 2 | 96.07492 | 0 | 0.6738027 | |
| ## 11 | 0 | 80.99633 | 0 | 0.7574574 | |
| ## 12 | 2 | 68.71062 | 0 | 0.6347914 | |
| ## 13 | 2 | 98.89793 | 0 | 0.5453331 | |
| ## 14 | 2 | 54.83150 | 0 | 0.7968360 | |
| ## 15 | 0 | 73.40559 | 0 | 0.5570022 | |
| ## 16 | 0 | 79.47569 | 0 | 0.6635424 | |
| ## 17 | 0 | 67.36563 | 0 | 0.4423808 | |
| ## 18 | 0 | 72.20612 | 0 | 0.7609989 | |
| ## 19 | 0 | 67.01432 | 0 | 0.7161740 | |
| ## 20 | 0 | 138.69347 | 1 | 0.6034935 | |
| ## | ShapeBFRStatusCh1 | ShapeLWRCh1 | ShapeLWRStatusCh1 | ShapeP2ACh1 | |
| ## 1 | 0 | 2.462450 | 0 | 1.883006 | |
| ## 2 | 2 | 1.468181 | 0 | 2.255810 | |
| ## 3 | 1 | 1.328408 | 0 | 1.272193 | |
| ## 4 | 0 | 2.826854 | 1 | 2.545840 | |
| ## 5 | 0 | 2.727127 | 1 | 2.018155 | |
| ## 6 | 0 | 1.043013 | 0 | 1.086278 | |
| ## 7 | 1 | 1.068329 | 0 | 1.091958 | |
| ## 8 | 0 | 2.406246 | 0 | 2.389160 | |
| ## 9 | 0 | 1.335671 | 0 | 1.919299 | |
| ## 10 | 0 | 1.698051 | 0 | 1.827275 | |
| ## 11 | 1 | 1.427439 | 0 | 1.197255 | |
| ## 12 | 0 | 1.313937 | 0 | 1.397364 | |
| ## 13 | 2 | 2.780739 | 1 | 3.071547 | |
| ## 14 | 1 | 1.487642 | 0 | 1.237124 | |
| ## 15 | 2 | 1.490323 | 0 | 1.588658 | |
| ## 16 | 0 | 1.430953 | 0 | 1.369114 | |
| ## 17 | 2 | 1.182277 | 0 | 2.160058 | |
| ## 18 | 1 | 1.720413 | 0 | 1.274781 | |
| ## 19 | 0 | 1.028672 | 0 | 1.099823 | |
| ## 20 | 0 | 1.528861 | 0 | 1.943913 | |
| ## | ShapeP2AStatusCh1 | SkewIntenCh1 | SkewIntenCh3 | SkewIntenCh4 | SkewIntenStatusCh1 |
| ## 1 | 0 | 0.45450484 | 0.46039340 | 1.2327736 | 0 |
| ## 2 | 0 | 0.39870467 | 0.61973079 | 0.5272631 | 0 |
| ## 3 | 0 | 0.47248709 | 0.97137879 | 0.3247065 | 0 |
| ## 4 | 1 | 0.88167138 | 0.99989280 | 0.6044399 | 1 |
| ## 5 | 0 | 0.51702723 | 1.17675337 | 0.9258123 | 0 |
| ## 6 | 0 | -0.70259526 | 0.15320233 | 0.6511667 | 2 |

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|-------|-------------------------|-------------------------|-------------------|-------------------|---|
| ## 7 | 0 | -0.81606076 | -0.22448186 | 0.1023062 | 2 |
| ## 8 | 1 | 0.66550394 | 0.01142938 | 1.6120078 | 0 |
| ## 9 | 0 | 0.59674364 | 1.51738349 | 0.4218934 | 0 |
| ## 10 | 0 | 0.58681762 | 0.40825422 | 0.6756981 | 0 |
| ## 11 | 0 | -0.56913821 | 0.76536204 | 0.8186378 | 2 |
| ## 12 | 0 | 0.54673722 | 1.43236317 | 0.7038007 | 0 |
| ## 13 | 1 | 0.63116945 | 0.18603538 | 1.2101592 | 0 |
| ## 14 | 0 | -0.08283974 | 0.08641559 | 2.0018820 | 0 |
| ## 15 | 0 | 0.44309238 | 0.55567549 | 0.1372252 | 0 |
| ## 16 | 0 | 0.25651828 | 1.07109399 | 1.0454462 | 0 |
| ## 17 | 0 | 0.41807947 | 0.88133481 | 0.7258839 | 0 |
| ## 18 | 0 | 0.27550348 | 0.77003268 | 0.6839946 | 0 |
| ## 19 | 0 | 5.27421649 | 0.17712659 | 0.2239078 | 1 |
| ## 20 | 0 | 0.66893477 | 1.93614592 | 1.6299375 | 0 |
| ## | SkewIntenStatusCh3 | SkewIntenStatusCh4 | SpotFiberCountCh3 | SpotFiberCountCh4 | |
| ## 1 | 0 | 0 | 1 | 4 | |
| ## 2 | 0 | 0 | 4 | 11 | |
| ## 3 | 0 | 0 | 2 | 6 | |
| ## 4 | 0 | 0 | 4 | 7 | |
| ## 5 | 0 | 0 | 1 | 7 | |
| ## 6 | 2 | 0 | 1 | 4 | |
| ## 7 | 2 | 0 | 0 | 4 | |
| ## 8 | 2 | 1 | 2 | 7 | |
| ## 9 | 0 | 0 | 1 | 11 | |
| ## 10 | 0 | 0 | 1 | 7 | |
| ## 11 | 0 | 0 | 1 | 4 | |
| ## 12 | 0 | 0 | 0 | 5 | |
| ## 13 | 2 | 0 | 0 | 6 | |
| ## 14 | 2 | 1 | 2 | 4 | |
| ## 15 | 0 | 0 | 1 | 4 | |
| ## 16 | 0 | 0 | 1 | 5 | |
| ## 17 | 0 | 0 | 4 | 4 | |
| ## 18 | 0 | 0 | 1 | 2 | |
| ## 19 | 2 | 0 | 2 | 2 | |
| ## 20 | 1 | 1 | 2 | 12 | |
| ## | SpotFiberCountStatusCh3 | SpotFiberCountStatusCh4 | TotalIntenCh1 | TotalIntenCh2 | |
| ## 1 | 0 | 0 | 2781 | 700 | |
| ## 2 | 1 | 1 | 24964 | 160997 | |
| ## 3 | 0 | 0 | 11552 | 47510 | |
| ## 4 | 1 | 0 | 5545 | 28869 | |
| ## 5 | 0 | 0 | 6603 | 30305 | |
| ## 6 | 0 | 0 | 53779 | 107680 | |
| ## 7 | 2 | 0 | 43950 | 100626 | |
| ## 8 | 0 | 0 | 4401 | 1135 | |
| ## 9 | 0 | 1 | 7593 | 41852 | |
| ## 10 | 0 | 0 | 6512 | 32638 | |
| ## 11 | 0 | 0 | 70614 | 157313 | |
| ## 12 | 2 | 0 | 4613 | 30855 | |
| ## 13 | 2 | 0 | 4111 | 1204 | |
| ## 14 | 0 | 0 | 7203 | 38161 | |
| ## 15 | 0 | 0 | 4340 | 30719 | |
| ## 16 | 0 | 0 | 14461 | 74259 | |
| ## 17 | 1 | 0 | 4743 | 15434 | |
| ## 18 | 0 | 0 | 64408 | 170105 | |

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|-------|---------------------|---------------------|---------------------|---------------------|
| ## 19 | 0 | 0 | 88725 | 148012 |
| ## 20 | 0 | 1 | 25911 | 119024 |
| ## | TotalIntenCh3 | TotalIntenCh4 | TotalIntenStatusCh1 | TotalIntenStatusCh2 |
| ## 1 | 1690 | 392 | 0 | 2 |
| ## 2 | 54675 | 128368 | 0 | 1 |
| ## 3 | 26344 | 43959 | 0 | 0 |
| ## 4 | 8042 | 8843 | 0 | 0 |
| ## 5 | 5569 | 11037 | 0 | 0 |
| ## 6 | 21234 | 57231 | 1 | 1 |
| ## 7 | 20929 | 46187 | 0 | 1 |
| ## 8 | 4136 | 373 | 0 | 2 |
| ## 9 | 6488 | 24325 | 0 | 0 |
| ## 10 | 7503 | 23162 | 0 | 0 |
| ## 11 | 15699 | 51325 | 1 | 1 |
| ## 12 | 3332 | 11466 | 0 | 0 |
| ## 13 | 3862 | 432 | 0 | 2 |
| ## 14 | 7911 | 24389 | 0 | 0 |
| ## 15 | 5548 | 17588 | 0 | 0 |
| ## 16 | 14474 | 23099 | 0 | 0 |
| ## 17 | 6265 | 17534 | 0 | 2 |
| ## 18 | 45045 | 113205 | 1 | 1 |
| ## 19 | 58224 | 120536 | 1 | 1 |
| ## 20 | 56359 | 154931 | 0 | 1 |
| ## | TotalIntenStatusCh3 | TotalIntenStatusCh4 | VarIntenCh1 | VarIntenCh3 |
| ## 1 | 0 | 2 | 12.47468 | 7.609035 |
| ## 2 | 1 | 1 | 18.80923 | 56.715352 |
| ## 3 | 0 | 0 | 17.29564 | 37.671053 |
| ## 4 | 0 | 2 | 13.81897 | 30.005643 |
| ## 5 | 0 | 0 | 15.40797 | 20.504288 |
| ## 6 | 0 | 0 | 115.00462 | 70.918444 |
| ## 7 | 0 | 0 | 93.38827 | 47.293662 |
| ## 8 | 0 | 2 | 12.41264 | 9.872296 |
| ## 9 | 0 | 0 | 11.79258 | 17.923023 |
| ## 10 | 0 | 0 | 11.46656 | 14.866582 |
| ## 11 | 0 | 0 | 67.84403 | 40.929727 |
| ## 12 | 0 | 0 | 13.92294 | 18.643027 |
| ## 13 | 0 | 2 | 12.26925 | 9.362472 |
| ## 14 | 0 | 0 | 21.24513 | 23.862052 |
| ## 15 | 0 | 0 | 12.32497 | 17.747143 |
| ## 16 | 0 | 0 | 20.95648 | 42.316360 |
| ## 17 | 0 | 0 | 20.58588 | 40.653090 |
| ## 18 | 1 | 1 | 100.07556 | 138.203909 |
| ## 19 | 1 | 1 | 201.36869 | 152.312103 |
| ## 20 | 1 | 1 | 21.19878 | 81.758387 |
| ## | VarIntenStatusCh1 | VarIntenStatusCh3 | VarIntenStatusCh4 | WidthCh1 |
| ## 1 | 0 | 2 | 2 | 10.64297 |
| ## 2 | 0 | 0 | 0 | 32.16126 |
| ## 3 | 0 | 0 | 0 | 21.18553 |
| ## 4 | 0 | 0 | 2 | 13.39283 |
| ## 5 | 0 | 0 | 0 | 13.19856 |
| ## 6 | 1 | 0 | 0 | 15.53666 |
| ## 7 | 1 | 0 | 0 | 14.90031 |
| ## 8 | 0 | 2 | 2 | 12.34625 |
| ## 9 | 0 | 0 | 0 | 24.23834 |

```

## 10          0          2          2 18.33625
## 11          0          0          0 19.81476
## 12          0          0          2 17.54686
## 13          0          2          2 12.49863
## 14          0          0          0 12.58091
## 15          0          0          2 17.66034
## 16          0          0          0 19.43055
## 17          0          0          0 17.39602
## 18          1          1          1 15.47200
## 19          1          1          1 20.69449
## 20          0          0          1 28.73863
##      WidthStatusCh1 XCentroid YCentroid
## 1          2         42         14
## 2          1        215        347
## 3          0        371        252
## 4          0        487        295
## 5          0        283        159
## 6          0        191        127
## 7          0        180        138
## 8          2        373        181
## 9          1        236        467
## 10         0        303        468
## 11         0        473         15
## 12         0        211        495
## 13         2        347        252
## 14         2        229        277
## 15         0        172        207
## 16         0        276        385
## 17         0        239        404
## 18         0        466        449
## 19         0         95         95
## 20         1         77        344

```

```

set.seed(125)

# split train and test
library(dplyr)
training <- dplyr::filter(seg1, Case == "Train")
testing <- dplyr::filter(seg1, Case == "Test")

# fit CART model with rpart
fit <- train(Class ~ ., method="rpart", data=training)

fit$finalModel

```

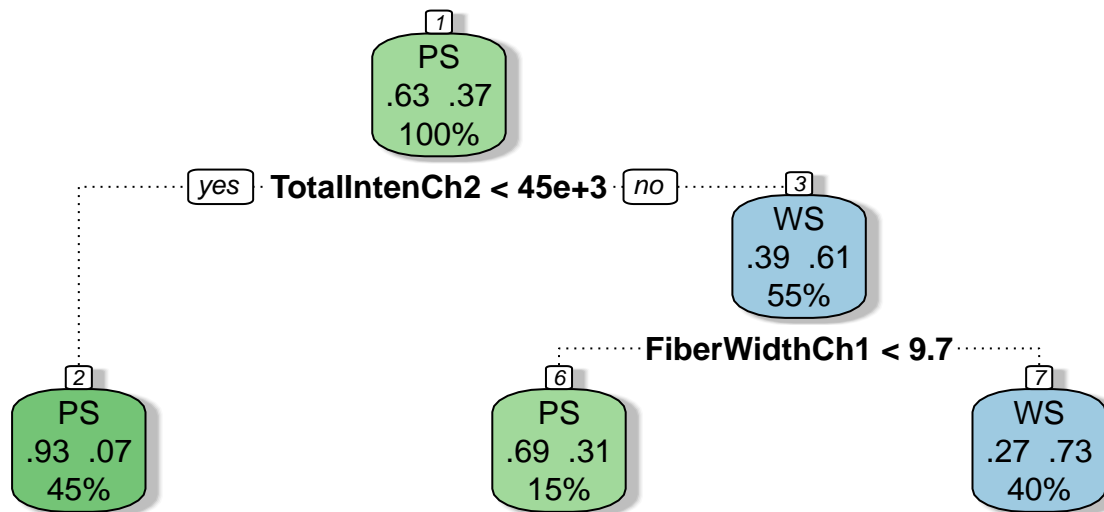
```

## n= 1009
##
## node), split, n, loss, yval, (yprob)
##      * denotes terminal node
##
## 1) root 1009 373 PS (0.63032706 0.36967294)
##   2) TotalIntenCh2< 45323.5 454 34 PS (0.92511013 0.07488987) *
##   3) TotalIntenCh2>=45323.5 555 216 WS (0.38918919 0.61081081)
##   6) FiberWidthCh1< 9.673245 154 47 PS (0.69480519 0.30519481) *

```

```
##      7) FiberWidthCh1>=9.673245 401 109 WS (0.27182045 0.72817955) *
```

```
fancyRpartPlot(fit$finalModel)
```



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Practical example 2

```
# Set seed for reproducibility
set.seed(1111)
```

Data processing- In this section the data is downloaded and manipulated. Some basic transformations

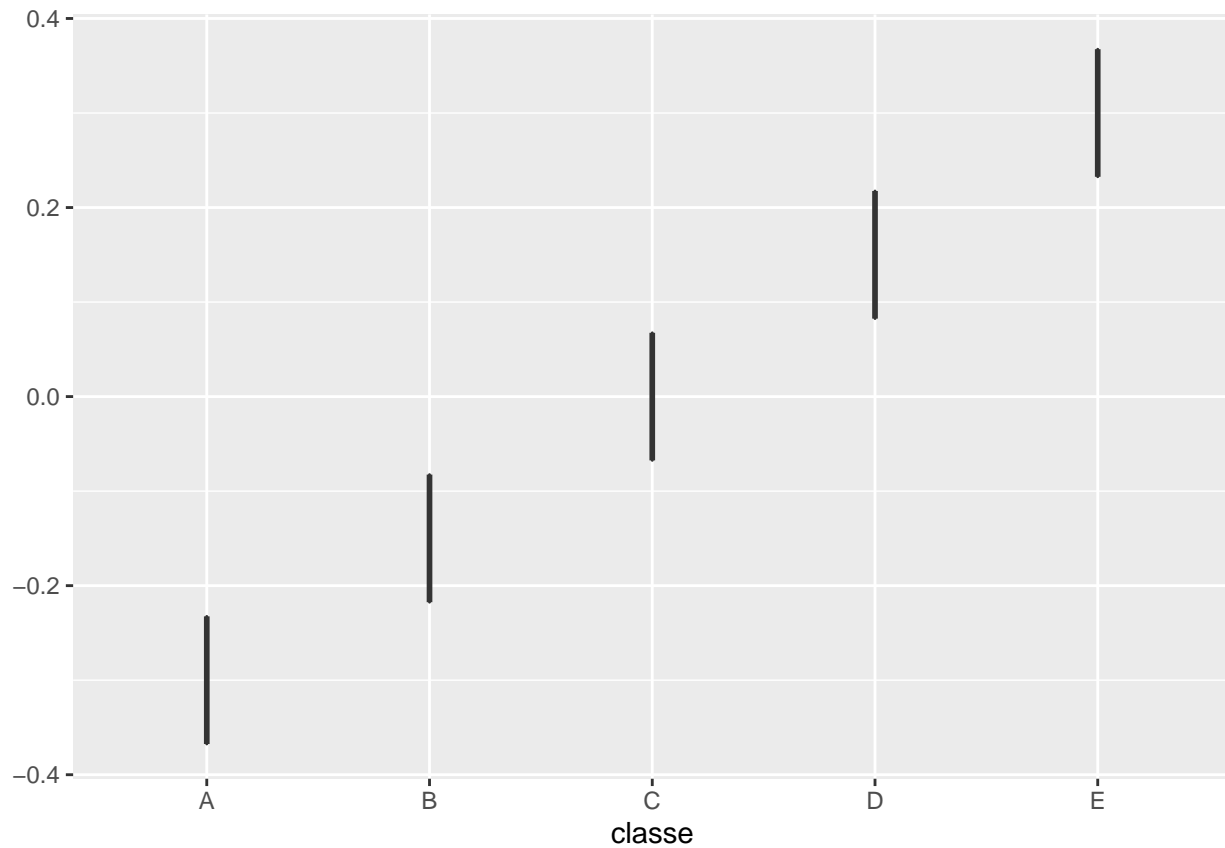
```
training <- read.csv("./data/pml-training.csv", na.strings=c("NA", "#DIV/0!", "")) ##Cleaning
testing <- read.csv("./data/pml-testing.csv", na.strings=c("NA", "#DIV/0!", ""))
training <- training[, colSums(is.na(training)) == 0]
testing <- testing[, colSums(is.na(testing)) == 0]
```

```
training <- training[, -c(1:7)] ##Subset for just the columns of interest.
testing <- testing[, -c(1:7)]
```

Cross-validation: In this section cross-validation will be performed by splitting the training data into

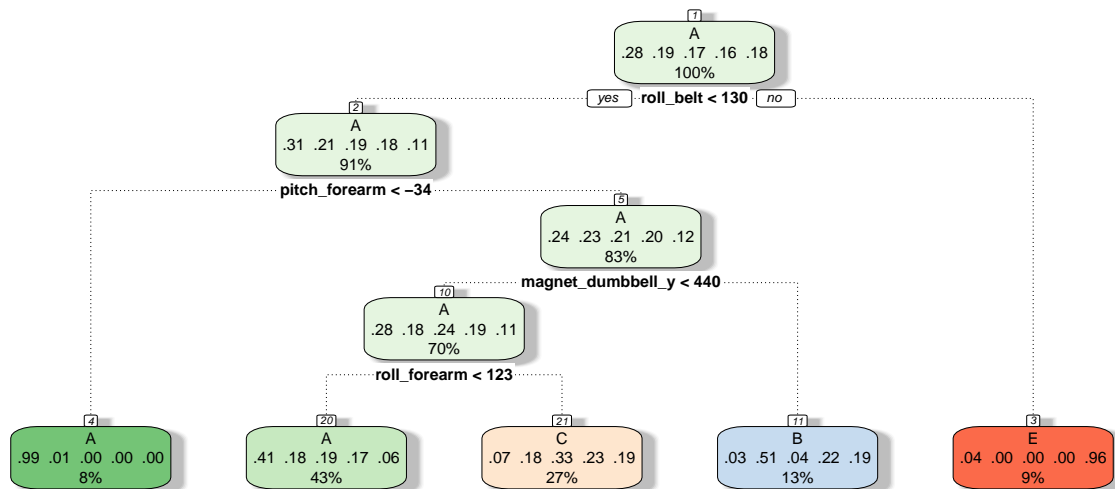
```
library(caret)
subSamples <- createDataPartition(y=training$classe, p=0.75, list=FALSE)
subTraining <- training[subSamples, ]
subTesting <- training[-subSamples, ]
```

```
library(ggplot2) ##some exploration
p1 <- ggplot(subTraining, aes(classe))
plot <- p1 + geom_boxplot()
plot
```



```
## Prediction models: In this section a decision tree and random forest will be used on the data.
library(rattle)
# Fit model
modFitDT <- train(classe ~ ., method="rpart", data=subTraining)
# Perform prediction
predictDT <- predict(modFitDT, subTesting, type = "prob")
# Plot result

fancyRpartPlot(modFitDT$finalModel)
```



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##This Random forest takes 1.5hrs to run, be careful!!!!!!!!!!

Fit model

```
library(caret)
```

```
library(randomForest)
```

```
modFit <- train(classe~ .,data=subTraining,method="rf")
```

```
modFit
```

```
## Random Forest
```

```
##
```

```
## 14718 samples
```

```
## 52 predictor
```

```
## 5 classes: 'A', 'B', 'C', 'D', 'E'
```

```
##
```

```
## No pre-processing
```

```
## Resampling: Bootstrapped (25 reps)
```

```
## Summary of sample sizes: 14718, 14718, 14718, 14718, 14718, ...
```

```
## Resampling results across tuning parameters:
```

```
##
```

```
## mtry Accuracy Kappa
```

```
## 2 0.9883892 0.9853054
```

```
## 27 0.9881484 0.9850017
```

```
## 52 0.9802377 0.9749918
```

```
##
```

```
## Accuracy was used to select the optimal model using the largest value.
```

```
## The final value used for the model was mtry = 2.
```

```
gTree <- getTree(modFit$finalModel,k=3) ##to view the individual trees, you can use this function, which
head(gTree, n=30)
```

```
##      left daughter right daughter split var split point status prediction
## 1         2         3         21 -270.50000      1          0
## 2         4         5         51 -247.50000      1          0
## 3         6         7         41  -32.95000      1          0
## 4         8         9         28   24.55794      1          0
## 5        10        11          3    1.78500      1          0
## 6        12        13         13 -312.50000      1          0
## 7        14        15         50 -434.50000      1          0
## 8        16        17         36   -2.50000      1          0
## 9        18        19         27   65.70858      1          0
## 10       20       21         48  235.50000      1          0
## 11       22       23          4   19.50000      1          0
## 12        0         0          0    0.00000     -1          1
## 13       24       25         52  787.50000      1          0
## 14       26       27          4   20.50000      1          0
## 15       28       29         36   26.50000      1          0
## 16       30       31         20    0.66500      1          0
## 17       32       33          9   32.00000      1          0
## 18        0         0          0    0.00000     -1          5
## 19        0         0          0    0.00000     -1          2
## 20       34       35         43   31.50000      1          0
## 21       36       37         35   -8.50000      1          0
## 22       38       39         45    0.98500      1          0
## 23        0         0          0    0.00000     -1          5
## 24       40       41         10   29.00000      1          0
## 25        0         0          0    0.00000     -1          2
## 26       42       43         52 -720.50000      1          0
## 27       44       45         36   61.00000      1          0
## 28       46       47         27   62.03453      1          0
## 29       48       49         14    2.19000      1          0
## 30       50       51         36 -144.50000      1          0
```

```
predictRF <- predict(modFit, subTesting, type = "prob")
```

```
# Perform prediction
```

```
predicts <- predict(modFit, testing, type="prob")
predicts
```

```
##      A      B      C      D      E
## 1 0.094 0.738 0.116 0.032 0.020
## 2 0.870 0.058 0.060 0.004 0.008
## 3 0.144 0.612 0.180 0.010 0.054
## 4 0.912 0.008 0.056 0.018 0.006
## 5 0.938 0.014 0.040 0.002 0.006
## 6 0.014 0.110 0.142 0.052 0.682
## 7 0.036 0.010 0.116 0.804 0.034
## 8 0.090 0.676 0.104 0.098 0.032
## 9 0.998 0.002 0.000 0.000 0.000
## 10 0.994 0.004 0.000 0.000 0.002
```

```
## 11 0.076 0.736 0.076 0.078 0.034
## 12 0.070 0.042 0.822 0.020 0.046
## 13 0.032 0.892 0.008 0.020 0.048
## 14 1.000 0.000 0.000 0.000 0.000
## 15 0.018 0.074 0.050 0.038 0.820
## 16 0.034 0.084 0.014 0.026 0.842
## 17 0.968 0.000 0.002 0.002 0.028
## 18 0.038 0.898 0.014 0.034 0.016
## 19 0.142 0.760 0.040 0.044 0.014
## 20 0.016 0.980 0.002 0.000 0.002
```

Forecasting

Traditionally independent variable over TIME