

COM662 Data Analytics

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Week 6 – Linear and Logistic Regression

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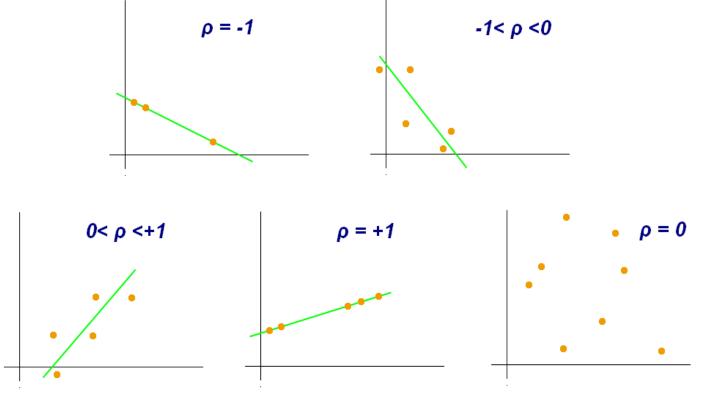
Week 6 Content

- Review correlation
- Linear regression
- Multiple regression
- Model fitness and significance
- Using regression as a predictive model
- Evaluating a linear regression model
- Logistic regression
- Application of logistic regression
 - Odds ratios
 - Binary classification/prediction



Correlation / association

- Correlation
 - Strength of association between variables



Regression Definition

- Regression models the relationship between one or many independent variables (features) with one dependent/target variable.
- Typically used to model continuous variables
- Multiple regression vs. simple linear regression



Regression Regression to the Mean

Heights of parents vs. heights of their children

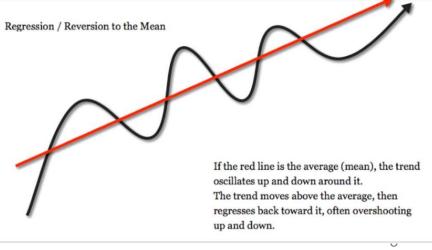
"It appeared from these experiments that the offspring did not tend to resemble their parents in size, but always to be more mediocre than they – to be smaller than the parents, if the parents were large; to be larger than the parents, if the parents were small."

Sir Francis Galton

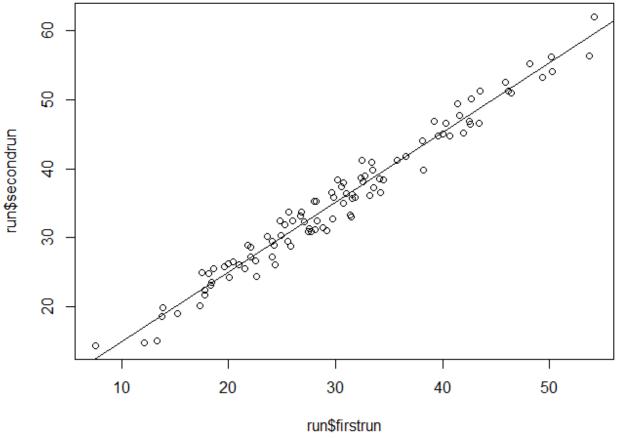
https://select-statistics.co.uk/blog/regression-to-the-mean-as-

relevant-today-as-it-was-in-the-1900s/





RegressionSimple Linear Regression





$$\hat{y} = b_0 + b_1 \mathbf{x}$$

Multiple Linear Regression

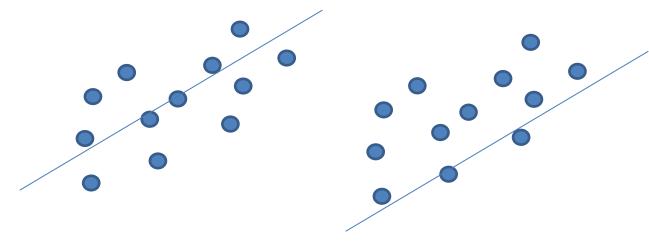
$$\hat{\mathbf{y}} = b\mathbf{X} + c\mathbf{Y} + d\mathbf{Z} + a$$

$$\hat{y} = \sum_{i=1}^{n} b_i \cdot X_i + b_0$$



Regression Ordinary Least Squares Regression

- Finds the regression line that has the least squared error between what is predicted and the actual values.
- OLS minimizes the sum of the squared residuals.
- It finds b₀ and b₁.

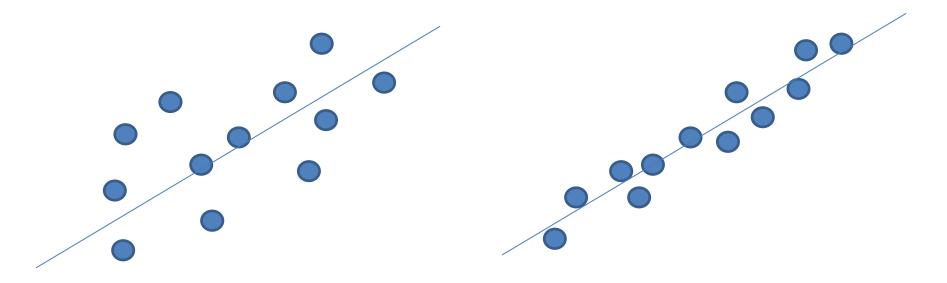




$$\hat{eta} = rac{\sum x_i y_i - rac{1}{n} \sum x_i \sum y_i}{\sum x_i^2 - rac{1}{n} (\sum x_i)^2} = rac{\mathrm{Cov}[x,y]}{\mathrm{Var}[x]} \ \hat{lpha} = \overline{y} - \hat{eta} \, \overline{x}$$

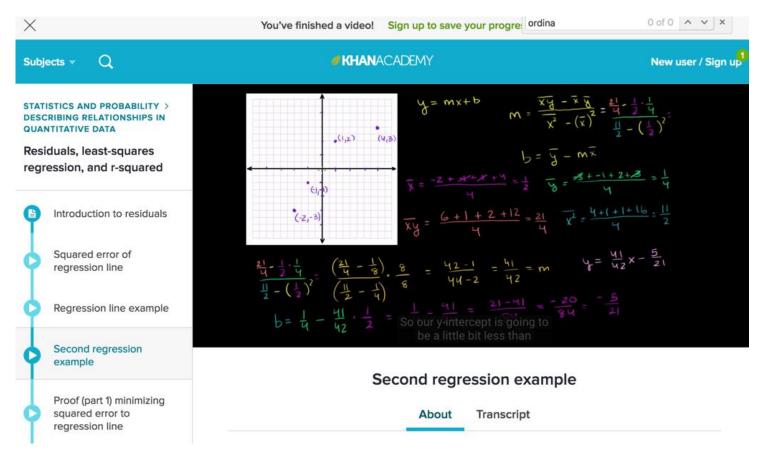
Regression Goodness of Fit

Definition: How the regression line fits the observed data





Exercise - watch online video



https://www.khanacademy.org/math/statistics-probability/describing-relationships-quantitative-data/residuals-least-squares-rsquared/v/second-regression-example

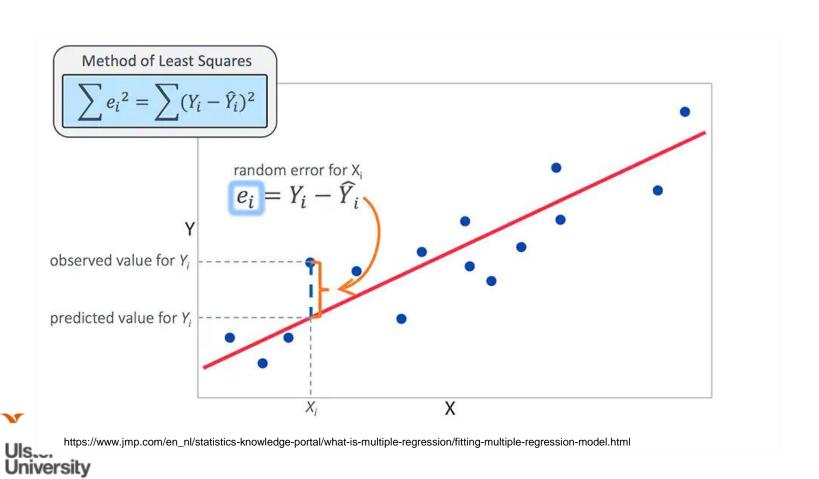


Regression Goodness of Fit

- How the regression line fits the observed data
- R Squared (Coefficient of determination)
 - R-squared is the % of the total variation which the model explains
 - R-squared is always between 0 and 100% or 0 and 1
 - R-squared is a statistical measure of how close the data are to the fitted regression line.



Goodness of Fit - Residuals absolute error

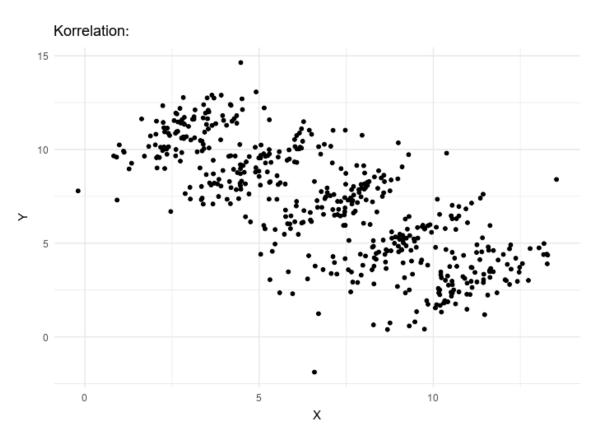


Statistical Significance in Regression - F Test

- Compares a model with no predictors (intercept-only model) to your model.
 - "Null hypothesis: The fit of the intercept-only model and your model are equal.
 - Alternative hypothesis: The fit of the intercept-only model is significantly reduced compared to your model."



RegressionSimpsons Paradox

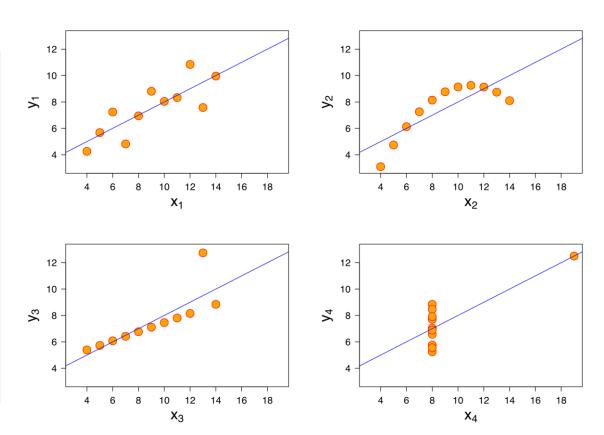




Regression Anscombe's quartet

Anscombe's quartet

7 tiloooiiibo o quartot							
ı		II		III		IV	
X	У	X	У	X	У	X	У
10.0	8.04	10.0	9.14	10.0	7.46	8.0	6.58
8.0	6.95	8.0	8.14	8.0	6.77	8.0	5.76
13.0	7.58	13.0	8.74	13.0	12.74	8.0	7.71
9.0	8.81	9.0	8.77	9.0	7.11	8.0	8.84
11.0	8.33	11.0	9.26	11.0	7.81	8.0	8.47
14.0	9.96	14.0	8.10	14.0	8.84	8.0	7.04
6.0	7.24	6.0	6.13	6.0	6.08	8.0	5.25
4.0	4.26	4.0	3.10	4.0	5.39	19.0	12.50
12.0	10.84	12.0	9.13	12.0	8.15	8.0	5.56
7.0	4.82	7.0	7.26	7.0	6.42	8.0	7.91
5.0	5.68	5.0	4.74	5.0	5.73	8.0	6.89



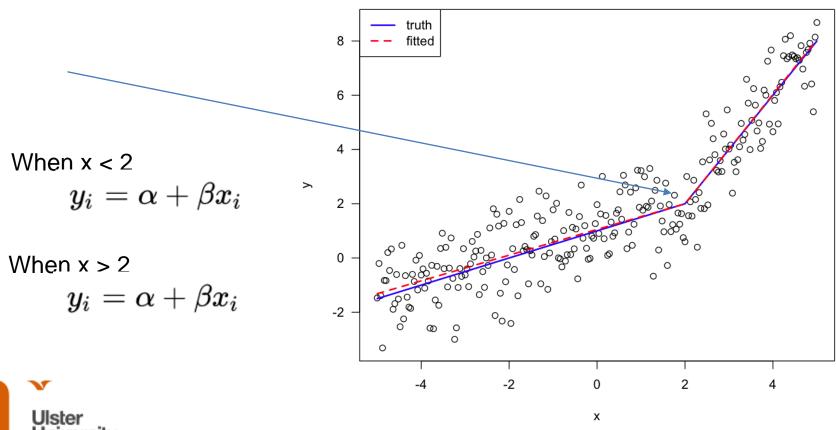
Anscombe, F. J. (1973). "Graphs in Statistical Analysis". American Statistician. 27 (1): 17–21. doi:10.1080/00031305.1973.10478966. JSTOR 2682899.

https://en.wikipedia.org/wiki/Anscombe%27s_quartet



RegressionSegmented regression

AKA piecewise regression / "broken-stick regression"



Regression Building a model using Multiple Regression

- Stepwise regression
 - Use a test for each independent variable (correlation)
- Model building using iteration
 - Backwards elimination
 - Forward selection



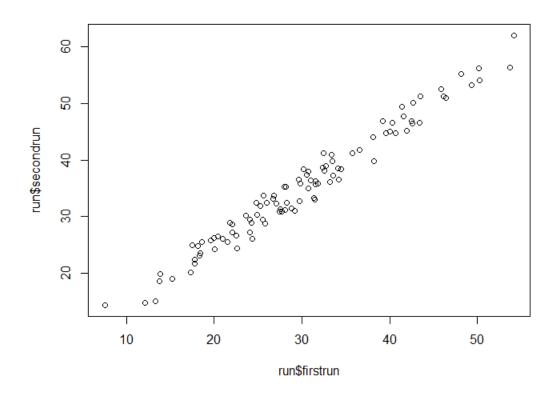
Regression Multicollinearity

 This is when multiple predictors are highly correlated meaning that coefficients maybe less interpretable



Using run.csv

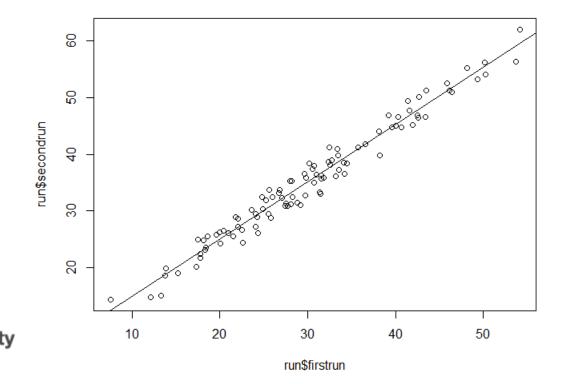
```
> run <- read.csv('run.csv')
> plot(run$firstrun,run$secondrun)
> |
```





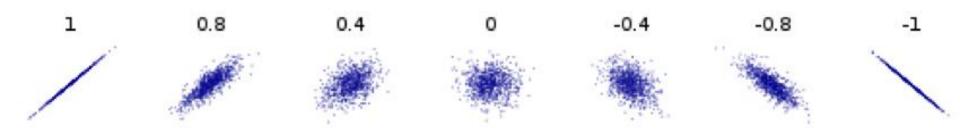
Ulster

```
> plot(run$firstrun,run$secondrun)
> reg1 <- lm(run$secondrun ~ run$firstrun)
> abline(reg1)
>
```



Pearson Correlation Coefficient

```
> r <- cor(run$firstrun,run$secondrun)
> r
[1] 0.9843855
> |
```



https://en.wikipedia.org/wiki/Correlation



Check if the Correlation is significant



p-value < 0.05 indicates the correlation is statistically significant

Regression Tutorial

- Multiple Regression
- help("~") tilde is used to separate the left- and righthand sides in a model formula.

```
lma <- lm(run$finalrun ~ firstrun + secondrun, data=run)
 summary(lma)
Call:
lm(formula = run$finalrun ~ firstrun + secondrun, data = run)
Residuals:
   Min
            1Q Median
-9.1691 -1.5454 0.1766 1.6946 6.8319
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) 6.1518 1.2025 5.116 1.60e-06 ***
           0.1764
                        0.1649 1.070
firstrun
                        0.1607 5.400 4.82e-07 ***
secondrun
            0.8679
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 2.873 on 96 degrees of freedom
Multiple R-squared: 0.9334,
                             Adjusted R-squared: 0.932
F-statistic: 672.6 on 2 and 96 DF, p-value: < 2.2e-16
```



Create a model and predict a value



- Build model using 80% of the rows
- Test on 20% of the rows

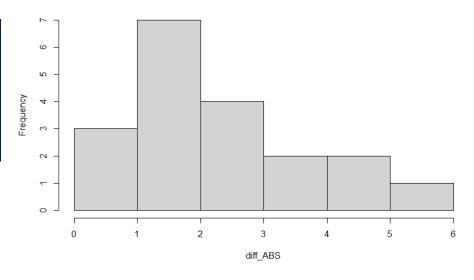
```
> set.seed(12345)
> run2 <- run[order(runif(99)),]
> lma <- lm(finalrun ~ firstrun + secondrun, data=run2[c(1:80),])
> testing <- run2[c(81:99),]
> predictions <- predict(lma, testing)</pre>
```



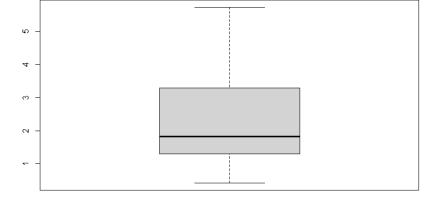
Regression Tutorial

```
> diff <- predictions - testing$finalrun
> diff_ABS <- abs(diff)
> mean(diff)
[1] -0.1306889
> hist(diff_ABS)
> boxplot(diff_ABS)
> |
```

Histogram of diff_ABS







 How many predicted distances of the final run were higher than the real distance of the final run?

```
> length(diff[diff>0])
[1] 10
```

 How many predicted distances were within 2km of the real distance of the final run?

```
> length(diff_ABS[diff_ABS<=2])
[1] 10
```



Regression Tutorial

```
> testing$prediction <- predictions
> testing$difference <- diff ABS
> testing
    X firstrun secondrun finalrun prediction difference
92 92 32.37726 38.73920 44.88548
                                   45.30739
                                             0.4219054
87 87 17.52331 24.99804 29.37527
                                   31.07447 1.6991964
52 52 41.91553 45.16510 51.27962
                                   52.54004 1.2604169
37 37 32.42804 41.16914 50.93965
                                  47.25107
                                             3.6885802
81 81 50.28730 54.16673 66.55377
                                  61.56143
                                             4.9923423
   2 41.60819 47.71729 55.34708
                                  54.50125
                                             0.8458281
    4 38.18542 39.78412 46.55384
                                   47.43082
                                             0.8769718
38 38 31.48147 33.00327 39.31016
                                   40.54661
                                             1.2364473
43 43 34.14534 38.57678 47.85165
                                   45.57167
                                             2.2799835
91 91 49.39089 53.19785 62.11803
                                   60.59146
                                             1.5265686
70 70 17.74187 21.66239 22.73644
                                   28.47046
                                             5.7340216
67 67 26.02081
               32.46219 42.79469
                                   38.90118
                                             3.8935087
20 20 25.81556
               28.77266 34.49935
                                   35.92144
                                             1.4220917
               50.13416 58.74417
77 77 42.71507
                                   56.66955
                                             2.0746192
51 51 17.27106
               20.12811 22.97570
                                   27.14557
                                             4.1698691
23 23 45.89486
               52.51884 60.62786
                                   59.27353
                                             1.3543277
65 65 25.63270
               33.79786 38.04635
                                   39.87699
                                             1.8306355
63 63 12.11494
               14.85375 24.69498
                                   21.80381
                                             2.8911652
10 10 32.69781 39.02388 43.19283
                                   45.60511
                                             2.4122794
```

Two new columns added

- https://archive.ics.uci.edu/ml/datasets/wine+quality
- Load the dataset

```
df red <- read.csv("winequality-red.csv", sep = ";")</pre>
> head(df red)
  fixed.acidity volatile.acidity citric.acid residual.sugar chlorides free.sulfur.dioxide total.sulfur.dioxide density
                             0.70
                                                                                                                       0.9978
            7.4
                                          0.00
                                                           1.9
                                                                    0.076
                                                                                            11
            7.8
                             0.88
                                                           2.6
                                                                    0.098
                                                                                                                   67 0.9968
                                          0.00
                                                                                            25
            7.8
                             0.76
                                          0.04
                                                           2.3
                                                                    0.092
                                                                                                                   54 0.9970
                                                                                            15
           11.2
                             0.28
                                          0.56
                                                           1.9
                                                                    0.075
                                                                                            17
                                                                                                                     0.9980
            7.4
                             0.70
                                          0.00
                                                           1.9
                                                                    0.076
                                                                                            11
                                                                                                                      0.9978
            7.4
                             0.66
                                          0.00
                                                                    0.075
                                                                                            13
                                                                                                                      0.9978
   pH sulphates alcohol quality
1 3.51
            0.56
                      9.4
2 3.20
            0.68
                      9.8
3 3.26
            0.65
                      9.8
4 3.16
                      9.8
            0.58
                                 6
5 3.51
            0.56
                      9.4
6 3.51
            0.56
                      9.4
```

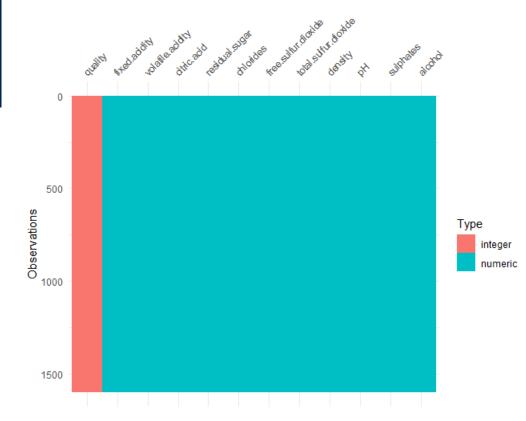


Regression Tutorial - Wine Quality Data Set

Check the missing value

```
> # check missing value
> sum(is.na(df_red))
[1] 0
> # visulisation of missing values
> library('visdat')
> vis_dat(df_red)
> |
```



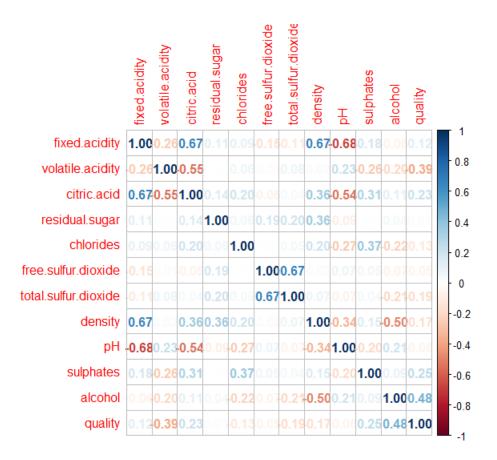


Regression Regression Tutorial - Wine Quality Data Set

Correlation analysis

```
> library('corrplot')
corrplot 0.92 loaded
> M <- cor(df_red)
> corrplot(M, method = 'number')
> |
```





- Modelling
 - Splitting Train Datasets and Test Datasets

```
> set.seed(1)
> sampleSize <- round(nrow(df_red)*0.8)
> idx <- sample(seq_len(sampleSize), size = sampleSize)
> 
> train_red <- df_red[idx,]
> test_red <- df_red[-idx,]
> |
```



- Modelling
 - Create linear regression model using all the features

```
> model red1 <- lm(quality ~ fixed.acidity + volatile.acidity + citric.acid + chlorides + free.sulfur.dioxide + total.sulfu
r.dioxide + density + pH + sulphates + alcohol, data = train red)
> summary(model red1)
Call:
lm(formula = quality ~ fixed.acidity + volatile.acidity + citric.acid +
    chlorides + free.sulfur.dioxide + total.sulfur.dioxide +
    density + pH + sulphates + alcohol, data = train red)
Residuals:
              10 Median
     Min
                                30
                                        Max
-2.67536 -0.38553 -0.06879 0.45454 1.97578
Coefficients:
                      Estimate Std. Error t value Pr(>|t|)
(Intercept)
                     1.510e+01 1.912e+01
                                            0.790 0.429935
fixed.acidity
                    1.547e-02 2.696e-02 0.574 0.566286
volatile.acidity
                    -1.057e+00 1.358e-01 -7.780 1.49e-14 ***
citric.acid
                    -1.774e-01 1.657e-01 -1.070 0.284621
chlorides
                    -1.779e+00 4.627e-01 -3.845 0.000126 ***
free.sulfur.dioxide 3.392e-03 2.480e-03 1.368 0.171691
total.sulfur.dioxide -3.645e-03 8.201e-04 -4.444 9.58e-06 ***
                    -1.102e+01 1.954e+01 -0.564 0.572664
density
                    -3.835e-01 2.010e-01 -1.908 0.056598 .
Нq
                    7.945e-01 1.217e-01 6.527 9.67e-11 ***
sulphates
alcohol
                     2.924e-01 2.462e-02 11.878 < 2e-16 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.6476 on 1268 degrees of freedom
Multiple R-squared: 0.3695, Adjusted R-squared: 0.3645
F-statistic: 74.3 on 10 and 1268 DF, p-value: < 2.2e-16
```

- Modelling
 - Create linear regression model using only one feature

```
> model red2 <- lm(quality ~ alcohol, data = train red)</pre>
> summary(model red2)
Call:
lm(formula = quality ~ alcohol, data = train red)
Residuals:
           10 Median
   Min
                            3Q
                                   Max
-2.8861 -0.4048 -0.1827 0.5582 2.5581
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) 1.81376
                       0.18792 9.652 <2e-16 ***
alcohol
            0.37021
                       0.01797 20.601 <2e-16 ***
Signif. codes: 0 \*** 0.001 \** 0.01 \*' 0.05 \.' 0.1 \' 1
Residual standard error: 0.7041 on 1277 degrees of freedom
Multiple R-squared: 0.2494,
                               Adjusted R-squared: 0.2489
F-statistic: 424.4 on 1 and 1277 DF, p-value: < 2.2e-16
```

- Modelling
 - Create linear regression model using top 5 most correlated features

```
model red3 <- lm(quality ~ alcohol + volatile.acidity + sulphates + citric.acid + density,</pre>
                 data = train red)
 summary (model red3)
Call:
lm(formula = quality ~ alcohol + volatile.acidity + sulphates +
   citric.acid + density, data = train red)
Residuals:
   Min
            10 Median
-2.7055 -0.4068 -0.0748 0.4949 2.2302
Coefficients:
                Estimate Std. Error t value Pr(>|t|)
               -10.20026 13.12402 -0.777
(Intercept)
                                             0.437
alcohol
                 volatile.acidity -1.26276 0.13187 -9.576 < 2e-16 ***
sulphates
                0.57332 0.11255 5.094 4.03e-07 ***
citric.acid
                -0.16117 0.13303 -1.212
                                             0.226
density
               12.75053 13.08709 0.974
                                             0.330
Signif. codes: 0 \*** 0.001 \** 0.01 \*' 0.05 \.' 0.1 \ ' 1
Residual standard error: 0.6601 on 1273 degrees of freedom
Multiple R-squared: 0.3423, Adjusted R-squared: 0.3397
F-statistic: 132.5 on 5 and 1273 DF, p-value: < 2.2e-16
```

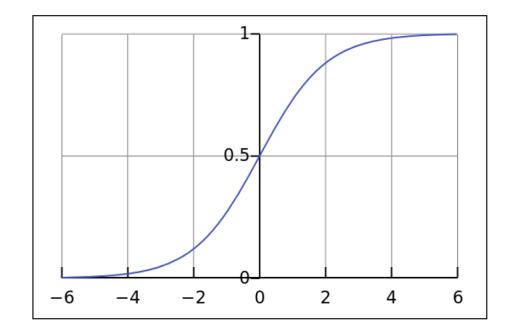
Logistic Regression

- A regression model where the dependent variable is nominal/categorical
 - Useful for feature engineering
 - Useful for binary classification
- Technology adoption
- Success/Unsuccess
- Pass/Fail
- Admitted or not admitted
- Useful for calculating the Odds Ratios for each independent variable, hence good for feature selection



Logistic Regression

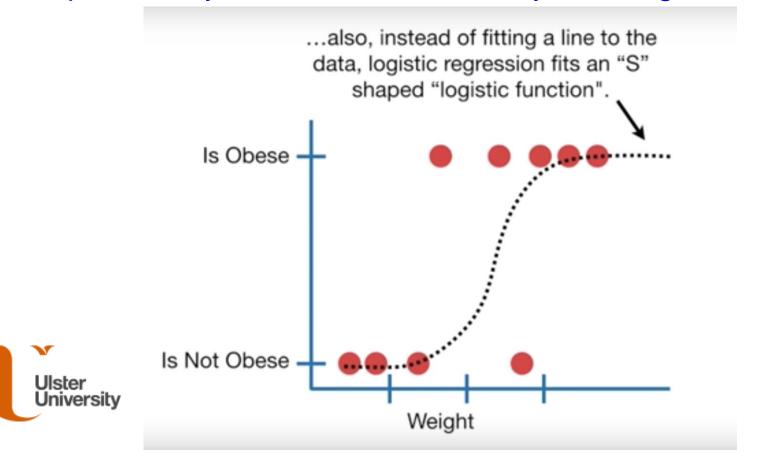
- Input: can take any value from a negative infinity to a positive infinity
- Output: values from 0 to 1 (a probability)



$$S(x)=rac{1}{1+e^{-x}}$$

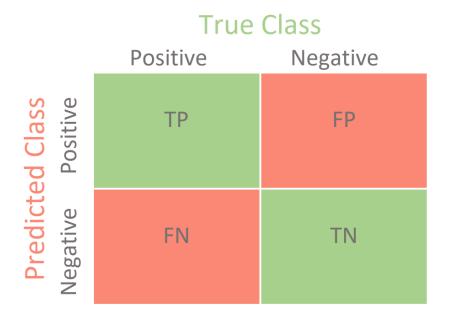


- StatQuest: Logistic Regression.
- https://www.youtube.com/watch?v=yIYKR4sgzI8



Logistic Regression Model evaluation

Confusion matrix



https://towards datascience.com/confusion-matrix-for-your-multi-class-machine-learning-model-ff9 a a 3bf7826



Logistic Regression

Exercise

Load data

```
> run <- read.csv('run.csv')
> head(run)
  X firstrun secondrun finalrun
1 1 46.47100 51.04981 62.91495
2 2 41.60819 47.71729 55.34708
3 3 40.29477 46.64771 56.34791
4 4 38.18542 39.78412 46.55384
5 5 34.50337 38.33918 45.68743
6 6 39.25199 46.94684 52.29005
> run$superrun = ifelse(run$finalrun >= 45,1,0)
> head(run)
  X firstrun secondrun finalrun superrun
1 1 46.47100 51.04981 62.91495
2 2 41.60819 47.71729 55.34708
3 3 40.29477 46.64771 56.34791
4 4 38.18542 39.78412 46.55384
5 5 34.50337 38.33918 45.68743
6 6 39.25199 46.94684 52.29005
```

Split training and test data

```
> # split train and test data
> set.seed(1)
> sampleSize <- round(nrow(run)*0.8)
> idx <- sample(seq_len(sampleSize), size = sampleSize)
> 
> train_run <- run[idx,]
> test_run <- run[-idx,]
> |
```



Create logistic regression model

```
# create logistic regression model
> logitModel <- glm(superrun ~ firstrun + secondrun, data=train run, family = binomial(link = 'logit'))
> summary(logitModel)
Call:
qlm(formula = superrun ~ firstrun + secondrun, family = binomial(link = "logit"),
    data = train run)
Deviance Residuals:
     Min
                1Q Median
                                             Max
                                    3Q
-1.63898 -0.15492 -0.00658 0.02821 1.89502
Coefficients:
            Estimate Std. Error z value Pr(>|z|)
(Intercept) -25.9130 8.4084 -3.082 0.00206 **
firstrun 0.5608 0.3281 1.709 0.08746 . secondrun 0.2117 0.2727 0.776 0.43760
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
    Null deviance: 104.903 on 78 degrees of freedom
Residual deviance: 24.515 on 76 degrees of freedom
AIC: 30.515
Number of Fisher Scoring iterations: 8
```

```
probabilities
          80
                                     82
                        81
                                                   83
                                                                 84
                                                                              85
9.999993e-01 9.999989e-01 1.876316e-03 4.603853e-03 3.790518e-01 1.787480e-04 4.775197e-02
                        88
                                     89
                                                   90
                                                                 91
          87
2.050390e-05 9.968462e-01 6.054316e-03 9.476916e-01 9.999978e-01 6.091173e-01 1.165638e-03
                        95
                                     96
                                                   97
          94
                                                                 98
1.892691e-02 1.762211e-01 8.104753e-01 3.039719e-01 3.043883e-01 7.838931e-05
```

```
> predicted <- as.numeric(probabilities > 0.5)
> predicted
 [1] 1 1 0 0 0 0 0 0 1 0 1 1 1 0 0 0 1 0 0
> |
```

```
> predicted <- as.factor(predicted)
> test_run$superrun <- as.factor(test_run$superrun)
> |
```



Model evaluation using confusion matrix

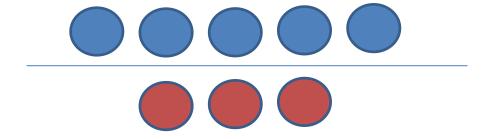
```
> library(caret)
Loading required package: lattice
> confusionMatrix(predicted, test run$superrun)
Confusion Matrix and Statistics
         Reference
Prediction 0 1
        0 11 2
        1 1 6
              Accuracy: 0.85
                95% CI : (0.6211, 0.9679)
   No Information Rate: 0.6
   P-Value [Acc > NIR] : 0.01596
                 Kappa : 0.6809
 Mcnemar's Test P-Value : 1.00000
           Sensitivity: 0.9167
           Specificity: 0.7500
         Pos Pred Value: 0.8462
        Neg Pred Value: 0.8571
             Prevalence: 0.6000
         Detection Rate: 0.5500
   Detection Prevalence: 0.6500
     Balanced Accuracy: 0.8333
       'Positive' Class: 0
```

- exp(coef(logitModel)) ## odds ratios only
- exp(cbind(OR = coef(logitModel), confit(logitModel)## odds ratios and 95% CI



Logistic Regression Another example

- What are the odds of winning the next football game?
 - From the last 8 games, if you won 5 and lost 3 then what are the odds of winning the next game?



$$5/3 = 1.66...$$
 times



Logistic Regression Odds ratios

- How do I interpret odds ratios in logistic regression?
 - https://stats.oarc.ucla.edu/stata/faq/how-do-i-interpretodds-ratios-in-logistic-regression/

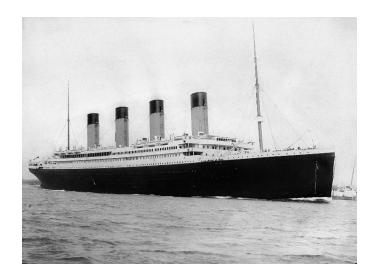


Logistic Regression To learn more - watch

- Stats quest:
 - Logistic regression overview
 - https://www.youtube.com/watch?v=yIYKR4sgzl8
 - Logistic regression coefficients
 - https://www.youtube.com/watch?v=vN5cNN2-HWE
 - Maximum likelihood
 - https://www.youtube.com/watch?v=BfKanl1aSG0
 - R squared and p values for logistic regression
 - https://www.youtube.com/watch?v=xxFYro8QuXA
 - Odds ratios from logit
 - https://www.youtube.com/watch?v=8nm0G-1uJzA

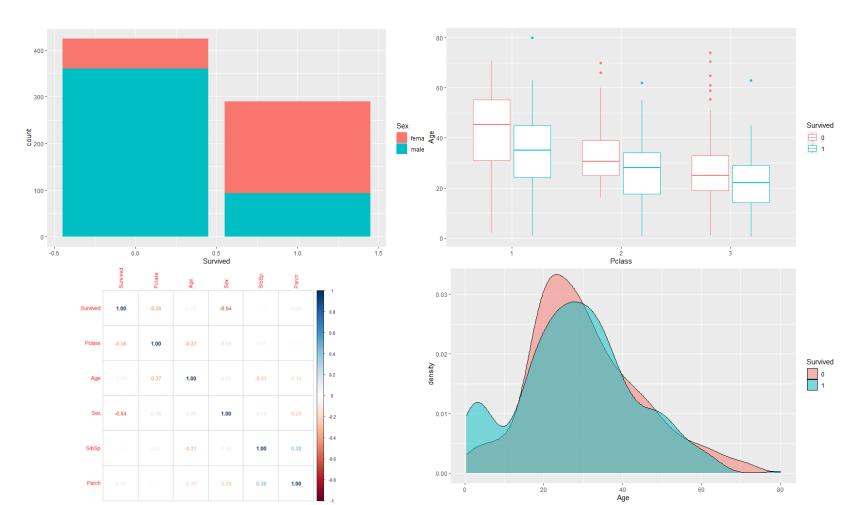


- Studying the Titanic passenger survival
- Titanic dataset from Kaggle https://www.kaggle.com/c/titanic
- build a logistic regression model for survival





Review data exploratory analysis in Week 3



Load the data

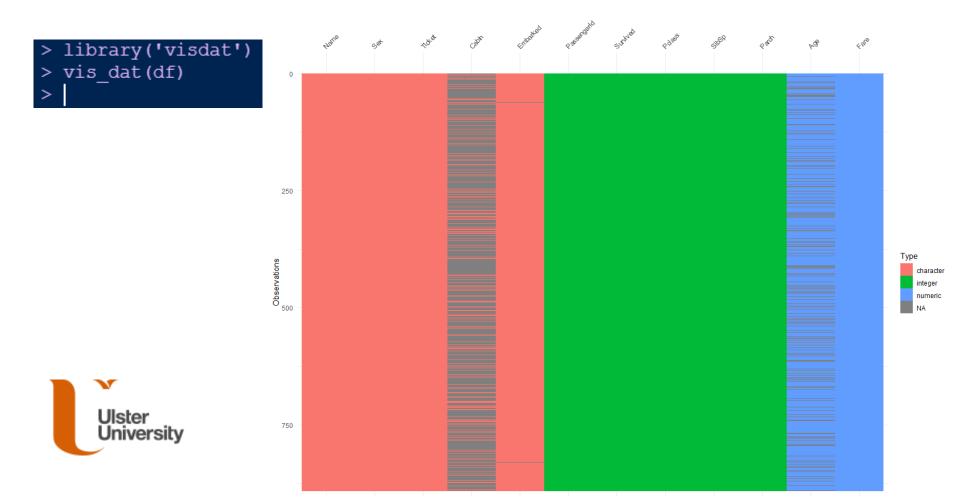
```
> df <- read.csv('train.csv', na.strings = '')  # empty values are read as NA values
> head(df)
 PassengerId Survived Pclass
                                                                                     Sex Age SibSp Parch
                                                                                                                   Ticket
                                                                                                                              Fare Cabin Embarked
                                                                             Name
                                                          Braund, Mr. Owen Harris
                           1 Cumings, Mrs. John Bradley (Florence Briggs Thayer) female
                                                                                                                 PC 17599 71.2833
                                                                                                                                                С
                                                           Heikkinen, Miss. Laina female
                                                                                                       0 STON/O2. 3101282 7.9250
                                     Futrelle, Mrs. Jacques Heath (Lily May Peel)
                                                         Allen, Mr. William Henry
                                                                                                                   373450 8.0500
                                                                                    male NA
                                                                                                                                                o
                                                                 Moran, Mr. James
                                                                                                                    330877 8.4583
```

Check missing values

```
> # check missing values
> sum(is.na(df$Cabin))
[1] 687
```

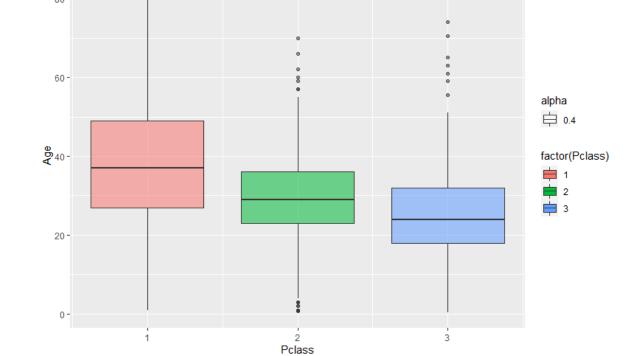


Missing values



Age vs Pclass

```
> library(ggplot2)
> ggplot(df,aes(Pclass,Age)) + geom_boxplot(aes(group=Pclass,fill=factor(Pclass),alpha=0.4))
Warning message:
Removed 177 rows containing non-finite values (stat_boxplot).
> |
```

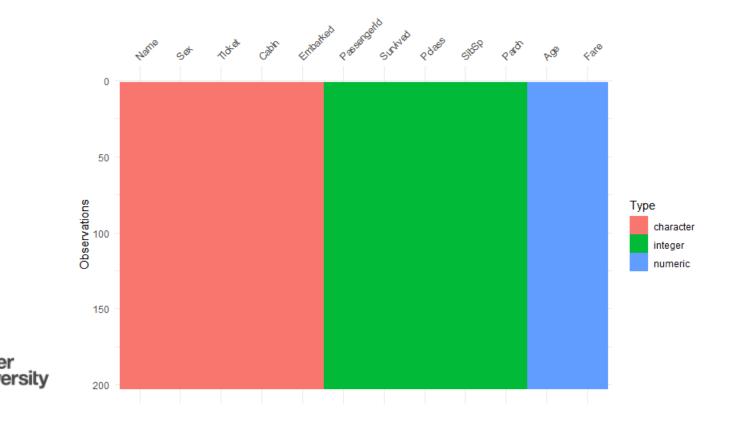




Impute missing age values

```
> # calculate average age for different Pclass
> avg P1 <- round(mean(df[df$Pclass==1,]$Age, na.rm = TRUE))
> avg P2 <- round(mean(df[df$Pclass==2,]$Age, na.rm = TRUE))
 avg P3 <- round(mean(df[df$Pclass==3,]$Age, na.rm = TRUE))</pre>
 impute age <- function(age, class) {</pre>
      out <- age
      for (i in 1:length(age)){
          if (is.na(age[i])){
              if (class[i] == 1){
                   out[i] <- avg P1
              else if (class[i] == 2){
                   out[i] <- avg P2
              }else{
                   out[i] <- avg P3
          }else{
              out[i]<-age[i]
      return (out)
  fixed.ages <- impute age(df$Age,df$Pclass)</pre>
 df$Age <- fixed.ages</pre>
 df <- na.omit(df) #Drop the rows with NA values
```

```
> # check if the imputation worked
> vis_dat(df)
> |
```



Select features

```
> df$Parch <- factor(df$Parch)</pre>
> library(dplyr)
                                                     > df$SibSp <- factor(df$SibSp)</pre>
                                                     >
Attaching package: 'dplyr'
The following objects are masked from 'package:stats':
    filter, lag
The following objects are masked from 'package:base':
    intersect, setdiff, setequal, union
> df <- select(df,-PassengerId,-Name,-Ticket,-Cabin)
> str(df)
'data.frame':
               202 obs. of 8 variables:
 $ Survived: int 1 1 0 1 1 1 1 0 1 1 ...
 $ Pclass : int 1 1 1 3 1 2 1 1 1 1 ...
 $ Sex : chr "female" "female" "male" "female" ...
 $ Age : num 38 35 54 4 58 34 28 19 38 49 ...
$ sibSp : int 1 1 0 1 0 0 0 3 1 1 ...
$ Parch : int 0 0 0 1 0 0 0 2 0 0 ...
$ Fare
         : num 71.3 53.1 51.9 16.7 26.6 ...
                  "C" "S" "S" "S"
$ Embarked: chr
 - attr(*, "na.action") = 'omit' Named int [1:689] 1 3 5 6 8 9 10 13 14 15 ...
  ..- attr(*, "names") = chr [1:689] "1" "3" "5" "6" ...
```

> df\$Survived <- factor(df\$Survived)</pre>

> df\$Pclass <- factor(df\$Pclass)</pre>

Build logistic regression model to predict survival

```
log.model <- qlm(formula=Survived ~ . , family = binomial(link='logit'),data = df)</pre>
> summary(log.model)
glm(formula = Survived ~ ., family = binomial(link = "logit"),
   data = df)
Deviance Residuals:
   Min
             10 Median
                                      Max
-2.7909 -0.7906 0.2852
                          0.5425
Coefficients:
             Estimate Std. Error z value Pr(>|z|)
(Intercept) 4.485693 0.969299 4.628 3.70e-06 ***
Pclass2
                      0.845046 0.140
             0.118245
                                         0.8887
Pclass3
            -1.478862 0.935923 -1.580
                                          0.1141
Sexmale
            -3.012731 0.506304 -5.950 2.67e-09 ***
Age
            -0.038646
                      0.015338 -2.520
                                          0.0118 *
SibSp1
           0.426687 0.417689 1.022
                                         0.3070
SibSp2
           0.886218
                      1.761204 0.503
                                         0.6148
SibSp3
            -0.906767
                      1.728729 -0.525
                                         0.5999
Parch1
           -0.319603
                      0.545424 -0.586
                                         0.5579
Parch2
           -0.652582
                      0.731659 -0.892
                                          0.3724
Parch4
           -13.825696 882.743802 -0.016
                                         0.9875
Fare
             0.001273
                      0.002925 0.435
                                         0.6633
                      1.639963 -1.146
Embarked0
            -1.878968
                                         0.2519
EmbarkedS
            -0.501598
                      0.434676 -1.154
                                        0.2485
Signif. codes: 0 \***' 0.001 \**' 0.01 \*' 0.05 \' 0.1 \' 1
(Dispersion parameter for binomial family taken to be 1)
   Null deviance: 258.07 on 201 degrees of freedom
Residual deviance: 173.92 on 188 degrees of freedom
AIC: 201.92
Number of Fisher Scoring iterations: 13
```

We can see clearly that Sex, Age are the most significant features. Which makes sense given the women and children first policy.

- Predicting using Test cases
 - Training and test split

```
> library(caTools)
> set.seed(101)
>
> split = sample.split(df$Survived, SplitRatio = 0.70)
>
> df_train = subset(df, split == TRUE)
> df_test = subset(df, split == FALSE)
> |
```



Build the model using training dataset

```
# build the model using training dataset
 log.model1 <- glm(formula=Survived ~ . , family = binomial(link='logit'),data = df train)</pre>
 summary(log.model1)
Call:
glm(formula = Survived ~ ., family = binomial(link = "logit"),
   data = df train)
Deviance Residuals:
             10 Median
-2.6092 -0.7143 0.2133 0.4894 2.3432
Coefficients:
             Estimate Std. Error z value Pr(>|z|)
(Intercept) 4.624e+00 1.192e+00 3.880 0.000104 ***
Pclass2
          1.841e+00 1.351e+00 1.362 0.173074
Pclass3
           -2.175e+00 1.186e+00 -1.834 0.066589 .
Sexmale
          -3.347e+00 6.903e-01 -4.849 1.24e-06 ***
Age
           -3.213e-02 1.930e-02 -1.664 0.096066 .
SibSp1
          4.689e-01 4.931e-01 0.951 0.341620
SibSp2
          1.472e+01 2.011e+03 0.007 0.994161
SibSp3
         -1.690e+01 3.956e+03 -0.004 0.996592
Parch1
          -7.259e-01 6.480e-01 -1.120 0.262605
Parch2
          -1.322e+00 9.968e-01 -1.326 0.184757
Parch4
           -1.724e+01 3.956e+03 -0.004 0.996523
Fare
          2.899e-03 3.408e-03 0.851 0.394898
EmbarkedO -1.170e+00 3.197e+00 -0.366 0.714289
EmbarkedS -7.758e-01 5.488e-01 -1.414 0.157502
Signif. codes: 0 \*** 0.001 \** 0.01 \*' 0.05 \'.' 0.1 \' 1
(Dispersion parameter for binomial family taken to be 1)
   Null deviance: 181.68 on 141 degrees of freedom
Residual deviance: 112.12 on 128 degrees of freedom
ATC: 140.12
Number of Fisher Scoring iterations: 16
```

Check the prediction accuracy

```
# Now check the prediction accuracy!
 probabilities <- predict(log.modell,newdata=df test,type='response')</pre>
 probabilities
                                                                            125
                       89
                                                 111
                                                                                          129
                                                                                                                                                231
4.417443e-01\ 5.875553e-07\ 2.882180e-01\ 2.976527e-01\ 9.908272e-01\ 1.498200e-01\ 8.105386e-01\ 8.870230e-01\ 5.203135e-01\ 5.024311e-01\ 9.687646e-01
         246
                      264
                                    285
                                                 293
                                                               306
                                                                            310
                                                                                          312
                                                                                                       326
                                                                                                                     332
                                                                                                                                  340
                                                                                                                                                342
9.999988e-01 3.133768e-01 3.441760e-01 8.805875e-01 5.143617e-01 9.786482e-01 1.000000e+00 9.793761e-01 2.934999e-01 3.010955e-01 5.689787e-07
                                        4.081726e-01
                                                                                                                         3.773614e-01 2.354377e-01
9.930039e-01 4.256300e-01
                          2.384780e-01
                                                      3.290564e-01 2.532270e-01 1.808523e-01 9.625446e-01
                                                                                                           2.759944e-01
                                    505
                                                 506
                                                               516
                                                                                          524
                                                                                                       537
9.730052e-01 9.328377e-01 9.730056e-01 8.150051e-01 2.868709e-01 9.903094e-01 9.341561e-01 2.956631e-01 9.392611e-01 1.000000e+00 9.618014e-01
                      586
                                    592
                                                 631
                                                               648
                                                                            708
                                                                                          711
                                                                                                       725
9.687893e-01 8.983150e-01 9.746426e-01 1.210536e-01 3.965905e-01 3.159569e-01 9.819478e-01 5.636863e-01 9.715035e-01 7.197754e-02 9.542876e-01
                                                 790
         764
                      766
                                    773
                                                               840
8.990071e-01 9.480654e-01 9.799236e-01 5.070712e-01 5.353868e-01
 # Now let's calculate from the predicted values:
 results <- ifelse(probabilities > 0.5,1,0)
        97 111 124 125 129 167 210 225 231 246 264 285 293 306 310 312 326 332 340 342 346 378 391
516 517 524 537 540 572 578 582 586 592 631 648 708 711 725 731 752 760 764 766 773 790 840
```



Check the prediction accuracy

```
> misClasificError <- mean(results != df_test$Survived)
> print(paste('Accuracy',1-misClasificError))
[1] "Accuracy 0.71666666666667"
>
> table(df_test$Survived, probabilities > 0.5)

    FALSE TRUE
    0    14    6
    1    11    29
>
```



Logistic Regression

Exercise

Confusion matrix



```
> predicted <- as.numeric(probabilities > 0.5)
> predicted <- as.factor(predicted)</pre>
> library(caret)
> confusionMatrix(predicted, df test$Survived)
Confusion Matrix and Statistics
          Reference
Prediction 0 1
         0 14 11
         1 6 29
               Accuracy: 0.7167
                 95% CI : (0.5856, 0.8255)
    No Information Rate: 0.6667
    P-Value [Acc > NIR] : 0.2495
                  Kappa: 0.4
 Mcnemar's Test P-Value: 0.3320
            Sensitivity: 0.7000
            Specificity: 0.7250
         Pos Pred Value: 0.5600
         Neg Pred Value: 0.8286
             Prevalence: 0.3333
         Detection Rate: 0.2333
   Detection Prevalence: 0.4167
      Balanced Accuracy: 0.7125
       'Positive' Class: 0
```

Statistical Model vs. ML algorithm What is the difference?

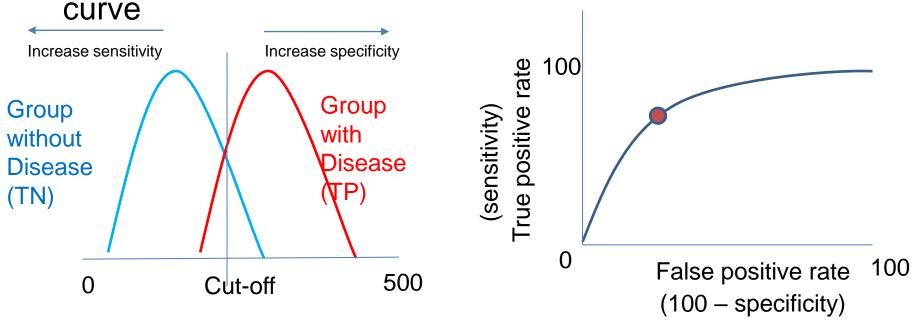
Read:

https://www.analyticsvidhya.com/blog/2015/07/difference-machine-learning-statistical-modeling/



ROC curve

An example of Receiver Operator Characteristic or ROC



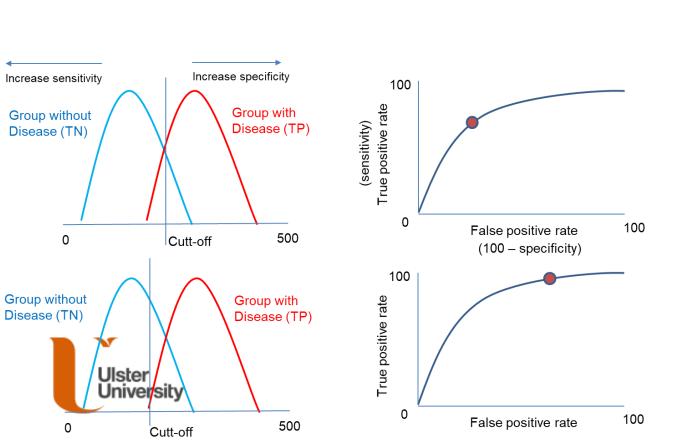


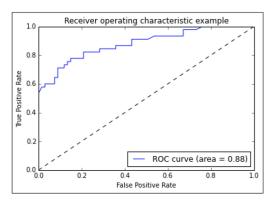
$$sensitivity = \frac{number\ of\ true\ positives}{number\ of\ true\ positives + number\ of\ false\ negatives}$$

$$\text{specificity} = \frac{\text{number of true negatives}}{\text{number of true negatives} + \text{number of false positives}}$$

ROC curve

An example of Receiver Operator Characteristic or ROC curve





Range	Category
0.90-1	This refers to excellent (A)
0.80-0.90	This refers to good (B)
0.70-0.80	This refers to fair (C)
0.60-0.70	This refers to poor (D)
0.50-0.60	This refers to fail (F)

Type 1 error

ROC curve

Receiver Operator Characteristic – Optional Exercise / Reading

- Play:
 - http://www.navan.name/roc/
- Watch:
 - https://www.youtube.com/watch?v=OAl6eAyP-yo
- Read:
 - http://people.inf.elte.hu/kiss/13dwhdm/roc.pdf



Further Reading

- Gareth, J., Daniela, W., Trevor, H. and Robert, T., 2013.
 An introduction to statistical learning: with applications in R. Springer. (Chapter 3)
- Lantz, Brett. Machine learning with R. Packt Publishing Ltd, 2013 (Chapter 6)
- https://stats.idre.ucla.edu/r/dae/logit-regression/
- https://www.scribbr.com/statistics/linear-regression-in-r/
- https://www.machinelearningplus.com/machinelearning/complete-introduction-linear-regression-r/

