HW3

Anubhay Shankar

2022-11-29

Let's generate a non-linear data-set on which Neural Networks outperforms Logistic Regression.

Dataset Description: The planar data is a **randomly generated non-linear dataset** which would need multiple decision boundaries to get an accurate classification or prediction.

```
planar dataset <- function(){ # A function makes it easier to modify the parameters, if required
set.seed(123) # To ensure reproducibility
m <- 800 # Total Length of the data frame
N <- m/2 # Sample size
D <- 2 # Dimensions
X <- matrix(0, nrow = m, ncol = D) # matrix where each row is a single example
Y <- matrix(0, nrow = m, ncol = 1) # label vector (0 for red, 1 for blue)
a <- 5 # Max value of the flower spread
for(j in 0:1){ # For loop to generate a random sample
      ix \leftarrow seq((N*j)+1, N*(j+1)) # range for sequence generation
      t \leftarrow seq(j*3.12,(j+1)*3.12,length.out = N) + rnorm(N, sd = 0.2) # petal inclination
      r \leftarrow a*sin(4*t) + rnorm(N, sd = 0.2) # radius
      X[ix,1] <- r*sin(t) # fills the first column of the X matrix
      X[ix,2] \leftarrow r*cos(t) \# fills the second column of the X matrix
      Y[ix,] <- j # will have value between 0 & 1
}
d <- as.data.frame(cbind(X, Y)) # Create a data-frame</pre>
names(d) <- c('X1','X2','Y') # Rename the columns accordingly
d
}
```

```
df <- planar_dataset()
df <- df[sample(nrow(df)), ] # Shuffle dataset
head(df)</pre>
```

```
## X1 X2 Y

## 740 2.053348 -3.3966571 1

## 730 -1.897771 1.4929127 1

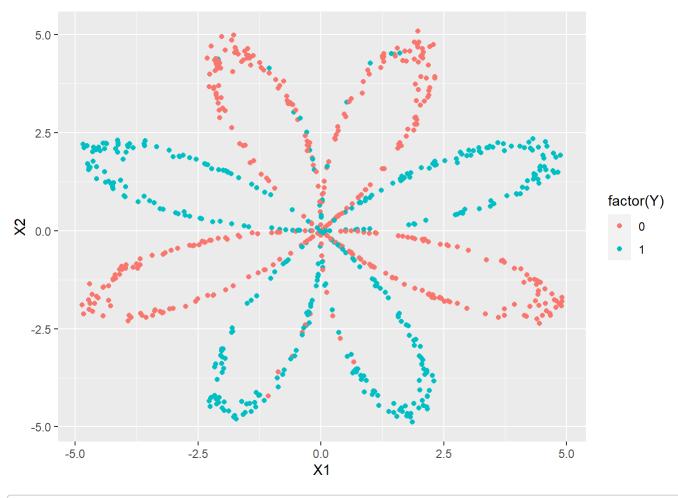
## 648 -3.592936 0.8797975 1

## 342 -1.210739 4.1583541 0

## 123 -1.155820 -1.0033961 0

## 700 -1.307497 1.0938884 1
```

```
ggplot(df, aes(x = X1, y = X2, color = factor(Y))) +
geom_point()
```



```
set.seed(123) # For reproducibility
index <- sample(1:nrow(df), 0.8 * nrow(df), replace = FALSE) # Do a 80-20 train-test split
train <- df[index,]
test <- df[-index,]
glimpse(train) # Get a quick glance of the data-frame structure</pre>
```

```
## Rows: 640
## Columns: 3
## $ X1 <dbl> -1.158604778, 0.812910525, -3.371047466, -0.155050665, -1.816881952...
## $ X2 <dbl> 4.09778115, -0.73494525, -2.01363648, 1.67141934, 2.62977897, 1.559...
## $ Y <dbl> 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 1, 1, 1, 1, 0, 0, 0...
```

```
glimpse(test)
```

```
## Rows: 160
## Columns: 3
## $ X1 <dbl> 2.05334818, -3.59293564, 4.72474337, 3.73321458, 1.55710728, 4.7002...
## $ X2 <dbl> -3.396657065, 0.879797525, 1.532745099, -1.992538553, 4.651756167, ...
## $ Y <dbl> 1, 1, 1, 0, 0, 0, 0, 1, 0, 0, 0, 1, 1, 1, 0, 0, 1, 0, 0, 0...
```

```
logit_train <- glm(Y ~ ., data = train, family = "binomial")
summary(logit_train) # Logistic regression using training set</pre>
```

```
##
## Call:
## glm(formula = Y ~ ., family = "binomial", data = train)
##
## Deviance Residuals:
##
      Min
                1Q Median
                                 3Q
                                         Max
## -1.6198 -1.2205 0.7633 1.1721
                                     1.5631
##
## Coefficients:
               Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) 0.095684 0.082008
                                    1.167
                                             0.243
## X1
               0.002088
                          0.032355 0.065
                                             0.949
## X2
              -0.214626
                          0.034320 -6.254 4.01e-10 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 886.60 on 639 degrees of freedom
## Residual deviance: 844.39 on 637 degrees of freedom
## AIC: 850.39
##
## Number of Fisher Scoring iterations: 4
```

```
logLik(logit_train) # Log likelihood of training set
```

```
## 'log Lik.' -422.1966 (df=3)
```

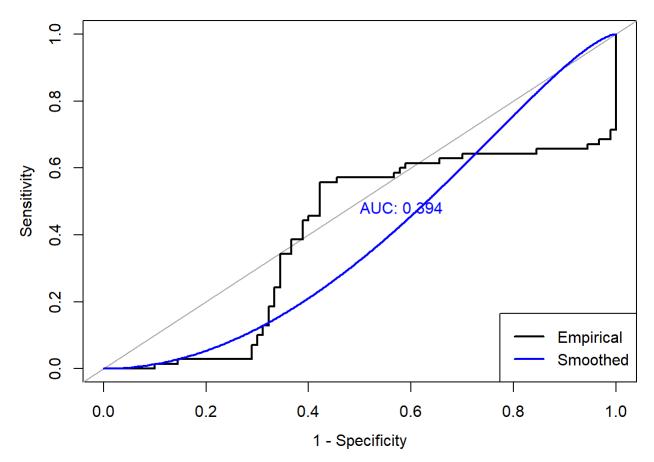
with(logit_train, pchisq(null.deviance - deviance, df.null - df.residual, lower.tail = FALSE)) # p-value of the training dataset

```
## [1] 6.826302e-10
```

```
pR2(logit_train)["McFadden"] # Logistic Regression equivalent of R-squared
```

```
## fitting null model for pseudo-r2
##
     McFadden
## 0.04760882
vif(logit_train) # Check for multicollinearity
##
         X1
## 1.000042 1.000042
predictions_test <- predict(logit_train, test, type = "response")</pre>
predict_binary_test <- ifelse(predictions_test > 0.5, 1, 0)
cftable_test <- table(predict_binary_test, test$Y)</pre>
cftable_test
##
## predict_binary_test 0 1
##
                      0 35 30
##
                      1 55 40
accuracy_test <- sum(diag(cftable_test))/sum(cftable_test)</pre>
accuracy_test
## [1] 0.46875
sensitivity_test <- cftable_test[1]/(cftable_test[1] + cftable_test[2])</pre>
sensitivity test
## [1] 0.3888889
specificity_test <- cftable_test[4]/(cftable_test[3] + cftable_test[4])</pre>
specificity_test
## [1] 0.5714286
ppv_test <- cftable_test[1]/(cftable_test[1] + cftable_test[3])</pre>
ppv_test
## [1] 0.5384615
npv_test <- cftable_test[4]/(cftable_test[2] + cftable_test[4])</pre>
npv_test
```

```
## [1] 0.4210526
error <- mean(predict_binary_test!=test$Y)</pre>
error
## [1] 0.53125
accuracy <- 1-error
accuracy
## [1] 0.46875
roc_test <- roc(test$Y,predictions_test)</pre>
## Setting levels: control = 0, case = 1
## Setting direction: controls > cases
plot.roc(roc_test, col=par("fg"),plot=TRUE,print.auc = FALSE, legacy.axes = TRUE, asp =NA)
roc_test
##
## Call:
## roc.default(response = test$Y, predictor = predictions_test)
## Data: predictions_test in 90 controls (test$Y 0) > 70 cases (test$Y 1).
## Area under the curve: 0.4049
plot.roc(smooth(roc_test), col="blue", add = TRUE,plot=TRUE,print.auc = TRUE, legacy.axes = TRUE
, asp = NA)
legend("bottomright", legend=c("Empirical", "Smoothed"),
       col=c(par("fg"), "blue"), lwd=2)
```



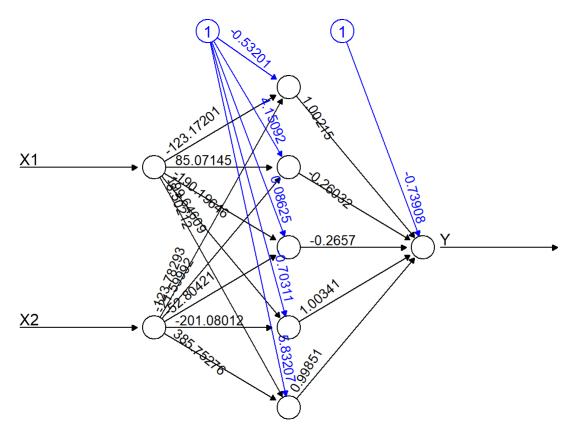
From the above observations we can conclude that Logistic Regression (LR) is not an appropriate model for this data. This is primarily due to the fact that Logistic Regression by default creates only one decision boundary. However, the Planar dataset is intentionally in a non-linear space, owing to the utilization of Sine and Cosine functions, hence, LR will have a horrible performance under such circumstances as shown by the AUC plot above.

```
seed <- 490
set.seed(seed)

planar_nn <- neuralnet(Y~., data = train, hidden = 5, rep = 100, stepmax = 10000) # Run a neural
network</pre>
```

Warning: Algorithm did not converge in 25 of 100 repetition(s) within the
stepmax.

plot(planar_nn, rep = 1) #Plot the NN structure with a single repetition



Error: 11.552722 Steps: 3852

```
predict <- compute(planar_nn, test[,c(1:2)]) # Predict the trained model on the test set

#head(predict$net.result) # View the converged results - OPTIONAL!
p <- predict$net.result #Store the predicted values

prediction <- ifelse(p>0.5,1,0) # Create the decision boundaries
pred_table <- table(prediction,test$Y) # Generate a confusion matrix
pred_table

##
## prediction 0 1
## 0 85 6</pre>
```

```
error_nn <- 1-sum(diag(pred_table))/sum(pred_table) # Error of the NN
error_nn
```

```
## [1] 0.06875
```

```
accuracy_nn <- 1-error_nn # Accuracy of the NN
accuracy_nn
```

##

1 5 64

```
## [1] 0.93125
```

From the above observations, it is clear that Neural Network (NN), accuracy ~94% vs. ~47%, vastly outperforms LR. This is primarily due to the non-linear structure of the data. Additionally, reduction or increase in the number of hidden layers has negligible impact on the resultant accuracy; however, it does increase the running time. Hence, to include more layers we can pair this with a few other techniques to see the improvement.

Let's now see a dataset where Logistic Regression (LR) performs better than Neural Network (NN).

Datset Description: The sinking of the Titanic is one of the most infamous shipwrecks in history.

On April 15, 1912, during her maiden voyage, the widely considered "unsinkable" RMS Titanic sank after colliding with an iceberg. Unfortunately, there weren't enough lifeboats for everyone onboard, resulting in the death of 1502 out of 2224 passengers and crew.

While there was some element of luck involved in surviving, it seems some groups of people were more likely to survive than others.

In this competition, you'll gain access to two similar datasets that include passenger information like name, age, gender, socio-economic class, etc. One dataset is titled "train.csv" and the other is titled "test.csv".

Train.csv will contain the details of a subset of the passengers on board (891 to be exact) and importantly, will reveal whether they survived or not, also known as the "ground truth".

The test.csv dataset contains similar information but does not disclose the "ground truth" for each passenger. It's your job to predict these outcomes.

Using the patterns you find in the train.csv data, predict whether the other 418 passengers on board (found in test.csv) survived.

```
train <- titanic_train # The Training Set
test <- titanic_test # The Testing Set
glimpse(train) # Analyze the metadata</pre>
```

```
## Rows: 891
## Columns: 12
## $ PassengerId <int> 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17,...
## $ Survived
                                                     <int> 0, 1, 1, 1, 0, 0, 0, 0, 1, 1, 1, 1, 0, 0, 0, 1, 0, 1, 0, 1...
                                                     <int> 3, 1, 3, 1, 3, 3, 1, 3, 3, 2, 3, 1, 3, 3, 3, 2, 3, 2, 3, 3...
## $ Pclass
## $ Name
                                                     <chr> "Braund, Mr. Owen Harris", "Cumings, Mrs. John Bradley (Fl...
                                                     <chr> "male", "female", "female", "female", "male", "m
## $ Sex
## $ Age
                                                     <dbl> 22, 38, 26, 35, 35, NA, 54, 2, 27, 14, 4, 58, 20, 39, 14, ...
                                                     <int> 1, 1, 0, 1, 0, 0, 0, 3, 0, 1, 1, 0, 0, 1, 0, 0, 4, 0, 1, 0...
## $ SibSp
## $ Parch
                                                     <int> 0, 0, 0, 0, 0, 0, 1, 2, 0, 1, 0, 0, 5, 0, 0, 1, 0, 0...
## $ Ticket
                                                     <chr> "A/5 21171", "PC 17599", "STON/O2. 3101282", "113803", "37...
## $ Fare
                                                     <dbl> 7.2500, 71.2833, 7.9250, 53.1000, 8.0500, 8.4583, 51.8625,...
                                                     <chr> "", "C85", "", "C123", "", "E46", "", "", "", "G6", "C...
## $ Cabin
## $ Embarked
                                                     <chr> "S", "C", "S", "S", "S", "Q", "S", "S", "S", "C", "S", "S"...
```

```
glimpse(test)
```

```
## Rows: 418
## Columns: 11
## $ PassengerId <int> 892, 893, 894, 895, 896, 897, 898, 899, 900, 901, 902, 903...
## $ Pclass
                <int> 3, 3, 2, 3, 3, 3, 3, 2, 3, 3, 1, 1, 2, 1, 2, 2, 3, 3, 3...
## $ Name
                <chr> "Kelly, Mr. James", "Wilkes, Mrs. James (Ellen Needs)", "M...
## $ Sex
                <chr> "male", "female", "male", "female", "female", "femal...
## $ Age
                <dbl> 34.5, 47.0, 62.0, 27.0, 22.0, 14.0, 30.0, 26.0, 18.0, 21.0...
                <int> 0, 1, 0, 0, 1, 0, 0, 1, 0, 2, 0, 0, 1, 1, 1, 1, 0, 0, 1, 0...
## $ SibSp
## $ Parch
                <int> 0, 0, 0, 0, 1, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0...
                <chr> "330911", "363272", "240276", "315154", "3101298", "7538",...
## $ Ticket
                <dbl> 7.8292, 7.0000, 9.6875, 8.6625, 12.2875, 9.2250, 7.6292, 2...
## $ Fare
                ## $ Cabin
                <chr> "Q", "S", "Q", "S", "S", "S", "Q", "S", "C", "S", "S", "S"...
## $ Embarked
```

colSums(is.na(train) | train == " ") # Check for NAs and blank values

```
## PassengerId
                    Survived
                                    Pclass
                                                   Name
                                                                  Sex
                                                                               Age
##
                                                       0
                                                                    0
                                                                               177
              0
                            0
                                         a
##
          SibSp
                       Parch
                                    Ticket
                                                                Cabin
                                                                          Embarked
                                                   Fare
##
                                                       0
                                                                    0
```

```
colSums(is.na(test) | test == " ")
```

```
## PassengerId
                      Pclass
                                      Name
                                                     Sex
                                                                   Age
                                                                              SibSp
##
                            0
                                          0
                                                        0
                                                                    86
                                                   Cabin
                                                             Embarked
##
          Parch
                      Ticket
                                      Fare
##
               0
                            0
                                          1
                                                        0
                                                                     0
```

```
set.seed(444)

titanic_glm <- glm(Survived ~ Sex, data = train_copy, family = 'binomial')
summary(titanic_glm)</pre>
```

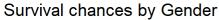
```
##
## Call:
## glm(formula = Survived ~ Sex, family = "binomial", data = train copy)
##
## Deviance Residuals:
       Min
                 10 Median
##
                                   3Q
                                           Max
## -1.6462 -0.6471 -0.6471
                               0.7725
                                        1.8256
##
## Coefficients:
##
               Estimate Std. Error z value Pr(>|z|)
                           0.1290
                                     8.191 2.58e-16 ***
## (Intercept)
                1.0566
                            0.1672 -15.036 < 2e-16 ***
## Sexmale
                -2.5137
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##
       Null deviance: 1186.7 on 890 degrees of freedom
## Residual deviance: 917.8 on 889 degrees of freedom
## AIC: 921.8
##
## Number of Fisher Scoring iterations: 4
```

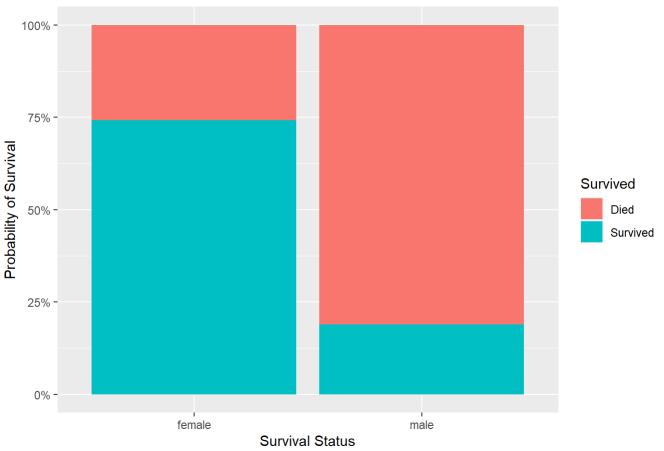
```
# Test for accuracy
predict_sex_survived <- predict(titanic_glm,newdata = test_copy,type = 'response')
# Since Survived can only be either 1 or 0, write if statement to round up of down the response
predict_sex_survived <- ifelse(predict_sex_survived>0.5,1,0)
error_1 <- mean(predict_sex_survived!=test_copy$Survived)
accuracy_1 <- 1-error_1
accuracy_1</pre>
```

[1] 0.7559809

```
prob_survival <- data.frame(prob_survival = titanic_glm$fitted.values,Survived = train_copy$Survived, Sex = train_copy$Sex)

ggplot(prob_survival, aes(fill = Survived, y = prob_survival, x = Sex)) +
    geom_bar(position = "fill", stat = "identity") +
    labs(x= "Survival Status", y = "Probability of Survival", title = "Survival chances by Gender"
) +
    scale_y_continuous(labels = scales::percent_format()) +
    scale_fill_discrete(labels = c("Died","Survived"))</pre>
```





logistic_complete <- glm(Survived~., data = train_copy, family = "binomial")
summary(logistic_complete)</pre>

```
##
## Call:
## glm(formula = Survived ~ ., family = "binomial", data = train_copy)
##
## Deviance Residuals:
##
      Min
                1Q Median
                                 3Q
                                         Max
##
  -2.6834 -0.6053 -0.4060
                             0.6202
                                      2.4785
##
## Coefficients:
##
                Estimate Std. Error z value Pr(>|z|)
## (Intercept) 16.633165 608.445775
                                    0.027 0.978191
## Pclass2
               -1.056168
                          0.304843 -3.465 0.000531 ***
## Pclass3
                          0.311508 -7.504 6.19e-14 ***
               -2.337544
## Sexmale
               -2.681168    0.201848    -13.283    < 2e-16 ***
               -0.043020 0.008119 -5.298 1.17e-07 ***
## Age
## SibSp
               ## Parch
               -0.099547
                          0.120556 -0.826 0.408955
## Fare
                0.002006
                          0.002463
                                     0.814 0.415414
## EmbarkedC
              -12.300751 608.445644 -0.020 0.983871
## EmbarkedQ
              -12.417556 608.445701 -0.020 0.983717
## EmbarkedS
              -12.718626 608.445631 -0.021 0.983323
##
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 1186.66 on 890 degrees of freedom
## Residual deviance: 779.97 on 880 degrees of freedom
## AIC: 801.97
##
## Number of Fisher Scoring iterations: 13
```

```
logistic_stepwise <- logistic_complete %>% stepAIC(direction = "both", trace = FALSE)
summary(logistic_stepwise)
```

```
##
## Call:
## glm(formula = Survived ~ Pclass + Sex + Age + SibSp, family = "binomial",
##
      data = train copy)
##
## Deviance Residuals:
##
      Min
               1Q
                   Median
                               3Q
                                      Max
## -2.7628 -0.5958 -0.4020
                           0.6177
                                   2,4872
##
## Coefficients:
              Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) 4.282319 0.421806 10.152 < 2e-16 ***
                       0.267854 -4.895 9.85e-07 ***
## Pclass2
            -1.311027
## Pclass3
             -2.547895
                       0.258793 -9.845 < 2e-16 ***
                       0.194536 -13.882 < 2e-16 ***
## Sexmale
             -2.700613
             ## Age
             ## SibSp
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 1186.66 on 890 degrees of freedom
## Residual deviance: 785.74 on 885 degrees of freedom
## AIC: 797.74
##
## Number of Fisher Scoring iterations: 5
```

```
AIC(logistic_complete,logistic_stepwise)
```

```
## df AIC
## logistic_complete 11 801.9660
## logistic_stepwise 6 797.7355
```

```
predict_logistic_test <- predict(logistic_stepwise,test_copy,type = "response")
predict_logistic_1 <- ifelse(predict_logistic_test>0.5,1,0)
cf_table_test <- table(predict_logistic_1,test_copy$Survived)
cf_table_test</pre>
```

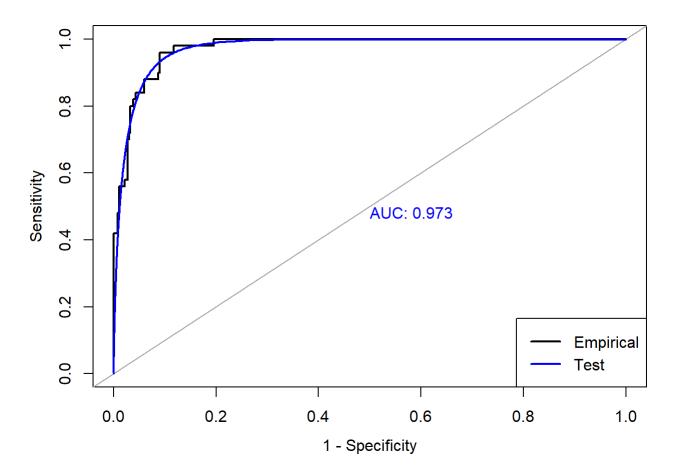
```
##
## predict_logistic_1 0 1
## 0 262 0
## 1 106 50
```

```
acc_test <- sum(diag(cf_table_test))/sum(cf_table_test)
acc_test</pre>
```

```
## [1] 0.7464115
sens_test <- cf_table_test[1]/(cf_table_test[1] + cf_table_test[2])</pre>
sens_test
## [1] 0.7119565
spec_test <- cf_table_test[4]/(cf_table_test[3] + cf_table_test[4])</pre>
spec test
## [1] 1
ppvtest <- cf_table_test[1]/(cf_table_test[1] + cf_table_test[3])</pre>
ppvtest
## [1] 1
npvtest <- cf_table_test[4]/(cf_table_test[2] + cf_table_test[4])</pre>
npvtest
## [1] 0.3205128
error_test <- mean(predict_logistic_1!=test_copy$Survived)</pre>
acc_test <- 1 - error_test</pre>
acc_test
## [1] 0.7464115
roc_test <- roc(test_copy$Survived,predict_logistic_test)</pre>
## Setting levels: control = 0, case = 1
## Setting direction: controls < cases
plot.roc(roc_test, col=par("fg"),plot=TRUE,print.auc = FALSE, legacy.axes = TRUE, asp =NA)
roc test
```

```
##
## Call:
## roc.default(response = test_copy$Survived, predictor = predict_logistic_test)
##
## Data: predict_logistic_test in 368 controls (test_copy$Survived 0) < 50 cases (test_copy$Survived 1).
## Area under the curve: 0.9743</pre>
```

```
plot.roc(smooth(roc_test), col="blue", add = TRUE,plot=TRUE,print.auc = TRUE, legacy.axes = TRUE
, asp =NA)
legend("bottomright", legend=c("Empirical", "Test"),
col=c(par("fg"), "blue"), lwd=2)
```



From the above observations it is clear that Logistic Regression does a great job in doing the requisite classifications. We'll now look at the performance of the Neural Networks on the same dataset below.

```
set.seed(777)

control <- trainControl(method = "repeatedcv", number = 10, repeats = 10)

logistic_bayesian <- train(Survived ~ FamilySize+CabinPos+Deck+Pclass+Sex+Age+SibSp+Parch+Fare+E
mbarked+Title+Mother+Child, data = train, method = "bayesglm", trControl = control, family = bin
omial(link = "logit"))

acc_bayesian <- paste0(round(max(logistic_bayesian$results$Accuracy),3)*100,'%')
cat('prediction accucary with bayesian logistic regression is ',acc_bayesian,'.\n')</pre>
```

```
## prediction accucary with bayesian logistic regression is 82.7% .
```

```
grid <- expand.grid(size = 1, decay = 0.01)

neuralnet <- train(Survived ~ FamilySize+CabinPos+Deck+Pclass+Sex+Age+SibSp+Parch+Fare+Embarked+
Title+Mother+Child, data = train, method = "nnet", trControl = control,maxit = 1000, trace = FAL
SE, tuneGrid = grid )

acc_nnet <- paste0(round(max(neuralnet$results$Accuracy),3)*100,'%')
cat('prediction accucary of neural net is ',acc_nnet,'.\n')</pre>
```

```
## prediction accucary of neural net is 82.9% .
```

With a simple Logistic Regression, we achieved an accuracy of ~77%. Though Neural Networks achieved an accuracy of ~83%, another form of Logistic Regression - Bayesian Logistic Regression managed to do the same. Hence, from the above observations it is clear that Logistic is better suited for this dataset than Neural Network.