Application of Neural Networks for Prediction of Authorization to Organ Donation

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Objective

Predict whether the family of a organ donor will say Yes/No to Organ Donation Methodology: Neural Network (Multilayer Perceptron)

In this project, no hidden layers were used. Linear basis function and sigmoidal activation function were used in the MLP

Specify the Input and Output Layer Size

```
library(ggplot2)
input_layer_size <- 8 # Excluding bias
output_layer_size <- 1</pre>
```

Loading data

```
r <- read.csv("C:/Users/pbagavan/SSIE519/mlclass-ex4/data.csv")

# include bias unit
bias <- rep(1, times=nrow(r))
r <- cbind(bias,r)

# Standardize age (AG) and Admission to Referral Time (AR)
r$AG <- ((r$AG*0.8)/95)+0.1
r$AR <- ((r$AR*0.08)/644.89)+0.1
r <- as.matrix(r)</pre>
```

Data Slicing

```
set.seed(1050)
index <- sample(1:nrow(r),round(0.70*nrow(r)))
train <- r[index,]
test <- r[-index,]</pre>
```

Training data

Preview table

head(round(train[1:5,],4)) bias AR AG Atele DSA_Def ODC RaceAfAm RaceAsian Timely Auth ## [1,] 1 0.1327 0.5716 0.9 0.1 0.1 0.1 0.1 0.1 0.9 ## [2,] 1 0.1025 0.5042 0.1 0.1 0.1 0.9 0.1 0.1 0.9 ## [3,] 1 0.1208 0.5716 0.1 0.1 0.1 0.1 0.1 0.1 0.9 ## [4.] 1 0.1000 0.7147 0.1 0.9 0.9 0.1 0.1 0.1 0.1 0.1 0.1 ## [5,] 0.1 0.9 1 0.1195 0.1000 0.1 0.1 0.1 # Storing the error in Epsilon_Store Epsilon_Store <- matrix(data=0,nrow=nrow(train),ncol=output_layer_size)</pre> Overall Error <- 0 # Input and output variables train_x <- train[,1:(input_layer_size+1)]</pre> train_Y <- train[,-(1:(input_layer_size+1))]</pre> # Initialize weights set.seed(500) w1 <- matrix(data=round((4*runif(input_layer_size+1,0,1)-2)/3,4), nrow=output layer size,ncol=input layer size+1, byrow = T)#t=1{1st Epoch, first step in epoch: zero out the c1 matrix} t=1 for(t in 1:465){ c1 <- matrix(data=0,nrow=output_layer_size,ncol=input_layer_size+1,</pre> bvrow = T) #i=1 {1st training example. First read in values for input layer activations} i <- 1 for(i in 1:nrow(train)){ # Forward propagate s <- train_x[i,] %*% t(w1) a <- 1/(1+exp(-s))# Backward propagate Epsilon <- a-train_Y[i]</pre> Epsilon_Store[i] <- Epsilon</pre> d <- Epsilon*a*(1-a)</pre> # Update c matrix c1 <- c1+t(train_x[i,])*d[1,1] #End of first epoch. Update weights $w1 \leftarrow w1 - 0.01*c1$ Overall_Error[t] <- 0.5*sum(Epsilon_Store^2)/input_layer_size</pre> }

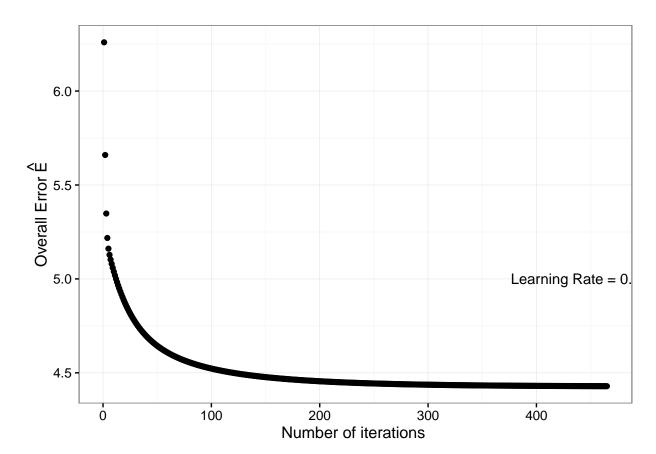
Input and out test variables

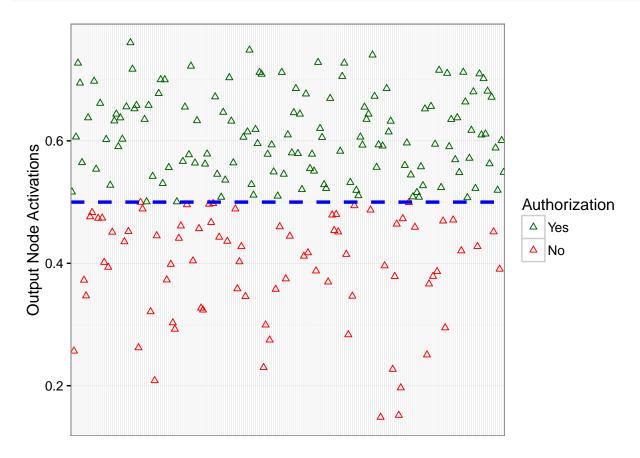
test_x <- test[,1:(input_layer_size+1)]
test_Y <- test[,-(1:(input_layer_size+1))]</pre>

```
# Save activations of output node
test_a <- rep(0, times=nrow(test))
i <- 1
for(i in 1:nrow(test)){
    # Forward propagate
    s <- test_x[i,] %*% t(w1)
    test_a[i] <- 1/(1+exp(-s))
}</pre>
```

Plotting activations

```
SampleNo <- 1:nrow(test_x)
OpActivations <- test_a
Authorization <- ifelse(OpActivations>=0.5, "darkgreen", "red")
plot_data <- as.data.frame(cbind(SampleNo,round(OpActivations,2), Authorization))
#Overall_Error
ggplot() + geom_point(aes(x=1:t,y=Overall_Error)) + theme_bw() +
    xlab("Number of iterations ")+ylab(expression(paste("Overall Error ",hat(E))))+
    annotate("text",x=440,y=5, label= "Learning Rate = 0.01")</pre>
```





Converting Auth from continuous to categorical variable

```
# Misclassification Rate
misclass <- round((sum(plot_data$category=="B"|plot_data$category=="C")/nrow(plot_data))*100,2)
misclass
## [1] 33.95</pre>
```

Threshold Distribution

```
df <- as.data.frame(cbind(test_a,test_Y))

plot_distribution <- function(df,threshold){
    v <- rep(NA, nrow(df))
    v <- ifelse(df$test_a >= threshold & df$test_Y == 0.9, "TP", v)
    v <- ifelse(df$test_a >= threshold & df$test_Y == 0.1, "FP", v)
    v <- ifelse(df$test_a < threshold & df$test_Y == 0.9, "FN", v)
    v <- ifelse(df$test_a < threshold & df$test_Y == 0.9, "FN", v)
    v <- ifelse(df$test_a < threshold & df$test_Y == 0.1, "TN", v)

df$pred_type <- v

ggplot(df, aes(x=test_Y, y=test_a))+
    geom_violin(fill=rgb(1,1,1,alpha = 0.6), color=NA)+
    geom_jitter(aes(color=pred_type),alpha=0.6)+
    geom_hline(yintercept=threshold,color="red",alpha=0.6)+
    scale_color_discrete(name=" ")+
    labs(title=sprintf("Threshold = %.2f", threshold))+
    xlab("Actual Value")+ylab("Predicted Value")
}</pre>
```

ROC Calculation

```
calculate_roc <- function(df, cost_of_fp, cost_of_fn, n=100) {
  tpr <- function(df, threshold) {
    sum(df$test_a >= threshold & df$test_Y == 0.9) / sum(df$test_Y == 0.9)
  }

fpr <- function(df, threshold) {
    sum(df$test_a >= threshold & df$test_Y == 0.1) / sum(df$test_Y == 0.1)
  }

cost <- function(df, threshold, cost_of_fp, cost_of_fn) {
    sum(df$test_a >= threshold & df$test_Y == 0.1) * cost_of_fp +
        sum(df$test_a <= threshold & df$test_Y == 0.1) * cost_of_fn
  }

roc <- data.frame(threshold = seq(0,1,length.out=n), tpr=NA, fpr=NA)
  roc$tpr <- sapply(roc$threshold, function(th) tpr(df, th))
  roc$cost <- sapply(roc$threshold, function(th) fpr(df, th))
  roc$cost <- sapply(roc$threshold, function(th) cost(df, th, cost_of_fp, cost_of_fn))</pre>
```

```
return(roc)
}
# ROC plotting
plot_roc <- function(roc, threshold, cost_of_fp, cost_of_fn) {</pre>
  library(gridExtra)
  library(grid)
  norm vec <- function(v) (v - min(v))/diff(range(v))</pre>
  idx_threshold = which.min(abs(roc$threshold-threshold))
  col_ramp <- colorRampPalette(c("green","orange","red","black"))(100)</pre>
  col_by_cost <- col_ramp[ceiling(norm_vec(roc$cost)*99)+1]</pre>
  p_roc <- ggplot(roc, aes(fpr,tpr)) +</pre>
    geom_line(color=rgb(0,0,1,alpha=0.3)) +
    geom_point(color=col_by_cost, size=4, alpha=0.5) +
    coord_fixed() +
    geom_line(aes(threshold, threshold), color=rgb(0,0,1,alpha=0.5)) +
    labs(title = sprintf(" ")) + xlab("1-Specificity") + ylab("Sensitivity") +
    geom_hline(yintercept=roc[idx_threshold, "tpr"], alpha=0.5, linetype="dashed") +
    geom_vline(xintercept=roc[idx_threshold,"fpr"], alpha=0.5, linetype="dashed")
  p_cost <- ggplot(roc, aes(threshold, cost)) +</pre>
    geom_line(color=rgb(0,0,1,alpha=0.3)) +
    geom_point(color=col_by_cost, size=4, alpha=0.5) +
    labs(title = sprintf(" ")) +
    geom_vline(xintercept=threshold, alpha=0.5, linetype="dashed")+
    xlab("Threshold")+ylab("Cost")
  sub_title <- sprintf("</pre>
                               threshold at %.2f - cost of FP = %d, cost of FN = %d", threshold, cost_of
  grid.arrange(p_roc, p_cost, ncol=2, sub=textGrob(sub_title, gp=gpar(cex=1), just="bottom"))
```

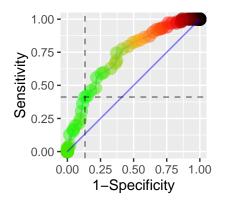
Cost-Specific Threshold Values

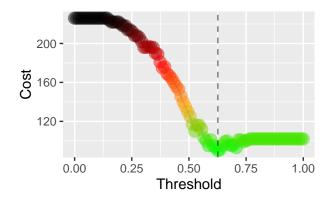
False Positives (FP): The model predicting incorrectly that the family will authorize when it actually does not

If I assume cost of 2 for FP and 1 for False Negatives (FN)

```
roc <- calculate_roc(df, 2,1,n=100)
Best_Threshold <- roc$threshold[which.min(roc$cost)]
Best_Threshold
## [1] 0.6262626

plot_roc(roc, Best_Threshold, 2, 1)</pre>
```





hreshold at 0.63 – cost of FP = 2, cost of FN = 1

plot_distribution(df, Best_Threshold)

