## Logistic regression algorithm implementation

```
For the below code:

* the feature_matrix is the matrix of features

* the labels is the vector of "true" classes

* the learn_rate is the learning rate for the gradient descent procedure

* the iters is the number of iterations to run gradient descent

* the betas is a vector of the coefficients

* the n is the number of predictors

* the m is the number of training examples (e.g., number of subjects)

The sigmoid logistic function:

sigmoid <- function(x){
    1/(1+exp(-x))}

}

The initialize_betas <- function(feature_matrix){
    betas <- as.matrix(rep(0, ncol(feature_matrix)))}

}
```

The cost function to calculate logistic loss:

```
cost <- function(betas, feature_matrix, labels){
  m <- nrow(feature_matrix)
  probs <- sigmoid(feature_matrix%*%betas)
  L <- (1/m)*sum( labels * log(probs) + (1-labels) * log( 1-probs ) )
}</pre>
```

The gradient descent function to calculate the average gradient:

```
gradient <- function(betas, feature_matrix, labels){
  n <- ncol(feature_matrix)
  probs <- sigmoid(feature_matrix%*%betas)
  dcost_dbeta <- t(feature_matrix) %*% (probs-labels)
  avg_gradient <- (1/n)*dcost_dbeta
}</pre>
```

The update\_betas function to use the gradient calculated above and learning rate to update the betas:

```
update_betas <- function(betas, feature_matrix, labels, learn_rate){
  gradient <- gradient(betas, feature_matrix, labels)
  betas <- betas - learn_rate*gradient
}</pre>
```

The logistic\_regression function combining all of the above:

```
logistic_regression <- function(feature_matrix, labels, learn_rate, iters, print_every){</pre>
  # add a column of ones at the beginning of the feature matrix to calculate the
  # intercept
  feature_matrix <- cbind(as.matrix(rep(1,nrow(feature_matrix))),feature_matrix)</pre>
  # save the cost to print
  cost_print <- c()</pre>
  # get the initial beta values
  betas <- initialize betas(feature matrix)</pre>
  for(i in 1:iters){
    betas <- update_betas(betas, feature_matrix, labels, learn_rate)</pre>
    cost <- cost(betas, feature_matrix, labels)</pre>
    cost_print[i] <- cost</pre>
    if(i %% print_every == 0){
      print(paste("Iteration#:",i))
      print(paste("Cost:",cost_print[i]))
  }
  return(list(betas = betas, final_cost = cost, all_costs = cost_print))
```

The predictor function to calculate predictions from betas obtained above:

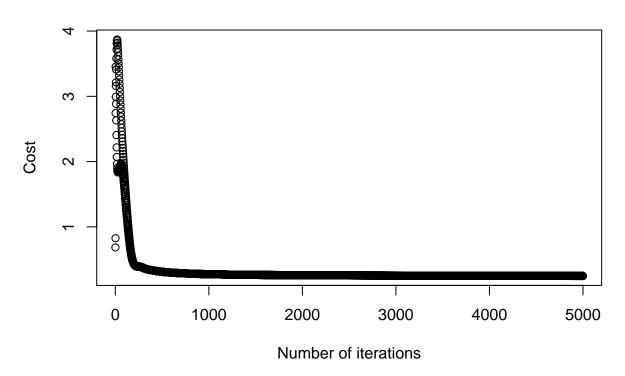
```
predictor <- function(feature_matrix, betas){
  preds <- sigmoid(feature_matrix%*%betas)
  preds
}</pre>
```

Train the model:

```
## [1] "Iteration#: 1000"
## [1] "Cost: -0.274708347977415"
## [1] "Iteration#: 2000"
## [1] "Cost: -0.259049322661902"
## [1] "Iteration#: 3000"
## [1] "Cost: -0.253871873882793"
## [1] "Iteration#: 4000"
## [1] "Cost: -0.250626029160736"
## [1] "Iteration#: 5000"
## [1] "Cost: -0.248113438653087"
```

Plot cost by iterations:

## **Cost versus iterations**

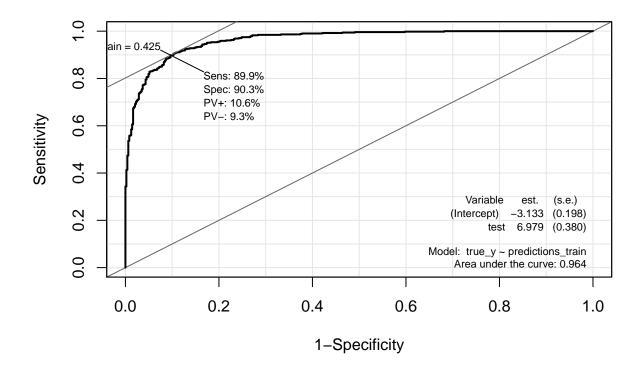


## Test the model:

```
# add a row of 1s to the train_data to conform to betas having an intercept term calculated
train_data_1 <- cbind(1, train_data)

# assess the predictions on the training data
predictions_train <- predictor(train_data_1, log_model$betas)

# find the optimal cutoff with an ROC curve
ROC(test = predictions_train, stat = true_y, plot = "ROC")</pre>
```



The best cutoff based on the ROC curve is 0.48 to maximize both sensitivity and specificity, 91.4% and 89.7%, respectively.

Make predictions using this cutoff value:

```
y_preds <- ifelse(predictions_train<0.425, 0, 1)
# print confusion matrix
table(true_y, y_preds)</pre>
```

```
## y_preds
## true_y 0 1
## 0 440 47
## 1 52 461
```

Make predictions on the test data:

```
# add a row of 1s to the test_data to conform to betas having an intercept term calculated
test_data_1 <- cbind(1, test_data)

predictions_test <- predictor(test_data_1, log_model$betas)

# using the same cutoff value that was found above on the train data
y_test <- ifelse(predictions_test<0.425, 0, 1)
table(y_test)</pre>
```

## y\_test ## 0 1 ## 479 521