## Problem Set #3

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- 1. (10 points) This question involves the Weekly dataset, which is included in the package ISLR2.
- (a) Fit a logistic regression with Direction as the response and the five lag variables plus Volume as predictors. Use the summary() to print the results.
- (b) For which of the predictors can you reject the null hypothesis H0: j = 0?
- (c) Compute the confusion matrix and overall fraction of correct predictions.
- (d) Repeat (c) using LDA. Which method performs better on this data?

```
# Load necessary libraries
library(ISLR2) # For the Weekly dataset
## Warning: package 'ISLR2' was built under R version 4.4.1
library(MASS)
               # For LDA (Linear Discriminant Analysis)
## Attaching package: 'MASS'
## The following object is masked from 'package:ISLR2':
##
##
      Boston
# Inspect the Weekly dataset
head(Weekly)
##
                 Lag2
                                             Volume Today Direction
    Year
           Lag1
                        Lag3
                               Lag4
                                     Lag5
## 1 1990 0.816 1.572 -3.936 -0.229 -3.484 0.1549760 -0.270
                                                               Down
Down
## 3 1990 -2.576 -0.270 0.816 1.572 -3.936 0.1598375 3.514
                                                                 Uр
                                                                 Uр
## 4 1990 3.514 -2.576 -0.270 0.816 1.572 0.1616300 0.712
## 5 1990 0.712 3.514 -2.576 -0.270 0.816 0.1537280 1.178
                                                                 Uр
## 6 1990 1.178 0.712 3.514 -2.576 -0.270 0.1544440 -1.372
                                                               Down
# (1a) Fit a logistic regression model
logistic_model <- glm(Direction ~ Lag1 + Lag2 + Lag3 + Lag4 + Lag5 + Volume,</pre>
                    data = Weekly, family = binomial)
# Print the summary of the logistic model
cat("(1a) Summary of Logistic Regression Model:\n")
```

## (1a) Summary of Logistic Regression Model: summary(logistic\_model) ## ## Call: ## glm(formula = Direction ~ Lag1 + Lag2 + Lag3 + Lag4 + Lag5 + ## Volume, family = binomial, data = Weekly) ## ## Coefficients: ## Estimate Std. Error z value Pr(>|z|)## (Intercept) 0.26686 0.08593 3.106 0.0019 \*\* -0.04127 0.02641 -1.563 0.1181 ## Lag1 ## Lag2 0.05844 0.02686 2.175 0.0296 \* ## Lag3 -0.01606 0.02666 -0.602 0.5469 ## Lag4 -0.02779 0.02646 -1.050 0.2937 -0.01447 0.02638 -0.549 0.5833 ## Lag5 -0.02274 0.03690 -0.616 0.5377 ## Volume ## ---## Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.05 '.' 0.1 ' ' 1 ## (Dispersion parameter for binomial family taken to be 1) ## ## Null deviance: 1496.2 on 1088 degrees of freedom ## Residual deviance: 1486.4 on 1082 degrees of freedom ## AIC: 1500.4 ## Number of Fisher Scoring iterations: 4 # (1b) Identify predictors for which the null hypothesis (HO: j = 0) can be rejected # Based on the p-values from the logistic regression summary cat("(1b) Predictors for which we can reject H0 (j = 0):\n") ## (1b) Predictors for which we can reject H0 (j = 0): # Extract p-values from the model summary and display significant ones logistic\_summary <- summary(logistic\_model)</pre> p\_values <- coef(logistic\_summary)[, 4] # p-values are in the 4th column # Print only the significant predictors (p-value < 0.05) significant\_predictors <- names(p\_values[p\_values < 0.05])</pre> print(significant\_predictors)

```
## [1] "(Intercept)" "Lag2"

# (1c) Compute confusion matrix and overall fraction of correct predictions for Logistic Regression

# Predict the probabilities using logistic regression

predicted_probs <- predict(logistic_model, type = "response")</pre>
```

```
# Convert probabilities to class labels (threshold of 0.5)
predicted_classes <- ifelse(predicted_probs > 0.5, "Up", "Down")
# Create confusion matrix for logistic regression
confusion_matrix <- table(Predicted = predicted_classes, Actual = Weekly$Direction)</pre>
cat("(1c) Confusion Matrix for Logistic Regression:\n")
## (1c) Confusion Matrix for Logistic Regression:
print(confusion_matrix)
##
            Actual
## Predicted Down Up
##
       Down 54 48
        Uр
              430 557
# Calculate the overall fraction of correct predictions (accuracy)
logistic_accuracy <- sum(diag(confusion_matrix)) / sum(confusion_matrix)</pre>
cat("Logistic Regression Accuracy: ", logistic_accuracy, "\n")
## Logistic Regression Accuracy: 0.5610652
# (1d) Perform Linear Discriminant Analysis (LDA) and repeat confusion matrix and accuracy calculation
# Fit the LDA model
lda_model <- lda(Direction ~ Lag1 + Lag2 + Lag3 + Lag4 + Lag5 + Volume, data = Weekly)</pre>
# Make predictions using LDA
lda_predictions <- predict(lda_model)</pre>
# Create confusion matrix for LDA
lda_confusion_matrix <- table(Predicted = lda_predictions$class, Actual = Weekly$Direction)</pre>
cat("(1d) Confusion Matrix for LDA:\n")
## (1d) Confusion Matrix for LDA:
print(lda_confusion_matrix)
            Actual
## Predicted Down Up
##
       Down 52 46
        Uр
              432 559
# Calculate the overall fraction of correct predictions for LDA (accuracy)
lda_accuracy <- sum(diag(lda_confusion_matrix)) / sum(lda_confusion_matrix)</pre>
cat("LDA Accuracy: ", lda_accuracy, "\n")
```

## LDA Accuracy: 0.5610652

```
# Compare the accuracy of Logistic Regression vs LDA
if (logistic_accuracy > lda_accuracy) {
    cat("Logistic Regression performs better with an accuracy of", logistic_accuracy, "\n")
} else {
    cat("LDA performs better with an accuracy of", lda_accuracy, "\n")
}
```

## LDA performs better with an accuracy of 0.5610652

Question #2

```
# Load necessary libraries
library(ggplot2)
```

## Warning: package 'ggplot2' was built under R version 4.4.1

```
# Load the dataset
setwd("C:/Users/ibach/OneDrive - Terillium/Desktop/Purdue MSBA/Machine Learning/HW 3")
data <- read.csv("logit_data.csv")
y <- data$grade
x <- data$hours</pre>
```

2A):

## Log-Likelihood Function

We take the natural logarithm of the likelihood function:

$$\ell(\beta \mid y) = \log L(\beta \mid y) = \sum_{i=1}^n \left( y_i \log(p(x_i;\beta)) + (1-y_i) \log(1-p(x_i;\beta)) \right)$$

Now, substitute the expansion for  $p(x_i; \beta)$  and  $1 - p(x_i; \beta)$ :

$$\ell(\beta \mid y) = \sum_{i=1}^n \left( y_i \log \left( \frac{1 + \exp(\beta_0 + \beta_1 x_i)}{\exp(\beta_0 + \beta_1 x_i)} \right) + (1 - y_i) \log \left( \frac{1}{1 + \exp(\beta_0 + \beta_1 x_i)} \right) \right)$$

Simplified, the log-likelihood function becomes:

$$\ell(\beta \mid y) = \sum_{i=1}^{n} (y_i(\beta_0 + \beta_1 x_i) - \log(1 + \exp(\beta_0 + \beta_1 x_i)))$$

Thus, the full log-likelihood function is written as:

$$\ell(\beta \mid y) = -\sum_{i=1}^n \log(1 + \exp(\beta_0 + \beta_1 x_i)) + \sum_{i=1}^n y_i (\beta_0 + \beta_1 x_i)$$

```
# (2a) Log-likelihood function
log_likelihood <- function(beta, y, x) {
  beta0 <- beta[1]
  beta1 <- beta[2]
  sum(-log(1 + exp(beta0 + beta1 * x)) + y * (beta0 + beta1 * x))
}</pre>
```

2B):

$$\ell(\beta \mid y) = \log L(\beta \mid y) = \sum_{i=1}^n \left( y_i \log(p(x_i;\beta)) + (1-y_i) \log(1-p(x_i;\beta)) \right)$$

Now, substitute the expansion for  $p(x_i; \beta)$  and  $1 - p(x_i; \beta)$ :

$$\ell(\beta \mid y) = \sum_{i=1}^n \left( y_i \log \left( \frac{1 + \exp(\beta_0 + \beta_1 x_i)}{\exp(\beta_0 + \beta_1 x_i)} \right) + (1 - y_i) \log \left( \frac{1}{1 + \exp(\beta_0 + \beta_1 x_i)} \right) \right)$$

Simplified, the log-likelihood function becomes:

$$\ell(\beta \mid y) = \sum_{i=1}^n \left( y_i (\beta_0 + \beta_1 x_i) - \log(1 + \exp(\beta_0 + \beta_1 x_i)) \right)$$

Thus, the full log-likelihood function is written as:

$$\ell(\beta \mid y) = -\sum_{i=1}^n \log(1 + \exp(\beta_0 + \beta_1 x_i)) + \sum_{i=1}^n y_i (\beta_0 + \beta_1 x_i)$$

```
# (2b) Gradient of the log-likelihood
gradient <- function(beta, y, x) {
  beta0 <- beta[1]
  beta1 <- beta[2]
  p <- 1 / (1 + exp(-(beta0 + beta1 * x)))
  grad_beta0 <- sum(y - p)
  grad_beta1 <- sum((y - p) * x)
  return(c(grad_beta0, grad_beta1))
}</pre>
```

2C):

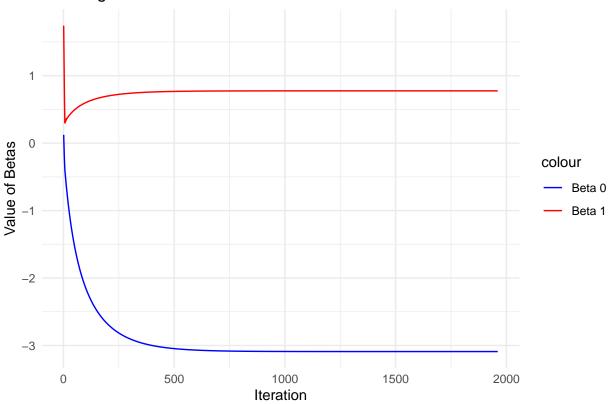
```
# (2c) Maximum likelihood estimates using glm
model <- glm(grade ~ hours, family = binomial(link = "logit"), data = data)
summary(model)</pre>
```

```
##
## Call:
## glm(formula = grade ~ hours, family = binomial(link = "logit"),
## data = data)
##
## Coefficients:
## Estimate Std. Error z value Pr(>|z|)
```

```
## (Intercept) -3.09053
                            0.36962 -8.361
                                               <2e-16 ***
                0.77508
                            0.07806
                                      9.930 <2e-16 ***
## hours
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
       Null deviance: 566.69 on 499 degrees of freedom
## Residual deviance: 236.73 on 498 degrees of freedom
## AIC: 240.73
##
## Number of Fisher Scoring iterations: 7
2D):
# (2d) Gradient ascent
gradient_ascent <- function(y, x, learning_rate = 0.001, epsilon = 1e-8, max_iter = 10000) {</pre>
  beta <- c(0, 0) # Initial values
  beta_history <- matrix(0, nrow = max_iter, ncol = 2)</pre>
  for (i in 1:max_iter) {
    grad <- gradient(beta, y, x)</pre>
    beta_new <- beta + learning_rate * grad</pre>
    beta_history[i, ] <- beta_new</pre>
    if (sqrt(sum((beta_new - beta)^2)) < epsilon) {</pre>
      beta_history <- beta_history[1:i, ]</pre>
      break
    }
    beta <- beta_new
 return(list(beta = beta, history = beta_history))
# Run gradient ascent
result <- gradient_ascent(y, x)
beta_estimates <- result$beta</pre>
beta_history <- result$history</pre>
2E):
# (2e) Predict probability of not getting an A for x = 10
x_new <- 10
p_{ext} = A < 1 / (1 + exp(-(beta_estimates[1] + beta_estimates[2] * x_new)))
p_not_getting_A <- 1 - p_getting_A</pre>
# Scatter plot of beta values
beta_df <- data.frame(iteration = 1:nrow(beta_history),</pre>
                      beta0 = beta_history[, 1],
                      beta1 = beta_history[, 2])
ggplot(beta_df, aes(x = iteration)) +
  geom_line(aes(y = beta0, color = "Beta 0")) +
  geom line(aes(y = beta1, color = "Beta 1")) +
 labs(title = "Convergence of Gradient Ascent for Beta Estimates",
```

```
x = "Iteration", y = "Value of Betas") +
scale_color_manual(values = c("Beta 0" = "blue", "Beta 1" = "red")) +
theme_minimal()
```

# Convergence of Gradient Ascent for Beta Estimates



# Output the probability that the student will not get an A p\_not\_getting\_A

## [1] 0.009374841