CMDA--3654

Homework 7

Eduardo Salvador

Due as a .pdf upload

Problem 1: [50 pts] Logistic Regression

Hermon Bumpus analyzed various characteristics of some house sparrows that were found on the ground after a severe winter storm in 1898. Some of the sparrows survived and some perished. The data on male sparrows in found in bumpus.csv are survival status (survived, perished), age (1 = adult, 2 = juvenile), the total length from tip of beak to tip of tail (in mm), the alar extent (length from tip to tip of the extended wings, in mm), the weight in grams, the length of the head in mm, the length of the humerus (arm bone, in inches), the length of the femur (thigh bones, in inches), the length of the tibio-tarsus (leg bone, in inches), the breadth of the skull in inches, and the length of the sternum in inches.

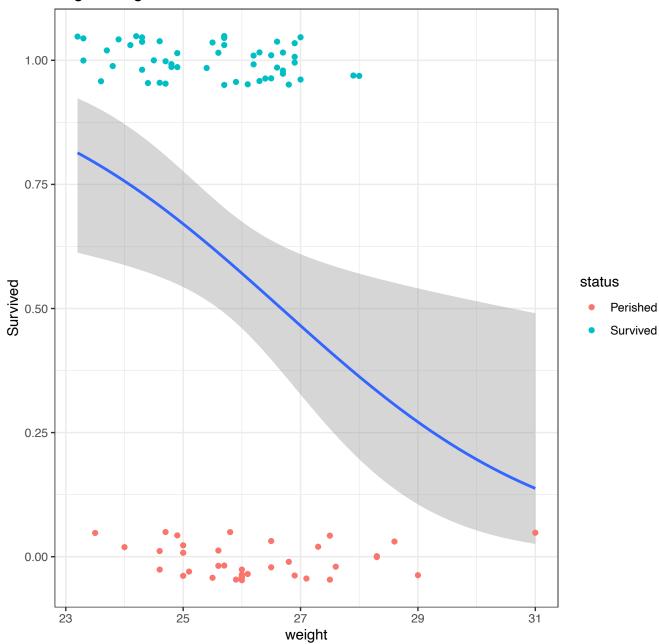
Analyze the data to see whether the probability of survival is associated with physical characteristics of the birds.

This would be consistent, according to Bumpus, with the theory of natural selection: those that survived did so because of some superior physical traits. Realize that (i) the sampling is from a population of grounded sparrows, and (ii) physical measurements and survival are both random.

a. Assuming that Weight is the only explanatory variable, fit a logistic regression model with Status as the response variable and answer the following questions.

```
library(tidyverse)
#Reading csv file
bump<-read.csv("/Users/eduardosalvador/Desktop/FINAL\ Spring\ Semester\ 2021/CMDA\ /Assignments/HW7/bumpus.csv
#Creating y variable of people who survived and those who have perished
bump <-bump %>%
    mutate(mystatus=if_else(status=="Survived",1,0))
#Creating regression model with ggplot
ggplot(bump,aes(x=weight,y=mystatus))+theme_bw()+ylab("Survived")+ggtitle("Logistic regression model with Stat
    geom_smooth(method="glm",method.args=list(family="binomial"))+geom_point(aes(color=status),position=position
```

Logistic regression model with Status



i. What is the probability a bird that weighs 25 grams survives?

#Creating fitting generalized linear model
bumpglm<-glm(mystatus~weight,data=bump,family=binomial)
head(bumpglm)</pre>

\$coefficients

(Intercept) weight 11.3201190 -0.4244016

\$residuals

1	2	3	4	5	6	7	8
1.397612	2.101072	2.101072	1.365255	1.335531	1.929158	1.414850	1.350078
9	10	11	12	13	14	15	16
1.271379	1.818079	1.818079	1.451600	1.582563	1.283144	1.661664	1.661664
17	18	19	20	21	22	23	24

1.929158	2.011470	1.308227	1.432835	2.756154	2.683182	1.720279	1.661664
25					30		
1.969441					2.011470		
33					38 -2.576863		40
1.634170					-2.576863 46		-2.045226 48
==					-2.330662		
49					2.330002		2.949027 56
					-2.330662		
57	58		60			63	64
-1.704036	-1.159401				1.451600	1.853546	1.381090
65	66	67	68	69	70	71	72
1.238936	2.011470	1.890550	2.101072	1.365255	2.148808	2.055320	1.471178
73			76			79	80
1.784086					-4.844644		-1.441416
81					86		
-3.310350	-1.766404	-2.511341	-1.372496	-3.034151	-1.704036	-2.330662	
****	_						
\$fitted.va		0	4	_		7	0
1	2				6		8
0.7155063	10				0.5183607 14		16
					0.7793359		
17	18				22		24
					0.3726918		
25	26				30		32
0.5077583	0.8136648	0.6018064			0.4971490		0.7719510
33	34			37		39	40
0.6119313	0.4653743	0.6979171	0.5183607	0.5605111	0.6119313	0.5813011	0.6219604
41	42	43	44	45	46	47	48
0.4029073	0.5915936			0.5183607	0.5709374	0.4548322	0.6609741
49	50			53		55	56
0.5709374					0.5709374		
57					62		64
					0.6888951		
65		67			70 0.4653743		
73	74	75	76	0.7324639	78	79	80
					0.7935865		
81	82		84	85	86	87	0.3002370
					0.4131580		
0.00.01.1	0.1000.02	0.0020001	0.2.2200	0.0.01200	0.1101000	0.0.000.1	
\$effects							
(Intercept	c) we:	ight					
-1.477207	73 2.4414	4522 0.89	957280 0	. 6474331	0.6349562	0.841382	25
0.667115	0.641	1371 0.60	054655 0	.8057024	0.8057024	0.681012	23
0.727577	75 0.6112	2022 0.7	543449 0	.7543449	0.8413825	0.867521	12
0.622902	23 0.6739	9782 1.09	949774 1	. 0733857	0.7737856	0.754344	19
0 054003	0 502	1002 0 71	542440 0	0171517	0 6474221	0 967501	10
0.854203	34 0.583	1223 0.73	543449 0	.8171517	0.6474331	0.867521	12
0.688230	0.6170	0120 O 7/	451209 O	.9106571	0.6739782	-1.159967	73
0.000230	0.01/	V120 U.1'	101200 0	. 0100011	0.0100102	1.103301	
-1.220138	32 -1.3069	9592 -1.2!	531900 -1	. 3260869	-1.0257537	-1.270533	25
							-
-1.455018	30 -1.2363	3991 -1.1	599673 -1	. 2363991	-1.0820472	-1.409199	91

```
-1.2363991 -1.3069592 -1.4317355 -1.5295544 -1.4317355 -1.2363991
 -0.9559097 -1.5295544 -1.0365033 -0.7249919 -0.9559097
                                                        0.6671153
 0.7362041 0.6810123 0.8171517 0.6538543 0.5886292
                                                       0.8675212
 0.6882306
 0.7946710 0.8542034 0.5886292 -1.7015995 -1.1193688 -1.8709810
 -1.1065901 -0.9281538 -1.5039064 -1.0587324 -1.2884483 -0.8924854
 -1.4317355 -1.0365033 -1.2363991
$R
           (Intercept)
                          weight
(Intercept) -4.417004 -114.334632
            0.000000 -5.752693
weight
$rank
[1] 2
summary(bumpglm)
glm(formula = mystatus ~ weight, family = binomial, data = bump)
Deviance Residuals:
        1Q Median 3Q
                                    Max
-1.7764 -1.2455 0.7607 1.0247
                                 1.4240
Coefficients:
          Estimate Std. Error z value Pr(>|z|)
(Intercept) 11.3201 4.5053 2.513 0.0120 *
          weight
---
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
   Null deviance: 118.01 on 86 degrees of freedom
Residual deviance: 111.25 on 85 degrees of freedom
AIC: 115.25
Number of Fisher Scoring iterations: 4
#Use predict function to get probability a bird that weights 25 grams
predict(bumpglm,newdata = data.frame(weight=25),type="response")
       1
0.6704185
#The probability a bird that weighs 25 grams survives is 67%.
ii. What is the probability a bird that weighs 30 grams survives?
#Use predict function to get probability a bird that weights 30 grams
predict(bumpglm,newdata = data.frame(weight=30),type="response")
```

```
1
0.1959299
```

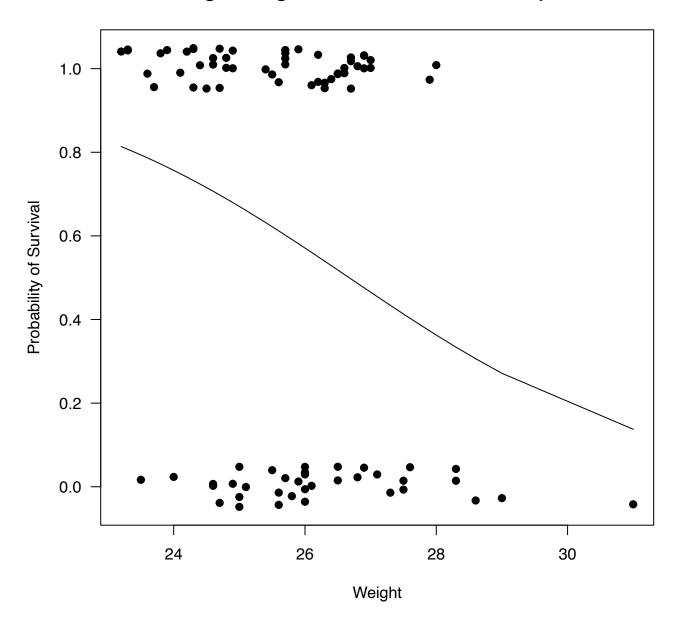
lines(ordw~sort(weight),data=bump)

#The probability a bird that weighs 30 grams survives is 19.59%.

iii. Plot the logistic regression model with a scatterplot of the observations. Make sure all plot elements m

#Creating scatter points with plot
plot(jitter(mystatus, amount = 0.05)~weight,data=bump,pch=19,las=1, main="Logistic regression model with a sca
#Creating line for regression model with lines function
ordw<-predict(bumpglm,data.frame(weight=sort(bump\$weight)),type = "response")

Logistic regression model with a scatterplot



iv. Suppose we come up with a classification rule that says we will consider a bird as having survived if the #The body weights would have to be 25.6 or lower.

b. Now consider using all of the physical characteristics as possible predictor variables in a logistic regression with Status as the response. Find the best subset of explanatory variables using stepAIC(). State the best model in terms of log-odds. Use this model for the remaining questions.

```
library (MASS)
#Creating fitting generalized linear model with all the variables
full\_model < -glm(mystatus - age+total\_length + alar\_extent + weight + head\_length + humerus\_length + femur\_length + tibio\_tarrelength + femur\_length + fe
#Creating fitting generalized linear model with null
null_model <- glm(mystatus ~ 1, data = bump, family= binomial)</pre>
#Finding best best subset of exploratory values using stepAIC()
best_model <- stepAIC(full_model, scope = list(lower = null_model, full_model), trace = 0, direction = "both")
summary (best_model)
Call:
glm(formula = mystatus ~ total_length + weight + humerus_length +
          sternum, family = binomial, data = bump)
Deviance Residuals:
         Min 1Q Median
                                                                              30
-2.2234 -0.5648 0.1540
                                                                     0.6094
                                                                                           2.2701
Coefficients:
                                    Estimate Std. Error z value Pr(>|z|)
 (Intercept)
                                      49.9861 18.4879 2.704 0.006857 **
total_length
                                       weight
                                       humerus_length 72.3327
                                                                  20.7640 3.484 0.000495 ***
                                                            11.7780 2.324 0.020101 *
                                       27.3775
sternum
 ---
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
 (Dispersion parameter for binomial family taken to be 1)
         Null deviance: 118.008 on 86 degrees of freedom
Residual deviance: 68.612 on 82 degrees of freedom
AIC: 78.612
Number of Fisher Scoring iterations: 6
      c. Is age group important? If so, how does the odds of survival change?
```

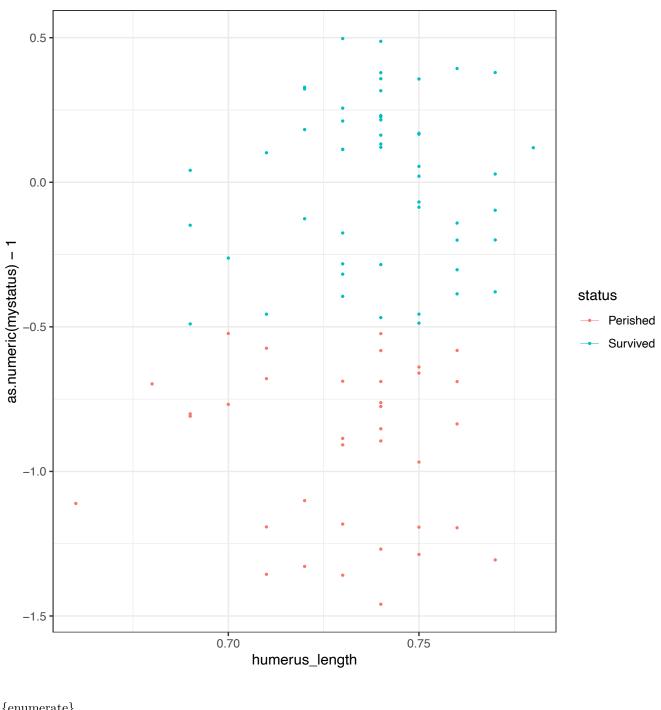
#It is NOT important in this case.

d. Is total length important? If so, if the total length is increased from 160 to 165 mm, and assuming everything else is held constant, what is change in odds of survival?

#total_length is important and the change in odds survival from a total length increaser from 160 to 165 would

e. Plot Status versus η the log-odds function and overlay the logistic regression curve.

```
#Created ggplot
ggplot(bump,aes(x=humerus_length, y=as.numeric(mystatus)-1))+theme_bw()+geom_smooth(aes(color=status),method =
```



 $\verb|\end{enumerate}|$

Problem 2: [50 pts] Classification using LDA, QDA, and SVM

Load the wine dataset from the rattle package in R.

Consider Type to be the response variable, and all other variables as features.

a. Describe the dataset in your own words, in 2-3 lines.

```
library(rattle)
data("wine")
view(wine)
```

#The wine data set includes 14 variables and 178 obs. Out of those variables, #the first one is type which consists of three types. The rest, are 13 chemical element variables.

b. Perform classification using LDA (linear discriminant analysis). Display the Confusion Matrix. Report the classification error rate.

```
library(caret)
library(klaR)
#The . allows to put all the variables, would've been beneficial to know earlier
LINDA<-lda(Type~.,data=wine)
#Predict value based on the input data
PLINDA<-predict(LINDA,data=wine)
#Confusion Matrix using LDA
confusionMatrix(PLINDA$class,wine$Type)</pre>
```

Confusion Matrix and Statistics

Reference

Prediction 1 2 3 1 59 0 0 2 0 71 0 3 0 0 48

Overall Statistics

Accuracy: 1

95% CI : (0.9795, 1)

No Information Rate : 0.3989 P-Value [Acc > NIR] : < 2.2e-16

Kappa: 1

Mcnemar's Test P-Value : NA

Statistics by Class:

	Class: 1	Class: 2	Class: 3
Sensitivity	1.0000	1.0000	1.0000
Specificity	1.0000	1.0000	1.0000
Pos Pred Value	1.0000	1.0000	1.0000
Neg Pred Value	1.0000	1.0000	1.0000
Prevalence	0.3315	0.3989	0.2697
Detection Rate	0.3315	0.3989	0.2697
Detection Prevalence	0.3315	0.3989	0.2697
Balanced Accuracy	1.0000	1.0000	1.0000

#The classification error rate is equal to 0.

c. Perform classification using QDA (quadratic discriminant analysis). Display the Confusion Matrix. Report the classification error rate.

```
#Performing QDA on all variables
quda<-qda(Type~.,data=wine)
#Predict value based on the input data and creating confusion matrix
confusionMatrix(predict(quda)$class,wine$Type)</pre>
```

Confusion Matrix and Statistics

Reference

Prediction 1 2 3

1 59 1 0

2 0 70 0

3 0 0 48

Overall Statistics

Accuracy : 0.9944

95% CI : (0.9691, 0.9999)

No Information Rate : 0.3989 P-Value [Acc > NIR] : < 2.2e-16

Kappa : 0.9915

Mcnemar's Test P-Value : NA

Statistics by Class:

	Class: 1	Class: 2	Class: 3
Sensitivity	1.0000	0.9859	1.0000
Specificity	0.9916	1.0000	1.0000
Pos Pred Value	0.9833	1.0000	1.0000
Neg Pred Value	1.0000	0.9907	1.0000
Prevalence	0.3315	0.3989	0.2697
Detection Rate	0.3315	0.3933	0.2697
Detection Prevalence	0.3371	0.3933	0.2697
Balanced Accuracy	0.9958	0.9930	1.0000

#The classification error rate is equal to almost 0, its accuracy is 0.994382

d. Perform classification using SVM (support vector machines). Display the Confusion Matrix. Report the classification error rate.

library(e1071)

#Creating a support vector machine of all variables in wine data

SUVM<-svm(Type~.,data=wine)</pre>

#Making confusion Matrix

confusionMatrix(predict(SUVM),wine\$Type)

Confusion Matrix and Statistics

Reference

Prediction 1 2 3

1 59 0 0

2 0 71 0

3 0 0 48

Overall Statistics

Accuracy : 1

95% CI : (0.9795, 1)

No Information Rate : 0.3989 P-Value [Acc > NIR] : < 2.2e-16

Kappa: 1

Mcnemar's Test P-Value : NA

Statistics by Class:

	Class: 1	Class: 2	Class: 3
Sensitivity	1.0000	1.0000	1.0000
Specificity	1.0000	1.0000	1.0000
Pos Pred Value	1.0000	1.0000	1.0000
Neg Pred Value	1.0000	1.0000	1.0000
Prevalence	0.3315	0.3989	0.2697
Detection Rate	0.3315	0.3989	0.2697
Detection Prevalence	0.3315	0.3989	0.2697
Balanced Accuracy	1.0000	1.0000	1.0000

#0

e. Rank the classification methods in your order of preference for this dataset, and justify your preference. (Hint: Note that the error rates should be calculated by cross-validation)

```
#The classification error rate was the same for LDA, and SVM, the QDA was a little higher #but by looking at the cross-validation, it can be cause by randomness in the data. #All 3 methods are good. My preference would be LDA since it is the one I used the most.
```

Problem 3: [20 pts Extra Credit]: More Logistic Regression

Load the mtcars data in R.

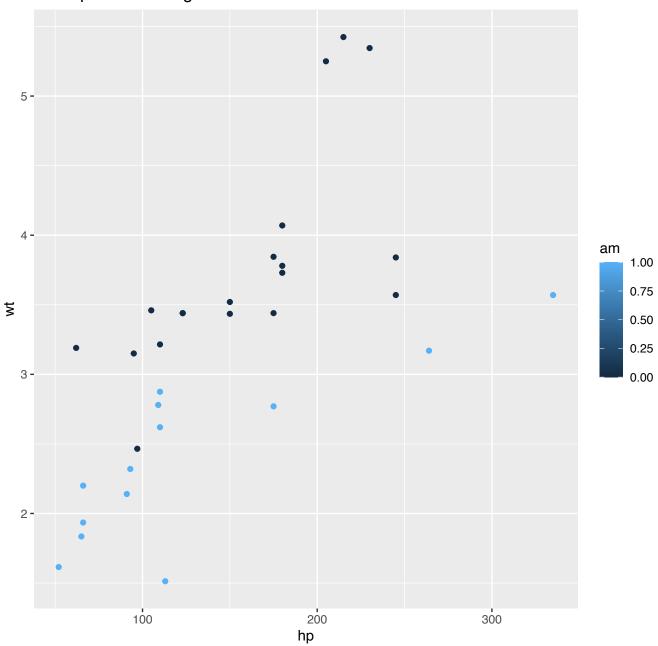
a. Describe the variable am in one sentence. We will consider am to be the response variable in the following questions.

#am variable is described as transmission where 1 is for manual and 0 for automatic

b. Construct a plot of hp (x-axis) and wt (y-axis), with different colors for automatic and manual transmission. From the plot, do you think automatic and manual transmission can be distinguished by weight and horsepower?

ggplot(mtcars,aes(x=hp,y=wt))+ggtitle("Horsepower vs Weight")+geom_point(aes(color=am))

Horsepower vs Weight



#By looking at the graph, I can say that cars with more weight and horsepower tend to be automatic.

c. Fit a logistic regression model with wt as the only feature. Using this model, explain whether heavier cars are more likely or less likely to have manual transmission. If weight increases by 1000 lbs, what is the change in odds of a car having manual transmission?

```
carlrmwt<-glm(am~wt,data=mtcars,family=binomial)
summary(carlrmwt)

Call:
glm(formula = am ~ wt, family = binomial, data = mtcars)

Deviance Residuals:
    Min     1Q     Median     3Q     Max
-2.11400     -0.53738     -0.08811     0.26055     2.19931</pre>
```

```
Estimate Std. Error z value Pr(>|z|)
(Intercept)
              12.040
                          4.510 2.670 0.00759 **
              -4.024
                          1.436 -2.801 0.00509 **
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
    Null deviance: 43.230 on 31 degrees of freedom
Residual deviance: 19.176 on 30 degrees of freedom
AIC: 23.176
Number of Fisher Scoring iterations: 6
#Heavier cars are less likkely to have manual transmissions because
#if weight increases by 1000 lbs, the change in odds of a car having
#manual transmission is around 98.21% and the odds in favor of manual transmission is of 0.0178813.
  d. Fit a logistic regression model with hp as the only feature. Using this model, explain whether cars with higher horsepower
     are more likely or less likely to have manual transmission. If horsepower increases by 100, what is the change in odds
     of a car having manual transmission?
carlrmhp<-glm(am~hp,data=mtcars,family=binomial)</pre>
summary(carlrmhp)
Call:
glm(formula = am ~ hp, family = binomial, data = mtcars)
Deviance Residuals:
              1Q
                   Median
                                         Max
                                      2.0379
-1.2955 -0.9968 -0.7818
                            1.1630
Coefficients:
             Estimate Std. Error z value Pr(>|z|)
                        0.915429
                                   0.848
                                             0.396
(Intercept) 0.776614
            -0.008117
                        0.006074 - 1.336
hp
                                             0.181
(Dispersion parameter for binomial family taken to be 1)
    Null deviance: 43.230 on 31 degrees of freedom
Residual deviance: 41.228 on 30 degrees of freedom
AIC: 45.228
Number of Fisher Scoring iterations: 4
# Higer hp cars have an odds in favor of manual transmission of 0.4441024
#which means they are less likely to have manual transmissions and for every
#100 increase in hp, the change in odds of a car having manual transmission decreases by 55.59%.
  e. If you had to choose between these two models, which one would you choose and why?
#Because the change of odds, I would chose the weight model since it is way higher
#than the change of odds of horsepower.
```

Coefficients: