

Lecture 06
2020 Spring Data-622
Logistic Regression
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Acknowledgements:
Generous support from IBM Power Systems Academic Initiative
IBM PSAI provides computing infrastructure for free

Script for Algorithms

Develop the Intuition

Understand the assumptions

Develop the mathematics

Run the algorithms

Learn to interpret the result/output

Predict using the model

Learn to determine the performance

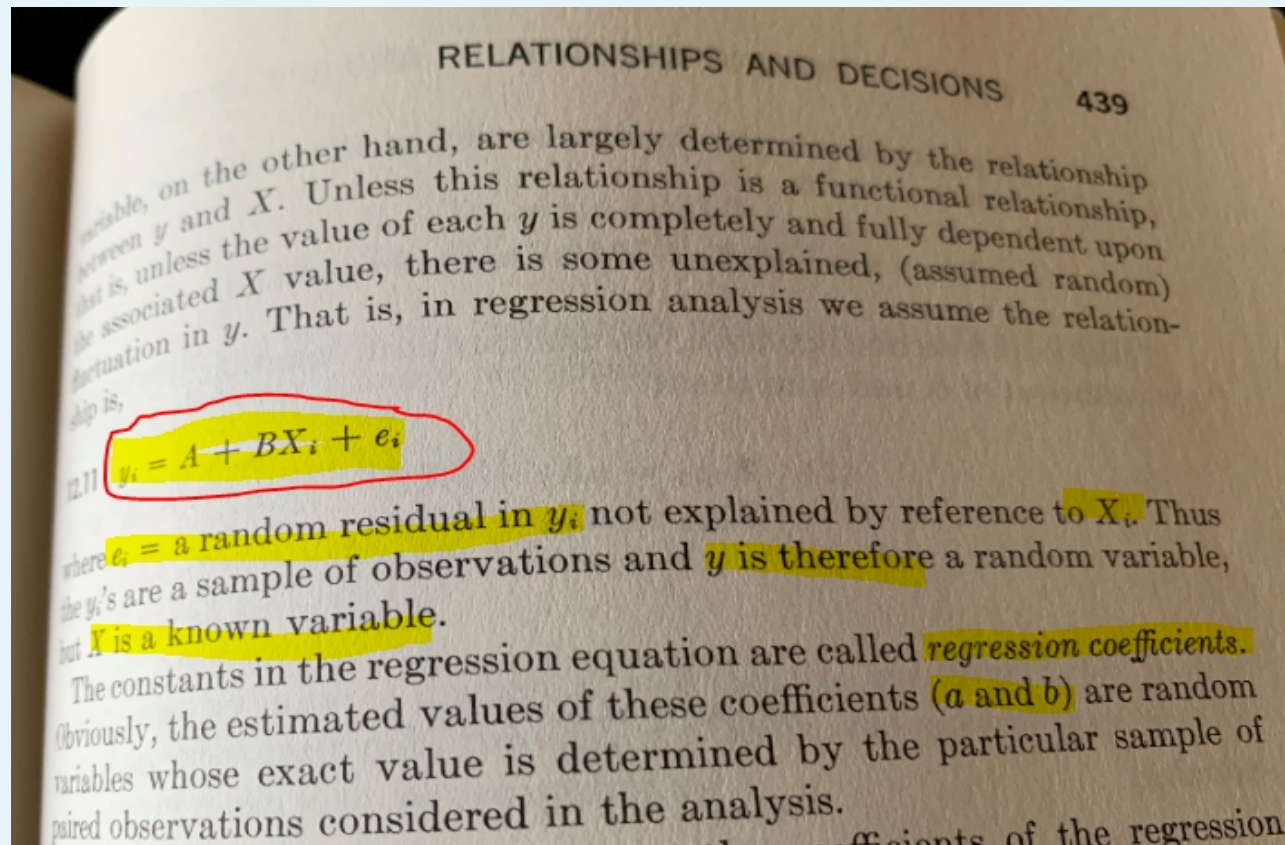
Distinguish training/testing error

Differentiate between overfitting/underfitting

Techniques to improve performance

OLS Model

Introduction to Quantitative Management – George J. Brabb



Need for a Logistic Function

Recall, $y = Xb + e$

In some scenarios, y , the DV that we wish to compute using observable X s is not continuous. It is dichotomous 1 or 0.

Consider, medical diagnostics, given some test results (X), physician has to determine if the patient is pregnant (YES or NO) or malignant or benign cancer.

Or Admissions committee, has to decide whether to admit or reject a student

Or a bank has to extend or decline credit to a customer.

There are so many instances where the dependent variable is dichotomous – can take one of two values. Binary Classification. When there are more than two, it is called Multi Class Classification.

Logistic Function

Let's understand how Logistic Regression works. For Linear Regression, where the output is a linear combination of input feature(s), we write the equation as:

$$\hat{Y} = \beta_0 + \beta_1 X + \epsilon$$

In Logistic Regression, we use the same equation but with some modifications made to \hat{Y} . Let's reiterate a fact about Logistic Regression: we calculate probabilities. And, probabilities always lie between 0 and 1. In other words, we can say:

1. The response value must be positive.
2. It should be lower than 1.

First, we'll meet the above two criteria. We know the exponential of any value is always a positive number. And, any number divided by number + 1 will always be lower than 1. Let's implement these two findings:

$$P(Y = 1|X) = \frac{e^{(\beta_0 + \beta_1 x)}}{e^{(\beta_0 + \beta_1 x)} + 1}$$

This is the logistic function.

As a function of x

<https://www.hackerearth.com/blog/wp-content/uploads/2017/01/equateimage.png>

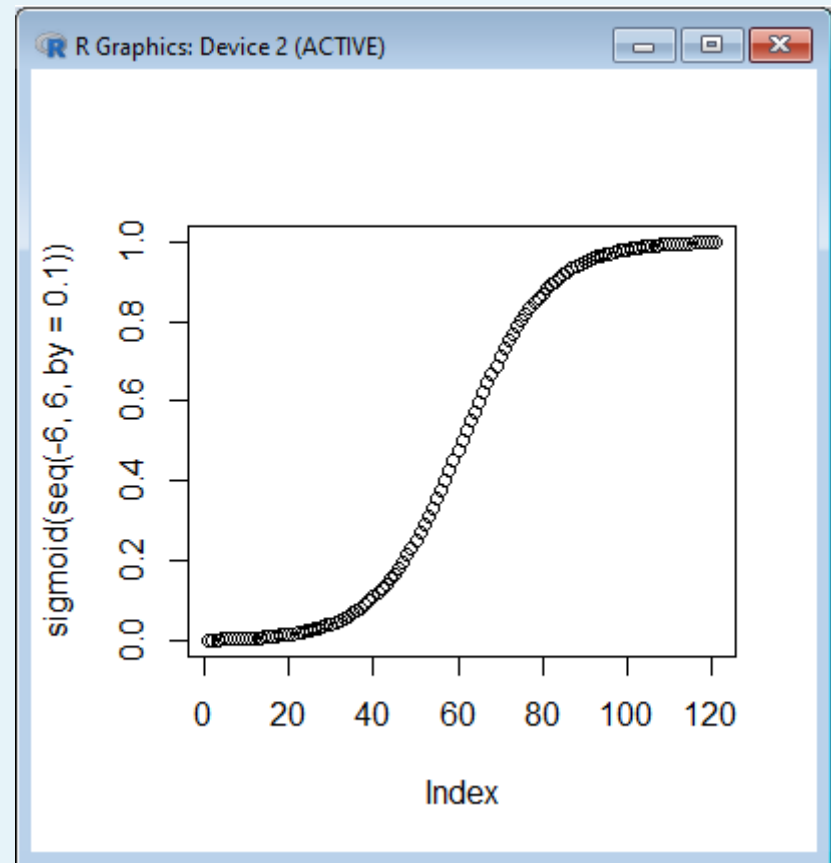
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$$\begin{aligned}\implies p(X) &= \frac{e^{(\beta_0 + \beta_1 x)}}{e^{(\beta_0 + \beta_1 x)} + 1} \\ \implies p(e^{(\beta_0 + \beta_1 x)} + 1) &= e^{(\beta_0 + \beta_1 x)} \\ \implies p \cdot e^{(\beta_0 + \beta_1 x)} + p &= e^{(\beta_0 + \beta_1 x)} \\ \implies p &= e^{(\beta_0 + \beta_1 x)} - p \cdot e^{(\beta_0 + \beta_1 x)} \\ \implies p &= e^{(\beta_0 + \beta_1 x)} (1 - p) \\ \implies \frac{p}{1 - p} &= e^{(\beta_0 + \beta_1 x)} \\ \implies \ln\left(\frac{p}{1 - p}\right) &= \beta_0 + \beta_1 x\end{aligned}$$

In search of a Transformation

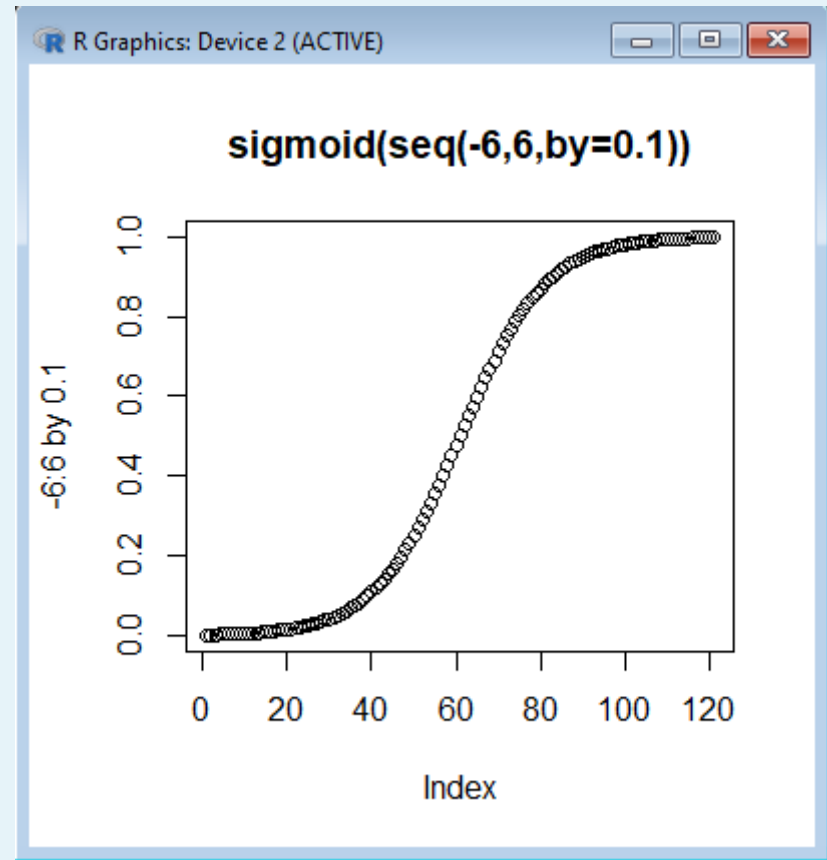
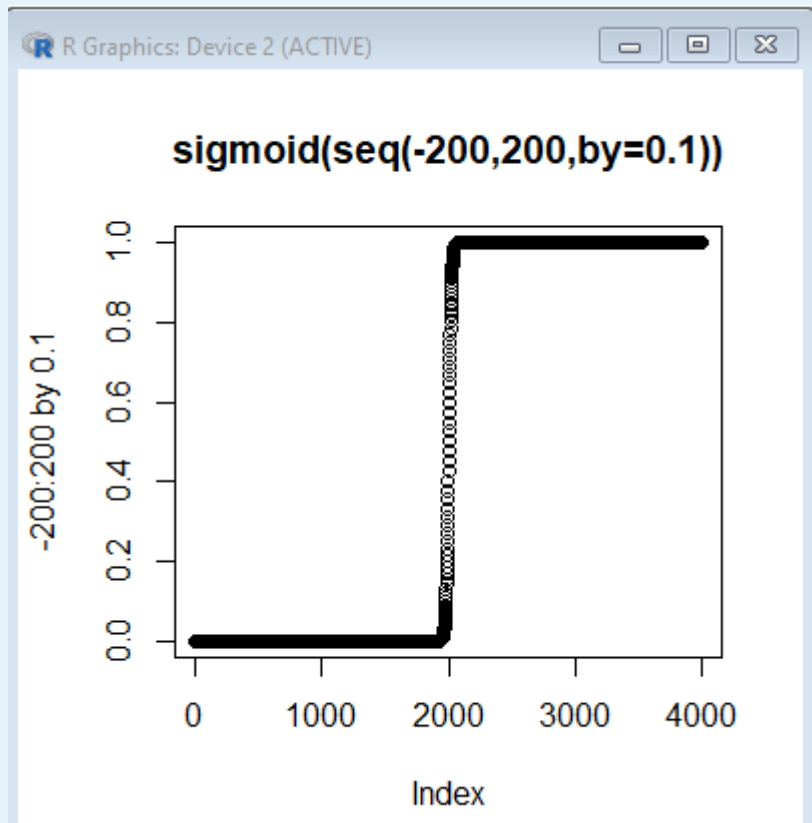
For binary classification, when the allowed values are either 0 or 1 we need a function that transforms X into one of two values.
One such function is the sigmoid function

```
vecp<-seq(-6,6,by=0.1)  
sigmoid <- function(x){ 1/(1+exp(-x))}  
plot(sigmoid(seq(-6,6,by=0.1)) )
```



In search of a Transformation

```
plot(sigmoid(seq(-200,200,by=0.1)),ylab="-200:200 by 0.1");title("sigmoid(seq(-200,200,by=0.1))")
```



```
plot(sigmoid(seq(-6,6,by=0.1)),ylab="-6:6 by 0.1");title("sigmoid(seq(-6,6,by=0.1))")
```


Logistic H: $\text{sigmoid} \leftarrow \text{function}(x) \{ 1/(1+\exp(-x)) \}$

Where $X = \sum \beta_i * X_i$

glm, the generalize linear model function in R computes the beta

```
logisticdata<-read.csv("https://pingax.com/wp-content/uploads/2013/12/data.csv")  
plot(logisticdata$score.1,logisticdata$score.2,col=as.factor(logisticdata$label),  
xlab="Score-1",ylab="Score-2") # view the data
```

Let us prepare data for training and testing

```
ccases1<-na.omit(logisticdata) #OR complete.cases , keep only complete observations
```

```
ccases2<-logisticdata[complete.cases(logisticdata),]
```

```
nrow(ccases1)
```

```
allidx<-1:nrow(ccases1)
```

```
set.seed(1313) # we are sampling for repeatability we set the seed
```

```
trainidx<-sample(allidx,round(0.7*nrow(ccases1)),replace=F)
```

```
traindata<-ccases1[trainidx,]
```

```
testdata<-ccases1[-trainidx,]
```

```
table(traindata$label)# check to make sure test and train are similarly distributed
```

```
table(testdata$label)
```

Running the model

we will run glm to generate
the model (betas) on
training data

```
glm.model<-  
glm(label~.,  
data=traindata,  
family='binomial')
```

```
summary(glm.model)
```

```
coef(glm.model)
```

The null deviance is 3 times
the deviance with the model.
So model is improving..

P-values are all small and we
can reject the NULL

The model is significant.

```
>  
> glm.model <-glm.model<-glm(label~.,data=traindata,family='binomial')  
>  
> summary(glm.model)  
  
Call:  
glm(formula = label ~ ., family = "binomial", data = traindata)  
  
Deviance Residuals:  
      Min       1Q   Median       3Q      Max   
-2.06587  -0.31121   0.02828   0.27766   1.82200  
  
Coefficients:  
              Estimate Std. Error z value Pr(>|z|)      
(Intercept) -22.70236     6.29030  -3.609  0.000307 ***  
score.1       0.18714     0.05128   3.649  0.000263 ***  
score.2       0.17713     0.05237   3.382  0.000718 ***  
---  
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
  
(Dispersion parameter for binomial family taken to be 1)  
  
    Null deviance: 92.360  on 69  degrees of freedom  
Residual deviance: 35.151  on 67  degrees of freedom  
AIC: 41.151  
  
Number of Fisher Scoring iterations: 7  
  
>  
> coef(glm.model)  
(Intercept)      score.1      score.2  
-22.7023627    0.1871382    0.1771322  
> |
```

Predicting using the model coeff

```
> computed.train.probabilities<-sigmoid(as.matrix(data.frame(intercept=1,traindata$score.1,traindata$score.2)) %*% betavec)
> computed.test.probabilities<-sigmoid(as.matrix(data.frame(intercept=1,testdata$score.1,testdata$score.2)) %*% betavec)
> computed.test.labels<-ifelse(computed.test.probabilities>0.5,1,0)
> computed.train.labels<-ifelse(computed.train.probabilities>0.5,1,0)
> table(computed.train.labels==traindata$label)
```

```
FALSE  TRUE
     9    61
```

```
> table(computed.test.labels==testdata$label)
```

```
FALSE  TRUE
     1    29
```

```
computed.train.probabilities<-sigmoid(as.matrix(
data.frame(intercept=1,traindata$score.1,traindata$score.2)) %*% betavec)
computed.test.probabilities<-sigmoid(as.matrix(
data.frame(intercept=1,testdata$score.1,testdata$score.2)) %*% betavec)
computed.test.labels<-ifelse(computed.test.probabilities>0.5,1,0)
computed.train.labels<-ifelse(computed.train.probabilities>0.5,1,0)
table(computed.train.labels==traindata$label)
table(computed.test.labels==testdata$label)
```

```
plot(1:70,computed.train.probabilities,col=ifelse(computed.train.probabilities>0.50,
'black','yellow'),xlab="obs",ylab="probabilities")
```

```
>
```

```
plot(1:30,computed.test.probabilities,col=ifelse(computed.test.probabilities>0.50,'
black','yellow'),xlab="obs",ylab="probabilities")
```

Predicting using predict

```
> estimated.train.proBABILITIES<-predict(glm.model,newdata=traindata[,c("score.1","score.2")],type='response')
> head(estimated.train.proBABILITIES)
      89      39      70      48      62      61
0.9998919382 0.2071082482 0.6438325823 0.9999825600 0.0004219775 0.9986386035
> estimated.train.labels<-ifelse(estimated.train.proBABILITIES>0.5,1,0)
> estimated.test.proBABILITIES<-predict(glm.model,newdata=testdata[,c("score.1","score.2")],type='response')
> estimated.test.labels<-ifelse(estimated.test.proBABILITIES>0.5,1,0)
> table(estimated.test.labels==computed.test.labels)

TRUE
 30
> table(estimated.train.labels==computed.train.labels)

TRUE
 70
> table(estimated.train.labels==traindata.labels)
Error in table(estimated.train.labels == traindata.labels) :
  object 'traindata.labels' not found
> table(estimated.train.labels==traindata.label)
Error in table(estimated.train.labels == traindata.label) :
  object 'traindata.label' not found
> table(estimated.train.labels==traindata$label)

FALSE  TRUE
   9    61
> table(estimated.test.labels==testdata$label)

FALSE  TRUE
   1    29
,
```

Predicting using predict

```
estimated.train.proBABILITIES<-  
predict(glm.model,newdata=traindata[,c("score.1","score.2")],type='response')  
head(estimated.train.proBABILITIES)  
estimated.train.labels<-ifelse(estimated.train.proBABILITIES>0.5,1,0)  
estimated.test.proBABILITIES<-  
predict(glm.model,newdata=testdata[,c("score.1","score.2")],type='response')  
estimated.test.labels<-ifelse(estimated.test.proBABILITIES>0.5,1,0)  
table(estimated.test.labels==computed.test.labels)  
table(estimated.train.labels==computed.train.labels)  
table(estimated.train.labels==traindata$label)  
table(estimated.test.labels==testdata$label)
```

Performance

```
require(ROCR)
glm_prediction<-prediction(estimated.test.probabilities,testdata$label)
glm_perf<-performance(glm_prediction,measure="tpr",x.measure="fpr")
glm_slot_fp<-slot(glm_prediction,"fp")
glm_slot_tp<-slot(glm_prediction,"tp")
glm_slot_tn<-slot(glm_prediction,"n.neg")
glm_slot_fn<-slot(glm_prediction,"n.pos")
glm_auc<-performance(glm_prediction,"auc")@y.values[[1]]

plot(unlist(glm_slot_fp)/unlist(glm_slot_tn),
unlist(glm_slot_tp)/unlist(glm_slot_fn),main="ROCR
Curve",xlab="FPR",ylab='TPR')
```

References

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<http://pingax.com/wp-content/uploads/2013/12/data.csv>

<https://towardsdatascience.com/understanding-logistic-regression-9b02c2aec102>

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