Lecture Nine – Confusion Matrix Checking the accuracy of the Logistic Regression Model Professor Esfandiari

A confusion matrix is a table that is often used to **describe the performance of a classification model** (or "classifier") on a set of test data for which the true values are known. The confusion matrix itself is relatively simple to understand, but the related terminology can be confusing.

We will start with an **example confusion matrix for a binary classifier**. In the following you are given the data for 200 patients that have diabetic. Based on a logistic regression model, we have made predictions. The following table presents the actual values and the predicted values.

Actual	Predicted		Colum
	No	Yes	totals
No	60	12	72
	TRUE NEGATIVE	FALSE POSITIVE	
		TYPE I ERROR	
Yes	8	120	128
	FALSE NEGATIVE	TRUE POSITIVE	
	TYPE II ERROR		
Row totals	68	132	200

Based on the prediction model,

Of the two hundred patients, 132 were classified as having the disease and 68 were classified as not having the disease.

Based on the actual results,

Of the two hundred patients, 128 actually had diabetic II and 72 did not.

We will now define, the different cells of the confusion matrix TRUE POSITIVE

True positive are the cases in which the patient actually has the disease and the model also predicts that the patient has the disease.

FALSE POSITIVE

False positive are the cases in which the patient actually has the disease, but, the model predicts that the patient does has the disease. **This is known as TYPE I ERROR.**

FALSE NEGATIVE

False negative are the cases in which the patient actually does not have the disease, but, the model predicts that the patient has the disease. THIS IS KNOWN AS TYPE II ERROR.

TRUE POSITIVE

False positive are the cases in which the patient actually has the disease and the model predicts that the patient has the disease.

Other Concepts Underlying the Model

ACCURACY

Overall, how well did the prediction based on the logistic model work?

$$Accuracy = \frac{\textit{True positive} + \textit{Ture negative}}{\textit{Total sample}} = \frac{120 + 60}{200} = 180/200 = 0.90$$

This means that in 90% of the cases the prediction based on the logistic model makes the correct classification.

MISCLASSIFICATION RATE - ERROR RATE

Overall, how often is the classification done by the prediction model wrong?

$$Misclassification \ Rate = \frac{false \ positive + false \ negative}{Total \ sample} = \frac{12 + 8}{200} = 20/200 = 0.10$$

Misclassification Rate = 1- accuracy rate = 1 - 0.90 = 0.10

This means that in 10% of the cases the prediction based on the logistic model makes the incorrect classification.

TRUE POSITIVE RATE - SENSITIVITY RATE

True positive rate: When the patient "actually has diabetic II", and the model also predicts "ves".

$$=$$
TP/actual yes $= 120/128 = 0.9375$

FALSE POSITIVE RATE – SENSITIVITY RATE

False positive rate: When the patient "does not have diabetic II", and model predicts "yes".

• **Specificity:** When it's actually no, how often does it predict no?

Specificity =
$$\frac{True\ No}{Actual\ No}$$
 = $60/72 = 0.833$

equivalent to 1 minus False Positive Rate = 1 - 0.167 = 0.833

• **Precision:** When it predicts yes, how often is it correct?

$$Precision = \frac{True\ positive}{Predicted\ yes} = 120/132 = 0.9090909$$

• Prevalence: How often does the yes condition actually occur in our sample?

actual yes/total =
$$128/200 = 0.64$$

Calculation of the Confusion Matrix using R

Using the diabetic data, we will create a logistic regression model for the prediction of having diabetic type II from age, smoking, and hypertension.

> m1<-glm(Diabetes.new~Age+SmokingStatus_NISTCode+HypertensionDX, family="binomial")
> summary(m1)

Coefficients:

	Estimate	Std. Error z value	Pr(> z)
(Intercept)	-3.380674	0.174617 -19.361	< 2e-16 ***
Age	0.025801	0.002736 9.430	< 2e-16 ***
SmokingStatus_NISTCodeFORMER	-0.355067	0.085528 -4.151	3.3e-05 ***
SmokingStatus_NISTCodeTRUE	-0.334447	0.130085 -2.571	0.0101 *
HypertensionDXyes	1.129735	0.090237 12.520	< 2e-16 ***

Signif. codes: 0 "*** 0.001 "** 0.01 "* 0.05". 0.1 " 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 4404.9 on 4402 degrees of freedom Residual deviance: 3953.5 on 4398 degrees of freedom

(5545 observations deleted due to missingness)

AIC: 3963.5

Number of Fisher Scoring iterations: 5

We will now use R to create the confusion matrix

R Codes

> library(nnet)

> m1<-

multinom(Diabetes.new~Age+HypertensionDX+SmokingStatus NISTCode,diabetic)

weights: 6 (5 variable) initial value 3051.927036 iter 10 value 1976.747412 final value 1976.731777 converged

R codes for the creation of the confusion matrix continued

- > p<-predict(m1,diabetic)
- > tab<-table(p,diabetic\$Diabetes.new)
- > tab

Predicted values	Actual Value		Row totals
	Does not have	Has diabetic	
	diabetic		
Does not have	3517	876	4393
diabetic	True negative	False Positive	
Has Diabetic	6	4	10
	False Negative	True positive	
Column Total	3523	880	4403

- **Accuracy:** Overall, how often is the classifier (predicted model) correct? (TP+TN)/total = (4+3517)/4403 = **0.799682**
- Misclassification Rate or error rate: Overall, how often is the classifier wrong?

(FP+FN)/total = (876+6)/4403 = 0.200318

equivalent to 1 minus Accuracy

• True Positive Rate: When it's actually yes, how often does it predict yes?

TP/actual yes = 4/880 = 0.004545455

also known as "Sensitivity"

• False Positive Rate: When it's actually no, how often does it predict yes?

FP/actual no = 876/3523 = 0.2486517

• **Specificity:** When it's actually no, how often does it predict no?

TN/actual no = 3517/3523 = 0.9982969

equivalent to 1 minus False Positive Rate

• Precision: When it predicts yes, how often is it correct?

TP/predicted yes = 4/10 = 0.4

• Prevalence: How often does the yes condition actually occur in our sample?

actual yes/total = 880/4403 = 0.1998637

CONCLUSION: Based on the confusion matrix given above, the accuracy rate is 80%. So, the predictive power of the model is good. You want most of the points to be on the diagonal; that is true positive and true negative. The model seems to be doing well with respect to true negative but not so well with respect to true positive.