For this pre live assignment we are going to be focusing on two main things, interpretation and verifying that logistic regression is a more general technique that has the previous methods as special cases. The data set we will explore is a coronary artery disease data set and the goal is to determine what risk factors would contribute to the likelihood of an individual having the disease or not. The data is below in the sas code.

Ca- coronary artery disease (1=Yes)

Sex- (1=Male, 0=Female)

ECG – categoral level for an echocardiogram result (0=low, 1=medium, 2=high)

Age – continuous, in years

1.      Use the proc freq and proc means procedures to explore if any relationships that exist between the response, disease status, and the potential risk factors.

2.     Fit a simple logistic regression using all 3 predictors and verify that the overall model is significant. We will discuss in live session that residual model checking for logistic regression is not very informative due to the categorical nature. One way to do this is with the Hosmer-Lemeshow lack of fit test. Obtain the statistic and p-value from the output. Does the result tell us that our model fits reasonable or not?

3.      Provide an interpretation of the regression coefficient for Age. What does the value represent in terms of the odds of having the disease?

4.      What is the odds ratio of a person with an Age=55  having coronary artery disease with respect to someone with who is 45years of age given they are both males and have the same ECG result.  (See 20.2.2 in text) Pay attention to the output you may need to do some “flipping” (reciprocal 1/OR).

5.      Consider a new patient comes into a doctors office and has measurement ECG=2, Age=50,Sex=Male.  What is the probability that this person has the disease? (Here we are trying to predict if the new person has the disease or not.)

6. Run a 2x2 analysis with CA and Sex and then run a logistic regression model just using Sex as an explanatory variable. Determine the odds ratio (disease:no disease) males vs females for the two analysis runs and verify that they are very close. This shows that logistic regression has the simple 2x2 table analysis as a special case when you have one categorical explanatory variable with just two levels.

Optional: It is not in the videos, but feel free to do a little google on the topic of ROC curves which we will discuss over the next couple of weeks. ROC stands for “Receiver Operating Characteristic”.

data coronary;

input sex ecg age ca @@ ;

cards;

0 0 28 0 1 0 42 1 0 1 46 0 1 1 45 0

0 0 34 0 1 0 44 1 0 1 48 1 1 1 45 1

0 0 38 0 1 0 45 0 0 1 49 0 1 1 45 1

0 0 41 1 1 0 46 0 0 1 49 0 1 1 46 1

0 0 44 0 1 0 48 0 0 1 52 0 1 1 48 1

0 0 45 1 1 0 50 0 0 1 53 1 1 1 57 1

0 0 46 0 1 0 52 1 0 1 54 1 1 1 57 1

0 0 47 0 1 0 52 1 0 1 55 0 1 1 59 1

0 0 50 0 1 0 54 0 0 1 57 1 1 1 60 1

0 0 51 0 1 0 55 0 0 2 46 1 1 1 63 1

0 0 51 0 1 0 59 1 0 2 48 0 1 2 35 0

0 0 53 0 1 0 59 1 0 2 57 1 1 2 37 1

0 0 55 1 1 1 32 0 0 2 60 1 1 2 43 1

0 0 59 0 1 1 37 0 1 0 30 0 1 2 47 1

0 0 60 1 1 1 38 1 1 0 34 0 1 2 48 1

0 1 32 1 1 1 38 1 1 0 36 1 1 2 49 0

0 1 33 0 1 1 42 1 1 0 38 1 1 2 58 1

0 1 35 0 1 1 43 0 1 0 39 0 1 2 59 1

0 1 39 0 1 1 43 1 1 0 42 0 1 2 60 1

0 1 40 0 1 1 44 1

;

run;quit;

\*Explore;

proc freq data=coronary;

tables sex\*ca ecg\*ca sex\*ecg\*ca / chisq relrisk;

run;quit;

proc means data=coronary;

class ca sex ecg;

types ca ca\*sex ca\*ecg;

var age;

run;

\*Simple proc logistic call to play around with;

proc logistic data=coronary ;

class sex ecg / param=ref;

model ca(event='1')= sex/ scale=none aggregate influence lackfit;

run;

\*Additional code for discussion in live session with additional options like effect plots and ROC curves.;

\*plots(only)=roc(id=obs);

proc logistic data=coronary ;

class sex ecg / param=ref;

model ca(event='1')= sex age ecg/ scale=none aggregate influence lackfit;

effectplot slicefit(sliceby=Sex plotby=ecg) / noobs;

run;

proc logistic data=coronary ;

class sex ecg / param=ref;

model ca(event='1')= sex ecg age sex\*ecg sex\*age ecg\*age age\*age/ selection=FORWARD start=3 scale=none details influence lackfit;

effectplot slicefit(sliceby=Sex plotby=ecg) / noobs;

run;

proc logistic data=coronary ;

class sex ecg;

model ca(event='1')= sex ecg age / scale=none ctable pprob=.5 aggregate lackfit;

ROC 'MainEffects' sex ecg age;

ROC 'Just Sex/AGE' sex age ;

roccontrast reference('Just Sex/AGE') / estimate e;

run;