

Appendix R code and output

Appendix 1. Anova table of model 2

```

              Response: target
              LR Chisq Df Pr(>Chisq)
sex           10.938  1 0.0009422 ***
cp            13.282  2 0.0013060 **
trestbps      4.592  1 0.0321253 *
chol          0.971  1 0.3243812
fbs           4.987  1 0.0255352 *
thalach       0.546  1 0.4598773
exang         4.806  1 0.0283614 *
slope        14.386  1 0.0001489 ***
ca           37.765  1 7.979e-10 ***
thal         10.674  1 0.0010867 **
exang:slope   6.779  1 0.0092261 **
sex:fbs       9.403  1 0.0021666 **
fbs:slope     10.029  1 0.0015413 **
exang:ca      10.005  1 0.0015608 **
trestbps:exang 6.184  1 0.0128931 *
thalach:ca    5.871  1 0.0153966 *
fbs:ca        2.729  1 0.0985694 .
thalach:exang 2.180  1 0.1398572
sex:thal      4.079  1 0.0434178 *
cp:chol       5.510  2 0.0635985 .

```

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

Appendix 2. odds ratio of final model

```

coefficient 95%CI(2.5%) 95%CI(97.5%) odds ratio
(Intercept)  "-1.96"    "-6.23"    " 2.22"    " 0.14"
sexM         " 0.14"    "-2.31"    " 2.38"    " 1.15"
cpasymptomatic " 1.43"    " 0.49"    " 2.43"    " 4.19"
cpnon_anginal "-0.14"    "-1.22"    " 0.91"    " 0.87"
trestbps     " 0.01"    "-0.01"    " 0.04"    " 1.01"
fbsTrue      " 6.10"    " 1.41"    "11.12"    "447.51"
exangYes     "-6.48"    "-13.83"   " 0.14"    " 0.00"
slopeup      "-0.85"    "-1.79"    " 0.07"    " 0.43"
ca           " 1.25"    " 0.69"    " 1.89"    " 3.49"
thalnormal   "-3.22"    "-5.83"    "-0.92"    " 0.04"
exangYes:slopeup "-2.34"   "-4.51"   "-0.41"    " 0.10"
sexM:fbsTrue  "-7.79"   "-13.28"  "-2.86"    " 0.00"
fbsTrue:slopeup "-7.29"   "-14.57"  "-1.56"    " 0.00"
exangYes:ca    " 2.43"    " 0.59"    " 4.73"    "11.41"
trestbps:exangYes " 0.06"    " 0.01"    " 0.12"    " 1.06"
fbsTrue:ca     " 1.80"    "-0.44"    " 4.58"    " 6.03"
sexM:thalnormal " 2.02"    "-0.44"    " 4.78"    " 7.52"

```

Appendix: Ready to run R code

```

setwd("~/Desktop")
#step1. take a look at the dataset
#read CSV file
heart1=read.table("project.1.data.1.csv",sep=";",head=T)
#preview of data
heart1[1:10,]
#check variable type
str(heart1)
#step2. Data cleaning
summary(heart1)
#change categorical variable to factor
heart1$sex[heart1$sex==1]="M"
heart1$sex[heart1$sex==0]="F"
heart1$sex=as.factor(heart1$sex)
heart1$fbs[heart1$fbs==1]="True"
heart1$fbs[heart1$fbs==0]="False"
heart1$fbs=as.factor(heart1$fbs)
heart1$cp[heart1$cp==1]="typical"
heart1$cp[heart1$cp==2]="atypical"
heart1$cp[heart1$cp==3]="non_anginal"
heart1$cp[heart1$cp==4]="ymptomatic"
heart1$cp=as.factor(heart1$cp)
heart1$exang[heart1$exang==1]="Yes"
heart1$exang[heart1$exang==0]="No"
heart1$exang=as.factor(heart1$exang)
heart1$restecg[heart1$restecg==0]="normal"
heart1$restecg[heart1$restecg==1]="ST_T_abnormal"
heart1$restecg[heart1$restecg==2]="probable or definite left ventricular hypertrophy"
heart1$restecg=as.factor(heart1$restecg)
heart1$thal[heart1$thal==3]="normal"
heart1$thal[heart1$thal==6]="fixed_defect"
heart1$thal[heart1$thal==7]="reversable_defect"
heart1$thal=as.factor(heart1$thal)
heart1$slope[heart1$slope==1]="up"
heart1$slope[heart1$slope==2]="flat"
heart1$slope[heart1$slope==3]="down"
heart1$slope=as.factor(heart1$slope)
heart1$target=as.factor(heart1$target)
str(heart1)
#step3. Visuals

```

Figure 2b.

```

#graph numeric variables vs. "target"
boxplot(age~target,data=heart1,xlab="Heart Disease",ylab="Age")
boxplot(trestbps~target,data=heart1,xlab="Heart Disease",ylab="resting blood pressure (in mm Hg) ")
boxplot(chol~target,data=heart1,xlab="Heart Disease",ylab="serum cholestoral in mg/dl")
boxplot(thalach~target,data=heart1,xlab="Heart Disease",ylab="maximum heart rate achieved")
boxplot(oldpeak~target,data=heart1,xlab="Heart Disease",ylab="ST depression induced by exercise relative to rest")
boxplot(ca~target,data=heart1,xlab="Heart Disease",ylab="number of major vessels(0-3)")
#graph categorical variable vs. "target"

```

Figure 1.

```

plot(heart1$target, xlab="heart disease", ylab="count")

```

```

spineplot(target~sex,data=heart1,ylab="Heart Disease",xlab="Sex")
spineplot(target~cp,data=heart1,ylab="Heart Disease",xlab="chest pain type")
spineplot(target~fbs,data=heart1,ylab="Heart Disease",xlab="fasting blood sugar > 120 mg/dl")
spineplot(target~restecg,data=heart1,ylab="Heart Disease",xlab="resting electrocardiographic")
spineplot(target~exang,data=heart1,ylab="Heart Disease",xlab="exercise induced angina")
spineplot(target~slope,data=heart1,ylab="Heart Disease",xlab="slope of the peak exercise ST segment")
spineplot(target~thal,data=heart1,ylab="Heart Disease",xlab="thal")

```

#Combine data with few observations

```
heart1=read.table("project.1.data.1.csv",sep="," ,head=T)
```

```

plot(heart1$sex)
heart1$sex[heart1$sex==1]="M"
heart1$sex[heart1$sex==0]="F"
heart1$sex=as.factor(heart1$sex)

```

```

plot(heart1$fbs)
heart1$fbs[heart1$fbs==1]="True"
heart1$fbs[heart1$fbs==0]="False"
heart1$fbs=as.factor(heart1$fbs)

```

#combine typical and atypical angina as one

```

plot(heart1$cp)
heart1$cp[heart1$cp==1]="angina"
heart1$cp[heart1$cp==2]="angina"
heart1$cp[heart1$cp==3]="non_anginal"
heart1$cp[heart1$cp==4]="asymptomatic"
heart1$cp=as.factor(heart1$cp)

```

```

plot(heart1$exang)
heart1$exang[heart1$exang==1]="Yes"
heart1$exang[heart1$exang==0]="No"
heart1$exang=as.factor(heart1$exang)

```

#combine ST_T_abnormal and "probable or definite left ventricular hypertrophy" as abnormal

```

plot(heart1$restecg)
heart1$restecg[heart1$restecg==0]="normal"
heart1$restecg[heart1$restecg==1]="abnormal"
heart1$restecg[heart1$restecg==2]="abnormal"
heart1$restecg=as.factor(heart1$restecg)

```

#combine fixed and reversable defect as defect

```

plot(heart1$thal)
heart1$thal[heart1$thal==3]="normal"
heart1$thal[heart1$thal==6]="defect"
heart1$thal[heart1$thal==7]="defect"
heart1$thal=as.factor(heart1$thal)

```

#combine flat and down together

```

plot(heart1$slope)
heart1$slope[heart1$slope==1]="up"
heart1$slope[heart1$slope==2]="flat | down"
heart1$slope[heart1$slope==3]="flat | down"
heart1$slope=as.factor(heart1$slope)
heart1$target=as.factor(heart1$target)

```

Figure 2a.

```
#graph categorical variable vs. "target"
plot(heart1$target, xlab="heart disease", ylab="count")
spineplot(target~sex,data=heart1,ylab="Heart Disease",xlab="Sex")
spineplot(target~cp,data=heart1,ylab="Heart Disease",xlab="chest pain type")
spineplot(target~fbs,data=heart1,ylab="Heart Disease",xlab="fasting blood sugar > 120 mg/dl")
spineplot(target~restecg,data=heart1,ylab="Heart Disease",xlab="resting electrocardiographic")
spineplot(target~exang,data=heart1,ylab="Heart Disease",xlab="exercise induced angina")
spineplot(target~slope,data=heart1,ylab="Heart Disease",xlab="slope of the peak exercise ST segment")
spineplot(target~thal,data=heart1,ylab="Heart Disease",xlab="Thalassemia")
```

#step 4.model selection

Model. 1

```
#model selection without interaction
fit.1=glm(target~1,data=heart1,family="binomial")
fit.2=glm(target~.,data=heart1,family="binomial")
select.1=step(fit.2, scope=list(lower=fit.1,upper=fit.2),direction="backward")
summary(select.1)
coef(select.1)
```

Model. 2

```
#model selection with interaction
fit.3=glm(target ~ (sex + cp + trestbps + chol + fbs + thalach + exang + slope + ca + thal)^2, family = "binomial",
data = heart1)
select.2=step(select.1, scope=list(lower=select.1,upper=fit.3),direction="forward")
summary(select.2)
coef(select.2)
```

Final model

```
library(car)# Anova
Anova(select.2) #based on this, exang not significant remove exang in fit.4
fit.final=glm(target ~ sex + cp + trestbps + fbs + exang + slope + ca + thal + exang:slope + sex:fbs + fbs:slope +
exang:ca + trestbps:exang + fbs:ca + sex:thal,
data=heart1, family=binomial)
summary(fit.final)
coef(fit.final)
confint(fit.final) #likelihood ratio CI
Anova(fit.final)
```

Final model. Odds ratio

```
oddsratio=cbind(coef(fit.final),confint(fit.final),exp(coef(fit.final)))
colnames(oddsratio)=c("coefficient", "95%CI(2.5%)", "95%CI(97.5.5%)", "odds ratio")
oddsratio=format(round(oddsratio,2), nsmall=2)
oddsratio
```

ROC AUC

#step 5. model evaluation: sensitivity/specificity/ROC/AUC

```
pi0=seq(1,0, length.out = 100)
nsample=nrow(heart1)
roc=NULL
```

```

cv.set.list=split(sample(nsample),rep(1:10,length=nsample)) #10-fold cross-validation method
for(k in 1:length(pi0))
{n_11=n_10=n_01=n_00=0
nsample=nrow(heart1)
for(i in 1:length(cv.set.list))
{
  training=heart1[-cv.set.list[[i]],]
  test=heart1[cv.set.list[[i]],]
  fit.training=glm(target ~ sex + cp + trestbps + fbs + exang + slope + ca + thal + exang:slope + sex:fbs +
fbs:slope +
                exang:ca + trestbps:exang + fbs:ca + sex:thal, data=training, family="binomial")
  for(j in 1:length(cv.set.list[[i]]))
  {if(predict(fit.training, test[j,], type="response")>=pi0[k])
  {Y.pred.1=1}else{Y.pred.1=0}
  if((test$target[j]==1)&(Y.pred.1==1))
  {n_11=n_11+1}
  if((test$target[j]==1)&(Y.pred.1==0))
  {n_10=n_10+1}
  if((test$target[j]==0)&(Y.pred.1==1))
  {n_01=n_01+1}
  if((test$target[j]==0)&(Y.pred.1==0))
  {n_00=n_00+1}
  }
}
sen=n_11/(n_10+n_11)
spe=n_00/(n_00+n_01)
roc=rbind(roc, c(1-spe, sen))
}
plot(roc, type="s",xlim=c(0,1), ylim=c(0,1), col="red", main="ROC curve", xlab="1-Specificity",
ylab="Sensitivity")
auc=sum(roc[-100, 2]*(roc[-1,1]-roc[-100,1]))
auc #AUC area under ROC curve
pred.accuracy=roc[,2]*sum(heart1$target==1)/nsample+(1-roc[,1])*sum(heart1$target==0)/nsample
z.1=which.max(pred.accuracy)
pi0[z.1] #best cut-off points
pred.accuracy[z.1] #best prediction accuracy

```

Appendix. output

```

> heart1$restecg[heart1$restecg==2]="probable or definite left ventricular hypertrophy"
> heart1$restecg=as.factor(heart1$restecg)
> heart1$thal[heart1$thal==3]="normal"
> heart1$thal[heart1$thal==6]="fixed_defect"
> heart1$thal[heart1$thal==7]="reversable_defect"
> heart1$thal=as.factor(heart1$thal)
> heart1$slope[heart1$slope==1]="up"
> heart1$slope[heart1$slope==2]="flat"
> heart1$slope[heart1$slope==3]="down"
> heart1$slope=as.factor(heart1$slope)
> heart1$target=as.factor(heart1$target)
> str(heart1)
'data.frame':   297 obs. of  14 variables:
 $ age   : int  63 67 67 37 41 56 62 57 63 53 ...

```

```

$ sex : Factor w/ 2 levels "F","M": 2 2 2 2 1 2 1 1 2 2 ...
$ cp : Factor w/ 4 levels "atypical","non_anginal",...: 3 4 4 2 1 1 4 4 4 4 ...
$ trestbps: int 145 160 120 130 130 120 140 120 130 140 ...
$ chol : int 233 286 229 250 204 236 268 354 254 203 ...
$ fbs : Factor w/ 2 levels "False","True": 2 1 1 1 1 1 1 1 2 ...
$ restecg : Factor w/ 3 levels "normal","probable or definite left ventricular hypertrophy",...: 2 2 2 1 2 1 2 1 2
2 ...
$ thalach : int 150 108 129 187 172 178 160 163 147 155 ...
$ exang : Factor w/ 2 levels "No","Yes": 1 2 2 1 1 1 1 2 1 2 ...
$ oldpeak : num 2.3 1.5 2.6 3.5 1.4 0.8 3.6 0.6 1.4 3.1 ...
$ slope : Factor w/ 3 levels "down","flat",...: 1 2 2 1 3 3 1 3 2 1 ...
$ ca : int 0 3 2 0 0 0 2 0 1 0 ...
$ thal : Factor w/ 3 levels "fixed_defect",...: 1 2 3 2 2 2 2 3 3 ...
$ target : Factor w/ 2 levels "0","1": 1 2 2 1 1 1 2 1 2 2 ...
> #step3. Visuals

```

Figure 2b

```

> #graph numeric variables vs. "target"
> boxplot(age~target,data=heart1,xlab="Heart Disease",ylab="Age")
> boxplot(trestbps~target,data=heart1,xlab="Heart Disease",ylab="resting blood pressure (in mm Hg) ")
> boxplot(chol~target,data=heart1,xlab="Heart Disease",ylab="serum cholestoral in mg/dl")
> boxplot(thalach~target,data=heart1,xlab="Heart Disease",ylab="maximum heart rate achieved")
> boxplot(oldpeak~target,data=heart1,xlab="Heart Disease",ylab="ST depression induced by exercise relative
to rest")
> boxplot(ca~target,data=heart1,xlab="Heart Disease",ylab="number of major vessels(0-3)")
> #graph categorical variable vs. "target"
> plot(heart1$target, xlab="heart disease", ylab="count")

```

Figure 1

```

> spineplot(target~sex,data=heart1,ylab="Heart Disease",xlab="Sex")
> spineplot(target~cp,data=heart1,ylab="Heart Disease",xlab="chest pain type")
> spineplot(target~fbs,data=heart1,ylab="Heart Disease",xlab="fasting blood sugar > 120 mg/dl")
> spineplot(target~restecg,data=heart1,ylab="Heart Disease",xlab="resting electrocardiographic")
> spineplot(target~exang,data=heart1,ylab="Heart Disease",xlab="exercise induced angina")
> spineplot(target~slope,data=heart1,ylab="Heart Disease",xlab="slope of the peak exercise ST segment")
> spineplot(target~thal,data=heart1,ylab="Heart Disease",xlab="thal")
>

```

```

> #Combine data with few observations
> heart1=read.table("project.1.data.1.csv",sep=" ",head=T)
>
> plot(heart1$sex)
> heart1$sex[heart1$sex==1]="M"
> heart1$sex[heart1$sex==0]="F"
> heart1$sex=as.factor(heart1$sex)
>
> plot(heart1$fbs)
> heart1$fbs[heart1$fbs==1]="True"
> heart1$fbs[heart1$fbs==0]="False"
> heart1$fbs=as.factor(heart1$fbs)
>
> #combine typical and atypical angina as one
> plot(heart1$cp)

```

```

> heart1$cp[heart1$cp==1]="angina"
> heart1$cp[heart1$cp==2]="angina"
> heart1$cp[heart1$cp==3]="non_anginal"
> heart1$cp[heart1$cp==4]="asymptomatic"
> heart1$cp=as.factor(heart1$cp)
>
> plot(heart1$exang)
> heart1$exang[heart1$exang==1]="Yes"
> heart1$exang[heart1$exang==0]="No"
> heart1$exang=as.factor(heart1$exang)
>
> #combine ST_T_abnormal and "probable or definite left ventricular hypertrophy" as abnormal
> plot(heart1$restecg)
> heart1$restecg[heart1$restecg==0]="normal"
> heart1$restecg[heart1$restecg==1]="abnormal"
> heart1$restecg[heart1$restecg==2]="abnormal"
> heart1$restecg=as.factor(heart1$restecg)
>
> #combine fixed and reversible defect as defect
> plot(heart1$thal)
> heart1$thal[heart1$thal==3]="normal"
> heart1$thal[heart1$thal==6]="defect"
> heart1$thal[heart1$thal==7]="defect"
> heart1$thal=as.factor(heart1$thal)
>
> #combine flat and down together
> plot(heart1$slope)
> heart1$slope[heart1$slope==1]="up"
> heart1$slope[heart1$slope==2]="flat | down"
> heart1$slope[heart1$slope==3]="flat | down"
> heart1$slope=as.factor(heart1$slope)
> heart1$target=as.factor(heart1$target)

```

Figure 2a

```

> #graph categorical variable vs. "target"
> plot(heart1$target, xlab="heart disease", ylab="count")
> spineplot(target~sex,data=heart1,ylab="Heart Disease",xlab="Sex")
> spineplot(target~cp,data=heart1,ylab="Heart Disease",xlab="chest pain type")
> spineplot(target~fbs,data=heart1,ylab="Heart Disease",xlab="fasting blood sugar > 120 mg/dl")
> spineplot(target~restecg,data=heart1,ylab="Heart Disease",xlab="resting electrocardiographic")
> spineplot(target~exang,data=heart1,ylab="Heart Disease",xlab="exercise induced angina")
> spineplot(target~slope,data=heart1,ylab="Heart Disease",xlab="slope of the peak exercise ST segment")
> spineplot(target~thal,data=heart1,ylab="Heart Disease",xlab="Thalassemia")

```

> **#step 4.model selection**

model 1 selection

```

> #model selection without interaction
> fit.1=glm(target~1,data=heart1,family="binomial")
> fit.2=glm(target~.,data=heart1,family="binomial")
> select.1=step(fit.2, scope=list(lower=fit.1,upper=fit.2),direction="backward")
Start: AIC=227.57
target ~ age + sex + cp + trestbps + chol + fbs + restecg + thalach +
  exang + oldpeak + slope + ca + thal

```

	Df	Deviance	AIC
- age	1	197.92	225.92
- oldpeak	1	198.85	226.85
- restecg	1	198.97	226.97
<none>		197.57	227.57
- fbs	1	199.71	227.71
- chol	1	199.75	227.75
- thalach	1	200.16	228.16
- exang	1	200.52	228.52
- trestbps	1	201.81	229.81
- slope	1	202.03	230.03
- sex	1	205.32	233.32
- thal	1	206.75	234.75
- cp	2	213.38	239.38
- ca	1	226.66	254.66

Step: AIC=225.92

target ~ sex + cp + trestbps + chol + fbs + restecg + thalach +
exang + oldpeak + slope + ca + thal

	Df	Deviance	AIC
- restecg	1	199.29	225.29
- oldpeak	1	199.30	225.30
- chol	1	199.90	225.90
<none>		197.92	225.92
- fbs	1	200.14	226.14
- thalach	1	200.17	226.17
- exang	1	200.96	226.96
- trestbps	1	201.81	227.81
- slope	1	202.24	228.24
- sex	1	206.19	232.19
- thal	1	207.01	233.01
- cp	2	214.17	238.17
- ca	1	227.54	253.54

Step: AIC=225.29

target ~ sex + cp + trestbps + chol + fbs + thalach + exang +
oldpeak + slope + ca + thal

	Df	Deviance	AIC
- oldpeak	1	200.60	224.60
<none>		199.29	225.29
- fbs	1	201.41	225.41
- thalach	1	201.61	225.61
- chol	1	201.96	225.96
- exang	1	202.41	226.41
- trestbps	1	203.65	227.65
- slope	1	204.41	228.41
- thal	1	207.71	231.71
- sex	1	208.79	232.79
- cp	2	215.53	237.53
- ca	1	229.65	253.65

Step: AIC=224.6

target ~ sex + cp + trestbps + chol + fbs + thalach + exang +
slope + ca + thal

	Df	Deviance	AIC
<none>		200.60	224.60
- fbs	1	202.92	224.92
- thalach	1	203.13	225.13
- chol	1	203.40	225.40
- exang	1	204.32	226.32
- trestbps	1	205.67	227.67
- thal	1	209.51	231.51
- slope	1	209.85	231.85
- sex	1	211.23	233.23
- cp	2	216.55	236.55
- ca	1	235.44	257.44

model 1

```
> summary(select.1)
```

Call:

```
glm(formula = target ~ sex + cp + trestbps + chol + fbs + thalach +  
exang + slope + ca + thal, family = "binomial", data = heart1)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-2.8956	-0.4994	-0.1579	0.4235	2.5855

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-3.802180	2.240935	-1.697	0.08975 .
sexM	1.550460	0.494219	3.137	0.00171 **
cpasymptomatic	1.380991	0.470005	2.938	0.00330 **
cpnon_anginal	-0.297578	0.507756	-0.586	0.55783
trestbps	0.022301	0.010149	2.197	0.02800 *
chol	0.006327	0.003736	1.693	0.09038 .
fbsTrue	-0.857185	0.570129	-1.503	0.13271
thalach	-0.015184	0.009723	-1.562	0.11838
exangYes	0.820650	0.423148	1.939	0.05245 .
slopeup	-1.223228	0.409763	-2.985	0.00283 **
ca	1.303097	0.252968	5.151	2.59e-07 ***
thalnormal	-1.171886	0.394730	-2.969	0.00299 **

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

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 409.95 on 296 degrees of freedom
Residual deviance: 200.60 on 285 degrees of freedom
AIC: 224.6

Number of Fisher Scoring iterations: 6

```
> coef(select.1)
```

(Intercept)	sexM	cpasymptomatic	cpnon_anginal	trestbps	chol	fbsTrue	thalach
-3.802180287	1.550459839	1.380991253	-0.297578161	0.022300661	0.006326923	-0.857185090	-0.015183944
exangYes	slopeup	ca	thalnormal				
0.820650053	-1.223227874	1.303096651	-1.171886079				

>

model 2 selection

```
> #model selection with interaction
> fit.3=glm(target ~ (sex + cp + trestbps + chol + fbs + thalach + exang + slope + ca + thal)^2, family =
"binomial", data = heart1)
> select.2=step(select.1, scope=list(lower=select.1,upper=fit.3),direction="forward")
Start: AIC=224.6
target ~ sex + cp + trestbps + chol + fbs + thalach + exang +
      slope + ca + thal
```

	Df	Deviance	AIC
+ exang:slope	1	196.84	222.84
+ sex:fbs	1	197.86	223.86
+ fbs:exang	1	198.11	224.11
+ sex:exang	1	198.58	224.58
+ chol:fbs	1	198.60	224.60
<none>		200.60	224.60
+ sex:thal	1	198.63	224.63
+ fbs:slope	1	198.66	224.66
+ cp:chol	2	196.92	224.92
+ trestbps:exang	1	199.00	225.00
+ chol:slope	1	199.15	225.15
+ sex:trestbps	1	199.55	225.55
+ trestbps:chol	1	199.62	225.62
+ thalach:ca	1	199.72	225.72
+ exang:thal	1	199.87	225.87
+ chol:ca	1	199.88	225.88
+ exang:ca	1	199.99	225.99
+ fbs:thalach	1	200.09	226.09
+ chol:exang	1	200.10	226.10
+ cp:trestbps	2	198.14	226.14
+ sex:chol	1	200.15	226.15
+ fbs:ca	1	200.17	226.17
+ cp:ca	2	198.26	226.26
+ trestbps:thal	1	200.29	226.29
+ thalach:thal	1	200.32	226.32
+ slope:thal	1	200.37	226.37
+ sex:slope	1	200.38	226.38
+ trestbps:slope	1	200.39	226.39
+ sex:thalach	1	200.40	226.40
+ cp:thalach	2	198.42	226.42
+ trestbps:ca	1	200.47	226.47
+ trestbps:fbs	1	200.50	226.50
+ chol:thal	1	200.50	226.50
+ slope:ca	1	200.50	226.50
+ chol:thalach	1	200.52	226.52
+ ca:thal	1	200.52	226.52
+ sex:ca	1	200.53	226.53
+ thalach:exang	1	200.58	226.58
+ fbs:thal	1	200.59	226.59
+ trestbps:thalach	1	200.60	226.60
+ sex:cp	2	198.60	226.60
+ thalach:slope	1	200.60	226.60
+ cp:exang	2	198.87	226.87

```

+ cp:slope      2  200.07 228.07
+ cp:fbs        2  200.17 228.17
+ cp:thal       2  200.18 228.18

```

Step: AIC=222.84

target ~ sex + cp + trestbps + chol + fbs + thalach + exang +
 slope + ca + thal + exang:slope

	Df	Deviance	AIC
+ sex:fbs	1	194.30	222.30
<none>		196.84	222.84
+ trestbps:exang	1	194.85	222.85
+ chol:fbs	1	195.11	223.11
+ fbs:exang	1	195.11	223.11
+ exang:ca	1	195.12	223.12
+ fbs:slope	1	195.16	223.16
+ cp:chol	2	193.34	223.34
+ chol:slope	1	195.46	223.46
+ sex:thal	1	195.50	223.50
+ sex:exang	1	195.63	223.63
+ trestbps:chol	1	195.94	223.94
+ chol:ca	1	196.01	224.01
+ sex:trestbps	1	196.02	224.02
+ exang:thal	1	196.08	224.08
+ chol:exang	1	196.13	224.13
+ thalach:exang	1	196.21	224.21
+ thalach:thal	1	196.28	224.28
+ thalach:ca	1	196.39	224.39
+ sex:chol	1	196.43	224.43
+ fbs:thalach	1	196.46	224.46
+ cp:thalach	2	194.52	224.52
+ trestbps:thal	1	196.54	224.54
+ cp:trestbps	2	194.54	224.54
+ fbs:ca	1	196.57	224.57
+ slope:ca	1	196.57	224.57
+ trestbps:slope	1	196.59	224.59
+ cp:ca	2	194.61	224.61
+ sex:slope	1	196.65	224.65
+ thalach:slope	1	196.68	224.68
+ sex:thalach	1	196.71	224.71
+ sex:ca	1	196.76	224.76
+ trestbps:ca	1	196.78	224.78
+ slope:thal	1	196.79	224.79
+ chol:thalach	1	196.79	224.79
+ cp:exang	2	194.80	224.80
+ trestbps:fbs	1	196.80	224.80
+ chol:thal	1	196.81	224.81
+ trestbps:thalach	1	196.83	224.83
+ ca:thal	1	196.83	224.83
+ fbs:thal	1	196.83	224.83
+ cp:slope	2	195.00	225.00
+ sex:cp	2	195.21	225.21
+ cp:fbs	2	196.17	226.17
+ cp:thal	2	196.45	226.45

Step: AIC=222.3

target ~ sex + cp + trestbps + chol + fbs + thalach + exang +

slope + ca + thal + exang:slope + sex:fbs

	Df	Deviance	AIC
+ fbs:slope	1	190.88	220.88
+ trestbps:exang	1	191.97	221.97
<none>		194.30	222.30
+ exang:ca	1	192.39	222.39
+ fbs:exang	1	192.62	222.62
+ chol:ca	1	192.76	222.76
+ sex:thal	1	192.88	222.88
+ cp:chol	2	191.22	223.22
+ sex:exang	1	193.26	223.26
+ chol:slope	1	193.43	223.43
+ thalach:ca	1	193.56	223.56
+ sex:chol	1	193.57	223.57
+ sex:trestbps	1	193.59	223.59
+ trestbps:chol	1	193.60	223.60
+ chol:exang	1	193.60	223.60
+ exang:thal	1	193.65	223.65
+ thalach:exang	1	193.71	223.71
+ thalach:thal	1	193.77	223.77
+ sex:slope	1	193.84	223.84
+ cp:trestbps	2	191.87	223.87
+ trestbps:thal	1	193.88	223.88
+ sex:ca	1	193.91	223.91
+ fbs:ca	1	193.91	223.91
+ trestbps:slope	1	193.95	223.95
+ fbs:thal	1	193.98	223.98
+ cp:thalach	2	192.01	224.01
+ cp:slope	2	192.02	224.02
+ slope:ca	1	194.05	224.05
+ chol:thal	1	194.13	224.13
+ sex:thalach	1	194.14	224.14
+ chol:fbs	1	194.14	224.14
+ fbs:thalach	1	194.16	224.16
+ thalach:slope	1	194.17	224.17
+ trestbps:fbs	1	194.25	224.25
+ trestbps:thalach	1	194.28	224.28
+ chol:thalach	1	194.28	224.28
+ cp:exang	2	192.29	224.29
+ cp:ca	2	192.30	224.30
+ trestbps:ca	1	194.30	224.30
+ slope:thal	1	194.30	224.30
+ ca:thal	1	194.30	224.30
+ sex:cp	2	192.50	224.50
+ cp:fbs	2	193.92	225.92
+ cp:thal	2	194.01	226.01

Step: AIC=220.88

target ~ sex + cp + trestbps + chol + fbs + thalach + exang +
slope + ca + thal + exang:slope + sex:fbs + fbs:slope

	Df	Deviance	AIC
+ exang:ca	1	187.51	219.51
<none>		190.88	220.88
+ fbs:exang	1	188.96	220.96
+ trestbps:exang	1	189.00	221.00

+ thalach:ca	1	189.28	221.28
+ sex:exang	1	189.49	221.49
+ sex:thal	1	189.50	221.50
+ chol:ca	1	189.65	221.65
+ cp:chol	2	187.78	221.78
+ sex:ca	1	189.80	221.80
+ chol:slope	1	189.81	221.81
+ cp:trestbps	2	187.82	221.82
+ sex:trestbps	1	189.92	221.92
+ sex:slope	1	189.98	221.98
+ cp:exang	2	188.09	222.09
+ sex:chol	1	190.16	222.16
+ exang:thal	1	190.17	222.17
+ fbs:ca	1	190.20	222.20
+ chol:exang	1	190.26	222.26
+ thalach:exang	1	190.26	222.26
+ trestbps:chol	1	190.27	222.27
+ thalach:thal	1	190.40	222.40
+ trestbps:thal	1	190.42	222.42
+ fbs:thal	1	190.50	222.50
+ trestbps:slope	1	190.68	222.68
+ trestbps:fbs	1	190.71	222.71
+ chol:thal	1	190.77	222.77
+ ca:thal	1	190.77	222.77
+ sex:thalach	1	190.77	222.77
+ chol:thalach	1	190.80	222.80
+ thalach:slope	1	190.81	222.81
+ chol:fbs	1	190.83	222.83
+ trestbps:ca	1	190.86	222.86
+ trestbps:thalach	1	190.86	222.86
+ fbs:thalach	1	190.87	222.87
+ slope:thal	1	190.87	222.87
+ slope:ca	1	190.88	222.88
+ cp:ca	2	189.06	223.06
+ cp:slope	2	189.07	223.07
+ cp:fbs	2	189.14	223.14
+ sex:cp	2	189.14	223.14
+ cp:thalach	2	189.35	223.35
+ cp:thal	2	190.69	224.69

Step: AIC=219.51

target ~ sex + cp + trestbps + chol + fbs + thalach + exang +
slope + ca + thal + exang:slope + sex:fbs + fbs:slope + exang:ca

	Df	Deviance	AIC
+ trestbps:exang	1	183.52	217.52
+ thalach:ca	1	185.36	219.36
<none>		187.51	219.51
+ chol:ca	1	185.75	219.75
+ fbs:ca	1	185.97	219.97
+ sex:trestbps	1	186.07	220.07
+ sex:exang	1	186.10	220.10
+ chol:slope	1	186.18	220.18
+ sex:thal	1	186.22	220.22
+ thalach:exang	1	186.30	220.30
+ cp:trestbps	2	184.32	220.32
+ cp:chol	2	184.37	220.37

+ sex:ca	1	186.47	220.47
+ chol:exang	1	186.55	220.55
+ cp:exang	2	184.55	220.55
+ exang:thal	1	186.64	220.64
+ sex:slope	1	186.65	220.65
+ trestbps:chol	1	186.66	220.66
+ sex:chol	1	186.80	220.80
+ trestbps:thal	1	186.85	220.85
+ fbs:exang	1	186.97	220.97
+ thalach:thal	1	187.19	221.19
+ trestbps:fbs	1	187.29	221.29
+ fbs:thal	1	187.33	221.33
+ cp:slope	2	185.35	221.35
+ trestbps:slope	1	187.39	221.39
+ chol:thal	1	187.40	221.40
+ chol:thalach	1	187.43	221.43
+ sex:thalach	1	187.44	221.44
+ thalach:slope	1	187.46	221.46
+ trestbps:thalach	1	187.47	221.47
+ slope:ca	1	187.47	221.47
+ trestbps:ca	1	187.47	221.47
+ slope:thal	1	187.49	221.49
+ fbs:thalach	1	187.49	221.49
+ chol:fbs	1	187.50	221.50
+ ca:thal	1	187.51	221.51
+ sex:cp	2	185.54	221.54
+ cp:thalach	2	185.99	221.99
+ cp:fbs	2	186.33	222.33
+ cp:ca	2	186.35	222.35
+ cp:thal	2	187.19	223.19

Step: AIC=217.52

target ~ sex + cp + trestbps + chol + fbs + thalach + exang +
 slope + ca + thal + exang:slope + sex:fbs + fbs:slope + exang:ca +
 trestbps:exang

	Df	Deviance	AIC
+ thalach:ca	1	181.17	217.17
+ fbs:ca	1	181.25	217.25
<none>		183.52	217.52
+ chol:slope	1	181.59	217.59
+ exang:thal	1	181.70	217.70
+ sex:thal	1	182.05	218.05
+ cp:chol	2	180.13	218.13
+ thalach:exang	1	182.18	218.18
+ chol:ca	1	182.30	218.30
+ trestbps:chol	1	182.69	218.69
+ sex:trestbps	1	182.95	218.95
+ sex:chol	1	182.96	218.96
+ sex:slope	1	183.02	219.02
+ cp:exang	2	181.05	219.05
+ sex:ca	1	183.06	219.06
+ trestbps:fbs	1	183.07	219.07
+ thalach:thal	1	183.19	219.19
+ trestbps:thal	1	183.28	219.28
+ sex:exang	1	183.32	219.32
+ sex:thalach	1	183.33	219.33

+ fbs:thal	1	183.36	219.36
+ trestbps:thalach	1	183.37	219.37
+ chol:exang	1	183.41	219.41
+ chol:thal	1	183.43	219.43
+ trestbps:ca	1	183.45	219.45
+ fbs:exang	1	183.48	219.48
+ fbs:thalach	1	183.49	219.49
+ slope:ca	1	183.50	219.50
+ chol:fbs	1	183.51	219.51
+ chol:thalach	1	183.51	219.51
+ slope:thal	1	183.51	219.51
+ trestbps:slope	1	183.51	219.51
+ thalach:slope	1	183.51	219.51
+ ca:thal	1	183.52	219.52
+ cp:slope	2	181.54	219.54
+ cp:ca	2	181.81	219.81
+ cp:thalach	2	182.04	220.04
+ cp:fbs	2	182.12	220.12
+ sex:cp	2	182.14	220.14
+ cp:trestbps	2	182.23	220.23
+ cp:thal	2	183.07	221.07

Step: AIC=217.17

target ~ sex + cp + trestbps + chol + fbs + thalach + exang +
 slope + ca + thal + exang:slope + sex:fbs + fbs:slope + exang:ca +
 trestbps:exang + thalach:ca

	Df	Deviance	AIC
+ fbs:ca	1	178.60	216.60
+ thalach:exang	1	178.78	216.78
<none>		181.17	217.17
+ sex:thal	1	179.18	217.18
+ cp:chol	2	177.32	217.32
+ chol:slope	1	179.45	217.45
+ chol:ca	1	179.58	217.58
+ exang:thal	1	179.75	217.75
+ trestbps:chol	1	180.47	218.47
+ sex:slope	1	180.53	218.53
+ sex:trestbps	1	180.60	218.60
+ sex:chol	1	180.76	218.76
+ fbs:thal	1	180.78	218.78
+ cp:slope	2	178.80	218.80
+ cp:thalach	2	178.81	218.81
+ sex:ca	1	180.82	218.82
+ trestbps:fbs	1	180.86	218.86
+ thalach:thal	1	180.87	218.87
+ sex:exang	1	180.90	218.90
+ cp:exang	2	178.91	218.91
+ trestbps:thal	1	181.00	219.00
+ chol:exang	1	181.02	219.02
+ chol:thal	1	181.07	219.07
+ slope:ca	1	181.09	219.09
+ chol:thalach	1	181.15	219.15
+ thalach:slope	1	181.15	219.15
+ trestbps:slope	1	181.15	219.15
+ sex:thalach	1	181.15	219.15
+ trestbps:thalach	1	181.15	219.15

+ ca:thal	1	181.16	219.16
+ fbs:exang	1	181.16	219.16
+ trestbps:ca	1	181.17	219.17
+ fbs:thalach	1	181.17	219.17
+ slope:thal	1	181.17	219.17
+ chol:fbs	1	181.17	219.17
+ cp:fbs	2	179.37	219.37
+ cp:trestbps	2	179.45	219.45
+ sex:cp	2	179.90	219.90
+ cp:ca	2	179.97	219.97
+ cp:thal	2	180.85	220.85

Step: AIC=216.6

target ~ sex + cp + trestbps + chol + fbs + thalach + exang +
 slope + ca + thal + exang:slope + sex:fbs + fbs:slope + exang:ca +
 trestbps:exang + thalach:ca + fbs:ca

	Df	Deviance	AIC
+ thalach:exang	1	176.14	216.14
<none>		178.60	216.60
+ sex:thal	1	176.67	216.67
+ exang:thal	1	176.80	216.80
+ fbs:exang	1	176.89	216.89
+ chol:ca	1	176.90	216.90
+ cp:chol	2	174.92	216.92
+ chol:slope	1	177.03	217.03
+ fbs:thal	1	177.52	217.52
+ trestbps:chol	1	177.71	217.71
+ cp:thalach	2	175.77	217.77
+ sex:slope	1	178.07	218.07
+ sex:chol	1	178.09	218.09
+ thalach:thal	1	178.10	218.10
+ sex:trestbps	1	178.18	218.18
+ cp:slope	2	176.23	218.23
+ trestbps:fbs	1	178.24	218.24
+ chol:thal	1	178.45	218.45
+ trestbps:thalach	1	178.48	218.48
+ chol:exang	1	178.48	218.48
+ sex:ca	1	178.51	218.51
+ fbs:thalach	1	178.51	218.51
+ cp:exang	2	176.54	218.54
+ trestbps:thal	1	178.56	218.56
+ sex:exang	1	178.56	218.56
+ slope:ca	1	178.57	218.57
+ chol:fbs	1	178.58	218.58
+ thalach:slope	1	178.58	218.58
+ slope:thal	1	178.59	218.59
+ trestbps:slope	1	178.59	218.59
+ trestbps:ca	1	178.59	218.59
+ ca:thal	1	178.60	218.60
+ chol:thalach	1	178.60	218.60
+ sex:thalach	1	178.60	218.60
+ cp:fbs	2	176.75	218.75
+ cp:trestbps	2	176.86	218.86
+ cp:ca	2	177.46	219.46
+ sex:cp	2	177.54	219.54
+ cp:thal	2	178.18	220.18

Step: AIC=216.14

target ~ sex + cp + trestbps + chol + fbs + thalach + exang +
 slope + ca + thal + exang:slope + sex:fbs + fbs:slope + exang:ca +
 trestbps:exang + thalach:ca + fbs:ca + thalach:exang

	Df	Deviance	AIC
+ sex:thal	1	174.11	216.11
<none>		176.14	216.14
+ exang:thal	1	174.16	216.16
+ chol:ca	1	174.31	216.31
+ chol:slope	1	174.68	216.68
+ cp:chol	2	172.68	216.68
+ fbs:exang	1	174.79	216.79
+ fbs:thal	1	174.90	216.90
+ cp:exang	2	173.13	217.13
+ trestbps:chol	1	175.13	217.13
+ trestbps:fbs	1	175.51	217.51
+ sex:trestbps	1	175.72	217.72
+ sex:chol	1	175.75	217.75
+ sex:slope	1	175.75	217.75
+ cp:slope	2	173.89	217.89
+ chol:fbs	1	175.95	217.95
+ thalach:slope	1	175.95	217.95
+ cp:trestbps	2	174.00	218.00
+ chol:thal	1	176.00	218.00
+ slope:thal	1	176.07	218.07
+ thalach:thal	1	176.07	218.07
+ trestbps:thalach	1	176.07	218.07
+ fbs:thalach	1	176.09	218.09
+ sex:ca	1	176.10	218.10
+ chol:exang	1	176.10	218.10
+ ca:thal	1	176.10	218.10
+ sex:thalach	1	176.11	218.11
+ slope:ca	1	176.12	218.12
+ chol:thalach	1	176.13	218.13
+ sex:exang	1	176.13	218.13
+ trestbps:ca	1	176.13	218.13
+ trestbps:thal	1	176.14	218.14
+ cp:fbs	2	174.14	218.14
+ trestbps:slope	1	176.14	218.14
+ cp:thalach	2	174.21	218.21
+ cp:ca	2	174.97	218.97
+ sex:cp	2	175.15	219.15
+ cp:thal	2	175.34	219.34

Step: AIC=216.11

target ~ sex + cp + trestbps + chol + fbs + thalach + exang +
 slope + ca + thal + exang:slope + sex:fbs + fbs:slope + exang:ca +
 trestbps:exang + thalach:ca + fbs:ca + thalach:exang + sex:thal

	Df	Deviance	AIC
+ cp:chol	2	168.60	214.60
+ chol:slope	1	171.72	215.72
<none>		174.11	216.11
+ exang:thal	1	172.25	216.25
+ fbs:thal	1	172.32	216.32

```

+ trestbps:chol  1  172.33 216.33
+ chol:ca       1  172.78 216.78
+ fbs:exang     1  172.84 216.84
+ sex:cp        2  170.92 216.92
+ sex:chol      1  173.22 217.22
+ trestbps:fbs  1  173.40 217.40
+ cp:exang      2  171.51 217.51
+ sex:thalach   1  173.53 217.53
+ sex:trestbps  1  173.62 217.62
+ thalach:thal  1  173.70 217.70
+ cp:fbs        2  171.72 217.72
+ cp:slope      2  171.81 217.81
+ thalach:slope 1  173.85 217.85
+ chol:fbs      1  173.87 217.87
+ trestbps:thalach 1  173.92 217.92
+ trestbps:ca   1  173.97 217.97
+ ca:thal       1  174.00 218.00
+ sex:slope     1  174.03 218.03
+ trestbps:thal 1  174.05 218.05
+ sex:ca        1  174.07 218.07
+ chol:exang    1  174.08 218.08
+ chol:thal     1  174.10 218.10
+ cp:trestbps   2  172.10 218.10
+ fbs:thalach   1  174.11 218.11
+ chol:thalach  1  174.11 218.11
+ sex:exang     1  174.11 218.11
+ slope:ca      1  174.11 218.11
+ trestbps:slope 1  174.11 218.11
+ slope:thal    1  174.11 218.11
+ cp:thalach    2  172.91 218.91
+ cp:ca         2  173.04 219.04
+ cp:thal       2  173.45 219.45

```

Step: AIC=214.6

target ~ sex + cp + trestbps + chol + fbs + thalach + exang +
 slope + ca + thal + exang:slope + sex:fbs + fbs:slope + exang:ca +
 trestbps:exang + thalach:ca + fbs:ca + thalach:exang + sex:thal +
 cp:chol

	Df	Deviance	AIC
<none>		168.60	214.60
+ fbs:thal	1	166.91	214.91
+ exang:thal	1	167.00	215.00
+ chol:slope	1	167.10	215.10
+ chol:ca	1	167.13	215.13
+ sex:thalach	1	167.24	215.24
+ fbs:exang	1	167.35	215.35
+ cp:slope	2	165.47	215.47
+ thalach:thal	1	167.90	215.90
+ trestbps:chol	1	167.94	215.94
+ ca:thal	1	168.15	216.15
+ cp:fbs	2	166.19	216.19
+ trestbps:fbs	1	168.25	216.25
+ cp:trestbps	2	166.37	216.37
+ sex:exang	1	168.42	216.42
+ trestbps:thal	1	168.44	216.44
+ thalach:slope	1	168.44	216.44

```

+ trestbps:ca    1 168.44 216.44
+ chol:fbs      1 168.47 216.47
+ sex:ca        1 168.47 216.47
+ sex:cp        2 166.47 216.47
+ sex:slope     1 168.51 216.51
+ cp:exang      2 166.53 216.53
+ sex:trestbps  1 168.54 216.54
+ sex:chol      1 168.56 216.56
+ slope:thal    1 168.56 216.56
+ chol:thal     1 168.56 216.56
+ trestbps:thalach 1 168.57 216.57
+ fbs:thalach   1 168.58 216.58
+ trestbps:slope 1 168.59 216.59
+ chol:exang    1 168.59 216.59
+ chol:thalach  1 168.60 216.60
+ slope:ca      1 168.60 216.60
+ cp:ca         2 166.91 216.91
+ cp:thal       2 167.97 217.97
+ cp:thalach    2 168.10 218.10

```

model 2

```
> summary(select.2)
```

Call:

```

glm(formula = target ~ sex + cp + trestbps + chol + fbs + thalach +
    exang + slope + ca + thal + exang:slope + sex:fbs + fbs:slope +
    exang:ca + trestbps:exang + thalach:ca + fbs:ca + thalach:exang +
    sex:thal + cp:chol, family = "binomial", data = heart1)

```

Deviance Residuals:

```

    Min      1Q  Median      3Q     Max
-3.2694 -0.4296 -0.0565  0.2711  2.5510

```

Coefficients:

```

              Estimate Std. Error z value Pr(>|z|)
(Intercept)   -0.748855   4.044172  -0.185 0.85310
sexM           -0.422105   1.575812  -0.268 0.78880
cpasymptomatic  4.231470   3.237593   1.307 0.19122
cpnon_anginal  6.775567   3.399967   1.993 0.04628 *
trestbps       0.005113   0.013687   0.374 0.70874
chol           0.020434   0.010570   1.933 0.05321 .
fbsTrue        6.101057   2.596119   2.350 0.01877 *
thalach       -0.031219   0.015716  -1.986 0.04699 *
exangYes      -12.229935   5.061918  -2.416 0.01569 *
slopeup       -0.789815   0.532386  -1.484 0.13793
ca            -3.955055   2.069777  -1.911 0.05602 .
thalnormal    -4.219209   1.669240  -2.528 0.01148 *
exangYes:slopeup -2.839590   1.144702  -2.481 0.01311 *
sexM:fbsTrue   -7.982742   2.772779  -2.879 0.00399 **
fbsTrue:slopeup -8.435324   3.451059  -2.444 0.01451 *
exangYes:ca     2.928729   1.096162   2.672 0.00754 **
trestbps:exangYes 0.064587   0.027683   2.333 0.01964 *
thalach:ca      0.034936   0.014104   2.477 0.01325 *
fbsTrue:ca      1.886769   1.291571   1.461 0.14406
thalach:exangYes 0.035525   0.024176   1.469 0.14172

```

```
sexM:thalnormal    3.067549  1.712458  1.791  0.07324 .
cpasymptomatic:chol -0.010954  0.012892 -0.850  0.39550
cpnon_anginal:chol -0.028377  0.013742 -2.065  0.03893 *
```

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

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 409.95 on 296 degrees of freedom
Residual deviance: 168.60 on 274 degrees of freedom
AIC: 214.6

Number of Fisher Scoring iterations: 8

```
> coef(select.2)
(Intercept)      sexM  cpasymptomatic  cpnon_anginal  trestbps      chol
-0.748855160 -0.422105336   4.231470256   6.775567135   0.005112785   0.020433762
  fbsTrue    thalach    exangYes      slopeup        ca      thalnormal
  6.101057188 -0.031218683  -12.229934801  -0.789814583  -3.955054944  -4.219209069
exangYes:slopeup  sexM:fbsTrue  fbsTrue:slopeup  exangYes:ca  trestbps:exangYes  thalach:ca
-2.839589972   -7.982741976   -8.435324047    2.928728815    0.064587296    0.034936465
  fbsTrue:ca  thalach:exangYes  sexM:thalnormal  cpasymptomatic:chol  cpnon_anginal:chol
  1.886769110   0.035524638    3.067549209   -0.010953740   -0.028376734
```

Anova test of model 2

```
> library(car)# Anova
> Anova(select.2) #based on this, exang not significant remove exang in fit.4
Analysis of Deviance Table (Type II tests)
```

Anova table

```
Response: target
      LR Chisq Df Pr(>Chisq)
sex      10.938  1 0.0009422 ***
cp       13.282  2 0.0013060 **
trestbps   4.592  1 0.0321253 *
chol       0.971  1 0.3243812
fbs       4.987  1 0.0255352 *
thalach    0.546  1 0.4598773
exang     4.806  1 0.0283614 *
slope    14.386  1 0.0001489 ***
ca       37.765  1 7.979e-10 ***
thal     10.674  1 0.0010867 **
exang:slope  6.779  1 0.0092261 **
sex:fbs     9.403  1 0.0021666 **
fbs:slope   10.029  1 0.0015413 **
exang:ca    10.005  1 0.0015608 **
trestbps:exang  6.184  1 0.0128931 *
thalach:ca   5.871  1 0.0153966 *
fbs:ca       2.729  1 0.0985694 .
thalach:exang  2.180  1 0.1398572
sex:thal     4.079  1 0.0434178 *
```

```
cp:chol      5.510 2 0.0635985 .
```

```
---
```

```
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

final model

```
> fit.final=glm(target ~ sex + cp + trestbps + fbs + exang + slope + ca + thal + exang:slope + sex:fbs + fbs:slope +
+
+       exang:ca + trestbps:exang + fbs:ca + sex:thal,
+       data=heart1, family=binomial)
> summary(fit.final)
```

Call:

```
glm(formula = target ~ sex + cp + trestbps + fbs + exang + slope +
    ca + thal + exang:slope + sex:fbs + fbs:slope + exang:ca +
    trestbps:exang + fbs:ca + sex:thal, family = binomial, data = heart1)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-3.4569	-0.4336	-0.0947	0.2612	2.4811

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-1.95595	2.13466	-0.916	0.35952
sexM	0.14207	1.17904	0.121	0.90409
cpasymptomatic	1.43154	0.49096	2.916	0.00355 **
cpnon_anginal	-0.14392	0.53785	-0.268	0.78901
trestbps	0.01199	0.01292	0.928	0.35343
fbsTrue	6.10370	2.41927	2.523	0.01164 *
exangYes	-6.48437	3.53362	-1.835	0.06650 .
slopeup	-0.84664	0.47333	-1.789	0.07366 .
ca	1.25061	0.30519	4.098	4.17e-05 ***
thalnormal	-3.21868	1.24334	-2.589	0.00963 **
exangYes:slopeup	-2.33809	1.03681	-2.255	0.02413 *
sexM:fbsTrue	-7.79417	2.59945	-2.998	0.00271 **
fbsTrue:slopeup	-7.29079	3.34994	-2.176	0.02953 *
exangYes:ca	2.43431	1.05663	2.304	0.02123 *
trestbps:exangYes	0.05952	0.02731	2.179	0.02931 *
fbsTrue:ca	1.79628	1.29162	1.391	0.16431
sexM:thalnormal	2.01785	1.32224	1.526	0.12699

```
---
```

```
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 409.95 on 296 degrees of freedom
Residual deviance: 181.36 on 280 degrees of freedom
AIC: 215.36

Number of Fisher Scoring iterations: 8

```
> coef(fit.final)
```

(Intercept)	sexM	cpasymptomatic	cpnon_anginal	trestbps	fbsTrue
-1.95594655	0.14207425	1.43154201	-0.14392298	0.01199279	6.10370348
exangYes	slopeup	ca	thalnormal	exangYes:slopeup	sexM:fbsTrue

```

-6.48437093 -0.84664308 1.25061443 -3.21867936 -2.33808641 -7.79417495
fbsTrue:slopeup exangYes:ca trestbps:exangYes fbsTrue:ca sexM:thalnormal
-7.29079329 2.43430601 0.05952450 1.79627544 2.01784581

```

```
> confint(fit.final) #likelihood ratio CI
```

```
Waiting for profiling to be done...
```

```

2.5 % 97.5 %
(Intercept) -6.226246301 2.22132173
sexM -2.308944600 2.38077245
cpasymptomatic 0.491543888 2.42788816
cpnon_anginal -1.215875696 0.90940084
trestbps -0.013093513 0.03802441
fbsTrue 1.413968216 11.12328113
exangYes -13.828832009 0.13770603
slopeup -1.793345893 0.07425921
ca 0.686246497 1.89046044
thalnormal -5.832627281 -0.91695987
exangYes:slopeup -4.513718453 -0.40889522
sexM:fbsTrue -13.280754706 -2.86420426
fbsTrue:slopeup -14.570415005 -1.56275404
exangYes:ca 0.593558905 4.72522190
trestbps:exangYes 0.008979013 0.11683091
fbsTrue:ca -0.442775296 4.57517758
sexM:thalnormal -0.441355023 4.77877217
> Anova(fit.final)

```

```
Analysis of Deviance Table (Type II tests)
```

```
Response: target
```

```

LR Chisq Df Pr(>Chisq)
sex 7.956 1 0.004792 **
cp 13.378 2 0.001245 **
trestbps 7.105 1 0.007688 **
fbs 3.854 1 0.049623 *
exang 4.702 1 0.030134 *
slope 17.300 1 3.191e-05 ***
ca 43.574 1 4.082e-11 ***
thal 12.267 1 0.000461 ***
exang:slope 5.745 1 0.016536 *
sex:fbs 9.482 1 0.002075 **
fbs:slope 7.251 1 0.007086 **
exang:ca 7.391 1 0.006556 **
trestbps:exang 5.421 1 0.019894 *
fbs:ca 2.316 1 0.128061
sex:thal 2.540 1 0.111023

```

```
---
```

```
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
>
```

```
>
```

```
> oddsratio=cbind(coef(fit.final),confint(fit.final),exp(coef(fit.final)))
```

```
Waiting for profiling to be done...
```

```
> colnames(oddsratio)=c("coefficient", "95%CI(2.5%)", "95%CI(97.5.5%)", "odds ratio")
```

```
> oddsratio=format(round(oddsratio,2), nsmall=2)
```

odds ratio

```
> oddsratio
```

```
coefficient 95%CI(2.5%) 95%CI(97.5.5%) odds ratio
```

```

(Intercept)  "-1.96"  "-6.23"  " 2.22"  " 0.14"
sexM         " 0.14"  "-2.31"  " 2.38"  " 1.15"
cpasymptomatic " 1.43"  " 0.49"  " 2.43"  " 4.19"
cpnon_anginal "-0.14"  "-1.22"  " 0.91"  " 0.87"
trestbps     " 0.01"  "-0.01"  " 0.04"  " 1.01"
fbsTrue      " 6.10"  " 1.41"  "11.12"  "447.51"
exangYes     "-6.48"  "-13.83" " 0.14"  " 0.00"
slopeup      "-0.85"  "-1.79"  " 0.07"  " 0.43"
ca           " 1.25"  " 0.69"  " 1.89"  " 3.49"
thalnormal   "-3.22"  "-5.83"  "-0.92"  " 0.04"
exangYes:slopeup "-2.34"  "-4.51"  "-0.41"  " 0.10"
sexM:fbsTrue  "-7.79"  "-13.28" "-2.86"  " 0.00"
fbsTrue:slopeup "-7.29"  "-14.57" "-1.56"  " 0.00"
exangYes:ca    " 2.43"  " 0.59"  " 4.73"  "11.41"
trestbps:exangYes " 0.06"  " 0.01"  " 0.12"  " 1.06"
fbsTrue:ca     " 1.80"  "-0.44"  " 4.58"  " 6.03"
sexM:thalnormal " 2.02"  "-0.44"  " 4.78"  " 7.52"

```

ROC curve

```

> #step 5. model evalution: sensitivity/specificity/ROC/AUC
> pi0=seq(1,0, length.out = 100)
> nsample=nrow(heart1)
> roc=NULL
> cv.set.list=split(sample(nsample),rep(1:10,length=nsample)) #10-fold cross-validation method
> for(k in 1:length(pi0))
+ {n_11=n_10=n_01=n_00=0
+ nsample=nrow(heart1)
+ for(i in 1:length(cv.set.list))
+ {
+   training=heart1[-cv.set.list[[i]],]
+   test=heart1[cv.set.list[[i]],]
+   fit.training=glm(target ~ sex + cp + trestbps + fbs + exang + slope + ca + thal + exang:slope + sex:fbs +
+ fbs:slope +
+   exang:ca + trestbps:exang + fbs:ca + sex:thal, data=training, family="binomial")
+   for(j in 1:length(cv.set.list[[i]]))
+   {if(predict(fit.training, test[j,], type="response")>=pi0[k])
+   {Y.pred.1=1}else{Y.pred.1=0}
+   if((test$target[j]==1)&(Y.pred.1==1))
+   {n_11=n_11+1}
+   if((test$target[j]==1)&(Y.pred.1==0))
+   {n_10=n_10+1}
+   if((test$target[j]==0)&(Y.pred.1==1))
+   {n_01=n_01+1}
+   if((test$target[j]==0)&(Y.pred.1==0))
+   {n_00=n_00+1}
+   }
+ }
+ sen=n_11/(n_10+n_11)
+ spe=n_00/(n_00+n_01)
+ roc=rbind(roc, c(1-spe, sen))
+ }
> plot(roc, type="s",xlim=c(0,1), ylim=c(0,1), col="red", main="ROC curve", xlab="1-Specificity",
ylab="Sensitivity")
> auc=sum(roc[-100, 2]*(roc[-1,1]-roc[-100,1]))

```

```
> auc #AUC area under ROC curve
[1] 0.9091697
> pred.accuracy=roc[,2]*sum(heart1$target==1)/nsample+(1-roc[,1])*sum(heart1$target==0)/nsample
> z.1=which.max(pred.accuracy)
> pi0[z.1] #best cut-off points
[1] 0.4949495
> pred.accuracy[z.1] #best prediction accuracy
[1] 0.8518519
>
```