# **Question 11.1**

Using the crime data set uscrime.txt from Questions 8.2, 9.1, and 10.1, build a regression model using:

- Stepwise regression
- Lasso
- Elastic net

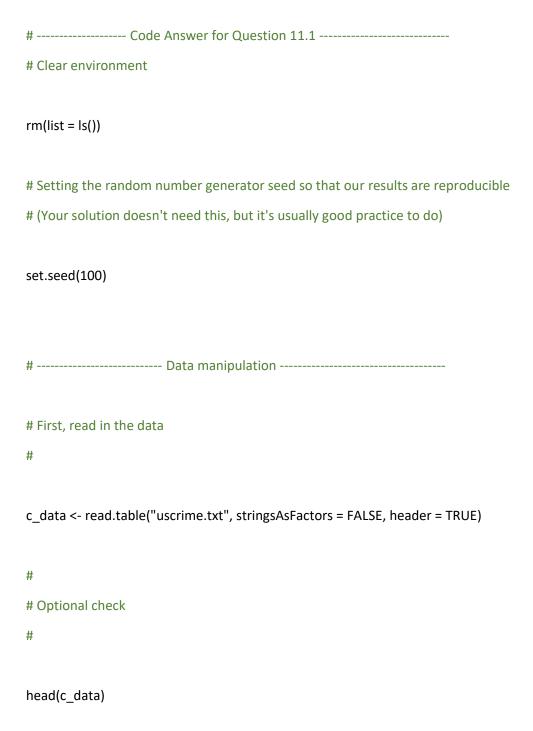
For Parts 2 and 3, remember to scale the data first – otherwise, the regression coefficients will be on different scales and the constraint won't have the desired effect.

For Parts 2 and 3, use the glmnet function in R.

#### Notes on R:

- For the elastic net model, what we called λ in the videos, glmnet calls "alpha"; you can get a range of results by varying alpha from 1 (lasso) to 0 (ridge regression) [and, of course, other values of alpha in between].
- In a function call like glmnet (x, y, family="mgaussian", alpha=1) the predictors x need to be in R's matrix format, rather than data frame format. You can convert a data frame to a matrix using as.matrix for example, x <- as.matrix(data[,1:n-1])
- Rather than specifying a value of T, glmnet returns models for a variety of values of T.

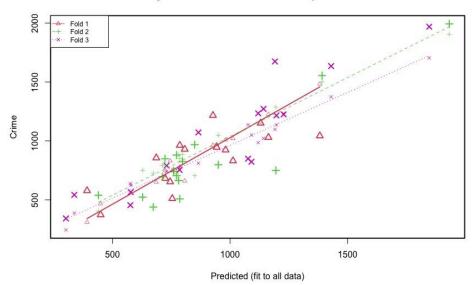
-----



```
# ------ Start Stepwise Regression -----
# Stepwise Regression using original variables and Cross Validation
# In backward stepwise regression.
#Scaling the data except the response variable and categorical
s_Data = as.data.frame(scale(c_data[,c(1,3,4,5,6,7,8,9,10,11,12,13,14,15)]))
s_Data <- cbind(c_data[,2],s_Data,c_data[,16]) # Add column 2 back in
colnames(s_Data)[1] <- "So"
colnames(s_Data)[16] <- "Crime"
library(caret)
# Perform 5 fold CV
ct_data <- trainControl(method = "repeatedcv", number = 5, repeats = 5)
ImFit_Step <- train(Crime ~ ., data = s_Data, "ImStepAIC", scope =</pre>
           list(lower = Crime~1, upper = Crime~.), direction = "backward",trControl=ct_data)
##Step: AIC=503.93
##.outcome ~ M + Ed + Po1 + M.F + U1 + U2 + Ineq + Prob
#
#Fitting a new model with the above 8 variables
mod_data = Im(Crime ~ M.F+U1+Prob+U2+M+Ed+Ineq+Po1, data = s_Data)
summary(mod_data)
```

cv.lm(c\_data,form.lm = formula(Crime  $\sim$  M.F+U1+Prob+U2+M+Ed+Ineq+Po1),m=3,dots=FALSE,plotit = TRUE,printit = TRUE)





##

##Residual standard error: 195.5 on 38 degrees of freedom

##Multiple R-squared: 0.7888, Adjusted R-squared: 0.7444

##F-statistic: 17.74 on 8 and 38 DF, p-value: 1.159e-10

#We got an Adjusted R-SQuared value = 0.7444 using the selected 8 variables using # Backward StepWise regression and Cross Validation

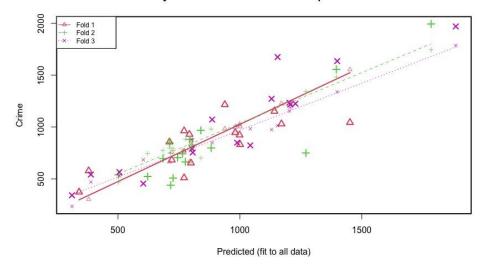
# Now let's use cross-validation twith only 47 data points, to see how good is the model # 47-fold cross-validation

#

SStot <- sum((c\_data\$Crime - mean(c\_data\$Crime))^2) totsse <- 0

```
for(i in 1:nrow(s_Data)) {
 mod_data_i = Im(Crime ~ M.F+U1+Prob+U2+M+Ed+Ineq+Po1, data = s_Data[-i,])
pred_i <- predict(mod_data_i,newdata=s_Data[i,])</pre>
totsse <- totsse + ((pred_i - c_data[i,16])^2)
}
R2_mod <- 1 - totsse/SStot
R2_mod
## 0.6676
# Notice that in the model above, the p-value for M.F is above 0.1.
# We might keep it in the model, because it's close to 0.1 and
# might be important. That's what we tested above.
# Or, we might remove it, and re-run the model without it.
# Let's see what happens if we do:
mod_data1 = Im(Crime ~ U1+Prob+U2+M+Ed+Ineq+Po1, data = s_Data)
summary(mod_data1)
cv.lm(s_Data,form.lm = formula(Crime ~ U1+Prob+U2+M+Ed+Ineq+Po1),m=3,dots=FALSE,plotit =
TRUE,printit = TRUE)
```

# Small symbols show cross-validation predicted values



#Residual standard error: 199.8 on 39 degrees of freedom

#Multiple R-squared: 0.7738, Adjusted R-squared: 0.7332

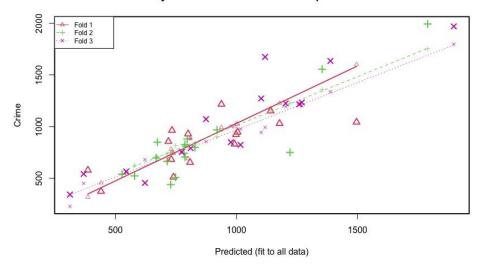
#F-statistic: 19.06 on 7 and 39 DF, p-value: 8.805e-11

# Now notice that U1 doesn't look significant... so we can take # it out too, and re-run the model.

mod\_data2 = Im(Crime ~ Prob+U2+M+Ed+Ineq+Po1, data = s\_Data)
summary(mod\_data2)

 $cv.lm(s\_Data,form.lm = formula(Crime ~ Prob+U2+M+Ed+Ineq+Po1), m=3, dots=FALSE, plot it = TRUE, print it = TRUE) \\$ 

### Small symbols show cross-validation predicted values



##

##Residual standard error: 200.7 on 40 degrees of freedom

##Multiple R-squared: 0.7659, Adjusted R-squared: 0.7307

##F-statistic: 21.8 on 6 and 40 DF, p-value: 3.42e-11

# This model looks good, so now let's see how it cross-validates:

```
SStot <- sum((c_data$Crime - mean(c_data$Crime))^2)

totsse <- 0

for(i in 1:nrow(s_Data)) {

mod_data2_i = lm(Crime ~ Prob+U2+M+Ed+Ineq+Po1, data = s_Data[-i,])

pred_i <- predict(mod_data2_i,newdata=s_Data[i,])

totsse <- totsse + ((pred_i - c_data[i,16])^2)

}

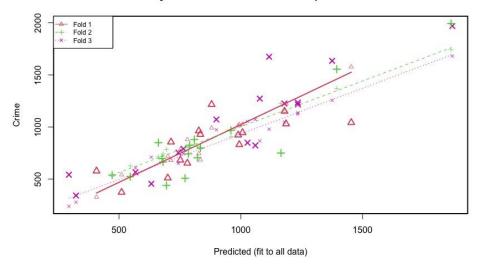
R3_mod <- 1 - totsse/SStot

R3_mod
```

## 0.666

```
# So, cross-validation shows that it's about the same whether
# we include M.F and U1 (0.668) or not (0.666). That gives some
# support to the idea that M.F and U1 really aren't significant.
# Since the quality is about the same, we should probably use the
# simpler model.
# ------ Lasso Regression ------
library(glmnet)
#building lasso Regression Model
XP=data.matrix(s_Data[,-16])
YP=data.matrix(s_Data$Crime)
lasso=cv.glmnet(x=as.matrix(s_Data[,-16]),y=as.matrix(s_Data$Crime),alpha=1,
        nfolds = 5,type.measure="mse",family="gaussian")
#Output the coefficients of the variables selected by lasso
coef(lasso, s=lasso$lambda.min)
#Fitting a new model with 9 variables
mod_lasso1 = Im(Crime ~So+M+Ed+Po1+M.F+NW+U2+Ineq+Prob, data = s_Data)
summary(mod_lasso1)
cv.lm(s_Data,form.lm = formula(Crime ~So+M+Ed+Po1+M.F+NW+U2+Ineq+Prob),m=3,dots=FALSE,plotit
= TRUE, printit = TRUE)
```

### Small symbols show cross-validation predicted values



##

##Residual standard error: 204.5 on 37 degrees of freedom

##Multiple R-squared: 0.775, Adjusted R-squared: 0.72

##F-statistic: 14.2 on 9 and 37 DF, p-value: 1.54e-09

#We obtain a slightly lower Adjusted R-SQuared value = 0.72 using the selected 9 variables using # Lasso regression and Cross Validation

# Now let's see how it cross-validates:

```
SStot <- sum((c_data$Crime - mean(c_data$Crime))^2)
totsse <- 0
for(i in 1:nrow(s_Data)) {
  mod_lasso1_i = lm(Crime ~ So+M+Ed+Po1+M.F+NW+U2+Ineq+Prob, data = s_Data[-i,])
  pred_i <- predict(mod_lasso1_i,newdata=s_Data[i,])
  totsse <- totsse + ((pred_i - c_data[i,16])^2)
}
LR3_mod <- 1 - totsse/SStot</pre>
```

```
LR3_mod
## 0.666
# based on above observation, three of the variables (So, M.F, and NW)
# are not significant. Let's remove them but
#It's exactly the same model we got above
# stepwis regression model!
# Please refer above R3_mod value
# ------ Elastic Net Regression-----
#We vary alpha in steps of 0.1 from 0 to 1 and calculate the resultant R-Squared values
R2=c()
for (i in 0:10) {
 mod_elastic = cv.glmnet(x=as.matrix(s_Data[,-16]),y=as.matrix(s_Data$Crime),
             alpha=i/10,nfolds = 5,type.measure="mse",family="gaussian")
 #The deviance(dev.ratio) shows the percentage of deviance explained,
 #(equivalent to r squared in case of regression)
 R2 = cbind(R2,mod_elastic$glmnet.fit$dev.ratio[which(mod_elastic$glmnet.fit$lambda ==
mod_elastic$lambda.min)])
}
```

```
##[,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9]
##[1,] 0.7062797 0.7535141 0.7386241 0.7402678 0.7603113 0.7734391 0.7886831 0.763631
0.7966378
##[,10] [,11]
##[1,] 0.7569592 0.7726187
#Best value of alpha
alpha_best = (which.max(R2)-1)/10
alpha_best
## 0.8
#Therefore we find that the best value of alpha may not lie somewhether between 0 and 1
#Lets build the model using this alpha value.
E_net=cv.glmnet(x=as.matrix(s_Data[,-16]),y=as.matrix(s_Data$Crime),alpha=alpha_best,
           nfolds = 5,type.measure="mse",family="gaussian")
#Output the coefficients of the variables selected by Elastic Net
coef(E_net, s=E_net$lambda.min)
```

# The Elastic Net selects 13 variables compared to 10 in Lasso and 8 in Step Wise.

# Next we compare how this new model performs

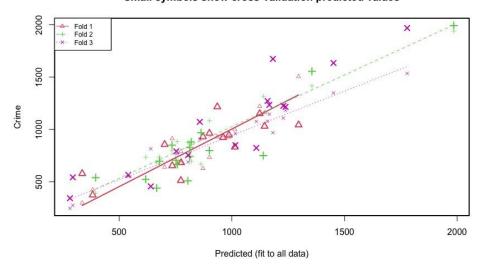
# compared to the Lasso and Step Wise models

 $mod_Elastic_net = Im(Crime \sim So+M+Ed+Po1+Po2+M.F+Pop+NW+U1+U2+Wealth+Ineq+Prob, data = s_Data)$ 

summary(mod\_Elastic\_net)

 $cv.lm(s\_Data,form.lm = formula(Crime \\ \sim So+M+Ed+Po1+Po2+M.F+Pop+NW+U1+U2+Wealth+Ineq+Prob), m=3, dots=FALSE, plotit = TRUE, printit = TRUE)$ 

### Small symbols show cross-validation predicted values



##

##Residual standard error: 204 on 33 degrees of freedom

##Multiple R-squared: 0.8005, Adjusted R-squared: 0.7219

##F-statistic: 10.19 on 13 and 33 DF, p-value: 4.088e-08

# The R-SQuared value is similar using Elastic Net and 13 variables. Therefore this method # may not be doing a good job as it selects 3 more variables for a similar RSquared value

# Now let's see how it cross-validates:

SStot <- sum((c\_data\$Crime - mean(c\_data\$Crime))^2) totsse <- 0

```
for(i in 1:nrow(s_Data)) {
 mod_Enet_i = Im(Crime ~ So+M+Ed+Po1+Po2+M.F+Pop+NW+U1+U2+Wealth+Ineq+Prob, data =
s Data[-i,])
 pred_i <- predict(mod_Enet_i,newdata=s_Data[i,])</pre>
totsse <- totsse + ((pred_i - c_data[i,16])^2)
}
ER5_mod <- 1 - totsse/SStot
ER5_mod
## 0.574
# That's a much worse cross-validated R-squared estimate. Why?
# As before, look at the p-values. Most of those variables'
# p-values seem to indicate that they're not significant.
# If we remove them all, we're left with M, Ed, Po1, U2, Ineq,
# and Prob.
# Does that look familiar? It should -- it's the same set of 6
# variables we were left with after removing insignificant ones
# from the Stepwise and Lasso models above!
# Before we quit, let's go back and use PCA on the variables,
# and then build Stepwise, Lasso, and Elastic Net models using
# the principal components.
# -----Implementing the above 3 models using Principal Component Analysis------
# Run PCA on matrix of scaled predictors
```

```
pca <- prcomp(c_data[,1:15], scale. = TRUE)</pre>
summary(pca)
# PCA PLOT
screeplot(pca, type="lines",col="blue")
var <- pca$sdev^2
propvar <- var/sum(var)</pre>
plot(propvar, xlab = "Principal Component", ylab = "Proportion of Variance Explained", ylim = c(0,1), type
= "b")
plot(cumsum(propvar), xlab = "Principal Component", ylab = "Cumulative Proportion of Variance
Explained",
  ylim = c(0,1), type = "b")
# For the purpose of this question, let us use all the PCs instead of the original variables
# and evaluate the performance of the 3 above models
# Creating a dataframe of response variable and PCs
#-----
PCcrime <- as.data.frame(cbind(pca$x, c_data[,16]))
colnames(PCcrime)[16] <- "Crime"
```

```
# ------ Stepwise Regression ------
# Stepwise Regression using PCs and Cross Validation
# In backward stepwise regression. Our lower model will have only the intercept
# and all variables in our full model.
library(caret)
# Now use the code below to perform 5 fold CV
ctrl <- trainControl(method = "repeatedcv", number = 5, repeats = 5)
set.seed(1)
ImFit_Step_PC <- train(Crime ~ ., data = PCcrime, "ImStepAIC", scope =</pre>
             list(lower = Crime~1, upper = Crime~.), direction = "backward",trControl=ctrl)
##Step: AIC=507.37
##.outcome ~ PC1 + PC2 + PC4 + PC5 + PC6 + PC7 + PC12 + PC14 +
##PC15
#Fitting a new model with these 9 PCS
mod1_Step_PC = Im(Crime ~ PC15+PC6+PC14+PC7+PC4+PC12+PC2+PC1+PC5, data = PCcrime)
summary(mod1_Step_PC)
```

```
##Residual standard error: 201.2 on 37 degrees of freedom
##Multiple R-squared: 0.7823, Adjusted R-squared: 0.7293
##F-statistic: 14.77 on 9 and 37 DF, p-value: 8.755e-10
#We obtain an Adjusted R-SQuared value = 0.729 using the selected 9PCs using
# Backward StepWise regression and Cross Validation. This is slightly lower
# than using the same method on the original variables
# Now let's see how it cross-validates:
SStot <- sum((c_data$Crime - mean(c_data$Crime))^2)</pre>
totsse <- 0
for(i in 1:nrow(PCcrime)) {
mod_lasso_i = lm(Crime ~ PC15+PC6+PC14+PC7+PC4+PC12+PC2+PC1+PC5, data = PCcrime[-i,])
pred_i <- predict(mod_lasso_i,newdata=PCcrime[i,])</pre>
totsse <- totsse + ((pred_i - PCcrime[i,16])^2)
}
RR2_mod <- 1 - totsse/SStot
RR2_mod
## 0.6311
# Notice that PC15 and PC6 were not significant in the model
# above. If we take them out, here's what we get:
mod_Step2_PC = Im(Crime ~ PC14+PC7+PC4+PC12+PC2+PC1+PC5, data = PCcrime)
summary(mod_Step2_PC)
```

```
##
##Residual standard error: 207 on 39 degrees of freedom
##Multiple R-squared: 0.757, Adjusted R-squared: 0.713
##F-statistic: 17.3 on 7 and 39 DF, p-value: 3.41e-10
# Now let's see how it cross-validates:
SStot <- sum((c_data$Crime - mean(c_data$Crime))^2)
totsse <- 0
for(i in 1:nrow(PCcrime)) {
mod_lasso_i = lm(Crime ~ PC14+PC7+PC4+PC12+PC2+PC1+PC5, data = PCcrime[-i,])
pred_i <- predict(mod_lasso_i,newdata=PCcrime[i,])</pre>
totsse <- totsse + ((pred_i - PCcrime[i,16])^2)
RR3_mod <- 1 - totsse/SStot
RR3_mod
## 0.627
# About the same as above, so the simpler model might be better to use.
# ------ Lasso Regression ------
library(glmnet)
```

#building lasso

```
XP=data.matrix(PCcrime[,-16])
YP=data.matrix(PCcrime$Crime)
lasso_PC=cv.glmnet(x=as.matrix(PCcrime[,-16]),y=as.matrix(PCcrime$Crime),alpha=1,
                                nfolds = 5,type.measure="mse",family="gaussian")
#Output the coefficients of the variables selected by lasso
coef(lasso_PC, s=lasso_PC$lambda.min)
#Fitting a new model with these 12 PCs compared to 10 original variables
mod_lPC = Im(Crime \sim PC1 + PC2 + PC3 + PC4 + PC5 + PC6 + PC7 + PC10 + PC12 + PC13 + PC14 + PC15, data = PC1 + PC10 + PC
PCcrime)
summary(mod_I_PC)
##
##Residual standard error: 202.1 on 34 degrees of freedom
##Multiple R-squared: 0.7981, Adjusted R-squared: 0.7269
##F-statistic: 11.2 on 12 and 34 DF, p-value: 1.408e-08
#We obtain a similar Adjusted R-SQuared value = 0.7269 using the selected 12 PCs instead
# of the 10 variables using Lasso regression
# Now let's see how it cross-validates:
SStot <- sum((c_data$Crime - mean(c_data$Crime))^2)</pre>
```

```
totsse <- 0
for(i in 1:nrow(PCcrime)) {
mod_lasso_i = lm(Crime ~ PC1+PC2+PC3+PC4+PC5+PC6+PC7+PC10+PC12+PC13+PC14+PC15, data =
PCcrime[-i,])
pred_i <- predict(mod_lasso_i,newdata=PCcrime[i,])</pre>
totsse <- totsse + ((pred i - PCcrime[i,16])^2)
LRR2_mod <- 1 - totsse/SStot
LRR2_mod
## 0.5857
# Looks worse. But notice that PCs 3, 6, 10, 13, and 15 do not
# appear to be significant. Let's take them out.
# When we do, we get the same model as when we use only significant
# variables from the stepwise PC model.
# ------ Elastic Net -----
#We vary alpha in steps of 0.1 from 0 to 1 and calculate the resultant R-Squared values
R2_PC=c()
for (i in 0:10) {
model = cv.glmnet(x=as.matrix(PCcrime[,-16]),y=as.matrix(PCcrime$Crime),
          alpha=i/10,nfolds = 5,type.measure="mse",family="gaussian")
#The deviance(dev.ratio) shows the percentage of deviance explained,
#(equivalent to r squared in case of regression)
```

```
R2_PC = cbind(R2_PC,model$glmnet.fit$dev.ratio[which(model$glmnet.fit$lambda ==
model$lambda.min)])
}
R2_PC
##
            [,2]
                  [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10] [,11]
      [,1]
##[1,] 0.7695465 0.7517182 0.7787271 0.8014505 0.749221 0.7857614 0.7590517 0.7981891
0.7635869 0.7937638 0.7940698
#Best value of alpha
alpha_best_PC = (which.max(R2_PC)-1)/10
alpha_best_PC
## 0.3
# An interesting observation after we use PCs instead of original variables. We observe that the best
# alpha value=0.3 which is slightly closer to a Lasso model. The R-Squared values are
# slightly higher here. Lets build the model using this alpha value.
Elastic_net_PC=cv.glmnet(x=as.matrix(PCcrime[,-16]),y=as.matrix(PCcrime$Crime),alpha=alpha_best,
             nfolds = 5,type.measure="mse",family="gaussian")
#Output the coefficients of the variables selected by Elastic Net
coef(Elastic_net_PC, s=Elastic_net_PC$lambda.min)
```

```
# The Elastic Net selects only 10 PCs compared to 12 in Lasso. Next we compare how this new model
performs
# compared to the Lasso model
mod_Elastic_net_PC = Im(Crime \sim PC1+PC2+PC3+PC4+PC5+PC6+PC7+PC12+PC14+PC15, data = PCcrime)
summary(mod_Elastic_net_PC)
##
##Residual standard error: 200 on 36 degrees of freedom
##Multiple R-squared: 0.7908, Adjusted R-squared: 0.7327
##F-statistic: 13.61 on 10 and 36 DF, p-value: 1.785e-09
# The R-SQuared value is slightly higher using Elastic Net and only 10 PCS compared to 12 PCs which
# was returned by Lasso. Elastic Net performs relatively better compared to Stepwise and Lasso
# Now let's see how it cross-validates:
SStot <- sum((c_data$Crime - mean(c_data$Crime))^2)</pre>
totsse <- 0
for(i in 1:nrow(PCcrime)) {
mod_lasso_i = lm(Crime ~ PC1+PC2+PC3+PC4+PC5+PC6+PC7+PC12+PC14+PC15, data = PCcrime[-i,])
pred_i <- predict(mod_lasso_i,newdata=PCcrime[i,])</pre>
totsse <- totsse + ((pred i - PCcrime[i,16])^2)
}
ER PC mod <- 1 - totsse/SStot
ER PC mod
```

# If we take out the seemingly-insignificant variables PC3,

# PC6, and PC15, we're left with the same model we had before

# after taking insignificant variables out of a PC model.