

Head and Neck Cancer Progression-free Survival Prediction Based on Texture Data Using LASSO (with R code)

In this project, our objective is to build a predictive model for head and neck cancer progressive-free survival (PFS), which is also our respond of interest. Our predictors are textures of fractional intravascular blood volume at baseline measurement or follow-ups.

In this project, the major difficulty is how to select the right variables. The raw data from the cancer texture is very large and most of those predictors are not as significant to our interest. The LASSO (Least Absolute Shrinkage and Selection Operator) is a regression method that involves penalizing the absolute size of the regression coefficients. As a result, we decided to use LASSO to select the variables and developed our own lasso function using R.

Data Manipulation

To begin with, data manipulation is performed before any regression process:

- (a) Remove completely missing columns.
- (b) Remove rows with completely missing predictors.
- (c) Remove columns with constant predictors.
- (d) Remove (second or later occurrence of) columns which are collinear (correlation 1) with another.
- (e) Calculate “baseline average” and “follow-up differences”

Table I shows the size of predictor matrix after each step of the data manipulation. The tolerance is chosen as 10^{-4} .

TABLE 1 Predictor Matrix Sizes

| | Observation Number | Predictor Number |
|-----------------|---------------------------|-------------------------|
| Raw Data | 33 | 238 |
| Step (a) | 33 | 235 |
| Step (b) | 29 | 235 |
| Step (c) | 29 | 235 |
| Step (d) | 29 | 219 |
| Step (e) | 29 | 150 |

Multiple Imputation

Multiple imputation is a statistical technique for analyzing incomplete data sets, that is, data sets for which some entries are missing. As a result, multiple imputation allows us to perform complete dataset analyses (which many/most statistical and machine learning procedures require), and obtain corresponding estimates of interest on each complete dataset. In this project, three imputed datasets are created through the multiple imputation process using the R library mi.

Please see the following references for multiple imputation.

- Su, Yu-Sung, et al. "Multiple imputation with diagnostics (mi) in R: Opening windows into the black box." *Journal of Statistical Software* 45.2 (2011): 1-31;
- Rubin, Donald B. "Multiple imputation after 18+ years." *Journal of the American Statistical Association* 91.434 (1996): 473-489.

Lasso via Modification to Least Angle Regression

The lasso algorithm proposed by Efron is realized through R (see the Appendix).

- Efron, Bradley, et al. "Least angle regression." *The Annals of statistics* 32.2 (2004): 407-499.

In order to test my LASSO algorithm, I compared my lasso algorithm actions with the actions in the library lars in R. In TABLE II, "Action 1" is acquired from my algorithm, and "Action 2" is acquired from the library function lars. They are exactly the same, and both take 110 iterations to gain the final results.

TABLE II Action Comparison

| Iter. | Action 1 | Action 2 |
|-------|----------|----------|-------|----------|----------|-------|----------|----------|-------|----------|----------|
| 1 | 90 | 90 | 31 | 130 | 130 | 61 | 64 | 64 | 91 | 63 | 63 |
| 2 | 127 | 127 | 32 | 132 | 132 | 62 | -63 | -63 | 92 | -17 | -17 |
| 3 | 96 | 96 | 33 | -118 | -118 | 63 | 58 | 58 | 93 | -63 | -63 |
| 4 | 55 | 55 | 34 | 99 | 99 | 64 | 62 | 62 | 94 | -124 | -124 |
| 5 | 86 | 86 | 35 | 131 | 131 | 65 | 124 | 124 | 95 | -118 | -118 |
| 6 | 105 | 105 | 36 | -79 | -79 | 66 | 63 | 63 | 96 | 124 | 124 |
| 7 | 84 | 84 | 37 | 116 | 116 | 67 | -17 | -17 | 97 | 50 | 50 |
| 8 | -86 | -86 | 38 | 118 | 118 | 68 | -105 | -105 | 98 | -124 | -124 |
| 9 | 56 | 56 | 39 | 79 | 79 | 69 | 78 | 78 | 99 | 16 | 16 |
| 10 | 146 | 146 | 40 | -133 | -133 | 70 | 105 | 105 | 100 | 48 | 48 |
| 11 | -84 | -84 | 41 | 145 | 145 | 71 | -94 | -94 | 101 | 3 | 3 |
| 12 | 75 | 75 | 42 | 96 | 96 | 72 | -67 | -67 | 102 | -15 | -15 |
| 13 | 84 | 84 | 43 | -84 | -84 | 73 | -105 | -105 | 103 | 142 | 142 |
| 14 | 94 | 94 | 44 | 41 | 41 | 74 | 118 | 118 | 104 | -44 | -44 |
| 15 | -96 | -96 | 45 | -79 | -79 | 75 | 147 | 147 | 105 | 15 | 15 |
| 16 | -56 | -56 | 46 | 49 | 49 | 76 | 120 | 120 | 106 | 104 | 104 |
| 17 | 99 | 99 | 47 | 67 | 67 | 77 | -124 | -124 | 107 | -16 | -16 |
| 18 | 78 | 78 | 48 | -145 | -145 | 78 | -147 | -147 | 108 | 118 | 118 |
| 19 | -99 | -99 | 49 | -41 | -41 | 79 | -118 | -118 | 109 | -3 | -3 |
| 20 | 79 | 79 | 50 | -78 | -78 | 80 | 17 | 17 | 110 | 145 | 145 |
| 21 | 69 | 69 | 51 | 44 | 44 | 81 | 15 | 15 | 111 | | |
| 22 | 133 | 133 | 52 | -118 | -118 | 82 | -131 | -131 | 112 | | |
| 23 | 51 | 51 | 53 | -55 | -55 | 83 | -99 | -99 | 113 | | |
| 24 | 54 | 54 | 54 | 118 | 118 | 84 | 144 | 144 | 114 | | |
| 25 | 62 | 62 | 55 | 76 | 76 | 85 | 117 | 117 | 115 | | |
| 26 | 137 | 137 | 56 | -118 | -118 | 86 | -127 | -127 | 116 | | |
| 27 | 118 | 118 | 57 | 62 | 62 | 87 | -63 | -63 | 117 | | |
| 28 | 126 | 126 | 58 | 17 | 17 | 88 | 124 | 124 | 118 | | |
| 29 | -62 | -62 | 59 | 63 | 63 | 89 | 99 | 99 | 119 | | |
| 30 | -137 | -137 | 60 | -62 | -62 | 90 | 118 | 118 | | | |

I also compared the β from my algorithm (β_1) with the β from the function lars in the library (β_2), see TABLE III. They are exactly the same.

TABLE III Action Comparison

| Predictor ID | β_1 | β_2 | Predictor ID | β_1 | β_2 |
|--------------|-------------|-------------|--------------|-------------|-------------|
| 1 | 0.1399669 | 0.1399669 | 15 | 0.3503249 | 0.3503249 |
| 2 | -0.15348659 | -0.15348659 | 16 | -0.05922694 | -0.05922694 |
| 3 | -1.47835984 | -1.47835984 | 17 | -0.03209639 | -0.03209639 |
| 4 | 0.25446971 | 0.25446971 | 18 | 0.40883653 | 0.40883653 |
| 5 | -0.14958195 | -0.14958195 | 19 | 0.12781619 | 0.12781619 |
| 6 | 0.95110139 | 0.95110139 | 20 | 0.03902563 | 0.03902563 |
| 7 | 1.11528411 | 1.11528411 | 21 | -0.48478189 | -0.48478189 |
| 8 | -1.32583124 | -1.32583124 | 22 | 0.3573575 | 0.3573575 |
| 9 | 0.67937989 | 0.67937989 | 23 | 0.16942768 | 0.16942768 |
| 10 | 0.2776513 | 0.2776513 | 24 | 0.71598979 | 0.71598979 |
| 11 | 0.61291334 | 0.61291334 | 25 | -0.1635122 | -0.1635122 |
| 12 | -0.24589955 | -0.24589955 | 26 | 0.16354518 | 0.16354518 |
| 13 | -0.13537225 | -0.13537225 | 27 | -0.0033004 | -0.0033004 |
| 14 | -0.09928269 | -0.09928269 | 28 | 0.2182681 | 0.2182681 |

- Figure 1 shows the sum of absolute beta value (t) vs. iteration
- Figure 2 shows my lasso path
- Figure 3 shows the lasso path from library lars
- Figure 4 shows \overline{err} (the apparent mean absolute predictive error rate)
- Figure 5 γ (the no information error rate)

sum of absolute beta Vs. iteration

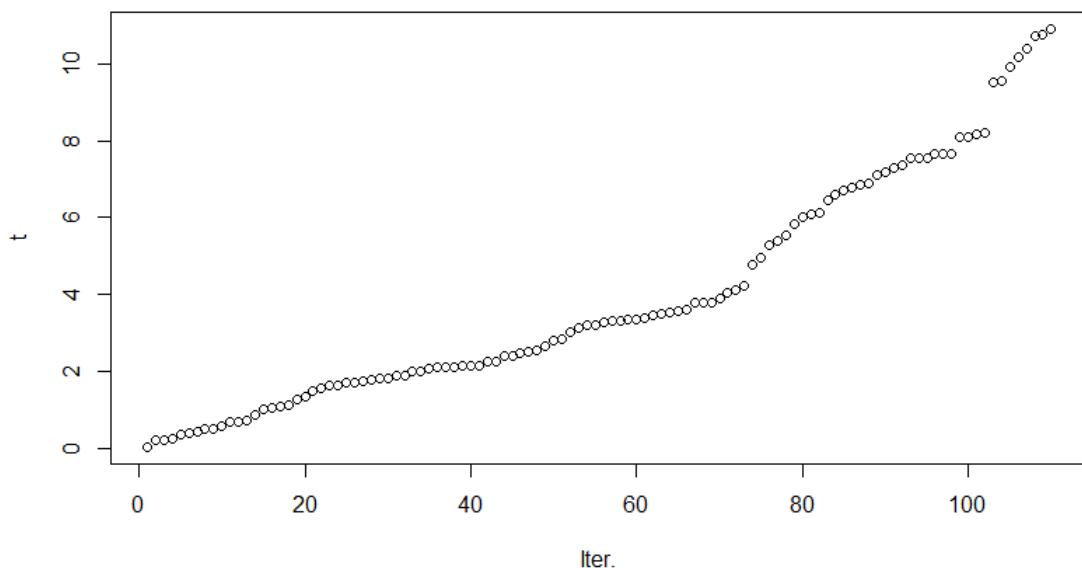


Figure 1 sum of absolute beta Vs. iteration

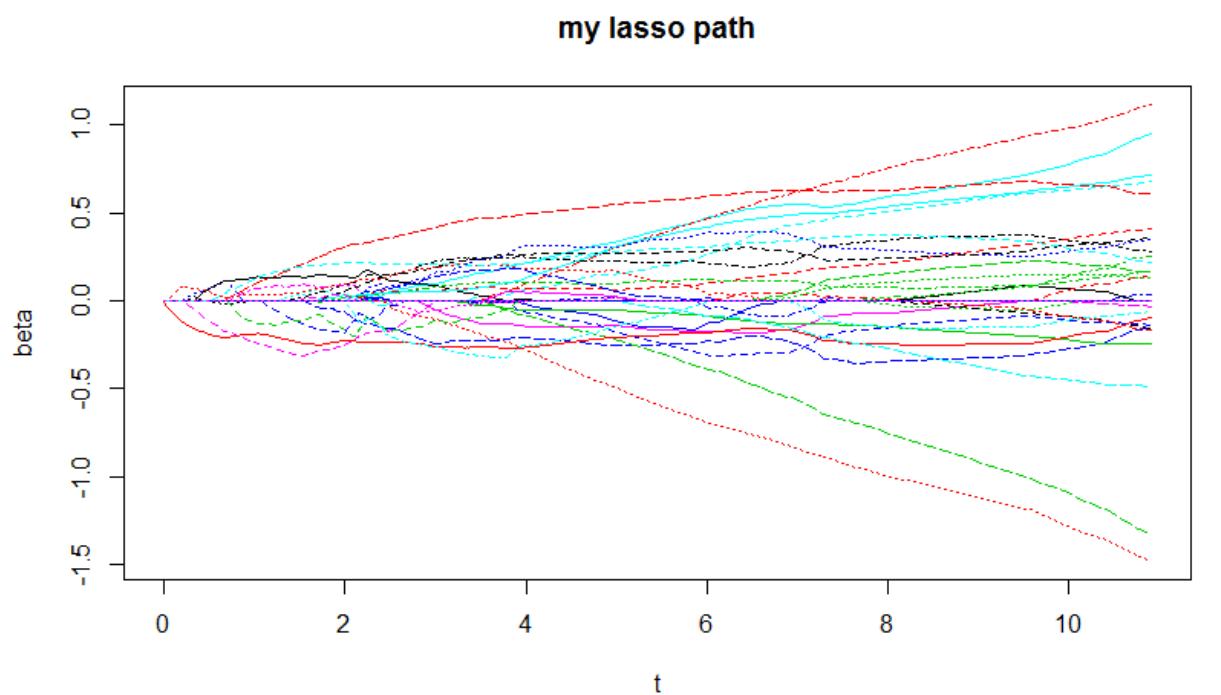


Figure 2 my lasso path

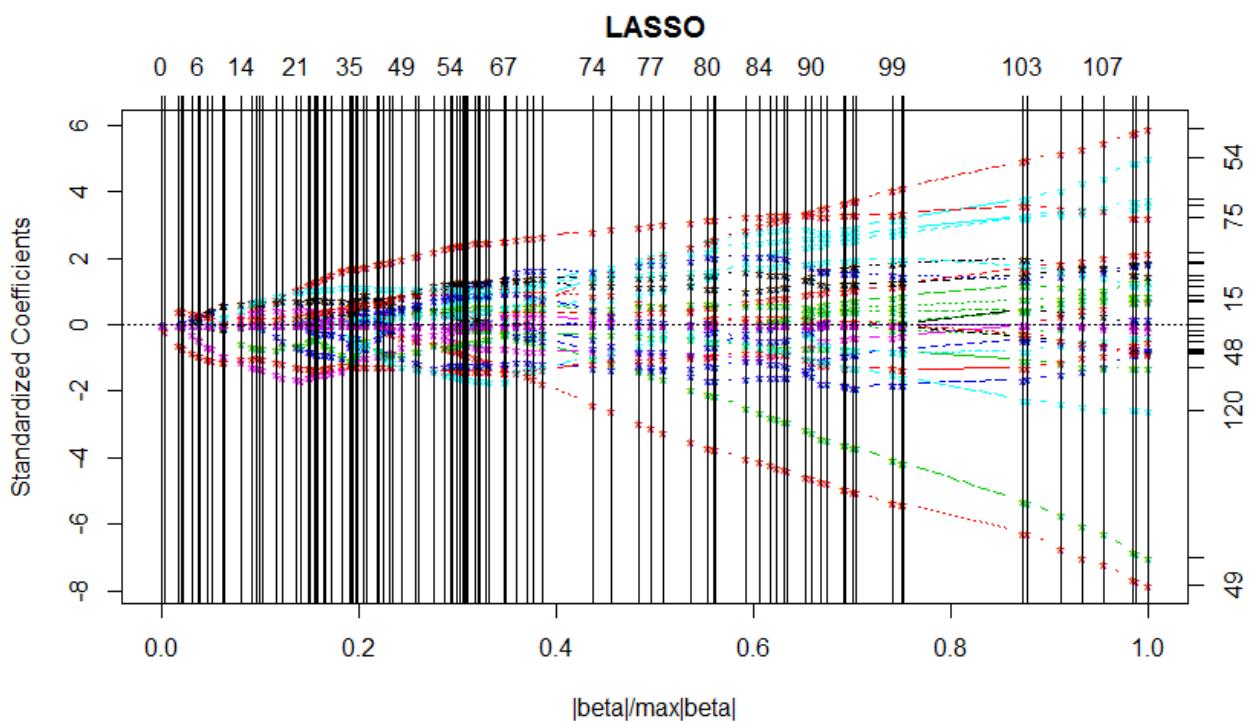


Figure 3 lasso path from lars

apparent mean absolute predictive error rate

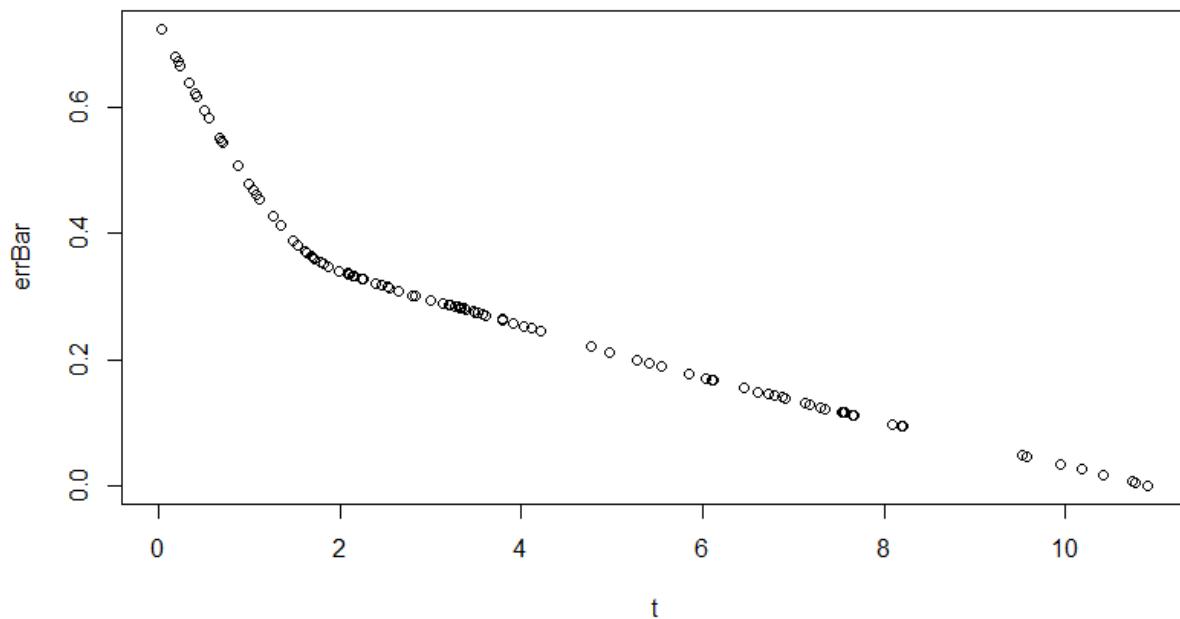


Figure 4 apparent mean absolute predictive error

no information error rate

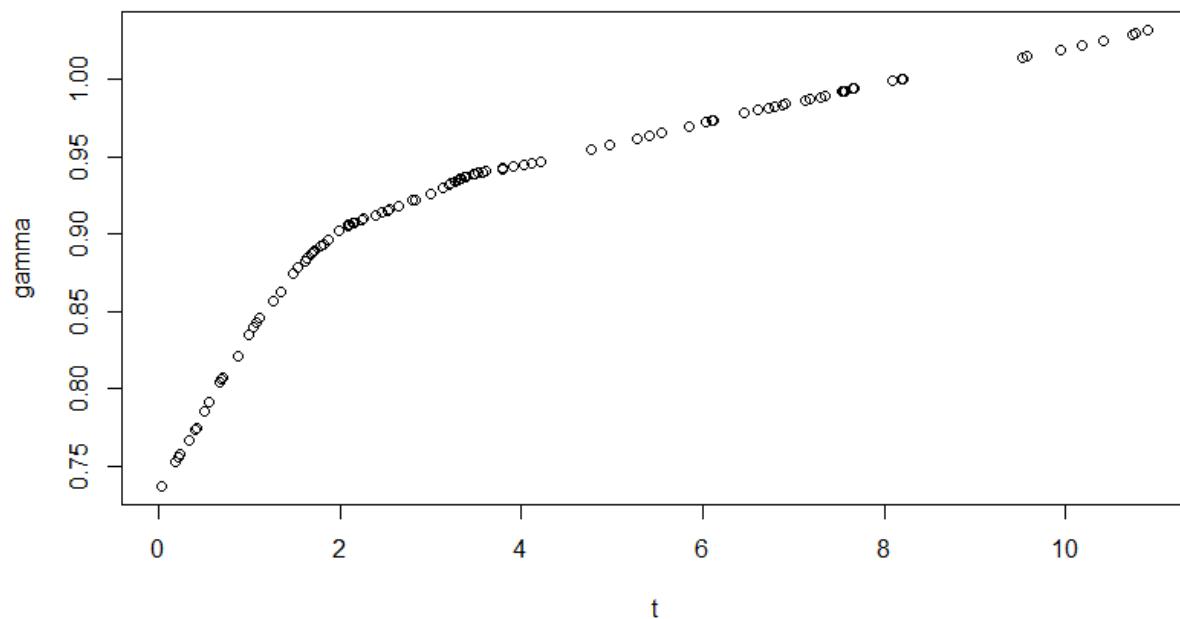


Figure 5 no information error

Lasso via Modification to Least Angle Regression

(a) Calculate the $\widehat{Err}^{(0.632+)}$ using \overline{err} , $\widehat{Err}^{(1)}$ and $\widehat{\gamma}$.

$$\widehat{Err}^{(0.632+)} = (1 - \widehat{w}) \times \overline{err} + \widehat{w} \times \widehat{Err}^{(1)}$$

where

$$\widehat{Err}^{(1)} = \frac{1}{n} \sum_{i=1}^n \frac{1}{|C^{-i}|} \sum_{b \in C^{-i}} Q(y_i, \widehat{\mu}^b(x_i))$$

$$Q(y_i, \widehat{\mu}^b(x_i)) = |(y_i - \overline{y^b}) - ((x_i - \overline{x^b}) / sd(x^b)) \times \boldsymbol{\beta}^b|$$

where $\overline{x^b}$ and $sd(x^b)$ is mean and standard deviation of x in the bootstrap b , and $\overline{y^b}$ is the mean of y in the bootstrap b .

(2) The average $\widehat{Err}^{(0.632+)}$ across three multiple imputation datasets is calculated, see Figure 6.

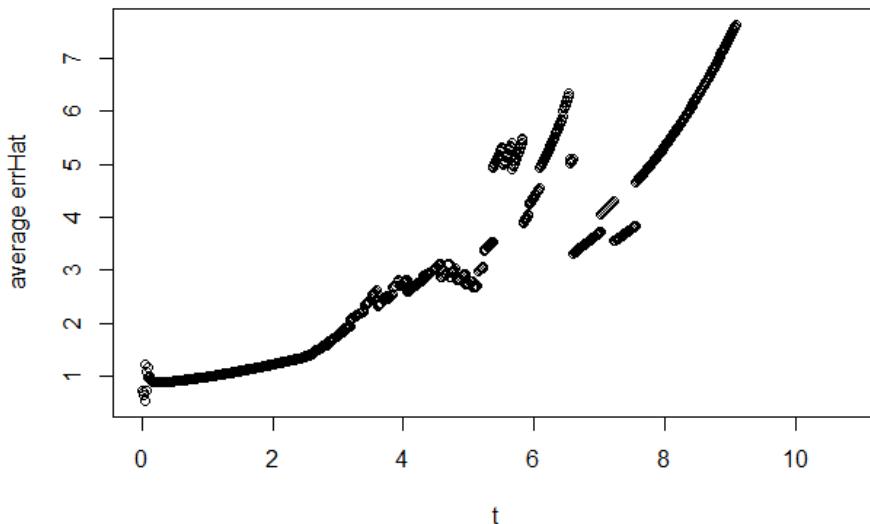


Figure 6 the average errHat

(3) Choose the best t value.

The best t is chosen as the one that minimize the average $\widehat{Err}^{(0.632+)}$. In this example, the best t value is chosen around 0.45, which corresponds to choosing around 7 predictors (depends on the MI dataset).

Problem 5 Visualization

(a) The lasso path and $\widehat{Err}^{(0.632+)}$ are calculated and plotted as the following figures:

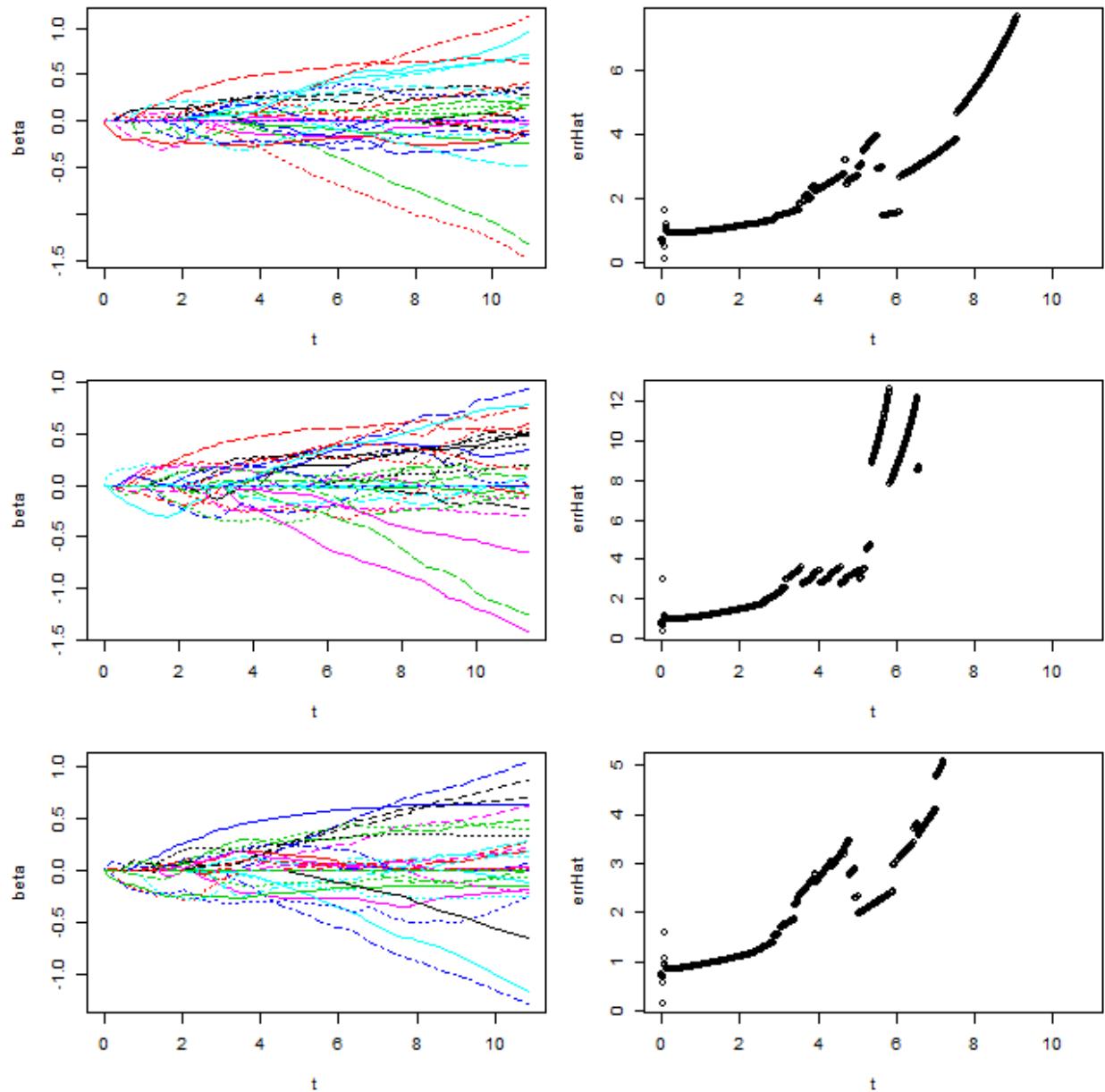


Figure 7 Lasso path and errHat from three MI datasets

(b) The average $\widehat{Err}^{(0.632+)}$ across three multiple imputation datasets is plotted in **Figure 6**.

Interpretation

In this example, we choose $t = 0.45$. The final model coefficients are calculated by averaging β from three multiple imputation datasets (β_1 , β_2 and β_3). In each multiple imputation model i , β_i is interpolated so that the t value is 0.45.

The final model is listed as follows:

| Predictor ID | Beta | Predictor Name |
|--------------|--------------|--|
| 55 | -0.125843705 | v1_Texture.Strength_average |
| 56 | -0.022846644 | v1_Maximum.Intensity_average |
| 75 | 0.01346343 | v1_High.Intensity.Large.Zone.Emphasis_average |
| 84 | 0.00146977 | v1_Sum.of.Squares.or.Variance_difference |
| 86 | 0.008499992 | v1_Sum.Variance_difference |
| 88 | 0.060127267 | v1_Difference.Variance_difference |
| 90 | -0.098805288 | v1_Information.Measure.of.Correlation1_difference |
| 96 | 0.018049183 | v1_Contrast.1_difference |
| 105 | 0.046300367 | v1_Sum.of.Squares.or.Variance.1_difference |
| 110 | -0.023923559 | v1_Information.Measure.of.Correlation1.1_difference |
| 114 | -0.062074634 | v1_Inverse.Difference.Moment.Normalized.1_difference |
| 127 | 0.03842584 | v1_Contrast.2_difference |
| 148 | 0.029746621 | v1_High.Intensity.Short.Zone.Emphasis_difference |

However, the above beta coefficients are based on several normalization processes. To begin with, we standardized the raw data. For example, the mean value of y (PFS) is 4.836. Then we calculate the baseline average and follow up difference as our model predictors. Again, we normalized the data before we could use lasso. Now, based on our normalized data, we ranked the importance of the predictors as follows:

| Beta | Predictor Name |
|---------|--|
| -0.1258 | v1_Texture.Strength_average |
| -0.0988 | v1_Information.Measure.of.Correlation1_difference |
| -0.0621 | v1_Inverse.Difference.Moment.Normalized.1_difference |
| 0.06013 | v1_Difference.Variance_difference |
| 0.0463 | v1_Sum.of.Squares.or.Variance.1_difference |
| 0.03843 | v1_Contrast.2_difference |
| 0.02975 | v1_High.Intensity.Short.Zone.Emphasis_difference |
| -0.0239 | v1_Information.Measure.of.Correlation1.1_difference |
| -0.0228 | v1_Maximum.Intensity_average |
| 0.01805 | v1_Contrast.1_difference |
| 0.01346 | v1_High.Intensity.Large.Zone.Emphasis_average |
| 0.0085 | v1_Sum.Variance_difference |
| 0.00147 | v1_Sum.of.Squares.or.Variance_difference |

If the coefficient is positive, the increase of the corresponding predictor will increase the PFS, vice versa. And the large the coefficient is the more correlated the predictor is with the PFS. For example, the predictor “v1_Texture.Strength_average” is negative. The increase of 1 unit of the predictor will lead the PFS to decrease 0.1258. Please note that the 1 unit of “v1_Texture.Strength_average” here is 1 unit after the normalization. To go back to the original value, we need to reverse our normalization process.

Appendix I Data Manipulation and Multiple Imputation Example

```
library(mi)
library(lars)

# Read in texture and response data
myXData=read.csv("Texture.csv",header=TRUE)
myYData=read.csv("TextureResponse.csv",header=TRUE)
# Restrict attention to particular variables
myLogical=sapply(attributes(myXData)$names,function(varname){
  if(varname %in% c("PatID","TimePoint","PFS")){
    return(TRUE)
  }else{
    return(((length(grep(pattern="v1_",varname))>0))&
      (!((length(grep(pattern="Xa",varname))>0)| 
        (length(grep(pattern="DeltaX",varname))>0))))
  }
})
myXData=myXData[,myLogical]

# Rearrange data
myXtemp=data.matrix(myXData[,6:dim(myXData)[2]])
patientIDsTemp=sort(unique(c(myXData$PatID,myYData$Pat.No)))
timePointsTemp=sort(unique(as.character(myXData$TimePoint)))
PFS=matrix(NA,length(patientIDsTemp),1)
predictors=matrix(NA,length(patientIDsTemp),(dim(myXtemp)[2])*length(timePointsTemp))
counter=1
for(pt in patientIDsTemp){
  index=which(myYData$Pat.No==pt)
  if(length(index)==1){
    PFS[counter]=log(myYData$PFS[index])
  }
  for(jj in 1:length(timePointsTemp)){
    index=which((myXData$TimePoint==timePointsTemp[jj])&
      (myXData$PatID==pt))
    if(length(index)==1){
      predictors[counter,(jj-1)*dim(myXtemp)[2]+1:(jj*dim(myXtemp)[2])]=
        myXtemp[index,]
    }
  }
  counter=counter+1
}
# normalize the predictors
predictorsStar=apply(predictors,2,function(xx){
  if(sum(!is.na(xx))>=2){
    return((xx-mean(xx,na.rm=TRUE))/sd(xx,na.rm=TRUE))
  }else{
    return(xx)
  }
})
```

```

# normalize the PFS mean = zero
myData=data.frame(PFS,predictorsStar)
myData$PFS=myData$PFS-mean(myData$PFS,na.rm=TRUE)

# Label columns
myNames=c("PFS",
         paste(attributes(myXData)$names[6:dim(myXData)[2]],"_baseline1",sep=""),
         paste(attributes(myXData)$names[6:dim(myXData)[2]],"_baseline2",sep=""),
         paste(attributes(myXData)$names[6:dim(myXData)[2]],"_followUp",sep=""))
attributes(myData)$names=myNames

#####
##### Data manipulation #####
#####

# Remove completely missing columns.
temp1 = logical(length = dim(myData)[2])
for(ii in 1:dim(myData)[2]){
  if(sum(abs(myData[,ii])>matrix(0.0001,dim(myData)[1],1), na.rm = TRUE)>0){
    temp1[ii] = TRUE
  }else{
    temp1[ii] = FALSE # missing columns
  }
}
myData1=myData[,temp1]

# Remove rows with completely missing predictors.
temp = logical(length = dim(myData1)[1])
for(ii in 1:dim(myData1)[1]){
  if(sum(abs(myData1[ii,2:dim(myData1)[2]])>matrix(0.0001,1,dim(myData1)[2]), na.rm = TRUE)>0){
    temp[ii] = TRUE
  }else{
    temp[ii] = FALSE # missing columns
  }
}
myData2=myData1[temp,]

# Remove columns with constant predictors.
temp2 = logical(length = dim(myData2)[2])
tempData = myData2
tempData[is.na(tempData)] = 0
for(ii in 1:dim(myData2)[2]){
  if(abs(abs(max(tempData[,ii]))+abs(min(tempData[,ii])))>0.0001){
    temp2[ii] = TRUE
  }else{
    temp2[ii] = FALSE # missing columns
  }
}
myData3=myData2[,temp2]

# Remove (second or later occurrence of) columns which are collinear (correlation 1) with another.
varmatrix = var(myData3[,-1], use ="pairwise.complete.obs")

```

```

temp = matrix(data = FALSE, nrow = dim(myData3)[2]-1, ncol = dim(myData3)[2]-1)
for(ii in 1:(dim(myData3)[2]-1)){
  for(jj in ii:(dim(myData3)[2]-1)){
    if (varmatrix[ii,jj]>0.9999&ii!=jj){
      temp[ii,jj] = TRUE
    }
  }
}
temp3 = colSums(temp)
temp3 = temp3==0
# sum(temp2)
temp3 = append(temp3,1,after = 0)
myData4 = myData3[,as.logical(temp3)]

#####
##### MI Algorithm #####
#####

myData = myData4
myInfo=mi.info(myData)
myInfo=update(myInfo,"type",as.list(rep("predictive-mean-matching",dim(myData)[2])))
myMIs=mi(myData,myInfo)
misStar=mi.completed(myMIs)

#DeletedNames = myNames[!temp1]
#temp = myNames[temp1]
#DeletedNames = rbind(as.matrix(DeletedNames),as.matrix(temp[as.logical(!temp3)]))

sampleNames = attributes(myXData)$names[6:dim(myXData)[2]]
temp = matrix(NA,dim(myData4)[1],dim(as.matrix(sampleNames))[1])
tempNames=paste(sampleNames,"_average",sep="")
attributes(temp)$names=tempNames
averageData = list(temp,temp,temp)
tempNames=paste(sampleNames,"_difference",sep="")
attributes(temp)$names=tempNames
differenceData = list(temp,temp,temp)

# compute the average and difference data
for(kk in 1:3){
  baseLine1Data = matrix(NA,dim(misStar[[1]])[1],dim(as.matrix(sampleNames))[1])
  baseLine2Data = matrix(NA,dim(misStar[[1]])[1],dim(as.matrix(sampleNames))[1])
  followUpData = matrix(NA,dim(misStar[[1]])[1],dim(as.matrix(sampleNames))[1])
  for(ii in 1:dim(as.matrix(sampleNames))[1]){
    if (length(grep(pattern = paste(sampleNames[ii],"_baseline1",sep=""),attributes(misStar[[1]])$names))!=0){
      baseLine1Data[,ii]=misStar[[kk]][grep(pattern =
      paste(sampleNames[ii],"_baseline1",sep=""),attributes(misStar[[1]])$names)]
    }
    if (length(grep(pattern = paste(sampleNames[ii],"_baseline2",sep=""),attributes(misStar[[1]])$names))!=0){
      baseLine2Data[,ii]=misStar[[kk]][grep(pattern =
      paste(sampleNames[ii],"_baseline2",sep=""),attributes(misStar[[1]])$names)]
    }
    if (length(grep(pattern = paste(sampleNames[ii],"_followUp",sep=""),attributes(misStar[[1]])$names))!=0){

```

```
followUpData[,ii]=misStar[[kk]][,grep(pattern = paste(sampleNames[ii],"_followUp",sep=""),attributes(misStar[[1]])$names)]
}
}
# compute the average value
for(ii in 1:dim(as.matrix(sampleNames))[1]){
  averageData[[kk]][,ii] = as.matrix(rowMeans(cbind(baseLine1Data[,ii],baseLine2Data[,ii])),na.rm = TRUE))
}
# compute the difference
differenceData[[kk]] = followUpData-averageData[[kk]]
}
```

Appendix II Lars Example

```

# Initialization
k = 0
A = matrix(0,dim(myX)[2],dim(myX)[2])
Ac = matrix(1,dim(myX)[2],dim(myX)[2])
beta = matrix(0,dim(myX)[2],dim(myX)[2])
XA = matrix(0,dim(myX)[1],k)
cBar = matrix(0,dim(myX)[2],dim(myX)[2])
u = matrix(0,dim(myX)[1],dim(myX)[2]+1)
betaA = matrix(0,dim(myX)[1],k)
#{min(dim(myX)[1]-1,dim(myX)[2])-1}
for(k in 1:(min(dim(myX)[1]-1,dim(myX)[2]-1))){
  # update cBar (for k-1)
  cBar[k,] = t(myX) * %*% (myY - u[,k])
  # find cBar for Ac elements
  cBarc = cBar[k,]
  cBarc[as.logical(A[k,])] = 0
  # update XA
  cat(which.max(abs(cBarc)),sep=" ",fill=TRUE)
  A[k:dim(myX)[2],which.max(abs(cBarc))] = 1
  Ac[k:dim(myX)[2],which.max(abs(cBarc))] = 0
  s = sign(cBar[k,as.logical(A[k,])])
  XA = myX[,as.logical(A[k,])]
  XA = XA * %diag(as.vector(s),length(s))
  # update cBar
  CBar = mean(abs(t(XA) * %*% (myY - u[,k])))
  # cat(abs(t(XA) * %*% (myY - u[,k])),sep=" ",fill=TRUE)
  # update angles
  angles =
  t(myX) * %*% XA * %*% solve(t(XA) * %*% XA,matrix(1,sum(A[k,]),
  1))
  temp = matrix(NA,dim(myX)[2],3)
  temp[,1] = (CBar - cBar[k,]) / (1 - angles)
  temp[,2] = (CBar + cBar[k,]) / (1 + angles)
  temp[as.logical(temp[,1] < 0),1] = +Inf
  temp[as.logical(temp[,2] < 0),2] = +Inf
  temp[,3] = pmin(temp[,1],temp[,2])
  temp[as.logical(A[k,]),3] = +Inf
  c = min(temp[,3])
  # update u
  u[,k+1] =
  c * XA * %*% solve(t(XA) * %*% XA) * %*% matrix(1,sum(A[,k]),1) + u[,k]
  #update betaA
  b = c * solve(t(XA) * %*% XA) * %*% matrix(1,sum(A[,k]),1)
  j = sum(A[,1:which.max(abs(cBarc))])
  betaA = betaA + b[-j]
  betaA = append(betaA,b[j],after=j-1)
  beta[k,as.logical(A[k,])] = betaA
}

# ordinary least square solution
s = sign(cBar[k,as.logical(A[k,])])
betaBar = solve(t(XA) * %*% XA) * %*% t(XA) * %*% (myY)
betaBarSign = s * betaBar

# caned function
object = lars(myX, myY, type = c("lars"))
# caned function dim(beta)[1]=dim(X)[1]+1
caned_beta = unique(object$beta[29,])[-1]
diff = cbind(betaBarSign, as.matrix(caned_beta))
cbind(unique(s * beta[,as.logical(A[,])]), caned_beta, betaBarSign)

```

Appendix III Lasso Example

```
#####
##### Lasso #####
#####

# Initialization
k = 0
A = matrix(0,dim(myX)[2],dim(myX)[2])
Ac = matrix(1,dim(myX)[2],dim(myX)[2])
beta = matrix(0,dim(myX)[2],dim(myX)[2])
XA = matrix(0,dim(myX)[1],k)
cBar = matrix(0,dim(myX)[2],dim(myX)[2])
u = matrix(0,dim(myX)[1],dim(myX)[2]+1)
betaA = matrix(0,dim(myX)[1],k)
jj = 0
action = matrix(0,dim(myX)[2],1)
numErr = 10^-12

S = matrix(0,dim(myX)[2],dim(myX)[2]) # record the sign
# run lasso
for(k in 1:dim(myX)[2]){
  # if jj needs to leave the active variable set XA
  if(jj!=0){
    cat(-jj,sep=" ",fill=TRUE)
    action[k,] = -jj
    # exclude jj from the active variable set XA
    A[k:dim(myX)[2],jj] = 0
    Ac[k:dim(myX)[2],jj] = 1
    cBar[k,] = t(myX)%%(myY-u[,k]) # which one to use u[,k] or u[,k-1]
    s = sign(cBar[,as.logical(A[,k])])
    XA = myX[,as.logical(A[,k])]
    XA = XA%*%diag(as.vector(s),length(s))
    # update cBar
    CBar = mean(abs(t(XA)%*%(myY-u[,k])))
    # cat(abs(t(XA)%*%(myY-u[,k])),sep=" ",fill=TRUE)
    # update angles
    angles = t(myX)%*%XA%*%solve(t(XA)%*%XA,matrix(1,sum(A[,k]),1))
    temp = matrix(NA,dim(myX)[2],3)
    temp[,1] = (CBar-cBar[,k,])/ (1-angles)
    temp[,2] = (CBar+cBar[,k,])/ (1+angles)
    temp[as.logical(temp[,1]<0),1] = +Inf
    temp[as.logical(temp[,2]<0),2] = +Inf
    temp[,3] = pmin(temp[,1],temp[,2])
    temp[as.logical(A[,k]),3] = +Inf
    c = min(temp[temp[,3]>numErr,3])
    # Lasso Modification
    if(k>1){
      bBar = solve(t(XA)%*%XA)%*%matrix(1,sum(A[,k]),1)
      d = matrix(0,dim(myX)[2],1)
      # d[as.logical(A[,k]),1]=s*bBar
      d[as.logical(A[,k]),1]=bBar
      betaOld = matrix(0,dim(myX)[2],1)
```

```

# fix betaA delete the zero elements
betaA = betaA[-sum(A[k-1,1:jj])]
betaOld[as.logical(A[k,]),1]=betaA
cLasso = matrix(0,dim(myX)[2],1)
cLasso[as.logical(A[k,]),1] = -betaOld[as.logical(A[k,]),1]/d[as.logical(A[k,]),1]
if(sum(cLasso>0)==0){
  cStar = Inf
}else{
  cStar = min(cLasso[cLasso>0])
}
if(cStar<c){
  c = cStar
  jj = which(cLasso==cStar) # the predictor that should leave the active variable set
}else{
  jj = 0
}
#
# update u
u[,k+1] = c*XA%*%solve(t(XA)%*%XA)%*%matrix(1,sum(A[k,]),1)+u[,k]
#update betaA
b = c*solve(t(XA)%*%XA)%*%matrix(1,sum(A[k,]),1)
betaA = betaA+b
beta[k,as.logical(A[k,])] = betaA
}else{
  # update cBar (for k-1)
  cBar[k,] = t(myX)%*%(myY-u[,k])
  # find cBar for Ac elements: cBarc
  cBarc = cBar[k,]
  cBarc[as.logical(A[k,])] = 0
  # update XA
  cat(which.max(abs(cBarc)),sep=" ",fill=TRUE)
  action[k,] = which.max(abs(cBarc))
  A[k:dim(myX)[2],which.max(abs(cBarc))] = 1
  Ac[k:dim(myX)[2],which.max(abs(cBarc))] = 0
  s = sign(cBarc[k,as.logical(A[k,])])
  XA = myX[,as.logical(A[k,])]
  XA = XA%*%diag(as.vector(s),length(s))
  # update cBar
  CBar = mean(abs(t(XA)%*%(myY-u[,k])))
  # cat(abs(t(XA)%*%(myY-u[,k])),sep=" ",fill=TRUE)
  # update angles
  angles = t(myX)%*%XA%*%solve(t(XA)%*%XA,matrix(1,sum(A[k,]),1))
  temp = matrix(NA,dim(myX)[2],3)
  temp[,1] = (CBar-cBarc[k,])/(1-angles)
  temp[,2] = (CBar+cBarc[k,])/(1+angles)
  temp[as.logical(temp[,1]<0),1] = +Inf
  temp[as.logical(temp[,2]<0),2] = +Inf
  temp[,3] = pmin(temp[,1],temp[,2])
  temp[as.logical(A[k,]),3] = +Inf
  c = min(temp[temp[,3]>numErr,3])
  # Lasso Modification
}

```

```

if(k>1){
  bBar = solve(t(XA) %*% XA) %*% matrix(1,sum(A[,]),1)
  d = matrix(0,dim(myX)[2],1)
  # d[as.logical(A[,]),1]=s*bBar
  d[as.logical(A[,]),1]=bBar
  betaOld = matrix(0,dim(myX)[2],1)
  betaOld[as.logical(A[,]),1]=betaA

  ## something is wrong here
  cLasso = matrix(0,dim(myX)[2],1)
  cLasso[as.logical(A[,]),1] = -betaOld[as.logical(A[,]),1]/d[as.logical(A[,]),1]
  if(sum(cLasso>0)==0){
    cStar = Inf
  }else{
    cStar = min(cLasso[cLasso>0])
  }
  if(cStar<c){
    c = cStar
    jj = which(cLasso==cStar) # the predictor that should leave the active variable set
  }else{
    jj = 0
  }
  # update u
  u[,k+1] = c*XA%*%solve(t(XA)%*%XA)%*%matrix(1,sum(A[,]),1)+u[,k]
  #update betaA
  b = c*solve(t(XA)%*%XA)%*%matrix(1,sum(A[,]),1)
  j = sum(A[,1:which.max(abs(cBar))])
  betaA = betaA+b[-j]
  betaA = append(betaA,b[jj],after=j-1)
  beta[,as.logical(A[,])]=betaA
}
# collect AMAPER and t and NIER
S[,] = sign(cBar[,])
# stop criteria
if (dim(XA)[2]==(min(dim(myX)[1]-1,dim(myX)[2]-1)) & jj==0){
  break
}
}

```

Appendix IV Bootstrap Example

```

KK = 1
myX = cbind(averageData[[KK]],differenceData[[KK]])
myXNames = rbind(as.matrix(paste(sampleNames, "_average",sep="")),as.matrix(paste(sampleNames, "_difference",sep="")))
##### attributes(myX)$names=myXNames
myX = myX[,colSums(is.na(myX))==0]
myPFS = misStar[[KK]][,1]
# check colinearity
# myXInfo=mi.info(myX)
# normalize the predictors
myX=apply(myX,2,function(xx){
  return((xx-mean(xx,na.rm=TRUE))/sd(xx,na.rm=TRUE))
})
# normalize the PFS mean = zero
myY=myPFS-mean(myPFS)

#####
# run lasso for the full data to get errBar and gamma #####
# Initialization
k = 0
A = matrix(0,dim(myX)[2],dim(myX)[2])
Ac = matrix(1,dim(myX)[2],dim(myX)[2])
beta = matrix(0,dim(myX)[2],dim(myX)[2])
XA = matrix(0,dim(myX)[1],k)
cBar = matrix(0,dim(myX)[2],dim(myX)[2])
u = matrix(0,dim(myX)[1],dim(myX)[2]+1)
betaA = matrix(0,dim(myX)[1],k)
jj = 0
action = matrix(0,dim(myX)[2],1)
numErr = 10^-12

S = matrix(0,dim(myX)[2],dim(myX)[2]) # record the sign
# run lasso
for(k in 1:dim(myX)[2]){
  # if jj needs to leave the active variable set XA
  if(jj!=0){
    cat(-jj,sep=" ",fill=TRUE)
    action[k,] = -jj
    # exclude jj from the active variable set XA
    A[k:dim(myX)[2],jj] = 0
    Ac[k:dim(myX)[2],jj] = 1
    cBar[k,] = t(myX)%*%(myY-u[,k]) # which one to use u[,k] or u[,k-1]
    s = sign(cBar[,as.logical(A[,k])])
    XA = myX[,as.logical(A[,k])]
    XA = XA%*%diag(as.vector(s),length(s))
    # update cBar
    CBar = mean(abs(t(XA)%*%(myY-u[,k])))
    # cat(abs(t(XA)%*%(myY-u[,k])),sep=" ",fill=TRUE)
    # update angles
    angles = t(myX)%*%XA%*%solve(t(XA)%*%XA,matrix(1,sum(A[,k]),1))
  }
}

```

```

temp = matrix(NA,dim(myX)[2],3)
temp[,1] = (CBar-cBar[k,])/(1-angles)
temp[,2] = (CBar+cBar[k,])/ (1+angles)
temp[as.logical(temp[,1]<0),1] = +Inf
temp[as.logical(temp[,2]<0),2] = +Inf
temp[,3] = pmin(temp[,1],temp[,2])
temp[as.logical(A[k,]),3] = +Inf
c = min(temp[temp[,3]>numErr,3])
# Lasso Modification
if(k>1){
  bBar = solve(t(XA) %*% XA) %*% matrix(1,sum(A[k,]),1)
  d = matrix(0,dim(myX)[2],1)
  # d[as.logical(A[k,]),1]=s*bBar
  d[as.logical(A[k,]),1]=bBar
  betaOld = matrix(0,dim(myX)[2],1)
  # fix betaA delete the zero elements
  betaA = betaA[-sum(A[k-1,1:jj])]
  betaOld[as.logical(A[k,]),1]=betaA
  cLasso = matrix(0,dim(myX)[2],1)
  cLasso[as.logical(A[k,]),1] = -betaOld[as.logical(A[k,]),1]/d[as.logical(A[k,]),1]
  if(sum(cLasso>0)==0){
    cStar = Inf
  }else{
    cStar = min(cLasso[cLasso>0])
  }
  if(cStar<c){
    c = cStar
    jj = which(cLasso==cStar) # the predictor that should leave the active variable set
  }else{
    jj = 0
  }
}
# update u
u[,k+1] = c*XA%*%solve(t(XA)%*%XA)%*%matrix(1,sum(A[k,]),1)+u[,k]
#update betaA
b = c*solve(t(XA) %*% XA) %*% matrix(1,sum(A[k,]),1)
betaA = betaA+b
beta[k,as.logical(A[k,])] = betaA
}else{
  # update cBar (for k-1)
  cBar[k,] = t(myX) %*% (myY-u[,k])
  # find cBar for Ac elements: cBarc
  cBarc = cBar[k,]
  cBarc[as.logical(A[k,])] = 0
  # update XA
  cat(which.max(abs(cBarc)),sep=" ",fill=TRUE)
  action[k,] = which.max(abs(cBarc))
  A[k:dim(myX)[2],which.max(abs(cBarc))] = 1
  Ac[k:dim(myX)[2],which.max(abs(cBarc))] = 0
  s = sign(cBarc[k,as.logical(A[k,])])
  XA = myX[,as.logical(A[k,])]
```

```

XA = XA%*%diag(as.vector(s),length(s))
# update cBar
CBar = mean(abs(t(XA)%*%(myY-u[,k])))
# cat(abs(t(XA)%*%(myY-u[,k])),sep=" ",fill=TRUE)
# update angles
angles = t(myX)%*%XA%*%solve(t(XA)%*%XA,matrix(1,sum(A[,]),1))
temp = matrix(NA,dim(myX)[2],3)
temp[,1] = (CBar-cBar[,k])/(1-angles)
temp[,2] = (CBar+cBar[,k])/(1+angles)
temp[as.logical(temp[,1]<0),1] = +Inf
temp[as.logical(temp[,2]<0),2] = +Inf
temp[,3] = pmin(temp[,1],temp[,2])
temp[as.logical(A[,]),3] = +Inf
c = min(temp[temp[,3]>numErr,3])
# Lasso Modification
if(k>1){
  bBar = solve(t(XA)%*%XA)%*%matrix(1,sum(A[,]),1)
  d = matrix(0,dim(myX)[2],1)
  # d[as.logical(A[,]),1]=s*bBar
  d[as.logical(A[,]),1]=bBar
  betaOld = matrix(0,dim(myX)[2],1)
  betaOld[as.logical(A[k-1]),1]=betaA

  ## something is wrong here
  cLasso = matrix(0,dim(myX)[2],1)
  cLasso[as.logical(A[,]),1] = -betaOld[as.logical(A[,]),1]/d[as.logical(A[,]),1]
  if(sum(cLasso>0)==0){
    cStar = Inf
  }else{
    cStar = min(cLasso[cLasso>0])
  }
  if(cStar<c){
    c = cStar
    jj = which(cLasso==cStar) # the predictor that should leave the active variable set
  }else{
    jj = 0
  }
}
# update u
u[,k+1] = c*XA%*%solve(t(XA)%*%XA)%*%matrix(1,sum(A[,]),1)+u[,k]
#update betaA
b = c*solve(t(XA)%*%XA)%*%matrix(1,sum(A[,]),1)
j = sum(A[,1:which.max(abs(cBar[,]))])
betaA = betaA+b[-j]
betaA = append(betaA,b[jj],after=j-1)
beta[,as.logical(A[,])]=betaA
}

# collect AMAPER and t and NIER
S[,k] = sign(cBar[,k])
# stop criteria
if (dim(XA)[2]==(min(dim(myX)[1]-1,dim(myX)[2]-1)) & jj==0){

```

```

break
}
}
# plot
errBar = matrix(NA,dim(myX)[2],1) # apparent mean absolute predictive error rate
gamma = matrix(NA,dim(myX)[2],1) # no information error rate
tAll = matrix(NA,dim(myX)[2],1)
for(kk in 1:k){
  tAll[kk,1] = sum(abs(beta[kk,]))
  XA = myX[,as.logical(A[kk,])]
  XA = XA%*%diag(as.vector(S[kk,as.logical(A[kk,])]),length(S[kk,as.logical(A[kk,])]))
  errBar[kk,1] = sum(abs(myY-XA%*%beta[kk,as.logical(A[kk,])]))/(dim(myX)[1])
  gamma[kk,1] = sum(abs(matrix(1,dim(myX)[1],1)%*%myY-
XA%*%as.matrix(beta[kk,as.logical(A[kk,])])%*%matrix(1,1,dim(myX)[1])))/(dim(myX)[1])^2
}

plot(tAll[1:k],xlab = "Iter.",ylab = "t")
title(main = "sum of absolute beta Vs. iteration")

plot(tAll[1:k],errBar[1:k],xlab = "t",ylab = "errBar")
title(main = "apparent mean absolute predictive error rate")

plot(tAll[1:k],gamma[1:k],xlab = "t",ylab = "gamma")
title(main = "no information error rate")

betaIndex = colSums(beta[1:k,])!=0
betaPlot = beta[1:k,betaIndex]*S[1:k,betaIndex]
matplot(rbind(0,as.matrix(tAll[1:k])),rbind(matrix(0,1,dim(betaPlot)[2]),betaPlot),type = "l", xlab ="t", ylab="beta")
title(main = "my lasso path")

#####
##### bootstrap algorithm #####
#####

# Perform Bootstrap Sampling
B = 25 # number of bootstraps
n = dim(myX)[1] # number of observations
p = dim(myX)[2] # number of predictors
Bl = matrix(sample.int(n,n*B,replace = TRUE),B,n)
UniBl = matrix(0,B,1)
myXB = array(0,dim=c(B,n,p))
myYB = array(0,dim=c(B,n))
Blc = matrix(0,B,n) # the unselected elements are 1 for each bootstrap
for (i in 1:B){
  temp = 1:29
  Blc[i,temp[!(1:29 %in% Bl[i,])]]=1
}
for(i in 1:B){
  myXB[i,,] = myX[Bl[i,],]
  myYB[i,] = myY[Bl[i,]]
  UniBl[i,] = length(unique(Bl[i,]))
}

```

```

# scale the data so that mean(y)=0, mean(x)=0 and var(x)=1
myYBMean = matrix(0,B,1)
myXBsd = matrix(0,B,p)
myXBMean = matrix(0,B,p)
for(i in 1:B){
  # mean(y)=0
  myYBMean[i,] = mean(myYB[i,])
  myYB[i,] = myYB[i,]-mean(myYB[i,])
  # mean(x)=0 and var(x)=1
  for (j in 1:p){
    myXBMean[i,j] = mean(myXB[i,,j])
    myXBsd[i,j] = sd(myXB[i,,j])
    myXB[i,,j] = (myXB[i,,j]-mean(myXB[i,,j]))/sd(myXB[i,,j])
  }
}
# run lesso for each bootstrap sample
betaB = array(NA,dim = c(B,p,p))
SB = array(NA,dim = c(B,p,p))
AcB = array(1,dim = c(B,p,p))
betaB = array(1,dim = c(B,p,p))
cBarB = array(1,dim = c(B,p,p))
tB = array(NA,dim = c(B,p))
errHatB = array(NA,dim = c(B,p,n))

for(bb in 1:B){
  cat(bb,sep=" ",fill=TRUE)
  ##### run lasso #####
  # Initialization
  k = 0
  A = matrix(0,p,p)
  Ac = matrix(1,p,p)
  beta = matrix(0,p,p)
  XA = matrix(0,n,k)
  cBar = matrix(0,p,p)
  u = matrix(0,n,p+1)
  betaA = matrix(0,n,k)
  jj = 0
  action = matrix(0,p,1)
  numErr = 10^-12
  S = matrix(0,p,p) # record the sign
  # run lasso
  for(k in 1:p){
    # if jj needs to leave the active variable set XA
    if(jj!=0){
      # cat(-jj,sep=" ",fill=TRUE)
      action[k,] = -jj
      # exclude jj from the active variable set XA
      A[k:p,jj] = 0
      Ac[k:p,jj] = 1
      cBar[k,] = t(myXB[bb,,])%*%(myYB[bb,]-u[,k]) # which one to use u[,k] or u[,k-1]
      s = sign(cBar[k,as.logical(A[k,])])
    }
  }
}

```

```

XA = myXB[bb,,][,as.logical(A[,k])]
XA = XA%*%diag(as.vector(s),length(s))
# update cBar
CBar = mean(abs(t(XA)%*%(myYB[bb,]-u[,k])))
# cat(abs(t(XA)%*%(myYB[bb,]-u[,k])),sep=" ",fill=TRUE)
# update angles
angles = t(myXB[bb,,])%*%XA%*%solve(t(XA)%*%XA,matrix(1,sum(A[,k]),1))
temp = matrix(NA,p,3)
temp[,1] = (CBar-cBar[,k])/(1-angles)
temp[,2] = (CBar+cBar[,k])/(1+angles)
temp[as.logical(temp[,1]<0),1] = +Inf
temp[as.logical(temp[,2]<0),2] = +Inf
temp[,3] = pmin(temp[,1],temp[,2])
temp[as.logical(A[,k]),3] = +Inf
c = min(temp[temp[,3]>numErr,3])
# Lasso Modification
if(k>1){
  bBar = solve(t(XA)%*%XA)%*%matrix(1,sum(A[,k]),1)
  d = matrix(0,p,1)
  # d[as.logical(A[,k]),1]=s*bBar
  d[as.logical(A[,k]),1]=bBar
  betaOld = matrix(0,p,1)
  # fix betaA delete the zero elements
  betaA = betaA[-sum(A[-k-1,1:jj])]
  betaOld[as.logical(A[,k]),1]=betaA
  cLasso = matrix(0,p,1)
  cLasso[as.logical(A[,k]),1] = -betaOld[as.logical(A[,k]),1]/d[as.logical(A[,k]),1]
  if(sum(cLasso>0)==0){
    cStar = Inf
  }else{
    cStar = min(cLasso[cLasso>0])
  }
  if(cStar<c){
    c = cStar
    jj = which(cLasso==cStar) # the predictor that should leave the active variable set
  }else{
    jj = 0
  }
  # update u
  u[,k+1] = c*XA%*%solve(t(XA)%*%XA)%*%matrix(1,sum(A[,k]),1)+u[,k]
  #update betaA
  b = c*solve(t(XA)%*%XA)%*%matrix(1,sum(A[,k]),1)
  betaA = betaA+b
  beta[k,as.logical(A[,k])] = betaA
}else{
  # update cBar (for k-1)
  cBar[,k] = t(myXB[bb,,])%*%(myYB[bb,]-u[,k])
  # find cBar for Ac elements: cBarc
  cBarc = cBar[,k]
  cBarc[as.logical(A[,k])] = 0
}

```

```

# update XA
# cat(which.max(abs(cBarc)),sep=" ",fill=TRUE)
action[k,] = which.max(abs(cBarc))
A[k:p,which.max(abs(cBarc))] = 1
Ac[k:p,which.max(abs(cBarc))] = 0
s = sign(cBar[k,as.logical(A[k,])])
XA = myXB[bb,,][,as.logical(A[k,])]
XA = XA%*%diag(as.vector(s),length(s))
# update cBar
CBar = mean(abs(t(XA)%*%(myYB[bb,]-u[,k])))
# cat(abs(t(XA)%*%(myYB[bb,]-u[,k])),sep=" ",fill=TRUE)
# update angles
angles = t(myXB[bb,,])%*%XA%*%solve(t(XA)%*%XA,matrix(1,sum(A[,]),1))
temp = matrix(NA,p,3)
temp[,1] = (CBar-cBar[k,])/(1-angles)
temp[,2] = (CBar+cBar[k,])/(1+angles)
temp[as.logical(temp[,1]<0),1] = +Inf
temp[as.logical(temp[,2]<0),2] = +Inf
temp[,3] = pmin(temp[,1],temp[,2])
temp[as.logical(A[,]),3] = +Inf
c = min(temp[temp[,3]>numErr,3])
# Lasso Modification
if(k>1){
  bBar = solve(t(XA)%*%XA)%*%matrix(1,sum(A[,]),1)
  d = matrix(0,p,1)
  # d[as.logical(A[,]),1]=s*bBar
  d[as.logical(A[,]),1]=bBar
  betaOld = matrix(0,p,1)
  betaOld[as.logical(A[k-1,]),1]=betaA

  ## something is wrong here
  cLasso = matrix(0,p,1)
  cLasso[as.logical(A[,]),1] = -betaOld[as.logical(A[,]),1]/d[as.logical(A[,]),1]
  if(sum(cLasso>0)==0){
    cStar = Inf
  }else{
    cStar = min(cLasso[cLasso>0])
  }
  if(cStar<c){
    c = cStar
    jj = which(cLasso==cStar) # the predictor that should leave the active variable set
  }else{
    jj = 0
  }
}
# update u
u[,k+1] = c*XA%*%solve(t(XA)%*%XA)%*%matrix(1,sum(A[,]),1)+u[,k]
#update betaA
b = c*solve(t(XA)%*%XA)%*%matrix(1,sum(A[,]),1)
j = sum(A[,1:which.max(abs(cBarc))])
betaA = betaA+b[-j]

```

```

betaA = append(betaA,b[j],after=j-1)
beta[k,as.logical(A[k,])] = betaA
}
S[k,] = sign(cBar[k,])
# stop criteria
if (dim(XA)[2]==UniBl[bb,]-1 & jj==0){
  break
}
}
# calculate indices
errHat = matrix(NA,p,n) # leave-one-out bootstrap error rate
# calculate errHat: leave-one-out bootstrap error rate
for(kk in 1:k){
  errHat[kk,as.logical(Blc[bb,])] = abs(((myY[as.logical(Blc[bb,])] - myYBMean[bb,1]) - as.matrix(((myX[as.logical(Blc[bb,])],-
  myXBMean[bb,])/myXBsd[bb,])[as.logical(A[kk,])]))%*%(S[kk,as.logical(A[kk,])] * beta[kk,as.logical(A[kk,])]))
}
tB[bb,k] = sum(abs(beta[kk,]))
errHatB[bb,,] = errHat
}

# Find the tmax and set the tGrid
tGridR = 1000
tGrid = seq(from = 0, to = max(max(tB,na.rm = TRUE),tAll,na.rm = TRUE), length.out = tGridR)
tBPlot = array(NA,dim = c(B,tGridR))
errBarPlot = array(NA, dim = c(tGridR))
gammaPlot = array(NA, dim = c(tGridR))
errHatBPlot = array(NA,dim = c(B,tGridR,n))
# calculate

# plot errBar
# adding values for t = 0
errBarPlot =
matrix(unlist(approx(rbind(0,as.matrix(tAll)),rbind(sum(abs(myY))/n,as.matrix(errBar)),tGrid,method="linear",rule=1)$y))
# plot(tGrid, errBarPlot)

# plot gamma
# adding values for t = 0
gammaPlot =
matrix(unlist(approx(rbind(0,as.matrix(tAll)),rbind(sum(abs(matrix(1,dim(myX)[1],1)%*%myY))/(dim(myX)[1])^2,as.matrix(gam-
ma)),tGrid,method="linear",rule=1)$y))
# plot(tGrid, gammaPlot)

# plot errHat
for(bb in 1:B){
  for(j in 1:n){
    if (is.na(errHatB[bb,1,j])==FALSE){
      # adding values for t = 0
      errHatBPlot[bb,,j] = matrix(unlist(approx(rbind(0,as.matrix(tB[bb,])),rbind(abs(myY[j]-
      myYBMean[bb,1]),as.matrix(errHatB[bb,,j])),tGrid,method="linear",rule=1)$y))
    }
  }
}

```

```

}

# plot(tGrid,errHatBPlot[1,,2])

errHatBPlus = matrix(NA,tGridR,n)
for(i in 1:n){
  cat(sum(Blc[,i]),fill = TRUE)
  # if we the i th observation has not been sampled in a draw
  if(sum(Blc[,i]>0)){
    CIndex = Blc[,i]==1

    if(sum(Blc[,i]>0)==1){
      errHatBPlus[,i] = errHatBPlot[as.logical(CIndex),,i]
    }else{
      errHatBPlus[,i] = colMeans(errHatBPlot[as.logical(CIndex),,i], na.rm = TRUE)
    }
  }
}
errHatBPlusPlot = rowMeans(errHatBPlus, na.rm =TRUE)
# plot(tGrid,errHatBPlusPlot)

# plot RHat
RHat = matrix(NA,tGridR,1)
RHat = (errHatBPlusPlot-errBarPlot)/(gammaPlot-errBarPlot)
# sum(is.na(errHatBPlusPlot))
# sum(is.na(errBarPlot))
# sum(is.na(errHatBPlusPlot-errBarPlot))
# plot(tGrid,RHat)

# plot ErrHatPlus
wHat = 0.632/(1-0.368*RHat)
# plot(tGrid,wHat)
ErrHatPlus = (1-wHat)*errBarPlot+wHat*errHatBPlusPlot
ErrHatPlus[as.logical(ErrHatPlus<0)]=NA
plot(tGrid,ErrHatPlus)

# plotbeta
betaBPlot = array(NA,dim =c(dim(betaPlot)[2],tGridR))
for(i in 1:dim(betaPlot)[2]){
  betaBPlot[i,] =
  matrix(unlist(approx(rbind(0,as.matrix(tAll[1:k,])),rbind(0,as.matrix(betaPlot[,i])),tGrid,method="linear",rule=1)$y))
}
matplot(tGrid,t(betaBPlot),type = "l", xlab ="t", ylab="beta")

# plot everything
dev.new(width = 6, hight = 6)
par(mfrow=c(2,2),cex=0.6)
# plot errBar
plot(tGrid, errBarPlot)
# plot gamma
plot(tGrid, gammaPlot)
# plot leave-one-out

```

```
plot(tGrid,errHatBPlusPlot)
# plot err0.632
plot(tGrid,ErrHatPlus)
```