

## Independent study with changes

```
df <- read.csv('C:/Users/17143/Desktop/Indepnent_Study/DIADEMwithProteomics_remission.csv', header = T
```

### Data cleaning

```
library(dplyr)

##
## Attaching package: 'dplyr'
##
## The following objects are masked from 'package:stats':
##
##     filter, lag
##
## The following objects are masked from 'package:base':
##
##     intersect, setdiff, setequal, union

library(tidyr)
sub<-df %>%select(which(colnames(df)=="remission"))
data <- df %>% mutate(ExSmoker = ifelse(ExSmoker == "No",0,1))
nums <- unlist(lapply(data, is.numeric))
data <- data[ , nums]
c <- c(2:11645)
data <- data[,c]
data[is.na(data)] <- 0
data <- data %>% select(-6786,-1282,-11614,-1246,-6785,-6781)
data <- data[,!grepl("hba1c",colnames(data))]
data <- data[,!grepl("HbA1c",colnames(data))]
data <- data[,!grepl("HOMA",colnames(data))]
data <- data[,!grepl("homa",colnames(data))]
data <- data[,!grepl("remission",colnames(data))]
data <- data[,!grepl("compremission",colnames(data))]
data <- data[,!grepl("_3",colnames(data))]
data <- data[,!grepl("_6",colnames(data))]
data <- data[,!grepl("_9",colnames(data))]
data<-data.frame(sub,data)
#data$x_12_0 <- data$x_12 - data$x_0
library(tidyselect)
df_12_0<-data %>% select(ends_with(c("_0","_12")))

df_12d<-df_12_0 %>% select(vars_select(names(df_12_0), ends_with("_12")))
df_0d<-df_12_0 %>% select(vars_select(names(df_12_0), ends_with("_0")))
#df_12d %>% select_if(not_all_na)
#df_0d %>% select_if(not_all_na)
df_12<-names(df_12d)
df_0<-names(df_0d)
#d=0
```

```

for(i in seq_along(df_0)) {
  ref<-substr(df_0[i],1,nchar(df_0[i])-2)
  #print(ref)

  ismatch <- sum(grepl(paste0("*", ref), df_12)) > 0
  #print(ismatch)
  name <-paste0(ref,"_12")
  if(ismatch) {
    if(any(names(df_12_0) == name)==TRUE){
      df_12_0[paste0(ref,"_12_0")] <- df_12_0[paste0(ref,"_12")] - df_12_0[paste0(ref,"_0")]
    }
  }
}

n<-(5428:7826)
df_12_0<-df_12_0[,n]
data<-data %>% select(!ends_with(c("_0","_12")))
data<-data.frame(data,df_12_0)
data <- as.data.frame(sapply(data, as.numeric))
data[is.na(data)] <- 0

```

## Train and Test data

```

smp_size <- 0.8*nrow(data)
train_ind <- sample(seq_len(nrow(data)),size=smp_size)
data_train <- (data[train_ind,])
data_test <- (data[-train_ind,])

```

## Lasso version of Logistic Regression

```
library(glmnet)
```

```

## Loading required package: Matrix
##
## Attaching package: 'Matrix'
## The following objects are masked from 'package:tidyr':
##
##   expand, pack, unpack
## Loaded glmnet 4.0-2
X_train=model.matrix(remission~.,data_train)[,-1]
Y_train=data_train$remission
X_te=model.matrix(remission~.,data_test)[,-1]
Y_te=data_test$remission

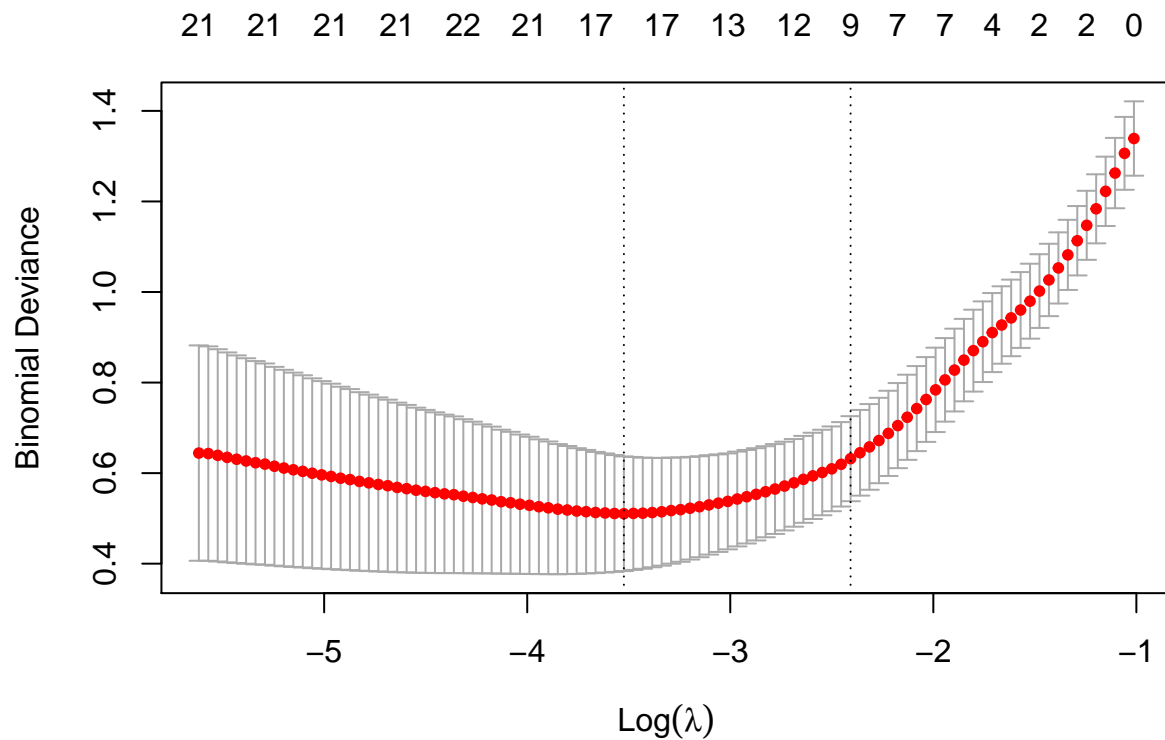
```

Different way to run lasso but doesn't seem to run accurately both error is 0

```

set.seed(123)
cv.lasso <- cv.glmnet(X_train, Y_train, alpha = 1, family = "binomial")
plot(cv.lasso)

```



```
cv.lasso$lambda.min
```

```
## [1] 0.02948501
```

```
cv.lasso$lambda.1se
```

```
## [1] 0.09004297
```

```
#coef(cv.lasso, cv.lasso$lambda.min)
```

```
#Using lambda.min
```

```
model <- glmnet(X_train, Y_train, alpha = 1, family = "binomial", lambda = cv.lasso$lambda.min)
```

```
# regression coefficients
```

```
coeffs<-coef(model)
```

```
coeffs<-as.data.frame(as.matrix(coeffs))
```

```
coeford<-coeffs[order(-coeffs$s0), , drop = FALSE]
```

```
row_sub = apply(coeford, 1, function(row) all(row !=0 ))
```

```
coef0<-coeford[row_sub,, drop = FALSE]
```

```
#View(coef0)
```

```
data.frame(coef0)
```

```
##                               s0
## (Intercept)                5.576954e+00
## cv_12_0                    4.721193e+00
## QUICKI_12_0                7.541397e-01
## LymphocyteAuto_12_0        5.859717e-01
## HADS_Depression3_12_0      1.173015e-01
## dneck120                   1.163967e-01
```

```

## Triglyceride_12_0      1.127194e-01
## Folate_12_0           8.101889e-02
## FT4_12_0             3.325813e-02
## BilirubinT_12_0      2.912162e-02
## IPAQ_1_12_0          1.943096e-02
## lbgi_12_0            1.363167e-02
## Sodium_12_0          7.586083e-04
## PRL_12_0             7.779636e-05
## Axis3MaxCounts_12_0  4.021943e-06
## HR_Standing_2_LAST   -3.765325e-02
## lacr_12_0            -9.869570e-02
## diabmedl             -1.392963e+00

probabilities <- model %>% predict(newx = X_te)
predicted.classes <- ifelse(probabilities > 0.5, "pos", "neg")
# Model accuracy
observed.classes <- data_test$remission
mean(predicted.classes == observed.classes)

## [1] 0

#Using lambda.1se
modell1 <- glmnet(X_train, Y_train, alpha = 1, family = "binomial", lambda = cv.lasso$lambda.1se)
# regression coefficients
coeffs1 <- coef(modell1)
coeffs1 <- as.data.frame(as.matrix(coeffs1))
coeford1 <- coeffs1[order(-coeffs1$s0), , drop = FALSE]
row_sub = apply(coeford1, 1, function(row) all(row != 0 ))
coef1 <- coeford1[row_sub, , drop = FALSE]
#View(coef1)
data.frame(coef1)

##                               s0
## (Intercept)      2.850405421
## cv_12_0          1.805469211
## LymphocyteAuto_12_0 0.249533560
## Folate_12_0       0.055981991
## dneck120          0.040377952
## Triglyceride_12_0  0.016155462
## FT4_12_0          0.009371377
## Sodium_12_0       0.006284426
## HR_Standing_2_LAST -0.015567546
## diabmedl         -0.705913113

probabilities <- modell1 %>% predict(newx = X_te)
predicted.classes <- ifelse(probabilities > 0.5, "pos", "neg")
observed.classes <- data_test$remission
mean(predicted.classes == observed.classes)

## [1] 0

```

## Random forest

```
library(randomForest)
```

```
## randomForest 4.6-14
```

```

## Type rfNews() to see new features/changes/bug fixes.

##
## Attaching package: 'randomForest'

## The following object is masked from 'package:dplyr':
##
##      combine

data$remission <- as.character(data$remission)
data$remission <- as.factor(data$remission)
fit <- randomForest(remission~., data)
print(fit)

##
## Call:
## randomForest(formula = remission ~ ., data = data)
##              Type of random forest: classification
##              Number of trees: 500
## No. of variables tried at each split: 53
##
##              OOB estimate of  error rate: 14.29%
## Confusion matrix:
##      0  1 class.error
## 0 18  9  0.33333333
## 1  1 42  0.02325581

imp<-importance(fit)
rfimp<-data.frame(imp)
rfimp<- rfimp[order(-rfimp$MeanDecreaseGini), , drop = FALSE]
row_sub = apply(rfimp, 1, function(row) all(row !=0 ))
rfimp<-rfimp[row_sub,, drop = FALSE]
data.frame(rfimp)

##              MeanDecreaseGini
## Folate_12_0      0.6949215781
## BEN              0.6732706046
## QUICKI_12_0      0.6445738774
## hba1c            0.4326952147
## 1ALT_12_0        0.3732597407
## LDLCalc_12_0     0.3613288126
## Cholesterol_12_0 0.3330551417
## Glucose_12_0     0.3194965142
## Urea_12_0        0.3109788516
## Hct_12_0         0.2855673469
## Fibro_ValidMeasuresNumber_12_0 0.2848778426
## MDRD4_12_0       0.2826000897
## Creatinine_12_0  0.2618182685
## CalciumCorr_12_0 0.2561314440
## VitD_12_0        0.2483706235
## dwaist120        0.2455817452
## SBP_Standing_Avg_12_0 0.2400449932
## FT4_12_0         0.2309291070
## DBP_Sitting_2_12_0 0.2272379814
## gmi_12_0         0.2232357232
## Fibro_CAP_med_12_0 0.2210682189

```

## ADP	0.2171457023
## SBP_Sitting_2_12_0	0.2116349498
## lvitD_12_0	0.2105950699
## TANITAidealBodyWeight_12_0	0.1956150835
## MPV_12_0	0.1927347403
## HR_Sitting_2_12_0	0.1912121052
## lAST_12_0	0.1911282916
## CKDEPI_12_0	0.1833670820
## TANITABoneMass_12_0	0.1790042723
## Albumin_12_0	0.1774163946
## HR_Sitting_1_12_0	0.1730144552
## HR_Standing_2_12_0	0.1708104013
## TANITATBW_12_0	0.1697073103
## Waistcircumference_12_0	0.1692776194
## min_sensor_12_0	0.1639926198
## DBP_Standing_2_12_0	0.1634344379
## nighttime_max_sens_glucose_12_0	0.1614362409
## MCV_12_0	0.1562447413
## TANITAMetabolicAge_12_0	0.1482540322
## SBP_Sitting_1_12_0	0.1445726372
## SBP_Sitting_Avg_12_0	0.1421247295
## DBP_Standing_1_12_0	0.1406801210
## daytime_max_sensor_glucose_12_0	0.1405903548
## MCHC_12_0	0.1368593798
## BilirubinT_12_0	0.1357907720
## percent_time_70_180_12_0	0.1357321621
## SBP_Standing_1_12_0	0.1320561565
## MaxLengthofSedentaryBreaks_1	0.1315668456
## Calorimetry_RMR_12_0	0.1286563746
## UricAcid_12_0	0.1264756470
## average_sensor_12_0	0.1250972020
## LymphocyteAuto_12_0	0.1230406625
## DBP_Sitting_1_12_0	0.1219302993
## daytime_avg_sensor_glucose_12_0	0.1203125603
## HR_Standing_1_12_0	0.1197537516
## dneck120	0.1186298427
## Potassium_12_0	0.1184840458
## d_weight_120	0.1165018201
## Neckcircumference_12_0	0.1153707211
## TotalMVPA_12_0	0.1153126289
## Calcium_12_0	0.1111131284
## Platelet_12_0	0.1110426086
## q1_sensor_12_0	0.1095589179
## VectorMagnitudeCPM_12_0	0.1075358064
## RD	0.1074793729
## BQC	0.1067292422
## VitB12_12_0	0.1064114817
## max_sensor_12_0	0.1052357222
## lPTH_12_0	0.1041075546
## TotalProtein_12_0	0.1039674762
## nighttime_min_sens_glucose_12_0	0.1036518162
## standard_deviation_12_0	0.1026460326
## PDWifavailable_12_0	0.1021893159
## TotalTimeinSedentaryBreaks_1	0.1009814270

## WBC_12_0	0.0997898019
## DBP_Standing_Avg_12_0	0.0995965413
## AGR2_12_0	0.0968379300
## BES	0.0955955936
## dfatmass120	0.0953339152
## Height_12_0	0.0938629761
## daytime_sd_12_0	0.0922292204
## median_sensor_12_0	0.0919560134
## lCPeptide_12_0	0.0880324324
## CDK2cyclinA_12_0	0.0858411385
## AUH	0.0853761039
## inLight_12_0	0.0846491448
## lInsulin_12_0	0.0846489488
## q3_sensor_12_0	0.0845349378
## IWQOL_DistressScaleAdj_12_0	0.0832180471
## average_auc_per_day_12_0	0.0829330169
## TANITAWeight_12_0	0.0826741408
## l2CDK2cyclinA_12_0	0.0823121499
## diabmedl	0.0822246679
## average_auc_180_12_0	0.0820961635
## HR_Sitting_2_LAST	0.0807080110
## SBP_Standing_2_12_0	0.0804908942
## AUQ	0.0803471789
## TANITAMuscleMass_12_0	0.0798923521
## StepsPerMinute_12_0	0.0796666017
## IWQOL_SexCount_12_0	0.0792815014
## AvgTimeperFreedson1998Bou	0.0759784636
## AVR	0.0750635035
## cv_12_0	0.0748465330
## HR_Standing_Avg_12_0	0.0743733170
## Hemoglobin_12_0	0.0739367784
## ADR	0.0738185015
## Axis3Counts_12_0	0.0735752993
## min_spent_over_250_day_12_0	0.0726631883
## nighttime_auc_12_0	0.0721770685
## NeutrophilAuto_12_0	0.0720053229
## day_night_sensor_ratio_12_0	0.0713342745
## Ficolin3_12_0	0.0676046225
## IWQOL_EsteemCount_12_0	0.0668345296
## HR_Sitting_Avg_12_0	0.0666085506
## l2CD23_12_0	0.0664263016
## percent_time_over_250_12_0	0.0660864596
## AUI	0.0660798183
## TANITAFatMass_12_0	0.0659613003
## Sodium_12_0	0.0655710389
## DMedLast	0.0646990583
## AUE	0.0642826304
## PolyUbiquitinK48_12_0	0.0626464722
## l2IL17sR_12_0	0.0620562149
## StepsCounts_12_0	0.0597127018
## Fibro_E_med_12_0	0.0593142225
## EMAP2_12_0	0.0593108956
## IWQOL_TotalCount_12_0	0.0589646328
## percent_time_over_200_12_0	0.0578850302

## 12a2Antiplasmin_12_0	0.0566658773
## IWQOL_TotalScaleAdj_12_0	0.0550713897
## AQX	0.0549618304
## TANITABMI_12_0	0.0544840336
## GPNMB_12_0	0.0543725325
## SMAC_12_0	0.0543024673
## min_spent_70_180_day_12_0	0.0540612224
## Semaphorin3E_12_0	0.0533998778
## 12GOR	0.0531220779
## RDWCV_12_0	0.0523391874
## IWQOL_PhysicalScaleAdj_12_0	0.0520334467
## Axis2CPM_12_0	0.0517003650
## pc1	0.0513720618
## AverageMVPAPerday_12_0	0.0509094665
## Fassoluble_12_0	0.0502585693
## VectorMagnitudeCounts_12_0	0.0501178659
## 12BASI_12_0	0.0493899638
## 12IL4_12_0	0.0492639222
## Hgb_12_0	0.0491961905
## HR_Standing_Avg_screening	0.0489460351
## min_spent_over_200_day_12_0	0.0483860789
## IWQOL_SexScaleAdj_12_0	0.0475661509
## estimated_a1c_12_0	0.0474868984
## DBP_Sitting_Avg_12_0	0.0472261905
## DPL	0.0466330143
## 12Mcl1_12_0	0.0458017510
## percent_time_70_180_day_12_0	0.0452659341
## IWQOL_WorkCount_12_0	0.0451973856
## CPeptide_12_0	0.0451755981
## 12EIZ	0.0448093027
## Diabetesmedno_LAST	0.0446919649
## 12TGFb3_12_0	0.0446270270
## HR_Standing_Avg_LAST	0.0440559997
## Enterokinase_12_0	0.0440558160
## TroponinIskeletalfasttwitc	0.0439422000
## D_steps120	0.0435266973
## Diabetesmedno_12_0	0.0432292631
## min_spent_over_200_night_12_0	0.0432113114
## insulinpmol_12_0	0.0431959541
## Axis2Counts_12_0	0.0423574602
## Axis3MaxCounts_12_0	0.0422123308
## CNTN2_12_0	0.0420073831
## hbgi_12_0	0.0416553137
## pc11	0.0415398562
## SECTM1_12_0	0.0414119309
## AUC	0.0412693017
## percent_time_70_180_night_12_0	0.0411820180
## 12SECTM1_12_0	0.0403609565
## Chloride_12_0	0.0403200000
## BMPRII_12_0	0.0396334362
## FSTL3_12_0	0.0395446083
## pc31	0.0389593563
## IP10_12_0	0.0385670831
## HR_Standing_1_LAST	0.0384748833



## Fucosyltransferase3_12_0	0.0384643077
## FCN1_12_0	0.0383741716
## l2Siglec3_12_0	0.0382392333
## FGA	0.0380789961
## percent_time_over_180_12_0	0.0380491837
## l2Tenascin_12_0	0.0379041789
## TotalTimeinFreedson1998Bo	0.0378724568
## REG4_12_0	0.0375603002
## HR_Standing_2_LAST	0.0373790557
## MMP12_12_0	0.0372583844
## HCE000342_12_0	0.0372207039
## l2MEPE_12_0	0.0371542064
## IWQOL_DistressCount_12_0	0.0371159030
## l2FOT	0.0368103437
## a2Antiplasmin_12_0	0.0366457132
## RSP04_12_0	0.0365087413
## MFGM_12_0	0.0364648221
## AUB	0.0360976801
## NrCAM_12_0	0.0360656056
## l1PAQ_WalkingMETs_12_0	0.0358588584
## MCH_12_0	0.0356947925
## GranzymeB_12_0	0.0356716115
## l2MICA_12_0	0.0351554069
## FETUB_12_0	0.0348033126
## l2IL18BPα_12_0	0.0343376238
## l2IP10_12_0	0.0343193348
## Light_12_0	0.0342567100
## EQ5D_Scale_12_0	0.0341718789
## IWQOL_PhysicalCount_12_0	0.0339805321
## AUJ	0.0337048359
## modd_12_0	0.0335960429
## l2EPI	0.0335885147
## l2MMP7_12_0	0.0333177076
## total_sensor_readings_12_0	0.0331549842
## TANITABMR_12_0	0.0329251701
## Axis3AverageCounts_12_0	0.0327788360
## l2Soggy1_12_0	0.0327028632
## proteinZinhibitor_12_0	0.0325948053
## AlkPhos_12_0	0.0325308933
## TSG6_12_0	0.0317313211
## Spondin1_12_0	0.0313032942
## l2ARSB_12_0	0.0312805452
## Axis1Counts_12_0	0.0312428571
## tst	0.0307678161
## AUP	0.0307638528
## l2CathepsinB_12_0	0.0307315609
## l2BAFF_12_0	0.0306092167
## nighttime_sd_12_0	0.0300462349
## HIF1α_12_0	0.0300087302
## JAG1_12_0	0.0297388289
## HDL_12_0	0.0296435795
## percent_time_over_180_night_12_0	0.0295860595
## ARSB_12_0	0.0295458432
## inMVPA_12_0	0.0294301626

## dmusclemass120	0.0293317350
## BTC_12_0	0.0292524868
## ROBO2_12_0	0.0291511438
## EPHB2_12_0	0.0291206614
## l2IFNb_12_0	0.0290778224
## PSMA_12_0	0.0289753212
## cJun_12_0	0.0286980035
## TRAILR4_12_0	0.0285878261
## WTLOSS_12_LAST	0.0285506200
## BAFF_12_0	0.0284968030
## Triglyceride_12_0	0.0282610723
## PLCG1_12_0	0.0281762293
## l2cJun_12_0	0.0279875476
## l2FCN1_12_0	0.0275814361
## min_spent_over_180_12_0	0.0272642259
## percent_time_over_250_night_12_0	0.0272014070
## Kallistatin_12_0	0.0271530438
## MSP_12_0	0.0270669090
## l2MRC2_12_0	0.0269580664
## adplepratio	0.0268245887
## Fibro_E_IQRE_med_12_0	0.0266693878
## l2Proteinase3_12_0	0.0266351203
## LRRT3_12_0	0.0265347001
## MnSOD_12_0	0.0261835765
## ITIheavychainH4_12_0	0.0260461686
## l2Endocan_12_0	0.0259625519
## l2DPL	0.0259125541
## CONA1_12_0	0.0258986648
## l2IL1F7_12_0	0.0258667784
## Tenascin_12_0	0.0258528809
## l1PAQ_ModerateMETs_12_0	0.0258494665
## percent_time_over_200_day_12_0	0.0257686346
## GGT_12_0	0.0253263204
## Transferrin_12_0	0.0253084321
## IL18Ra_12_0	0.0253082389
## j_index_12_0	0.0251554389
## IWQOL_EsteemScale_12_0	0.0251519581
## UCreatinine_12_0	0.0251221435
## min_spent_70_180_12_0	0.0251221435
## l2IFNA7_12_0	0.0250775510
## l2PGCB_12_0	0.0249181225
## l2BCMA_12_0	0.0249169608
## TopoisomeraseI_12_0	0.0248762257
## BGH3_12_0	0.0248204770
## l2CD97_12_0	0.0246873068
## VEGFD_12_0	0.0245445378
## l2OPG_12_0	0.0243366581
## l2IgA_12_0	0.0240596491
## l2CAMK1D_12_0	0.0239878184
## HDF	0.0239795250
## TimeToEat_Dinner	0.0238734556
## GFAP_12_0	0.0238471033
## Periostin_12_0	0.0237996461
## CD39_12_0	0.0236290137

## MMP8_12_0	0.0236237162
## TrkB_12_0	0.0234704762
## ARR	0.0233154837
## min_spent_over_200_12_0	0.0232640409
## Renin_12_0	0.0232569745
## l2PCI_12_0	0.0231814059
## conga_1_12_0	0.0231606061
## AUD	0.0231093027
## Proteindisulfideisomerase_12_0	0.0230792891
## HistoneH2Az_12_0	0.0230623974
## l2Kallistatin_12_0	0.0230486654
## C9_12_0	0.0230062125
## l2CNTN2_12_0	0.0229821991
## DNG	0.0229812121
## BNP32_12_0	0.0226455782
## l2CHKB_12_0	0.0224303461
## l2a2HSGlycoprotein_12_0	0.0222220817
## l2RAP_12_0	0.0221525641
## CCL28_12_0	0.0220144668
## l2TPSG1_12_0	0.0220101800
## daytime_min_sensor_glucose_12_0	0.0219978614
## HR_Standing_1_screening	0.0218236701
## l2MATN3_12_0	0.0217805528
## l2Renin_12_0	0.0214953047
## Neckcircumference_screening	0.0214701419
## d_lhdl120	0.0212909091
## Axis2MaxCounts_12_0	0.0212004421
## OX40Ligand_12_0	0.0210920941
## GFRa3_12_0	0.0210730159
## EosinophilAuto_12_0	0.0210483342
## Endocan_12_0	0.0210360674
## CgA_12_0	0.0208894737
## l2EMAP2_12_0	0.0208830958
## NKp46_12_0	0.0207149439
## PEDF_12_0	0.0205796818
## TANITAidealBodyWeight_LAST	0.0205000000
## GDNF_12_0	0.0202806023
## ProteasenexinI_12_0	0.0199047619
## PACAP38_12_0	0.0198395903
## GIIE_12_0	0.0198328762
## EGVEGF_12_0	0.0197608634
## l2LYPD3_12_0	0.0196462585
## GIB_12_0	0.0195528652
## LRRK2_12_0	0.0195460660
## CATZ_12_0	0.0195147739
## IPAQ_4_Trunc_12_0	0.0194809039
## BOC_12_0	0.0194026488
## nighttime_avg_sens_glucose_12_0	0.0193968711
## l2TLR2_12_0	0.0193925926
## IGFBP2_12_0	0.0191925170
## PDE7A_12_0	0.0191842105
## TIMD3_12_0	0.0190309681
## l2bEndorphin_12_0	0.0186994296
## ATS15_12_0	0.0186970760

## 12FSTL1_12_0	0.0186852002
## Plasmin_12_0	0.0185617667
## GGE	0.0185333333
## Semaphorin3A_12_0	0.0184063647
## GPC6_12_0	0.0183655706
## 12CCL28_12_0	0.0183627706
## MMP10_12_0	0.0182554589
## LIN7B_12_0	0.0181656140
## d_lpth120	0.0180100906
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## HADS_TotalDepression_12_0	0.0179978210
## 12TLR4MD2complex_12_0	0.0179299867
## XG	0.0178374194
## CAMK1D_12_0	0.0177948402
## HR_Sitting_1_screening	0.0177625596
## TACI_12_0	0.0175848596
## HADS_Depression3_12_0	0.0175704949
## DBP_Sitting_2_LAST	0.0175671088
## 12FAM3D_12_0	0.0173571429
## MATN3_12_0	0.0173251068
## 12IL6sRa_12_0	0.0172400412
## min_spent_over_120_12_0	0.0172324831
## p27Kip1_12_0	0.0171818182
## HGFA_12_0	0.0171664661
## 12LRRK2_12_0	0.0170666667
## NRX3B_12_0	0.0170479063
## HCE004331_12_0	0.0170475996
## Moderate_12_0	0.0170000000
## Osteopontin_12_0	0.0169588235
## HOC	0.0169182046
## Thrombin_12_0	0.0169032141
## QUACKI	0.0168690154
## AMPKa1b1g1_12_0	0.0168638662
## 12GA7331protein_12_0	0.0167978918
## Galectin8_12_0	0.0167555556
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## 12TAFI_12_0	0.0166532880
## 12Enterokinase_12_0	0.0166352201
## 12IL8_12_0	0.0166254100
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## 12MRCKB_12_0	0.0165942857
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## ZI	0.0165517241
## gp130soluble_12_0	0.0163902439
## 12BCAM_12_0	0.0163878788
## IFNg_12_0	0.0163809524
## 12IFNaA_12_0	0.0162722222
## 12GGE	0.0162249912
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## Fibro_E_IQR_12_0	0.0160173913
## 12eIF5_12_0	0.0160000000
## HR_Sitting_Avg_screening	0.0159079070

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## FGFR4_12_0	0.0158237861
## 12gp130soluble_12_0	0.0157792208
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## FAM3B_12_0	0.0154691877
## StepsAverageCounts_12_0	0.0153903509
## 12CHL1_12_0	0.0153757576
## 12IFN10_12_0	0.0153682680
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## 12EGVEGF_12_0	0.0151223135
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## Galectin2_12_0	0.0145281385
## av_awakening	0.0145174973
## GRN_12_0	0.0145174973
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## 12Aminoacylase1_12_0	0.0144927536
## TANITAFFM_12_0	0.0144857143
## 12IL5_12_0	0.0144510272
## Elafin_12_0	0.0144414716
## 12ADAM9_12_0	0.0144180563
## IgA_12_0	0.0143675214
## percent_time_over_200_night_12_0	0.0143624694
## EBP2_12_0	0.0143490355
## RNaseH1_12_0	0.0143307758
## FN14_12_0	0.0143163399
## TCCR_12_0	0.0142930403
## 12CSH_12_0	0.0142725490
## GHC2_12_0	0.0141493268
## MIS_12_0	0.0141415275
## MIP5_12_0	0.0140698413
## Vitronectin_12_0	0.0140480726

## 12LCK_12_0	0.0140394737
## IL10_12_0	0.0139777427
## 12ANGL4_12_0	0.0139428571
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## 12CJQ	0.0138400000
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## IL10Rb_12_0	0.0136362319
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## 12EphrinA3_12_0	0.0135168961
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## bEndorphin_12_0	0.0134603175
## 12WIF1_12_0	0.0134472727
## 12TIG2_12_0	0.0134292374
## 12MIP1a_12_0	0.0133983648
## 12Transferrin_12_0	0.0133981800
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## 12SonicHedgehog_12_0	0.0133329227
## NormScale_40_12_0	0.0133082632
## GFRA1_12_0	0.0133025641
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## FAM3D_12_0	0.0132452620
## Ckine_12_0	0.0132452620
## 12TopoisomeraseI_12_0	0.0132452620
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## PUR8_12_0	0.0130987013
## IL1F6_12_0	0.0130909091
## 12FGA	0.0130769231
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## GOT1_12_0	0.0130614192
## 12IL22RA1_12_0	0.0130089127
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## 12GNS_12_0	0.0129562998
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## 12IDS_12_0	0.0128866978
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## 12BMPER_12_0	0.0124102564
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## 12FSTL3_12_0	0.0122656450

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## l2PPAC_12_0	0.0115555556
## l2CRDL1_12_0	0.0115348837
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## Sphingosinekinase1_12_0	0.0115238095
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## l2STX1a_12_0	0.0112390244
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## l2Myostatin_12_0	0.0112130623
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## IWQOL_Sex1_12_0	0.0110402477
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## EphrinA3_12_0	0.0106430155
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## Notch1_12_0	0.0104034537
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## l2AnnexinV_12_0	0.0103434343
## CNTFRalpha_12_0	0.0103404255
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## IPAQ_TotalMETs_Trunc_12_0	0.0102857143
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## EphrinA5_12_0	0.0102466793
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## l2CNTFRalpha_12_0	0.0102023810
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## l2NANOG_12_0	0.0101587302
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## IL3_12_0	0.0077943309
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## CAMK1_12_0	0.0076518519
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## l2STAB2_12_0	0.0076097561
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## 12CD59_12_0	0.0074722222
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## Fasligandsoluble_12_0	0.0073914824
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## 12CSK_12_0	0.0073333333
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## 12HCE003300_12_0	0.0073310105
## KininogenHMW_12_0	0.0073204633
## VEGFsR3_12_0	0.0073083004
## MIA_12_0	0.0073083004
## ANGL4_12_0	0.0073083004
## 12Carbonicanhydrase9_12_0	0.0073083004
## 12FABPL_12_0	0.0072906404
## TIMP3_12_0	0.0072857143
## PACAP27_12_0	0.0072770563
## 12Lymphotoxina2b1_12_0	0.0072770563
## KX	0.0072727273
## Transketolase_12_0	0.0072727273
## 12Flt3ligand_12_0	0.0072727273
## 12Macrophagescavengerreceptor_12_0	0.0072727273
## ERBB2_12_0	0.0072603517
## BSSP4_12_0	0.0072603517
## TGFbRII_12_0	0.0072428571
## Eotaxin_12_0	0.0072428571
## 12Thrombin_12_0	0.0072428571
## SLIK1_12_0	0.0072403101
## TANITAFFM_LAST	0.0072380952
## BASI_12_0	0.0072380952

## l2Angiopoietin4_12_0	0.0072285714
## HXK1_12_0	0.0072245153
## C4b_12_0	0.0072245153
## CD97_12_0	0.0072245153
## l2FGF4_12_0	0.0072173730
## Glucagon_12_0	0.0072052632
## RS3_12_0	0.0072052632
## l2ECM1_12_0	0.0072052632
## ABL	0.0072000000
## TroponinI_12_0	0.0072000000
## sRANKL_12_0	0.0071637427
## HCE003183_12_0	0.0071428571
## IgE_12_0	0.0071382504
## CYTF_12_0	0.0071130435
## IFNgR1_12_0	0.0071111111
## l2PCadherin_12_0	0.0071069767
## AUK	0.0071024390
## C2_12_0	0.0070877005
## l2ENC	0.0070750000
## SMOC1_12_0	0.0070533333
## l2DNG	0.0070283401
## l2PlGF_12_0	0.0070144928
## transcriptionfactorMLR1isofo	0.0070103343
## l2GML	0.0070000000
## l2OSM_12_0	0.0069938534
## MIF_12_0	0.0069731868
## ABL1_12_0	0.0069498708
## l2BTK_12_0	0.0069428571
## l2EDAR_12_0	0.0069333333
## IL19_12_0	0.0069264069
## ATS1_12_0	0.0069018718
## TLR4_12_0	0.0068612403
## l2TGFbRIII_12_0	0.0068571429
## Siglec9_12_0	0.0068571429
## DcR3_12_0	0.0068571429
## l2AMG02_12_0	0.0068571429
## TANITAidealBodyWeight_screen	0.0068404558
## l2VEGFD_12_0	0.0067969231
## DLL1_12_0	0.0067914068
## avg_excur_over_140_per_day_12_0	0.0067820513
## percent_time_under_70_night_12_0	0.0067820513
## BDNF_12_0	0.0067820513
## l2CAMK2A_12_0	0.0067339130
## NumberofEpochs_12_0	0.0066818182
## l2GFra3_12_0	0.0066666667
## HSP70protein8_12_0	0.0066562771
## TCTP_12_0	0.0066304762
## l2Cardiotrophin1_12_0	0.0066176471
## IWQOL_Physical6_12_0	0.0066149871
## STAT1_12_0	0.0066149871
## CYTD_12_0	0.0065666667
## ADAM9_12_0	0.0065614035
## AveragelengthofSedentaryBrea	0.0065454545
## LCMT1_12_0	0.0065379208

## lIPAQ_VigorousMETs_Trunc_12_0	0.0064962406
## dipwalkingmetsg120	0.0064000000
## SOST_12_0	0.0064000000
## GranzymeH_12_0	0.0064000000
## NAGK_12_0	0.0064000000
## l2Artemin_12_0	0.0064000000
## annexinVI_12_0	0.0063333333
## lIPAQ_ModerateMETs_Trunc_12_0	0.0063116883
## Peroxiredoxin6_12_0	0.0062696296
## CLM6_12_0	0.0062485353
## Angiopoietin4_12_0	0.0062222222
## JAMC_12_0	0.0062222222
## UFC1_12_0	0.0062222222
## TS_12_0	0.0062222222
## l2CTACK_12_0	0.0062222222
## l2AIF1_12_0	0.0062051282
## l2CalcineurinBa_12_0	0.0062000000
## Duration_Month	0.0060000000
## pc4	0.0060000000
## EQ5D_Mobility_12_0	0.0060000000
## Macrophagescavengerreceptor_12_0	0.0060000000
## PSA6_12_0	0.0060000000
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## EphrinA4_12_0	0.0057142857
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## KI3L2_12_0	0.0057142857
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## l2ISLR2_12_0	0.0057142857
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## PAK7_12_0	0.0056666667
## l2SLIK1_12_0	0.0056000000
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## QORL1_12_0	0.0053333333
## l2ULBP3_12_0	0.0053333333
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## Peroxiredoxin1_12_0	0.0052424242
## IDS_12_0	0.0051428571
## d_ldl120	0.0050066667
## PlGF_12_0	0.0049855072
## WISP3_12_0	0.0049855072
## min_spent_over_140_12_0	0.0049777778
## l2FQB	0.0049696970
## Lymphotoxina2b1_12_0	0.0049696970

## GNS_12_0	0.0049696970
## CarbonicanhydraseI_12_0	0.0048761905
## MMP2_12_0	0.0048333333
## l2GranzymeB_12_0	0.0048000000
## l2LKHA4_12_0	0.0047659479
## l2bFGF_12_0	0.0047619048
## TANITABoneMass_LAST	0.0047407407
## l2DQS	0.0047086957
## l2IL17E_12_0	0.0046855960
## PDL2_12_0	0.0046582850
## TNFsRI_12_0	0.0046507177
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## PIAS4_12_0	0.0046153846
## CD38_12_0	0.0045961240
## EPI_12_0	0.0045414414
## l2IL17B_12_0	0.0045333333
## l2CaMKKalpha_12_0	0.0042458002
## l2FETUB_12_0	0.0040427783
## CathepsinS_12_0	0.0039215686
## FCRL3_12_0	0.0039215686
## l2YKL40_12_0	0.0039200000
## ER_12_0	0.0039183673
## AMNLS_12_0	0.0039183673
## BQB	0.0039166667
## EQ5D_Pain_12_0	0.0039166667
## STAB2_12_0	0.0039148936
## l2TMA_12_0	0.0039148936
## l2BPI_12_0	0.0039148936
## min_spent_over_250_12_0	0.0039130435
## NSF1C_12_0	0.0039130435
## dipaqtottrunc120	0.0039111111
## l2GRB2relatedadapterprotein2_1	0.0039111111
## pTEN_12_0	0.0039111111
## ADAM12_12_0	0.0039111111
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## calreticulin_12_0	0.0039111111
## LEG9_12_0	0.0039111111
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## l2HCE000414_12_0	0.0039111111
## l2TGFb1_12_0	0.0039111111
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## Galectin4_12_0	0.0039090909
## FYN_12_0	0.0039090909
## GSTA3_12_0	0.0039090909
## l2TLR4_12_0	0.0039090909
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## l2FYN_12_0	0.0039090909
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## CD48_12_0	0.0039069767
## CoagulationFactorIX_12_0	0.0039069767
## SEPR_12_0	0.0039069767
## l2UCRP_12_0	0.0039069767
## l2TACI_12_0	0.0039069767

## 12Siglec7_12_0	0.0039069767
## 12IL19_12_0	0.0039069767
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## NCK1_12_0	0.0039047619
## Tropomyosin1alphachain_12_0	0.0039047619
## 12BMP1_12_0	0.0039047619
## 12HFJ	0.0039024390
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## M2PK_12_0	0.0039024390
## CDON_12_0	0.0039024390
## NXPH1_12_0	0.0039024390
## 12IgE_12_0	0.0039024390
## SBP_Sitting_1_LAST	0.0039000000
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## IWQOL_Distress4_12_0	0.0039000000
## Neurotrophin5_12_0	0.0039000000
## min_spent_over_180_day_12_0	0.0038974359
## LRIG3_12_0	0.0038974359
## PGRPS_12_0	0.0038974359
## 12Lumican_12_0	0.0038974359
## 12ApoB_12_0	0.0038974359
## 12JDO	0.0038947368
## CRP_12_0	0.0038947368
## JAK2_12_0	0.0038947368
## NLGNX_12_0	0.0038947368
## 12ATS13_12_0	0.0038947368
## 12AggreCAN_12_0	0.0038947368
## 12a2Macroglobulin_12_0	0.0038947368
## 12pTEN_12_0	0.0038947368
## IFN10_12_0	0.0038918919
## NogoReceptor_12_0	0.0038918919
## 12CKMB_12_0	0.0038918919
## C5b6Complex_12_0	0.0038888889
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## ApoE2_12_0	0.0038888889
## lacr_12_0	0.0038888889
## 12MAPK2_12_0	0.0038888889
## 12Prekallikrein_12_0	0.0038888889
## STRATIFIN_12_0	0.0038857143
## TRAILR1_12_0	0.0038857143
## 12HER	0.0038823529
## IWQOL_Physical10_12_0	0.0038823529
## HCE004359_12_0	0.0038823529
## TSP2_12_0	0.0038823529
## 12FLRT2_12_0	0.0038823529
## 12GV_12_0	0.0038823529
## Height_screening	0.0038787879
## HADS_Depression7_12_0	0.0038787879
## 12ApoptosisregulatorBclW_12_0	0.0038787879
## CJQ	0.0038709677
## CLC1B_12_0	0.0038709677
## TNFa_12_0	0.0038709677

## HFA	0.0038666667
## NRP1_12_0	0.0038666667
## l2ICOS_12_0	0.0038666667
## TANITAWeight_LAST_CERNER	0.0038620690
## Tpo_12_0	0.0038620690
## l2PTK6_12_0	0.0038620690
## PDGFRA_12_0	0.0038571429
## AggreCAN_12_0	0.0038571429
## CaMKKalpha_12_0	0.0038571429
## PSelectin_12_0	0.0038571429
## Waistcircumference_screening	0.0038400000
## FCV	0.0038400000
## CoagulationFactorX_12_0	0.0038400000
## IF4A3_12_0	0.0038400000
## IL1sR9_12_0	0.0038400000
## LBP_12_0	0.0038333333
## ATS13_12_0	0.0038333333
## PHI_12_0	0.0038333333
## ANP_12_0	0.0038333333
## dipaqsitting120	0.0038260870
## NKp30_12_0	0.0038260870
## RAP_12_0	0.0038260870
## EMR2_12_0	0.0038260870
## NR1D1_12_0	0.0038260870
## l2C9_12_0	0.0038260870
## C5_12_0	0.0038181818
## FGF10_12_0	0.0038181818
## l2sICAM3_12_0	0.0038181818
## l2HEF	0.0038095238
## CD177_12_0	0.0038095238
## CoagulationFactorXI_12_0	0.0038095238
## LSAMP_12_0	0.0038095238
## BPI_12_0	0.0038095238
## CSRP3_12_0	0.0038095238
## l2Galectin3_12_0	0.0038095238
## l2OLR1_12_0	0.0038095238
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## calgranulinB_12_0	0.0038000000
## l2C5_12_0	0.0038000000
## l2PARC_12_0	0.0038000000
## l2Periostin_12_0	0.0038000000
## ASG	0.0037894737
## AUA	0.0037894737
## HLY	0.0037894737
## IFNA7_12_0	0.0037894737
## WFKN2_12_0	0.0037894737
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## l2dopadecarboxylase_12_0	0.0037894737
## CyclinB1_12_0	0.0037777778
## l2GMY	0.0037750363
## Prekallikrein_12_0	0.0037707391
## ROR1_12_0	0.0037662745
## UB2G2_12_0	0.0037662745



## MPIF1_12_0	0.0037647059
## FUT5_12_0	0.0037647059
## CRTAM_12_0	0.0037647059
## l2CyclophilinA_12_0	0.0037647059
## IWQOL_Physical1_12_0	0.0037568027
## BCMA_12_0	0.0037568027
## l2FCN2_12_0	0.0037568027
## ICOS_12_0	0.0037500000
## MBL_12_0	0.0037500000
## SLAF7_12_0	0.0037500000
## uPA_12_0	0.0037465310
## annexinII_12_0	0.0037465310
## LuxAverageCounts_12_0	0.0037353535
## daytime_auc_12_0	0.0037353535
## PARC_12_0	0.0037353535
## d_lvitd120	0.0037333333
## l2GMZ	0.0037333333
## dopadecarboxylase_12_0	0.0037333333
## PSME3_12_0	0.0037333333
## HCE001796_12_0	0.0037293869
## MMP13_12_0	0.0037293869
## CathepsinH_12_0	0.0037293869
## l2EPHA3_12_0	0.0037231451
## l2C2_12_0	0.0037166086
## l2HCK_12_0	0.0037166086
## l2PSelectin_12_0	0.0037142857
## MRCKB_12_0	0.0037025641
## GP114_12_0	0.0036950067
## Gastrinreleasingpeptide_12_0	0.0036950067
## l2IL12_12_0	0.0036950067
## l2LYVE1_12_0	0.0036923077
## EFNB2_12_0	0.0036786787
## IR_12_0	0.0036786787
## Fibrinogengchaindimer_12_0	0.0036666667
## FMP	0.0036605042
## CarbonicAnhydraseX_12_0	0.0036401515
## TANITAVisceralFatRating_scre	0.0036363636
## CLC4K_12_0	0.0036363636
## CD244_12_0	0.0036363636
## GDF11_12_0	0.0036214178
## a1Antitrypsin_12_0	0.0036065306
## l2EIX	0.0036000000
## IL23R_12_0	0.0036000000
## l2ITAC_12_0	0.0036000000
## l2GDF118_12_0	0.0035986395
## inSedentary_12_0	0.0035818686
## FBLN3_12_0	0.0035729469
## Cripto_12_0	0.0035555556
## totalminbed	0.0035539112
## l2CathepsinV_12_0	0.0035266667
## CATF_12_0	0.0035219512
## l2IL1F8_12_0	0.0035072464
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## FCG2A_12_0	0.0035000000

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## l2GFRa2_12_0	0.0035000000
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## SGTA_12_0	0.0034650980
## C3d_12_0	0.0034632035
## TANITAMuscleMass_screening	0.0034285714
## TotalMeds_12_0	0.0034285714
## SMAD3_12_0	0.0034285714
## Hemopexin_12_0	0.0034285714
## CDC37_12_0	0.0034285714
## LaminB1_12_0	0.0034285714
## SCGFalpha_12_0	0.0034285714
## Myokinasehuman_12_0	0.0034285714
## CPNE1_12_0	0.0034285714
## Noggin_12_0	0.0034285714
## l2LAG1_12_0	0.0034285714
## l2LRP8_12_0	0.0034285714
## l2HSP40_12_0	0.0034285714
## l2Sortingnexin4_12_0	0.0034285714
## TSLPR_12_0	0.0033600000
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## PROTOCOLFOLLOWED_ACTIVE	0.0033333333
## CJO	0.0033333333
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## Axis1MaxCounts_12_0	0.0033333333
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## ApoL1_12_0	0.0033333333
## S100A4_12_0	0.0033333333
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## GA7331protein_12_0	0.0033333333
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## 12IOA	0.0032000000
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## 12BMP14_12_0	0.0032000000
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## 12ETHE1_12_0	0.0032000000
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## HCE004333_12_0	0.0030000000
## FGF18_12_0	0.0030000000
## BGN_12_0	0.0030000000
## FCG3B_12_0	0.0030000000
## LCK_12_0	0.0030000000
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## 12PAK7_12_0	0.0030000000
## 12FN13_12_0	0.0030000000
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## IHP	0.0026666667
## l2EOY	0.0026666667
## HADS_Depression2_12_0	0.0026666667
## HADS_Anxiety7_12_0	0.0026666667
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## Somatostatin28_12_0	0.0026666667
## l1PAQ_VigorousMETs_12_0	0.0026666667
## l1PAQ_WalkingMETs_Trunc_12_0	0.0026666667
## l2B7_12_0	0.0026666667
## l2CalpainI_12_0	0.0026666667
## l2Ubiquitin1_12_0	0.0026666667
## l2Karyopherina2_12_0	0.0026666667
## l2PCNA_12_0	0.0026666667
## l2Galectin2_12_0	0.0026666667
## l2Thrombospondin1_12_0	0.0026666667
## l2PDGFCC_12_0	0.0026666667
## l2SKP1_12_0	0.0026666667
## BEU	0.0020000000
## l2BNX	0.0020000000
## l2CPZ	0.0020000000
## avg_excur_over_200_per_day_12_0	0.0020000000
## HMG1_12_0	0.0020000000
## GPC2_12_0	0.0020000000
## Chitotriosidase1_12_0	0.0020000000
## l2CDK1cyclinB_12_0	0.0020000000
## IL1sRI_12_0	0.0019230769
## l2SOD_12_0	0.0019111111
## ESM	0.0019090909
## PSME1_12_0	0.0019024390
## l2EDA_12_0	0.0019024390
## l2IL22_12_0	0.0018947368
## l2IL18Rb_12_0	0.0009166667

## CART

### Classification

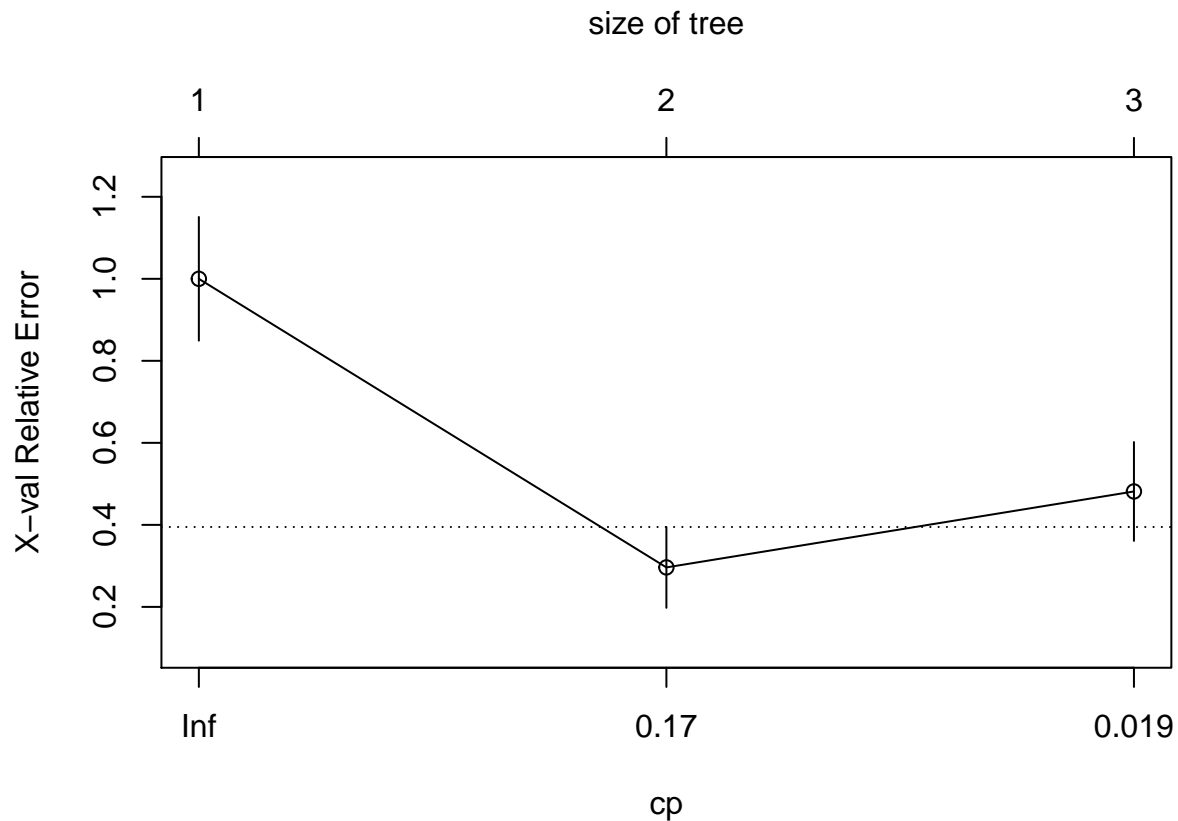
```
library(rpart)
library(rpart.plot)

fit1 <- rpart(remission~., method="class", data=data)
```

```
printcp(fit1)
```

```
##
## Classification tree:
## rpart(formula = remission ~ ., data = data, method = "class")
##
## Variables actually used in tree construction:
## [1] daytime_min_sensor_glucose_12_0 Folate_12_0
##
## Root node error: 27/70 = 0.38571
##
## n= 70
##
##      CP nsplit rel error  xerror   xstd
## 1 0.777778      0  1.00000 1.00000 0.150835
## 2 0.037037      1  0.22222 0.29630 0.098589
## 3 0.010000      2  0.18519 0.48148 0.120503
```

```
plotcp(fit1)
```



```
summary(fit1)
```

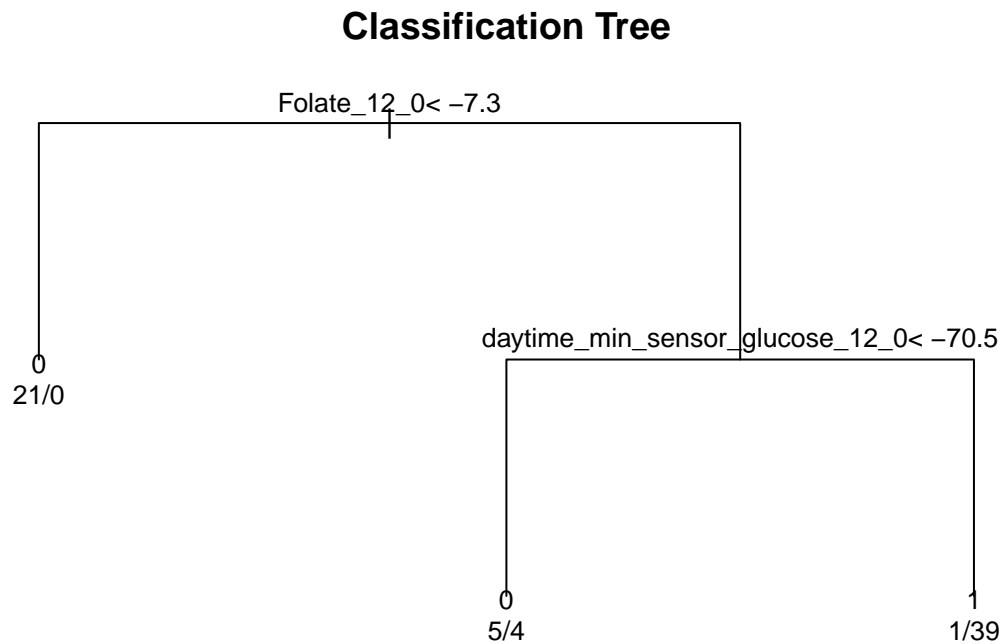
```
## Call:
## rpart(formula = remission ~ ., data = data, method = "class")
##   n= 70
##
##      CP nsplit rel error  xerror   xstd
## 1 0.777778      0  1.00000 1.00000 0.150835
## 2 0.037037      1  0.22222 0.29630 0.098589
## 3 0.010000      2  0.18519 0.48148 0.120503
```

```

## 1 0.77777778      0 1.0000000 1.0000000 0.15083542
## 2 0.03703704      1 0.2222222 0.2962963 0.09858891
## 3 0.01000000      2 0.1851852 0.4814815 0.12050257
##
## Variable importance
##           Folate_12_0           Albumin_12_0
##                17                14
##           Calcium_12_0           LAST_12_0
##                14                14
##           lvitD_12_0           QUICKI_12_0
##                14                14
## daytime_min_sensor_glucose_12_0      estimated_a1c_12_0
##                3                2
##                gmi_12_0 nighttime_avg_sens_glucose_12_0
##                2                2
##           average_sensor_12_0           q1_sensor_12_0
##                2                2
##
## Node number 1: 70 observations,      complexity param=0.7777778
##   predicted class=1   expected loss=0.3857143   P(node) =1
##   class counts:      27    43
##   probabilities: 0.386 0.614
##   left son=2 (21 obs) right son=3 (49 obs)
##   Primary splits:
##     Folate_12_0 < -7.3      to the left,   improve=22.64082, (0 missing)
##     LAST_12_0   < -1.191506 to the left,   improve=19.68123, (0 missing)
##     BEN         < 0.003126954 to the left,   improve=18.28681, (0 missing)
##     hba1c       < 2.4       to the left,   improve=18.28681, (0 missing)
##     Urea_12_0   < -2.6      to the left,   improve=18.28681, (0 missing)
##   Surrogate splits:
##     Calcium_12_0 < -2.235    to the left,   agree=0.943, adj=0.81, (0 split)
##     Albumin_12_0 < -37       to the left,   agree=0.943, adj=0.81, (0 split)
##     QUICKI_12_0  < -0.1013723 to the left,   agree=0.943, adj=0.81, (0 split)
##     lvitD_12_0   < -1.641707 to the left,   agree=0.943, adj=0.81, (0 split)
##     LAST_12_0    < -1.191506 to the left,   agree=0.943, adj=0.81, (0 split)
##
## Node number 2: 21 observations
##   predicted class=0   expected loss=0   P(node) =0.3
##   class counts:      21    0
##   probabilities: 1.000 0.000
##
## Node number 3: 49 observations,      complexity param=0.03703704
##   predicted class=1   expected loss=0.122449   P(node) =0.7
##   class counts:      6    43
##   probabilities: 0.122 0.878
##   left son=6 (9 obs) right son=7 (40 obs)
##   Primary splits:
##     daytime_min_sensor_glucose_12_0 < -70.5      to the left,   improve=4.136168, (0 missing)
##     Triglyceride_12_0                < -1.075    to the left,   improve=3.292517, (0 missing)
##     percent_time_70_180_night_12_0 < -20.91778 to the left,   improve=3.292517, (0 missing)
##     modd_12_0                        < -30.2217   to the left,   improve=3.292517, (0 missing)
##     SBP_Standing_Avg_LAST             < 111.75    to the left,   improve=3.128698, (0 missing)
##   Surrogate splits:
##     estimated_a1c_12_0                < -4.05     to the left,   agree=0.959, adj=0.778, (0 split)

```

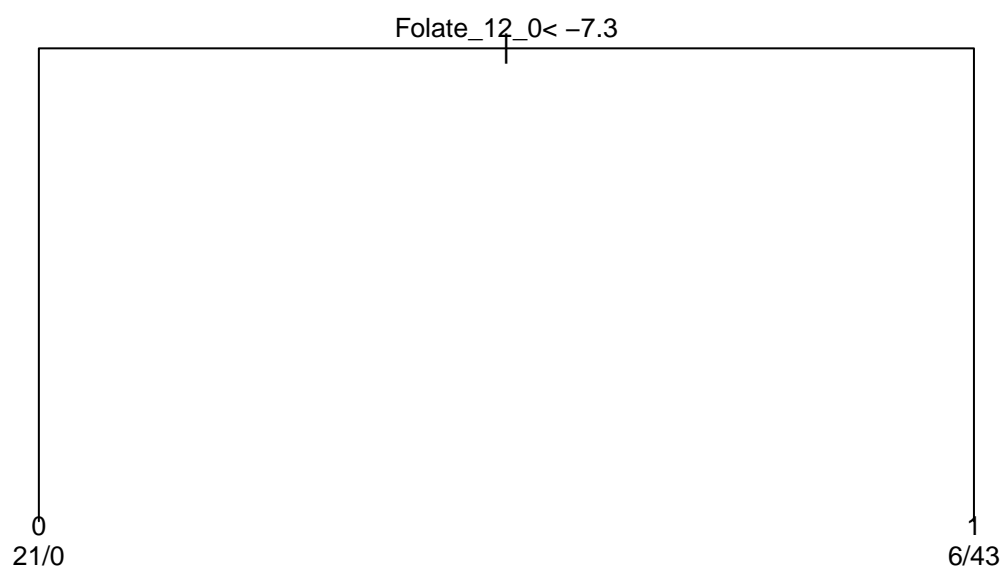
```
##      gmi_12_0 < -2.85      to the left, agree=0.959, adj=0.778, (0 split)
##      nighttime_avg_sens_glucose_12_0 < -99.7621 to the left, agree=0.959, adj=0.778, (0 split)
##      average_sensor_12_0 < -108.4068 to the left, agree=0.939, adj=0.667, (0 split)
##      q1_sensor_12_0 < -96      to the left, agree=0.939, adj=0.667, (0 split)
##
## Node number 6: 9 observations
##   predicted class=0 expected loss=0.4444444 P(node) =0.1285714
##   class counts:      5      4
##   probabilities: 0.556 0.444
##
## Node number 7: 40 observations
##   predicted class=1 expected loss=0.025 P(node) =0.5714286
##   class counts:      1     39
##   probabilities: 0.025 0.975
{plot(fit1, uniform=TRUE, main="Classification Tree")
text(fit1, use.n=TRUE, xpd=TRUE, cex=.8)}
```



```
pfit<- prune(fit1, cp= fit1$cptable[which.min(fit1$cptable[, "xerror"]), "CP"])

# plot the pruned tree
{plot(pfit, uniform=TRUE, main="Pruned Classification Tree")
text(pfit, use.n=TRUE, xpd=TRUE, cex=.8)}
```

## Pruned Classification Tree



## Regression

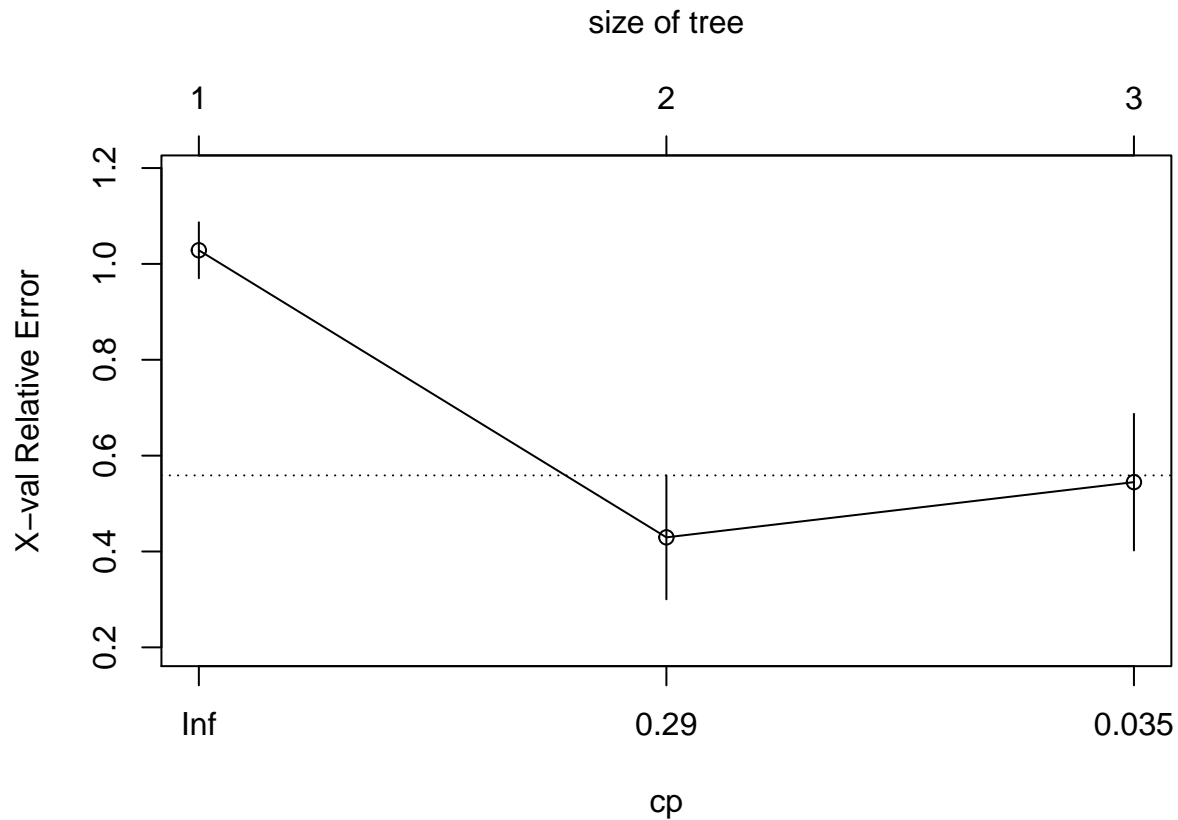
```
fit <- rpart(remission~., method="anova", data=data)
```

```
printcp(fit)
```

```
##
## Regression tree:
## rpart(formula = remission ~ ., data = data, method = "anova")
##
## Variables actually used in tree construction:
## [1] daytime_min_sensor_glucose_12_0 Folate_12_0
##
## Root node error: 16.586/70 = 0.23694
##
## n= 70
##
##      CP nsplit rel error  xerror   xstd
## 1 0.68254     0  1.00000 1.02849 0.058388
## 2 0.12469     1  0.31746 0.42950 0.129265
## 3 0.01000     2  0.19277 0.54472 0.142489
```

```
plotcp(fit)
```





```
summary(fit)
```

```
## Call:
## rpart(formula = remission ~ ., data = data, method = "anova")
##   n= 70
##
##           CP nsplit rel error   xerror   xstd
## 1 0.6825397     0 1.0000000 1.0284876 0.05838835
## 2 0.1246907     1 0.3174603 0.4295023 0.12926500
## 3 0.0100000     2 0.1927696 0.5447211 0.14248928
##
## Variable importance
##           Folate_12_0           Albumin_12_0
##                17                14
##           Calcium_12_0           lAST_12_0
##                14                14
##           lvitD_12_0           QUICKI_12_0
##                14                14
## daytime_min_sensor_glucose_12_0 estimated_a1c_12_0
##                3                2
##                gmi_12_0 nighttime_avg_sens_glucose_12_0
##                2                2
##           average_sensor_12_0           q1_sensor_12_0
##                2                2
##
## Node number 1: 70 observations,   complexity param=0.6825397
```

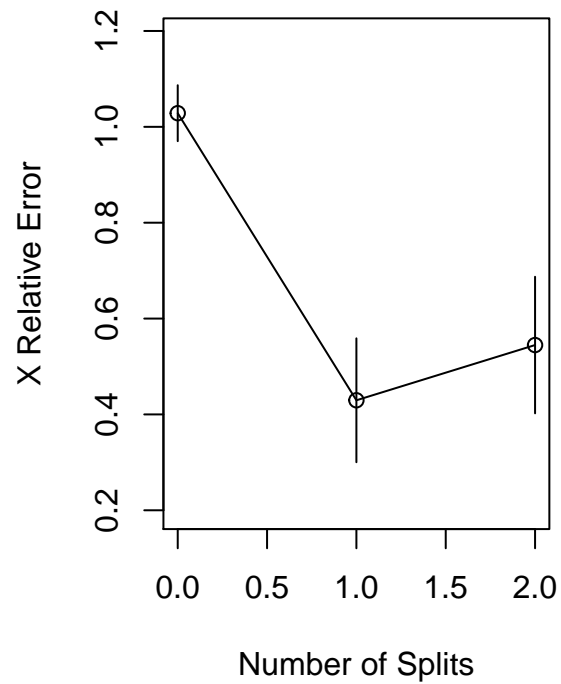
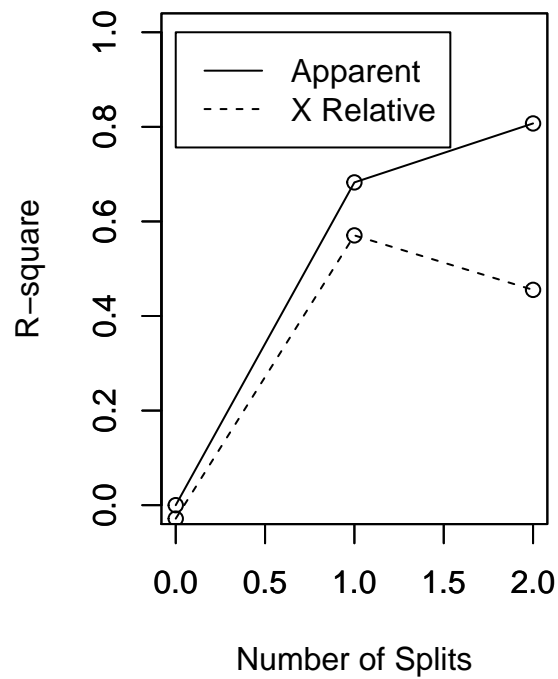
```

## mean=1.614286, MSE=0.2369388
## left son=2 (21 obs) right son=3 (49 obs)
## Primary splits:
## Folate_12_0 < -7.3 to the left, improve=0.6825397, (0 missing)
## lAST_12_0 < -1.191506 to the left, improve=0.5933188, (0 missing)
## BEN < 0.003126954 to the left, improve=0.5512821, (0 missing)
## hbaif < 2.4 to the left, improve=0.5512821, (0 missing)
## Urea_12_0 < -2.6 to the left, improve=0.5512821, (0 missing)
## Surrogate splits:
## Calcium_12_0 < -2.235 to the left, agree=0.943, adj=0.81, (0 split)
## Albumin_12_0 < -37 to the left, agree=0.943, adj=0.81, (0 split)
## QUICKI_12_0 < -0.1013723 to the left, agree=0.943, adj=0.81, (0 split)
## lvitD_12_0 < -1.641707 to the left, agree=0.943, adj=0.81, (0 split)
## lAST_12_0 < -1.191506 to the left, agree=0.943, adj=0.81, (0 split)
##
## Node number 2: 21 observations
## mean=1, MSE=0
##
## Node number 3: 49 observations, complexity param=0.1246907
## mean=1.877551, MSE=0.1074552
## left son=6 (9 obs) right son=7 (40 obs)
## Primary splits:
## daytime_min_sensor_glucose_12_0 < -70.5 to the left, improve=0.3927756, (0 missing)
## Triglyceride_12_0 < -1.075 to the left, improve=0.3126615, (0 missing)
## percent_time_70_180_night_12_0 < -20.91778 to the left, improve=0.3126615, (0 missing)
## modd_12_0 < -30.2217 to the left, improve=0.3126615, (0 missing)
## estimated_a1c_12_0 < -3.25 to the left, improve=0.2971051, (0 missing)
## Surrogate splits:
## estimated_a1c_12_0 < -4.05 to the left, agree=0.959, adj=0.778, (0 split)
## gmi_12_0 < -2.85 to the left, agree=0.959, adj=0.778, (0 split)
## nighttime_avg_sens_glucose_12_0 < -99.7621 to the left, agree=0.959, adj=0.778, (0 split)
## average_sensor_12_0 < -108.4068 to the left, agree=0.939, adj=0.667, (0 split)
## q1_sensor_12_0 < -96 to the left, agree=0.939, adj=0.667, (0 split)
##
## Node number 6: 9 observations
## mean=1.444444, MSE=0.2469136
##
## Node number 7: 40 observations
## mean=1.975, MSE=0.024375
par(mfrow=c(1,2))
rsq.rpart(fit)

##
## Regression tree:
## rpart(formula = remission ~ ., data = data, method = "anova")
##
## Variables actually used in tree construction:
## [1] daytime_min_sensor_glucose_12_0 Folate_12_0
##
## Root node error: 16.586/70 = 0.23694
##
## n= 70
##
## CP nsplit rel error xerror xstd

```

```
## 1 0.68254      0  1.00000 1.02849 0.058388
## 2 0.12469      1  0.31746 0.42950 0.129265
## 3 0.01000      2  0.19277 0.54472 0.142489
```

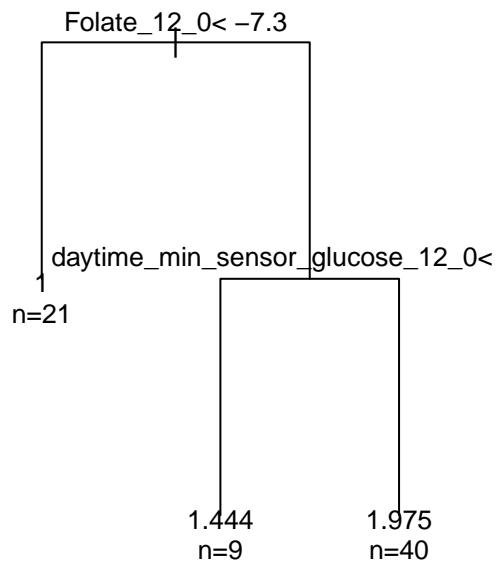


```
{plot(fit, uniform=TRUE, main="Regression Tree")
text(fit, use.n=TRUE, xpd=TRUE, cex=.8)}

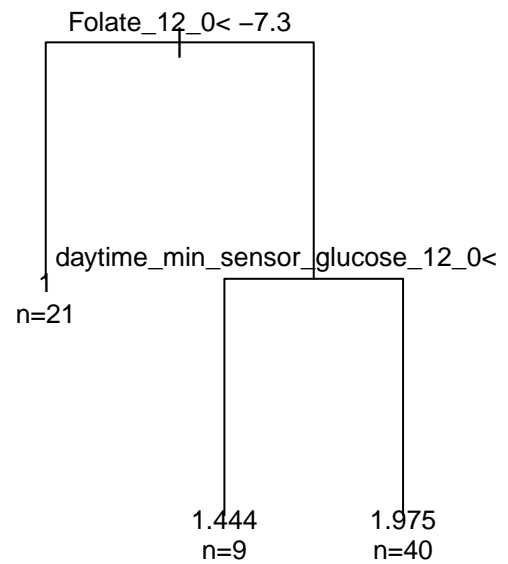
pfit<- prune(fit, cp= fit1$cptable[which.min(fit1$cptable[, "xerror"]), "CP"])

# plot the pruned tree
{plot(pfit, uniform=TRUE, main="Pruned Regression Tree")
text(pfit, use.n=TRUE, xpd=TRUE, cex=.8)}
```

## Regression Tree



## Pruned Regression Tree



## Multiple Linear Regression

```
fit<- lm(remission~., data=data)
```

```
## Warning in model.response(mf, "numeric"): using type = "numeric" with a factor
## response will be ignored
```

```
## Warning in Ops.factor(y, z$residuals): '-' not meaningful for factors
```

```
summ<-summary(fit)
```

```
## Warning in Ops.factor(r, 2): '^' not meaningful for factors
```

```
#summ
```

```
coeffs<-coefficients(fit)
```

```
coeffsord <- coeffs[order(-coeffs)]
```

```
#coeffsord
```

```
coeffsord<-na.omit(coeffsord)
```

```
data.frame(coeffsord)
```

```
##
## (Intercept)                coeffsord
## Mother_Cancer              4.740726e+00
## Father_Cancer              4.341466e+00
## TANITAFM_screening         4.329283e+00
## ExSmoker                   4.159041e+00
## TANITAWeight_screening     3.295370e+00
## TANITABMR_screening        3.042602e+00
```

## Father_Diabetes	1.872362e+00
## CurrentSmoker_Shisha	1.251051e+00
## SBP_Standing_Avg_screening	7.967440e-01
## Father_Obesity	4.323048e-01
## HR_Sitting_2_screening	4.008988e-01
## Takeaway_PerWeek	3.322471e-01
## Dates_Quantity	3.134188e-01
## SBP_Sitting_2_screening	2.922432e-01
## Mother_CVD	2.863922e-01
## Soda_PerWeek	2.333353e-01
## PhysicianDuration_121	2.293297e-01
## DBP_Sitting_1_screening	1.932015e-01
## DBP_Standing_1_screening	1.247072e-01
## DBP_Sitting_2_screening	1.048951e-01
## LatestAlbCr	9.593252e-02
## TANITATBW_screening	9.325901e-02
## HR_Standing_1_screening	7.934664e-02
## TANITAImpedance_screening	6.086061e-02
## SBP_Sitting_1_screening	6.018763e-02
## DieticianDuration_121	5.170409e-02
## TimeToEat_Breakfast	5.156961e-02
## Mother_Obesity	3.992159e-02
## Coffee_PerWeek	3.549674e-02
## Hipcircumference_screening	3.006949e-02
## Neckcircumference_screening	3.560605e-03
## TrainerDuration_121	1.439771e-03
## O0	4.338848e-04
## KX	-2.430382e-04
## Waistcircumference_screening	-7.510443e-04
## HG	-1.123688e-03
## RD	-1.561101e-03
## HR_Sitting_1_screening	-5.399085e-03
## EQ	-7.188664e-03
## UZ	-1.200850e-02
## TimeToEat_Lunch	-1.457020e-02
## TimeToEat_Dinner	-3.674659e-02
## Tea_PerWeek	-9.962502e-02
## TANITAMetabolicAge_screening	-1.128635e-01
## DBP_Standing_2_screening	-1.478293e-01
## Juice_PerWeek	-1.667087e-01
## HR_Standing_2_screening	-1.807813e-01
## Dates_PerWeek	-1.916388e-01
## Father_CVD	-2.009244e-01
## SBP_Standing_2_screening	-2.104589e-01
## DBP_Sitting_Avg_screening	-3.045556e-01
## VA	-3.353568e-01
## TANITAVisceralFatRating_scre	-3.694895e-01
## SBP_Sitting_Avg_screening	-3.723838e-01
## HR_Sitting_Avg_screening	-3.753063e-01
## Mother_Hypertension	-4.488260e-01
## SBP_Standing_1_screening	-4.796831e-01
## WX	-4.820194e-01
## NeutrophilAutoAbsoluteNeu	-5.718729e-01
## TANITAFatMass_screening	-9.192848e-01

## VC	-1.087770e+00
## Height_screening	-1.132343e+00
## Father_Hypertension	-1.386737e+00
## CurrentSmoker_Cigarettes	-1.603185e+00
## TANITAMuscleMass_screening	-1.658964e+00
## TANITAidealBodyWeight_screen	-1.862862e+00
## Mother_Diabetes	-2.652432e+00
## TANITABMI_screening	-4.902927e+00
## BK	-1.285434e+01