Independent study

data <- read.csv('C:/Users/17143/Desktop/Indpendent_Study/DIADEMwithProteomics_remission.csv', header =</pre>

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
data <- data %>% mutate(ExSmoker = ifelse(ExSmoker == "No",0,1))
nums <- unlist(lapply(data, is.numeric))</pre>
data <- data[ , nums]</pre>
c \leftarrow c(2:11645)
data <- data[,c]</pre>
data[is.na(data)] <- 0
data <- data %>% select(-6786,-1282,-11614,-1246,-6785,-6781)
data <- data[,!grepl("hba1c",colnames(data))]</pre>
data <- data[,!grepl("HbA1c",colnames(data))]</pre>
data <- data[,!grepl("HOMA",colnames(data))]</pre>
data <- data[,!grepl("homa",colnames(data))]</pre>
data <- data[,!grepl("remission_",colnames(data))]</pre>
data <- data[,!grepl("compremission",colnames(data))]</pre>
```

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,])</pre>
```

Lasso version of Logistic Regression

```
library(glmnet)

## Loading required package: Matrix

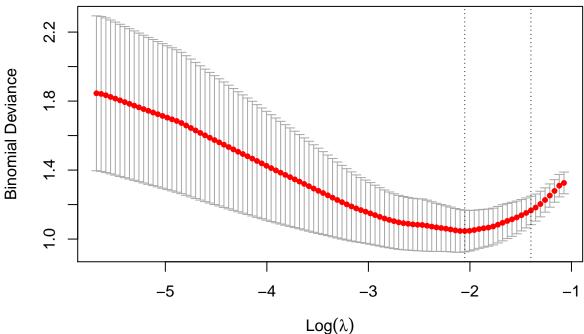
## Loaded glmnet 4.0-2
```

```
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)</pre>
```

32 31 30 31 33 32 31 32 28 26 19 12 8 4 2 1



```
cv.lasso$lambda.min
## [1] 0.1286703
cv.lasso$lambda.1se

## [1] 0.2467781
#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]</pre>
```

```
View(coef0)
data.frame(coef0)
## QUICKI_12
                                  3.175564e+00
## gmihba1_12
                                 4.664018e-01
## HDL_12
                                  4.074800e-01
## IWQOL_EsteemAVG_12
                                 1.668712e-01
## dneck120
                                 6.668647e-02
## percent_time_70_180_12
                                 2.166279e-03
## UricAcid 12
                                 1.718520e-03
## percent_time_70_180_night_12 1.751711e-04
## IWQOL EsteemScale 12
                                1.219488e-04
## DMedLast
                                -1.987887e-16
## diabmedl
                                -1.104585e-04
## Diabetesmedno_LAST
                                -1.491953e-04
## Diabetesmedno_12
                                -9.002303e-02
## Diabetesmedno_6
                                -2.259538e-01
## (Intercept)
                                -1.257286e+00
probabilities <- model %>% predict(newx = X_te)
predicted.classes <- ifelse(probabilities > 0.5, "pos", "neg")
# Model accuracy
observed.classes <- data_test$remission</pre>
mean(predicted.classes == observed.classes)
## [1] 0
#Using lambda.1se
model1 <- glmnet(X_train, Y_train, alpha = 1, family = "binomial", lambda = cv.lasso$lambda.1se)
# regression coefficients
coeffs1<-coef(model1)</pre>
coeffs1<-as.data.frame(as.matrix(coeffs1))</pre>
coeford1<-coeffs1[order(-coeffs1$s0), , drop = FALSE]</pre>
row_sub = apply(coeford1, 1, function(row) all(row !=0))
coef1<-coeford1[row_sub,, drop = FALSE]</pre>
View(coef1)
data.frame(coef1)
##
## QUICKI_12
                2.9742686387
## UricAcid_12 0.0008080566
## (Intercept) -0.1900546703
probabilities <- model1 %>% predict(newx = X_te)
predicted.classes <- ifelse(probabilities > 0.5, "pos", "neg")
observed.classes <- data_test$remission</pre>
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)</pre>
data$remission <- as.factor(data$remission)</pre>
fit <- randomForest(remission~., data)</pre>
print(fit)
##
## Call:
   randomForest(formula = remission ~ ., data = data)
##
##
                  Type of random forest: classification
##
                         Number of trees: 500
## No. of variables tried at each split: 107
##
           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)</pre>
rfimp<-data.frame(imp)
rfimp<- rfimp[order(-rfimp$MeanDecreaseGini), , drop = FALSE]</pre>
row sub = apply(rfimp, 1, function(row) all(row !=0))
rfimp<-rfimp[row_sub,, drop = FALSE]</pre>
View(rfimp)
data.frame(rfimp)
                                    MeanDecreaseGini
## Creatinine_12
                                         0.3018207939
## percent_time_70_180_night_12
                                         0.2818320882
## VitD_12
                                         0.2335704282
## WTLOSS_12
                                         0.2314719945
## EosinophilAuto_12
                                         0.2234741106
## Cholesterol_12
                                         0.2117557885
## AST 12
                                         0.2093061419
## FT4 12
                                         0.2092592301
## EQ5D Anxiety 12
                                         0.2060071552
## BilirubinT_12
                                         0.2018839493
## lvitD 12
                                         0.2010481385
## VitB12_12
                                         0.1976154042
## BEN
                                         0.1974306408
## Fibro_E_med_12
                                         0.1952207643
## 1AST 12
                                         0.1939994293
## FBG_12
                                         0.1870033905
## wtloss_0_12
                                         0.1841749096
## Urea_12
                                         0.1727274357
## ALT 12
                                         0.1677857588
```

0.1617295342

dwaist120

## Fibro_E_IQRE_med_12	0.1603504865
## 1PTH_12	0.1589444015
## 1CPeptide_12	0.1542560538
## SBP_Standing_1_12	0.1516111986
## ADP	0.1485126245
## X_12	0.1469666962
## Ferritin_12	0.1435804901
## Potassium_12	0.1403109753
## Chloride_12	0.1381084791
## IWQOL_Esteem5_12	0.1369773346
## Folate_12	0.1364975646
## d_weight_120	0.1271648859
## EQ5D_Scale_12	0.1252458637
## Calcium_12	0.1205832342
## EQ5D_Pain_12	0.1155423981
## BasophilAuto_12	0.1150628197
## TotalProtein_12	0.1124386948
## dfatmass120	0.1118960277
## IWQOL_TotalAVG_12	0.1086268208
<pre>## Waistcircumference_12</pre>	0.1083731712
## hbaf	0.1063066437
## CalciumCorr_12	0.1046559788
## lbgi_12	0.1040617864
## HDL_12	0.1033667269
## IWQOL_TotalScale_12	0.1033486508
## HR_Sitting_1_12	0.1031565501
## Axis3CPM_12	0.1018687775
## q1_sensor_12	0.1004834473
## Axis2AverageCounts_12	0.1003668682
## TANITABMR_12	0.0996092057
## Light_12	0.0991982705
## VectorMagnitudeMaxCounts_12	0.0985374739
## EQ5D_Mobility_12	0.0980589831
## IWQOL_PhysicalScaleAdj_12	0.0967223583
## AUC	0.0961401521
## Glucose_12	0.0939463285
## VectorMagnitudeCPM_12	0.0936117549
## 1ALT_12	0.0907260361
## excursions_over_120_12	0.0893692919
## IWQOL_WorkCount_12	0.0889268707
## CRP_12	0.0876618314
## HR_Standing_1_12	0.0867921789
## lferritin_12	0.0860142534
## IWQOL_Physical2_12	0.0859432252
## GGT_12	0.0854679164
## AVR	0.0851180046
## TANITAIdealBodyWeight_12	0.0832663946
## DQ5D_Selfcare_12	0.0827744631
## EQ5D_Activities_12	0.0826829932
## IWQOL_WorkScaleAdj_12	0.0826776335
## Chloride_6	0.0813876418
## inMVPA_12	0.0807976463
## AUI	0.0800263719
## 1HDL 9	0.0799232912
-	

	StepsAverageCounts_12	0.0798455988
	standard_deviation_12	0.0796192326
	IWQOL_Physical1_12	0.0790526984
	DBP_Standing_1_12	0.0779970295
	12IL17BR_3	0.0761133108
##	TANITAVisceralFatRating_12	0.0757634721
	AUD	0.0751359742
	QUICKI_6	0.0742108352
	DBP_Standing_2_12	0.0741251729
	TANITAWeight_12	0.0738550073
	SBP_Sitting_1_12	0.0711561077
	IPAQ_Sitting_12	0.0706340647
	Axis3MaxCounts_12	0.0704839601
	inLight_12	0.0703143597
	TANITABoneMass_12	0.0698880797
	IL17E_3	0.0682377932
	lIPAQ_TotalMETs_12	0.0666000590
	lInsulin_6	0.0665020106
	q3_sensor_12	0.0664460583
	hbgi_12	0.0659053638
	MPV_12	0.0648478114
	Calorimetry_RMR_12	0.0645130532
##	insulinpmol_9	0.0642840336
	00	0.0634781454
	Axis1AverageCounts_12	0.0632092707
	Neckcircumference_12	0.0629946863
	12IL34_3	0.0628368858
##	percent_time_over_180_0	0.0624424007
##	ADS	0.0607815518
	ADJ	0.0607373139
	percent_time_70_180_12	0.0603688312
	IWQOL_Esteem2_12	0.0598095056
	HR_Standing_2_12	0.0595962240
##	ALT_3	0.0592681341
	12Galectin4_3	0.0592204329
	num_days_good_data_12	0.0587714286
	TotalSedentaryBouts_12	0.0586893895
	LDLCalc_12	0.0586298962
	TANITAMuscleMass_12	0.0583776002
	Triglyceride_12	0.0576417151
	Albumin_12	0.0576161838
	Moderate_12	0.0568924959
	IWQOL_EsteemAVG_12	0.0567351145
	UMALBRatio_12	0.0555085714
	Glucose_6	0.0552458694
	IWQOL_TotalCount_12	0.0546031746
	MCV_12	0.0543182885
	LuxAverageCounts_12	0.0538657580
	total_auc_12	0.0537841125
	TANITAFFM_12	0.0535599845
	Insulin_12	0.0530357143
	12RANK_3	0.0528557626
	Platelet_12	0.0528228571
##	HR_Standing_Avg_12	0.0525417136

## SBP_Standing_Avg_12	0.0515945962
## PKCZ_3	0.0495918367
## HR_Sitting_3_12	0.0495827177
## Hct_12	0.0494444232
## IWQOL_Work4_12	0.0492255848
## median_sensor_12	0.0483246201
## Axis1CPM_12	0.0479846528
<pre>## day_night_sensor_ratio_12</pre>	0.0474575407
## UricAcid_12	0.0472628571
## DBP_Sitting_Avg_12	0.0470263305
## AUN	0.0469070437
## TANITAFatMass_9	0.0467972476
## BQC	0.0460178149
## IWQOL_Physical8_12	0.0459784698
## HR_Standing_1_0	0.0455754667
## HCE003300_3	0.0455736264
## 12phosphoglyceratekinase1_6	0.0449241137
## VitD_9	0.0448193939
## IPAQ_6_Trun_12	0.0446444298
## MCHC_12	0.0445823129
## 12NovH_3	0.0445333333
## 12DPL	0.0445122877
## 12ARTS1_3	0.0442514286
## SBP_Sitting_3_12	0.0442405046
## IWQOL_Esteem7_12	0.0439314286
## Urea_9	0.0438608220
## IPAQ_TotalMETs_12	0.0435590216
## lcrp_9	0.0435590216
## IWQOL_EsteemCount_12	0.0434603175
## nighttime_sd_12	0.0434194888
## SBP_Sitting_2_12	0.0432565940
## 12Trypsin2_3	0.0432000000
## LymphocyteAuto_12	0.0431092437
## SMOC1 3	0.0431092437
## QUICKI_12	0.0431092437
## ROBO3_3	0.0430951012
## Fibro_E_IQR_12	0.0425671075
## Sodium_9	0.0424779388
## j_index_12	0.0422323308
## MonocyteAuto_12	0.0420943915
## 12KLRF1 6	0.0420761905
## IWQOL_DistressCount_12	0.0420622222
## gmi_12	0.0417794585
## 12NANOG_3	0.0417794585
## DCSIGN_3	0.0414398681
## HDL_9	0.0414268908
## TotalSedentaryBreaks_12	0.0414228571
## daytime_sd_0	0.0413731302
## lcrp_12	0.0413063335
## min_spent_over_180_12	0.041000000
## TANITABMI_9	0.0411463203
## QUICKI_3	0.0411400200
## DBP_Standing_1_9	0.0407403029
## nighttime_avg_sens_glucose_12	0.0406352403
	5.5100002100

## HR_Standing_1_LAST	0.0405313668
## RDWCV_12	0.0402336508
## HDL_0	0.0396439394
## kcals_12	0.0394909430
## 12EF1beta_3	0.0394651515
## 12sTie1_3	0.0388065542
## OSM_6	0.0387292208
## UricAcid_6	0.0387108590
## 12LIN7B_6	0.0385714286
## Fibro_CAP_IQR_12	0.0384881563
## LAG3 3	0.0382015996
## GITR_3	0.0381921501
## LDLCalc_9	0.0381197700
## percent_time_over_140_12	0.0376414754
## Waistcircumference_6	0.0375669841
## EPOR 6	0.0375669841
## IWQOL_WorkScale_12	0.0374857143
## NANOG_3	0.0374010582
## 1IPAQ_TotalMETs_Trun_12	0.0374010382
## UCreatinine_9	0.0372173493
_	
## Fibro_E_IQRE_med_6	0.0371159030
## ERBB4_3	0.0368970072
## IGFI_3	0.0368703934
## inSedentary_12	0.0368556064
## percent_time_70_180_day_12	0.0367714286
## C1r_6	0.0367682382
## KREM2_3	0.0366149660
## 1AST_3	0.0364850959
## min_sensor_12	0.0361869313
## IWQOL_Physical7_12	0.0359863946
## 12LEG9_3	0.0359863946
## DBP_Standing_Avg_9	0.0358914233
## ARY	0.0358400000
## CPeptide_6	0.0356835979
## UMicroalb_12	0.0355032468
## 12PTHrP_0	0.0354470978
## IL1R4_6	0.0352149946
## IL8_3	0.0351673469
## GGT_9	0.0351620771
## Height_12	0.0348923077
## IWQOL_Esteem1_12	0.0348923077
## Insulin_9	0.0348048060
## RD	0.0347657143
## HR_Sitting_Avg_0	0.0346599895
## total_sensor_readings_12	0.0342857143
## IWQOL_Physical3_12	0.0341702656
## NogoReceptor_6	0.0341536508
## 120PG_3	0.0338800000
## 12CLC4K_3	0.0338800000
## StepsPerMinute_3	0.0336620647
## MDRD4_12	0.0336561404
## MDRD4_9	0.0336119658
## RBC_12	0.0336008658
## IWQOL_DistressScaleAdj_12	0.0334955423
· =	

	wtloss_0_9	0.0332921325
	12MAPK5_6	0.0332893475
	MBL_12	0.0332247619
	Averagekcalsperday_12	0.0331523810
	average_auc_per_day_12	0.0329456243
	SBP_Standing_1_6	0.0329063246
	IWQOL_Distress1_12	0.0329063246
	BilirubinT_9	0.0327788360
	HR_Sitting_Avg_3	0.0327690378
	VectorMagnitudeMaxCounts_3	0.0326859023
	12PH_3	0.0325814536
	12PHI_6	0.0325520534
	IWQOL_EsteemScale_12	0.0324521893
	Albumin_9	0.0322432140
	HR_Sitting_2_12	0.0321511971
	Endocan_3	0.0320380952
	modd_12	0.0317698853
	nighttime_min_sens_glucose_12	0.0316749917
	Potassium_3	0.0316483516
	TSH_12	0.0315584416
	COMPLETED12M	0.0315584416
	CathepsinS_6	0.0314903097
	12Eotaxin2_3	0.0314201299
	HR_Sitting_2_0	0.0313848767
##	12GREM1_6	0.0313003175
	max_sensor_0	0.0312169082
##	TANITAFatMass_12	0.0311537415
	PDE7A_3	0.0311537415
##	CD47_3	0.0311074197
	12SIG14_3	0.0309892587
	12FABP_3	0.0309428571
	diabmedl	0.0307318478
	HIV2Rev_3	0.0306884058
##	HR_Standing_1_9	0.0306342160
##	ERBB2_3	0.0305873196
	TANITABMI_12	0.0304295482
	Angiotensinogen_3	0.0302482116
	IWQOL_Physical4_12	0.0302120259
	HPG_3	0.0302120259
##	JAG1_0	0.0302109091
	AlkPhos_9	0.0301474274
	12ERBB1_3	0.0300787213
	12EDA_3	0.0298666667
	MRCKB_0	0.0296817453
##	CKDEPI_9	0.0296435795
##	Calcium_9	0.0296228571
	efficeincy	0.0295140147
##	KIRR3_0	0.0294617792
##	12ILT2_6	0.0294617383
##	12EPOR_6	0.0293486471
##	DBP_Sitting_3_12	0.0293261456
##	12kallikrein5_0	0.0293125541
	DBP_Sitting_2_9	0.0292640693
##	NumberofEpochs_3	0.0292163338

## TANITAMusc	leMass_9	0.0291432974
## 12SDF1_6		0.0290303433
## IWQOL_Work	AVG_12	0.0289821468
## modd_0		0.0289725927
## IL1RAcP_3		0.0289598344
## 12HCE00034	2_0	0.0288688172
## wtloss3_0		0.0286754460
## nighttime	max_sens_glucose_12	0.0285714286
## IgA_0	9 -	0.0284731485
## Fibro_E_IQ	R 3	0.0284424832
## Axis2MaxCo	-	0.0283942857
## 12FGF9_3		0.0283102041
## BSP 0		0.0281955556
## IL17RC_3		0.0280941210
	over_200_night_0	0.0280472050
## IWQOL_Tota		0.0278146032
## pc5	ibcalendj_12	0.0276140032
## cv 12		0.0277341615
## CV_12 ## IL17D_3		0.0276468272
## 1E17D_3 ## DPK		0.0275427937
## DHH 3		0.0275427937
## 12H6ST1 3		0.0275093027
-		
## 12ENA78_3		0.0274293233
## IL17B_0	050 1 0	0.0273147845
	over_250_day_0	0.0272605301
-	Freedson1998Bou	0.0269968969
## Axis1CPM_0		0.0268585659
## ApoB_3		0.0265863169
## OSM_3		0.0265722743
## 12PIK3CAPI	_	0.0264727273
## 12RNaseH1_	12	0.0264208683
## RDWCV_3		0.0264142857
## MCH_9		0.0261120879
## Trypsin2_0		0.0260980392
## min_spent_	70_180_12	0.0260923550
## FGF8A_6		0.0259453416
## min_spent_		0.0258705549
## IWQOL_Phys		0.0256530021
## auc_over_1	_	0.0254417701
## gmihba1_12		0.0254021978
## C3b_6		0.0252457143
## 12CD177_3		0.0252457143
## 12MATN2_0		0.0251101466
## 12ZAP70_3		0.0250914729
## WTLOSS_12_	LAST	0.0250381618
## 12RSP04_12		0.0248579019
## 12FactorH_	6	0.0248083857
## Time_12		0.0247714286
## CD27_0		0.0245805036
## EMAP2_0		0.0244577589
## 12tPA_0		0.0243599331
## 1ALT_9		0.0240790801
## average_se	nsor_0	0.0240593381
## 1PTH_6	-	0.0236680933
_		

## 12pTEN		0 .	.0234782609
## wtloss		0 .	.0233232175
## 12TACI	_6	0 .	.0232133256
## 12Grob	g_6	0 .	.0231580827
## min_sp	ent_over_180_night_0	0 .	.0231092200
## HIV2Re	v_0	0 .	.0230658979
## SIRT2_	0	0 .	.0229340294
## d_weig	ht_30	0 .	.0228299657
## 12Alph	aenolase_3	0 .	.0227781936
## DMedLa	st	0 .	.0227396428
## C8_0		0 .	.0227045455
## Tropon	inIskeletalfasttwitc	0 .	.0225894599
_	tiplasmin_0	0 .	.0225408163
## TGFb1_	6	0.	.0223940788
## 12Fact	orH 12	0.	.0222244462
## Time_3	_	0.	.0220835350
## Rb 6		0.	.0220020274
## FT4_9		0.	.0218272464
## 12CHST	6 3	0.	.0217666667
## MIA_O	-	0.	.0215485788
## EPHA3_	0	0.	.0215238095
## JAML1_		0.	.0214308943
## JAG1 3			.0213794967
## Ckine		0	.0213747262
## Axis2C			.0213186207
## cMyc_0	_		.0211314286
## 12ADAM			.0211038961
## TANITA	_		.0210380952
## NOTC2_	_		.0210173737
## NKp30_			.0210168961
	rd_deviation_0		.0208814192
	ekcalsperhour_12		.0206741201
## HR_Sit	<u>-</u>		.0206398349
## SDF1_6	9		.0205714286
_	ting_3_LAST		.0205512746
## C2_3	CIUE_O_LMDI	-	.0205312740
## 02_5 ## 12IL1R	AcP 6		.0204861776
## Rb_0	ACI_O		.0203636364
## 12BMP7	0		.0202365950
## daytim			.0202505950
## daytim			.0199980471
## 12DCSI			.0199584576
	_		.0199504570
## 12SEPR	_		.0196522096
## MSPR_0			
	stimulatedPDE_6		.0197676190
## GFRa3_			.0195222433
## AREG_0			.0194400020
## dmuscl	emass120		.0194181145
## MCH_0	1: 0		.0193050712
## Proper	_		.0192000000
## kallik	-		.0190939405
## 12IL8_			.0190924959
## inMVPA	_		.0190855731
## 12a1An	titrypsin_0	0.	.0190400458

## TECK_O	0.0190272364
## 12MFGM_12	0.0189943666
## nighttime_sd_0	0.0189729576
## MMP2_12	0.0189342857
## average_auc_per_day_0	0.0188833183
## inMVPA_O	0.0188558935
## 12TroponinT_6	0.0188053391
## 12IL5Ra_12	0.0187500000
## 1ALT_3	0.0185818566
## 12IL1F7_0	0.0185444444
## 12Aggrecan_0	0.0184666167
## Averagekcalsperday_3	0.0182430155
## 12IMB1_12	0.0182400000
## percent_time_over_250_0	0.0182237563
## 12Semaphorin6A_12	0.0182101122
## LivinB_0	0.0182052632
## PDK1 0	0.0182052632
## LDLR_3	0.0182050622
## 12PAFAH_3	0.0181313040
## 12NID2_0	0.0180846561
## HCE000342_3	0.0180333952
## HR_Standing_1_screening	0.0180236912
## 12SCGFbeta_0	0.0180230912
_	0.0179658653
## Diabetesmedno_12	0.0179036063
## 12SET_12	0.0179356061
## 12CSK21_0	0.0179259259
## 12sLeptinR_6	
## SPTA2_6	0.0178775744
## PESC_6	0.0178598550
## SLIK1_12	0.0178571429
## MIP5_0	0.0178503775
## 12ghrelin_12	0.017777778
## CathepsinH_0	0.0177509549
## LGMN_O	0.0177141674
## RDWCV_O	0.0176989967
## 12PSP_0	0.0176075650
## Triglyceride_3	0.0175837321
## Diabetesmedno_LAST	0.0175192858
## 12FTCD_3	0.0175157895
## 12PSP_6	0.0175108616
## Sulphonylurea_0	0.0175075988
## 12LRP8_3	0.0174743187
## MDC_0	0.0174618182
## FGF23_6	0.0174545455
## CONA1_3	0.0174409938
## sqrtghrelin_0	0.0174180165
## 1HDL_12	0.0173779904
## 12GFAP_12	0.0173051338
## 12Testican1_3	0.0173008625
## 12ERBB2_6	0.0172972973
## estimated_a1c_12	0.0172902963
## 12DLC8_0	0.0172386364
## QUICKI_9	0.0172023810
## 12IL17F_0	0.0171521336

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	12TF_6	0.0170086682
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##	12HistoneH12_0	0.0169477241
##	12EGVEGF_12	0.0169312169
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	12MK13_12	0.0164131231
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	12EDA_0	0.0162331081
	12Chk2_6	0.0161910015
	ALT_9	0.0161538462
	12IFNlambda2_6	0.0161333333
	KI3S1_6	0.0161293743
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	TCPTP_3	0.0160535117
	Artemin_0	0.0160190476
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	12FGR_0	0.0158142857
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	12CDK2cyclinA_12	0.0156829837
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	12Lactoferrin_6	0.0153035923
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	12EGVEGF_0	0.0151439259
	Hgb_12	0.0151250000
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## Neurotrophin3_0	0.0150146628
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## IFNlambda2_0	0.0149345455
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## 12AMPKa1b1g1_6	0.0149286379
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## 12sRANKL_0	0.0147917834
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## IMDH1_0	0.0147272727
## 12Leptin_6	0.0147272727
## 12GPDA_0	0.0147200000
## 12IL17RC_3	0.0146994774
## IF4A3_0	0.0146938776
## aldolaseA_3	0.0146938776
## SBP_Sitting_1_6	0.0146866930
## HR_Sitting_Avg_12	0.0146403181
## 12HGH_12	0.0146352543
<pre>## percent_time_over_250_12</pre>	0.0146086957
## 12GX_3	0.0145833333
## 12ITIheavychainH4_3	0.0145832831
<pre>## a1Antitrypsin_6</pre>	0.0145777778
## HCG_6	0.0145714286
## StepsMaxCounts_3	0.0145185185
## 12EMAP2_0	0.0144420493
## AurorakinaseA_12	0.0144390244
## 12TGFb1_6	0.0144000000
## 12HSP70protein8_3	0.0143013431
## SPD_0	0.0142857143
## Notch3_0	0.0142281481
## HR_Sitting_1_LAST	0.0142145079
## 12FGF17_3	0.0141949897
## inModerate_12	0.0141842105
## IWQOL_TotalScale_0	0.0141493268
## 12ERK1_3	0.0141243221
## HR_Sitting_1_3	0.0140833333
## 12PIK3CAPIK3R1_3	0.0140586953
## KI3S1_3	0.0140489510
## IL2sRa_0	0.0140455006
## 12ING1_0	0.0139808407
## 12SNP25_6	0.0139758125
## SIG14_3	0.0139706239
## Trypsin_0	0.0139653637
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## 12JAG2_3	0.0138888889
## Gelsolin_12	0.0138857143
## ERBB2_6	0.0138775758
## carbonicanhydraseII_6	0.0138753388
## 1IPAQ_ModerateMETs_Trun_12	0.0137931034
## Rb_3	0.0137619048
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## 12SMAD3_3	0.0137142857
## 12HCC1_3	0.0137142857
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## SBP_Standing_2_6
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## bFGF_6
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## 12FactorB 6
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## TNR4 6
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## MDC 6
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## 12NXPH1_3
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## GPNMB O
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## 12C9_12
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## Fasligandsoluble_3
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## 12DSC2_3
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## 12ALCAM_3
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## 12IgG_0
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## WBC 12
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## estimated_a1c_0
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## 12BQC
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## 12FCAR_6
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## 12NSF1C 0
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## 12Ficolin3_3
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## CD97 0
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## Properdin_0
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## 12CD30_0
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## HSP90ab_0
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## hnRNPA2B1_0
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## 12Cardiotrophin1_0
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## CoagulationFactorVII_0
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## 12PDE5A_3
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## 1HDL_0
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## TRAILR1_3
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## 12SHPS1 6
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## GIB 0
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## lInsulin 0
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## EF1beta_3
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## d_hr120
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## 12HXK1_0
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## 12GMCSF 12
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## ADR
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## 12Soggy1_0
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## MMP13_3
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## gmihba1_0
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## SSRP1_0
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## ARMEL_O
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## CD63_0
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## 1AST_0
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## lnghrelin_0
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## percent_time_over_180_night_12
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## 12IL18BPa 0
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## IL17BR 6
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## 12FractalkineCX3CL1_0
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## RASA1_3
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## 12IgD 6
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## 12B7H1 12
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## 12TMA 0
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## FGF12 0
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## 12IGFBP3 0
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## 12CoagulationFactorV_6
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## 1ALT_0
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## 12LRRK2_0
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## 12IL12Rb1_6
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## 12KI3S1_6
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## IPAQ_3_12
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## 12MBL_0
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## percent_time_over_200_0
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## WNK3 3
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## 12WISP3_12
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## Cadherin6 6
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## HDL_3
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## min_spent_over_180_0
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## HR_Standing_Avg_9
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## percent_time_over_200_day_12
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## 12STAT1 0
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## 12ThrombopoietinReceptor_0
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## MCV_O
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## TNR4_0
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## 12Alphaenolase_0
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## BilirubinT_3
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## 12CYTT_0
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## DGE
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## average_sensor_12
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## 12TSP4_0
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## HR_Standing_2_LAST
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## LRRK2 0
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## 12ADAMTS5 6
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## C2 0
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## HCE003300 0
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## C9_6
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## TANITABMI_screening
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## SPARCL1 6
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## OPG 3
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## Secretin_3
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## 12Rb_3
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## resistin_12
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## 12RAP_0
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## NXPH1_6
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## PHI_3
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## 12RXFP1_3
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## 12HCE003167_0
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## TACI_6
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## Diabetesmedno 6
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##	12IL8_0	0.0114461318
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##	12IL7_6	0.0112800000
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##	12ProteinC_0	0.0109090909
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##	12IL1RAcP_0	0.0108724528
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##	q3_sensor_0	0.0107675785

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	12PAK7_3	0.0107236479
##	12IL17D_3	0.0107236479
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##	12HAO_12	0.0107017189
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	FYN_3	0.0106730159
	12MDC_0	0.0106730159
##	TXD12_3	0.0106666667
	12S100A7_3	0.0106666667
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	12I309_12	0.0104347826
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	12RELT_6	0.0102722222
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##	12hnRNPAB_3	0.0093605442
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	WBC_9	0.0090685714
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	CLC4K_0	0.0090000000
	SCGFbeta_12	0.0090000000
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	PAPPA O	0.0089629630
	TGFb3_0	0.0088944444
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	CalcineurinBa 0	0.0088888889
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	12EGF 6	0.0088494983
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	12HIV2Rev_3	0.0087272727
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	12BCAM_0	0.0085838298
	12Discoidindomainreceptor2_0	0.0085815603
	12CNTN2_3	0.0085468115
	Cardiotrophin1_6	0.0084992026
##	IWQOL_Physical2_0	0.0084521739

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## 12MMP17 3
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## 12PDGFBB_3
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## 12TNR4_3	0.0075897436
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## 12HistoneH2Az_3	0.0074285714
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## 12NG36_3	0.0073504125
## 12CD59_6	0.0073504125
## lcrp_6	0.0073504125
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## 12RSP04_6	0.0073369565
## 12LRRK2_12	0.0073369565
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## 12IFNb_6	0.0073333333
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## 12EphB4_6	0.0073147392
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## 12IL4 6	0.0072930233
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## 12IL17sR_6	0.0072428571
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## 12TPSB2_0	0.0072380952
## 12DRG1_6	0.0072380952
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## NXPH1_3
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## 12TopoisomeraseI 3
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## 12CarbonicanhydraseVII 3
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## 1MIC1_0
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## MMP10_12
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## percent_time_over_250_night_12
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## 12GV_3
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## 12PDE9A 6
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	cJun_6	0.0062769231
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## SBP_Standing_Avg_6
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## Duration_Month
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## CDK8cyclinC_0
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## B7H1_12
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## sLeptinR_12
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## 12HistoneH2Az_0
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## 12IgA_3
## 12CDK2cyclinA_3
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## 12MOZ_6
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## 12sLRP1_6
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## insulinpmol 6
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## Proteindisulfideisomerase_0
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## Calcineurin 0
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## IgD_0
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## LG3BP 0
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## GV_6
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## 12SonicHedgehog 0
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## 12BMP1 0
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## 12HGFA 0
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## 12GPC5_3
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## 12IL18BPa_6
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## 12HPG_12
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## pc11
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## HR_Standing_1_6
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## SMAD3_3
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## LY86_3
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## FCG2A_12
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## 12IL4 0
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## 12NET4 3
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## 12RSP02 6
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## 12Elafin_6
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## IPAQ_7_6
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## HSP70_0
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## IF4G2 0
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## PBEF 0
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## Semaphorin6A_0
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## CD5L_3
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## TSG6_3
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## CAMK2A_6
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## GOR
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## LY9_12
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## Transferrin_12
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## 12CD97_0
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## 12IL19_0
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## 12BOC_3
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## 12ProteindisulfideisomeraseA3_6
## 12EphrinA4_12
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## HR_Standing_Avg_3	0.0039000000
## Glucose_3	0.0039000000
## IPAQ_Sitting_6_LOCF	0.0039000000
## IL16_0	0.0039000000
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## LYN_3	0.0039000000
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## HR_Sitting_Avg_screening	0.0038974359
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## min_spent_over_200_12	0.0038974359
## EphrinB3_3	0.0038974359
## DLRB1_3	0.0038974359
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## IF4A3_6	0.0038974359
## IGFIsR_12	
## RANK_12	0.0038974359
## 12CyclophilinA_0	0.0038974359
## 12PH_6	0.0038974359
## IL22BP_0	0.0038947368
## CYTN_O	0.0038947368
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## JAK2_12	0.0038947368
## 12CD22_0	0.0038947368
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## 12IL15Ra_0	0.0038947368
## 12TGFb2_0	0.0038947368
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## 12BPI_12	0.0038947368
## KEAP1_0	0.0038918919
## PSD7_12	0.0038918919
## 12BNP32_0	0.0038918919
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## 12CSF1_3	0.0038918919
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## 12IL1sRI_0	0.0038888889
## 12CystatinC_6	0.0038888889
## 12Properdin_12	0.0038888889

	12ApoD_12	0.0038888889
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	SGTA_3	0.0038857143
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	12EphrinA2_6	0.0038857143
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	Myokinasehuman_3	0.0038823529
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##	12ActivinRIB_0	0.0038823529
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##	12RAC3_12	0.0038823529
##	TANITATBW_9	0.0038787879
	IgE_0	0.0038787879
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##	12PRL_3	0.0038787879
##	UricAcid_0	0.0038750000
##	FGFR2_0	0.0038750000
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	NET1_3	0.0038620690
##	120steopontin_0	0.0038620690
##	12EphA1_12	0.0038620690
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##	bEndorphin_0	0.0038571429
##	OLR1_O	0.0038571429
##	NKp44_6	0.0038571429
##	IDE_6	0.0038571429
##	<pre>1IPAQ_ModerateMETs_6</pre>	0.0038571429
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	Tropomyosin4_0	0.0038461538
	Azurocidin_3	0.0038461538
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## 12TrkA_12	0.0038400000
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## MAPK2_6	0.0038333333
## 12FN13_0	0.0038333333
## 12EFNB1_12	0.0038333333
## TimeToEat_Lunch	0.0038260870
## insulinpmol_12	0.0038260870
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## CHM	0.0038260870
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## IL17RD_6	0.0038260870
## SORC2_6	0.0038260870
## 12PLPP_0	0.0038260870
## 12IL1a_6	0.0038260870
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## UCreatinine_12	0.0038181818
## max_sensor_12	0.0038181818
## EMR2_0	0.0038181818
## TCCR_0	0.0038181818
## NogoReceptor_3	0.0038181818
## TGFb2_6	0.0038181818
## XG	0.0038095238
## CJQ	0.0038095238
## Thyroglobulin_0	0.0038095238
## DR6_0	0.0038095238
## IL17RD_3	0.0038095238
## HCE000342_6	0.0038095238
## S100A12_6	0.0038095238
## DAPK2_12	0.0038095238
## PDGFAA_12	0.0038095238
## 12ER_0	0.0038095238
## 12FSH_0	0.0038095238
## 12CNTN2_0	0.0038095238
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## 12Trefoilfactor2_3	0.0038095238
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## 12ADAMTS4_12	0.0038095238
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## inLight_3	0.0038000000
## Freedson1998Bouts_6	0.0038000000
## TGFbRIII_0	0.0038000000
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## SDF1_12	0.0038000000
## 12NUDC3_0	0.0038000000
## 12ActivinAB_3	0.0038000000
## 12PARC_12	0.0038000000
## 12Testican2_12	0.0038000000
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## ZNRF3_0	0.0037894737
## BLC_0	0.0037894737
<u>-</u> -	2.300.002101

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## Cadherin2 0
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## CTAPIII 3
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## Cadherin12 6
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## FAM107B_12
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## lIPAQ_WalkingMETs_Trun_12
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## 12HINT1 12
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## CoagulationFactorX_0
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## CoagulationFactorIXab_6
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## 12FCN1 0
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## 12bECGF_3
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## 12F1t3_12
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## CHL1_0
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## percent_time_over_180_day_12
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## CKDEPI_12
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## 12LTBP4_3
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## 12sRAGE_3
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## 12Mcl1_6
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## 12IL17B 12
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## 12CATF_12
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## TFPI 0
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## 12UBE2N_0
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## 12Properdin_3
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## 12Peroxiredoxin5_6
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## d weight 60
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## 12PCSK7 3
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## 12PGCB 12
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## prostaticbindingprotein_0
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## BLC_12
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## 12ADAMTS5_0
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## IFNg_3
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## SonicHedgehog_12
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## LymphocyteAuto_6
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## 12CAMK1_6
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## MPV_9
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## NET1 12
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## 12LGMN 12
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## 12LG3BP 12
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## 12IL18Rb_3
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## IL3 0
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## HSP60 3
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## MAPK5 6
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## 12ULBP2_0
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## MICA_3
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## ThyroxineBindingGlobulin_12
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## 12CONA1_3
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## Fucosyltransferase3_0
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## B7_12
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## 12RXFP1_0
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## 12MO2R1_6
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## TGFb1_0
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## 12ASAHL 0
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## 12ALK1 3
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## Creatinine_9	0.0036923077
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## 12IGFBP7_0	0.0036786787
## 12IL1R4_0	0.0036666667
<pre>## insulinpmol_0</pre>	0.0036506239
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## 12TNFSF18_12	0.0036506239
## SMAD2_6	0.0036363636
## min_spent_70_180_night_12	0.0036219512
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## PDE4D_6	0.0036000000
## 12MK08_3	0.0036000000
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## CD48_0	0.0035818686
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## 12HCK_0	0.0035612536
## LIN7B_0	0.003555556
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## 12HAI1_12	0.003555556
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## 12Thymidinekinase_6	0.0035266667
## 120PG_12	0.0035266667
## 12AnnexinV_0	0.0035102564
## 12LaminB1_12	0.0035072464
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## 12C8_12	0.0035000000
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## 12IL15Ra_12	0.0034420168
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## TANITAMetabolicAge_6	0.0034285714
## SHC1_0	0.0034285714
## SECTM1_3	0.0034285714
## SPINT2_3	0.0034285714
## PSA2_3	0.0034285714
## EGFRvIII_3	0.0034285714
## MIP5_6	0.0034285714
## 12EphrinA5_3	0.0034285714
## 12MIC1_3	0.0034285714
## 12Glucagon_3	0.0034285714
## 12NogoReceptor_3	0.0034285714
## 12RXFP1_6	0.0034285714
## 12HBEGF_6	0.0034285714
## 12WISP1_12	0.0034285714
## C5_0	0.0034105263
## 12Semaphorin6A_3	0.0034105263
## MPV_6	0.0033333333
· - ·	2.2.20000000

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## Axis3Counts_0
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## Mcl1_0
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## B7 0
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## TNFSF18_0
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## ProteinC 0
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## Spondin1 0
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## IntegrinaVb5 0
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## PARK7_0
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## CathepsinG_3
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## PERL_3
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## NterminalproBNP_3
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## HIF1a_12
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## NKG2D_12
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## hnRNPK_12
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## 12BCMA_0
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## 12ADAMTS4_0
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## 12MRC2_0
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## 12transcriptionfactorMLR1isofo
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## 12MK11_3
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## 12DAN 12
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## 12IRE
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## 12RBP_12
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## 12TRAILR2_12
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## TANITAMuscleMass_screening
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## IWQOL_Sex4_0
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## IWQOL_Physical11_12
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## excursions_over_140_0
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## S100A4 0
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## MK08 0
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## MDM2 0
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## GRN_O
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## HCE000414 3
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## MSPR_3
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## MP2K4 3
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## ABL2 3
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## SHC1 3
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## RUXF_3
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## INGR2_3
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## ApoptosisregulatorBclW_6
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## PF4_6
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## ART_6
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## ER_6
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## Nectinlikeprotein2_6
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## EphA5_6
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## CyclophilinA_6
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## Kras_6
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## CarbonicanhydraseVII_12
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## 12VEGF_0	0.0032000000
## 12GA7331protein_0	0.0032000000
## 12PAI1_0	0.0032000000
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## 12Neurotrophin3_0	0.0032000000
## 12Ku70_3	0.0032000000
## 12PTP1C_3	0.0032000000
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## 12CD48_6	0.0032000000
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## DBP_Standing_2_3	0.0030000000
## DBP_Sitting_2_12	0.0030000000
## IWQOL_Esteem2_0	0.0030000000
## Time_0	0.0030000000
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## Marapsin_0	0.0030000000
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## sRAGE_3	0.0030000000
## C6_3	0.0030000000
## CK2A1B_3	0.0030000000
## BMP7 6	0.0030000000
## ASAH2_6	0.0030000000
## CarbonicanhydraseXIII_6	0.0030000000
## SPD 6	0.0030000000
## PDE3A_6	0.0030000000
## PACAP38_6	0.0030000000
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## GFRa1_12	0.0030000000
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## 12Carbonicanhydrase9_0	0.0030000000
## 12Cytochromec_3	0.0030000000
## 12HIPK3 3	0.0030000000
## 12Caspase3_3	0.0030000000
## 12Secretin_3	0.003000000
## 12CTAPIII_3	0.0030000000
## 1201AF111_3 ## 12MIS_3	0.0030000000
## 12M15_5 ## 12PKCZ_6	0.003000000
## 12PKCZ_6 ## 12MMP17_6	0.003000000
## 12Artemin_6	0.003000000
## 12Mrtemin_6 ## 12Ubiquitin_6	0.003000000
"" TSONIAUTOTHO	0.0030000000

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	PTH_12	0.0026666667
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	AUE	0.0026666667
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	excursions_under_70_12	0.0026666667
	DBP_Standing_Avg_LAST	0.0026666667
	NormScale_1_0	0.0026666667
	Vitronectin_0	0.0026666667
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	CoagulationFactorXa_0	0.0026666667
	IDUA_O	0.0026666667
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	CTGF_3	0.0026666667
	Galectin2_3	0.0026666667
	BLC_3	0.0026666667
	discoidindomainreceptor1_3	0.0026666667
	JAML1_3	0.0026666667
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	PYY_6	0.0026666667
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	12LSAMP_3	0.0026666667
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	12MMEL2_3	0.0026666667
	12SHPS1_3	0.0026666667
	12BAD_3	0.0026666667
	12WISP3_3	0.0026666667
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	12HIDH_6	0.0026666667
	12TRAILR1_6	0.0026666667
##	12HCG_6	0.0026666667
	12GP1BA_6	0.0026666667
	12PDE7A_6	0.0026666667
	12IFNgR1_6	0.0026666667
##	12GDF9_12	0.0026666667

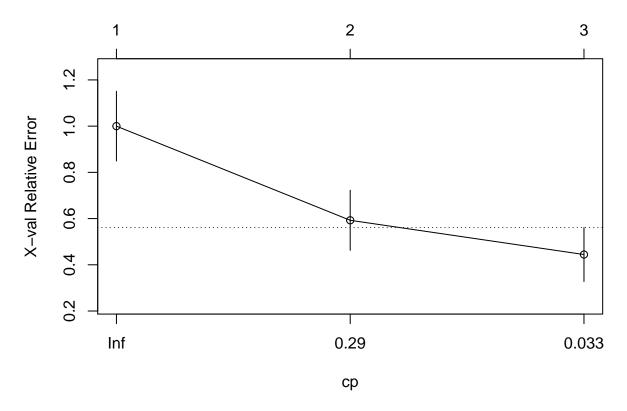
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## 12KPCT 12
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## Caspase3_0
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## ENP
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## 12MK08_0
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## 12GPDA 3
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## 12CD30Ligand_0
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## IFNgR1_0
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## SIRT2_3
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## TSG6_0
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## CaMKKalpha_0
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## FLI_0
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## Axis1CPM_3
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## IPAQ_TotalMETs_Trun_12_LOCF
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```

CART

Classification

```
library(rpart)
library(rpart.plot)
fit1 <- rpart(remission~., method="class", data=data)</pre>
printcp(fit1)
## Classification tree:
## rpart(formula = remission ~ ., data = data, method = "class")
##
## Variables actually used in tree construction:
## [1] Glucose_12 HDL_12
## Root node error: 27/70 = 0.38571
##
## n= 70
##
          CP nsplit rel error xerror
## 1 0.74074
              0 1.00000 1.00000 0.15084
## 2 0.11111
                  1 0.25926 0.59259 0.13012
                  2 0.14815 0.44444 0.11679
## 3 0.01000
plotcp(fit1)
```



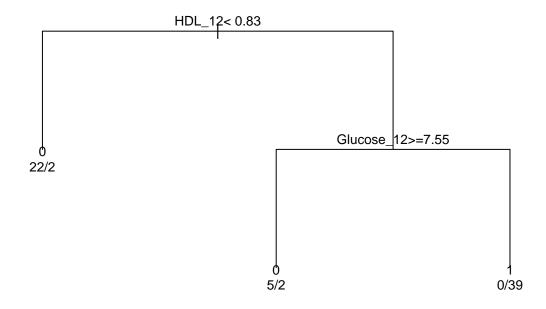


summary(fit1)

```
## Call:
## rpart(formula = remission ~ ., data = data, method = "class")
##
##
            CP nsplit rel error
                    0 1.0000000 1.0000000 0.1508354
## 1 0.7407407
## 2 0.1111111
                    1 0.2592593 0.5925926 0.1301200
## 3 0.0100000
                    2 0.1481481 0.4444444 0.1167863
##
##
   Variable importance
##
                  HDL_12
                                    dfatmass120
                                                        Fibro_E_med_12
##
                      16
                                             13
                                                                    13
##
        SBP_Sitting_1_12
                                      Height_12 Waistcircumference_12
##
                      13
                                             12
                                                                    12
##
                  FBG_12
                                     Glucose_12
                                                                  hbaf
##
                        5
                                              5
                                                                     3
##
               Glucose 3
                                      Glucose_6
                                                                 IL4 0
                        3
                                               3
##
                                                                     3
##
                                     complexity param=0.7407407
## Node number 1: 70 observations,
##
     predicted class=1 expected loss=0.3857143 P(node) =1
##
       class counts:
                         27
                               43
##
      probabilities: 0.386 0.614
     left son=2 (24 obs) right son=3 (46 obs)
##
```

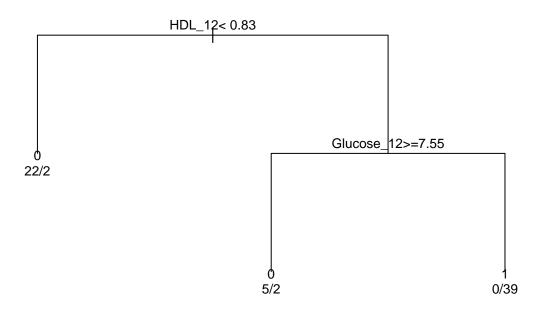
```
##
     Primary splits:
##
                         < 0.83
                                    to the left, improve=20.59172, (0 missing)
         HDL 12
##
         BilirubinT 12
                         < 4.2
                                    to the left, improve=19.68123, (0 missing)
         TotalProtein_12 < 66
                                                  improve=19.68123, (0 missing)
##
                                    to the left,
##
         UricAcid_12
                         < 97.5
                                    to the left,
                                                  improve=19.68123, (0 missing)
##
         Cholesterol 12 < 3.05
                                                  improve=19.68123, (0 missing)
                                    to the left,
##
     Surrogate splits:
##
         SBP_Sitting_1_12
                               < 104.5
                                          to the left, agree=0.943, adj=0.833, (0 split)
##
         Fibro E med 12
                               < 2.533333 to the left, agree=0.943, adj=0.833, (0 split)
##
         dfatmass120
                               < -0.55
                                          to the right, agree=0.943, adj=0.833, (0 split)
##
         Height_12
                               < 75
                                          to the left, agree=0.929, adj=0.792, (0 split)
                                          to the left, agree=0.929, adj=0.792, (0 split)
##
         Waistcircumference_12 < 40.95
##
## Node number 2: 24 observations
##
     predicted class=0 expected loss=0.08333333 P(node) =0.3428571
##
       class counts:
                        22
                               2
##
      probabilities: 0.917 0.083
##
                                      complexity param=0.1111111
## Node number 3: 46 observations,
     predicted class=1 expected loss=0.1086957 P(node) =0.6571429
##
##
       class counts:
                         5
                              41
##
     probabilities: 0.109 0.891
##
     left son=6 (7 obs) right son=7 (39 obs)
     Primary splits:
##
##
         Glucose 12
                          < 7.55
                                     to the right, improve=6.055901, (0 missing)
##
         FBG_12
                          < 7.55
                                     to the right, improve=6.055901, (0 missing)
##
                          < 7.2
                                     to the right, improve=4.468599, (0 missing)
         Glucose_3
##
         average_sensor_0 < 201.1093 to the right, improve=3.535754, (0 missing)
##
         estimated_a1c_0 < 8.65
                                     to the right, improve=3.535754, (0 missing)
##
     Surrogate splits:
##
         FBG_12
                   < 7.55
                              to the right, agree=1.000, adj=1.000, (0 split)
##
         hbaf
                   < 6.55
                              to the right, agree=0.957, adj=0.714, (0 split)
##
         Glucose_3 < 6.95
                              to the right, agree=0.935, adj=0.571, (0 split)
##
                              to the right, agree=0.935, adj=0.571, (0 split)
         Glucose_6 < 7.7
##
         IL4 0
                  < 408.7
                              to the left, agree=0.935, adj=0.571, (0 split)
##
## Node number 6: 7 observations
##
     predicted class=0 expected loss=0.2857143 P(node) =0.1
##
       class counts:
                         5
                               2
##
      probabilities: 0.714 0.286
##
## Node number 7: 39 observations
##
    predicted class=1 expected loss=0 P(node) =0.5571429
##
       class counts:
                         0
      probabilities: 0.000 1.000
{plot(fit1, uniform=TRUE, main="Classification Tree")
text(fit1, use.n=TRUE, xpd=TRUE, cex=.8)}
```

Classification Tree



```
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)}</pre>
```

Pruned Classification Tree

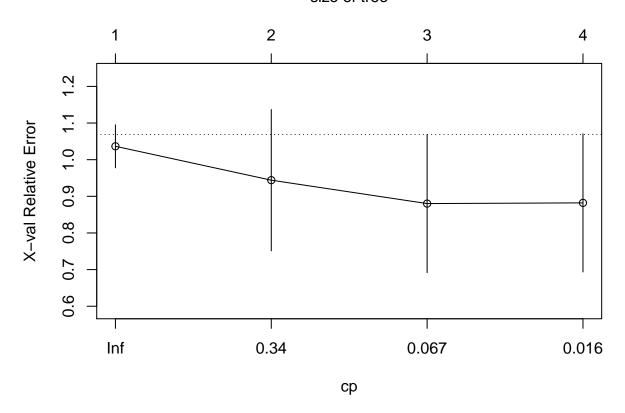


Regression

```
fit <- rpart(remission~., method="anova", data=data)</pre>
printcp(fit)
##
## Regression tree:
## rpart(formula = remission ~ ., data = data, method = "anova")
##
## Variables actually used in tree construction:
## [1] DBP_Standing_2_screening Glucose_12
                                                   HDL_12
## Root node error: 16.586/70 = 0.23694
##
## n= 70
##
##
          CP nsplit rel error xerror
## 1 0.620767 0 1.00000 1.03657 0.058676
```

plotcp(fit)



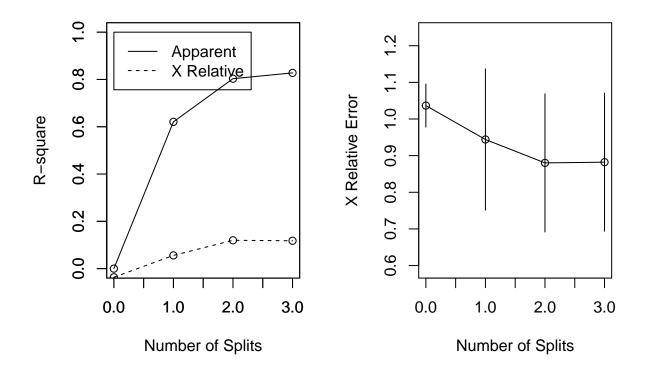


summary(fit)

```
## Call:
## rpart(formula = remission ~ ., data = data, method = "anova")
##
     n=70
##
##
              CP nsplit rel error
                                      xerror
## 1 0.62076671
                      0 1.0000000 1.0365692 0.05867597
## 2 0.18256376
                      1 0.3792333 0.9440776 0.19288556
## 3 0.02440425
                      2 0.1966695 0.8802024 0.18855559
## 4 0.01000000
                      3 0.1722653 0.8821346 0.18848687
##
## Variable importance
##
                        HDL_12
                                                {\tt dfatmass120}
##
                             15
                                                          13
##
                Fibro_E_med_12
                                           SBP_Sitting_1_12
##
                             13
##
                     Height_12
                                     {\tt Waistcircumference\_12}
##
                             12
                                                          12
                        FBG_12
##
                                                 Glucose_12
##
                              4
##
                          {\tt hbaf}
                                                  Glucose_3
##
                              3
                                                           3
##
                     Glucose_6
                                                       IL4_0
```

```
##
##
     DBP_Standing_2_screening DBP_Standing_Avg_screening
##
##
## Node number 1: 70 observations,
                                       complexity param=0.6207667
     mean=1.614286, MSE=0.2369388
##
     left son=2 (24 obs) right son=3 (46 obs)
##
##
     Primary splits:
##
         HDL 12
                         < 0.83
                                    to the left,
                                                   improve=0.6207667, (0 missing)
##
         BilirubinT_12
                         < 4.2
                                    to the left,
                                                   improve=0.5933188, (0 missing)
##
         TotalProtein_12 < 66
                                     to the left,
                                                   improve=0.5933188, (0 missing)
##
                         < 97.5
                                                   improve=0.5933188, (0 missing)
         UricAcid_12
                                    to the left,
         Cholesterol_12 < 3.05
                                                   improve=0.5933188, (0 missing)
##
                                    to the left,
##
     Surrogate splits:
##
         SBP_Sitting_1_12
                               < 104.5
                                           to the left, agree=0.943, adj=0.833, (0 split)
##
         Fibro_E_med_12
                               < 2.533333 to the left,
                                                         agree=0.943, adj=0.833, (0 split)
##
         dfatmass120
                               < -0.55
                                           to the right, agree=0.943, adj=0.833, (0 split)
##
                               < 75
                                           to the left, agree=0.929, adj=0.792, (0 split)
         Height_12
##
                                           to the left, agree=0.929, adj=0.792, (0 split)
         Waistcircumference_12 < 40.95
##
## Node number 2: 24 observations,
                                       complexity param=0.02440425
     mean=1.083333, MSE=0.07638889
##
     left son=4 (17 obs) right son=5 (7 obs)
##
##
     Primary splits:
##
         DBP_Standing_2_screening < 76
                                              to the right, improve=0.2207792, (0 missing)
##
         UricAcid_0
                                   < 281.5
                                              to the right, improve=0.2207792, (0 missing)
##
                                              to the right, improve=0.2207792, (0 missing)
         day_night_sensor_ratio_0 < 2.25</pre>
##
         IgA_0
                                   < 3212.7
                                              to the right, improve=0.2207792, (0 missing)
##
         LPPL_0
                                              to the right, improve=0.2207792, (0 missing)
                                   < 637.65
##
     Surrogate splits:
##
         DBP_Standing_Avg_screening < 79.75
                                                to the right, agree=0.958, adj=0.857, (0 split)
##
         SBP_Sitting_1_screening
                                    < 127.5
                                                to the right, agree=0.917, adj=0.714, (0 split)
##
         SBP_Sitting_2_screening
                                     < 121.5
                                                to the right, agree=0.917, adj=0.714, (0 split)
##
                                                to the right, agree=0.917, adj=0.714, (0 split)
         DBP_Standing_1_screening
                                     < 78.5
##
         SBP_Sitting_Avg_screening < 125.5
                                                to the right, agree=0.917, adj=0.714, (0 split)
##
## Node number 3: 46 observations,
                                       complexity param=0.1825638
##
     mean=1.891304, MSE=0.09688091
     left son=6 (7 obs) right son=7 (39 obs)
##
##
     Primary splits:
##
         Glucose 12
                          < 7.55
                                      to the right, improve=0.6794425, (0 missing)
                                      to the right, improve=0.6794425, (0 missing)
##
         FBG_12
                          < 7.55
##
         Glucose 3
                          < 7.2
                                      to the right, improve=0.5013550, (0 missing)
##
         average_sensor_0 < 201.1093 to the right, improve=0.3966944, (0 missing)
##
         estimated_a1c_0 < 8.65
                                      to the right, improve=0.3966944, (0 missing)
##
     Surrogate splits:
##
         FBG_12
                   < 7.55
                              to the right, agree=1.000, adj=1.000, (0 split)
##
         hbaf
                   < 6.55
                              to the right, agree=0.957, adj=0.714, (0 split)
##
         Glucose_3 < 6.95
                              to the right, agree=0.935, adj=0.571, (0 split)
##
         Glucose_6 < 7.7
                              to the right, agree=0.935, adj=0.571, (0 split)
##
                              to the left, agree=0.935, adj=0.571, (0 split)
         IL4_0
                   < 408.7
##
## Node number 4: 17 observations
     mean=1, MSE=0
```

```
##
## Node number 5: 7 observations
    mean=1.285714, MSE=0.2040816
##
## Node number 6: 7 observations
##
   mean=1.285714, MSE=0.2040816
## Node number 7: 39 observations
   mean=2, MSE=0
par(mfrow=c(1,2))
rsq.rpart(fit)
##
## Regression tree:
## rpart(formula = remission ~ ., data = data, method = "anova")
## Variables actually used in tree construction:
## [1] DBP_Standing_2_screening Glucose_12
                                                      HDL_12
## Root node error: 16.586/70 = 0.23694
##
## n= 70
##
          CP nsplit rel error xerror
## 1 0.620767 0 1.00000 1.03657 0.058676
## 2 0.182564 1 0.37923 0.94408 0.192886
## 3 0.024404
                2 0.19667 0.88020 0.188556
             3 0.17227 0.88213 0.188487
## 4 0.010000
```



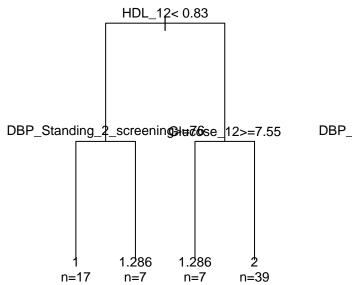
```
{plot(fit, uniform=TRUE, main="Regression Tree")
text(fit, use.n=TRUE, xpd=TRUE, cex=.8)}

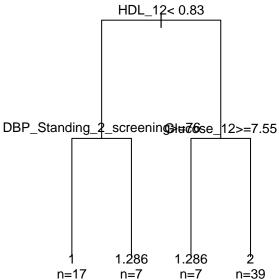
pfit<- prune(fit, cp= fit1$cptable[which.min(fit1$cptable[,"xerror"]),"CP"])

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

Regression Tree

Pruned Regression Tree





SVM

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)</pre>
```

##		coeffsord
##	(Intercept)	1907.5132020
##	TANITAFFM_O	225.0528983
##	TANITAFFM_screening	63.3263269
##	TANITAFatMass_0	57.2404458
##	TANITABMR_O	26.2820557

	TANITATBW_screening	21.0277334
	TANITABMI_O	20.9776533
	TANITABMR_screening	17.8787612
	Mother_Cancer	16.2537069
	Father_Hypertension	14.0111450
##	TANITAIdealBodyWeight_screen	12.9576645
##	_ · · · · · · · · · · · · · · · · ·	10.2340697
##	TANITAVisceralFatRating_scre	10.2109495
	ExSmoker	9.7999856
##	CurrentSmoker_Cigarettes	8.4651516
##	Mother_Hypertension	7.7865994
##	TANITAWeight_screening	7.3327687
##	Mother_Obesity	3.7944617
##	Dates_Quantity	3.4093357
##	TANITAMetabolicAge_screening	2.6693243
##	Father_Obesity	2.2123729
##	TimeToEat_Breakfast	1.7811497
##	DBP_Sitting_2_screening	1.7394594
##	Neckcircumference_0	1.4402402
##	HR_Standing_1_screening	1.3647995
##	Waistcircumference_screening	1.1205229
##	Father_CVD	0.8665498
##	HR_Sitting_3_screening	0.8636234
##	LatestAlbCr	0.8567369
##	SBP_Sitting_1_screening	0.7176876
##	DBP_Standing_1_screening	0.6517275
##	Takeaway_PerWeek	0.6365369
##	Soda_PerWeek	0.6018107
##	Dates_PerWeek	0.3031706
##	SBP_Sitting_3_screening	0.2881038
##	DBP_Standing_2_screening	0.2401073
##	Coffee_PerWeek	0.1561715
##	Tea_PerWeek	0.1178284
##	TANITAImpedance_screening	0.1032210
	TimeToEat_Lunch	-0.2557238
##	SBP_Standing_1_screening	-0.3118365
##	Waistcircumference_0	-0.4571104
##	DBP_Sitting_3_screening	-0.4937112
	SBP_Sitting_2_screening	-0.5099673
	SBP_Sitting_Avg_screening	-0.5788230
	SBP_Standing_2_screening	-0.5878252
	HR_Sitting_2_screening	-0.6677836
	TimeToEat_Dinner	-0.8034717
	Neckcircumference_screening	-0.8650100
	HR_Sitting_1_screening	-0.8715946
	DBP_Sitting_1_screening	-0.9539603
	Juice_PerWeek	-0.9763528
	HR_Standing_2_screening	-1.2253485
	Hipcircumference_screening	-1.2566155
	TANITAMetabolicAge_0	-1.8676782
	TANITAFatMass_screening	-3.8747837
	Height_0	-4.6271985
	CurrentSmoker_Shisha	-5.9791696
	Father_Diabetes	-6.0845759
	-	

##	Mother_Diabetes	-6.9235629
##	TANITAVisceralFatRating_0	-11.2919537
##	Height_screening	-11.5902615
##	TANITATBW_O	-21.0509717
##	Father_Cancer	-24.5504862
##	TANITABMI_screening	-36.5704339
##	TANITAWeight_0	-56.0416969
##	BK	-74.0482636
##	TANITAMuscleMass_screening	-89.3995749
##	EQ	-110.4355206
##	TANITAMuscleMass_0	-165.0274222