

FILTER

2023-07-22

```
Assignment <- read.csv("C:/Users/User/Downloads/chd.csv", header = TRUE)
Assignment$adiposity = NULL

library(dplyr)

## Warning: package 'dplyr' was built under R version 4.3.1
##
## 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(ggplot2)

## Warning: package 'ggplot2' was built under R version 4.3.1

library(tidyr)

## Warning: package 'tidyr' was built under R version 4.3.1

# Fit the logistic regression model
model <- glm(chd ~ ., family = binomial(link = 'logit'), data = Assignment)

# Predict probabilities
probabilities <- predict(model, type = "response")

# Predicted classes based on a threshold of 0.5
predicted.classes <- ifelse(probabilities > 0.5, "pos", "neg")

# Select only numeric predictors
mydata <- Assignment %>%
  dplyr::select_if(is.numeric)

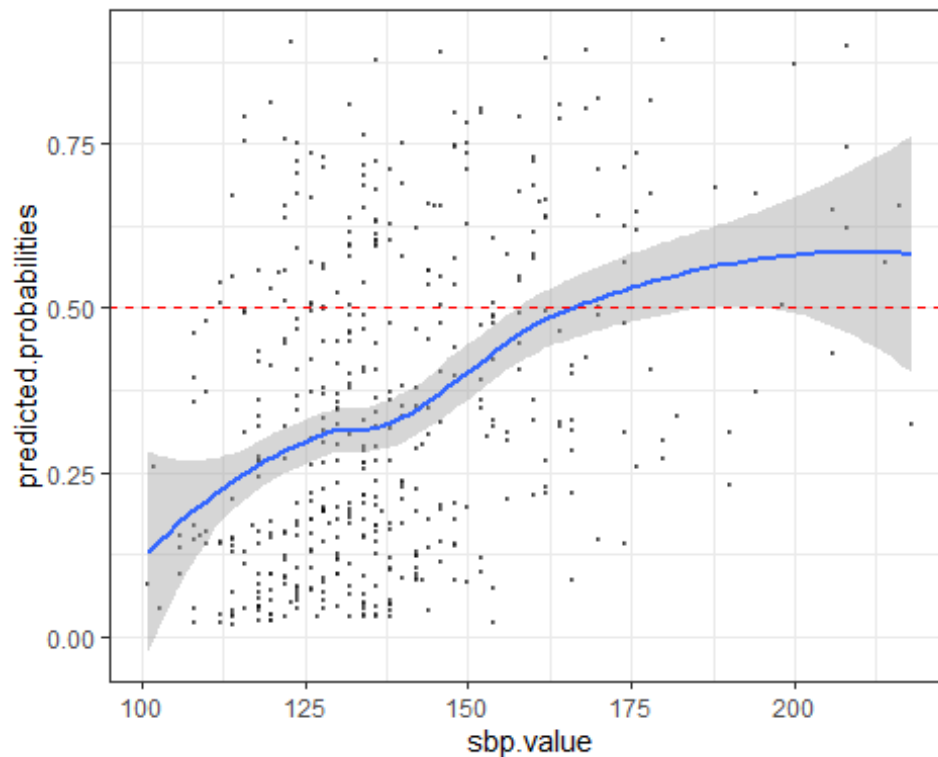
# Add logit values to the dataframe
mydata$logit <- probabilities

# Gather the data for plotting
mydata <- mydata %>%
  gather(key = "predictors", value = "predictor.value", -logit)
```

```
#####sbp#####
# Filter only the "sbp" predictor for the plot
mydata_sbp <- mydata %>%
  filter(predictors == "sbp")

# Plot the graph for "sbp" predictor
ggplot(mydata_sbp, aes(predictor.value, logit)) +
  geom_point(size = 0.5, alpha = 0.5) +
  labs(x="sbp.value", y="predicted.probabilities") +
  geom_smooth(method = "loess") +
  theme_bw() + geom_hline(yintercept = 0.5, linetype = "dashed", color =
"red")

## `geom_smooth()` using formula = 'y ~ x'
```

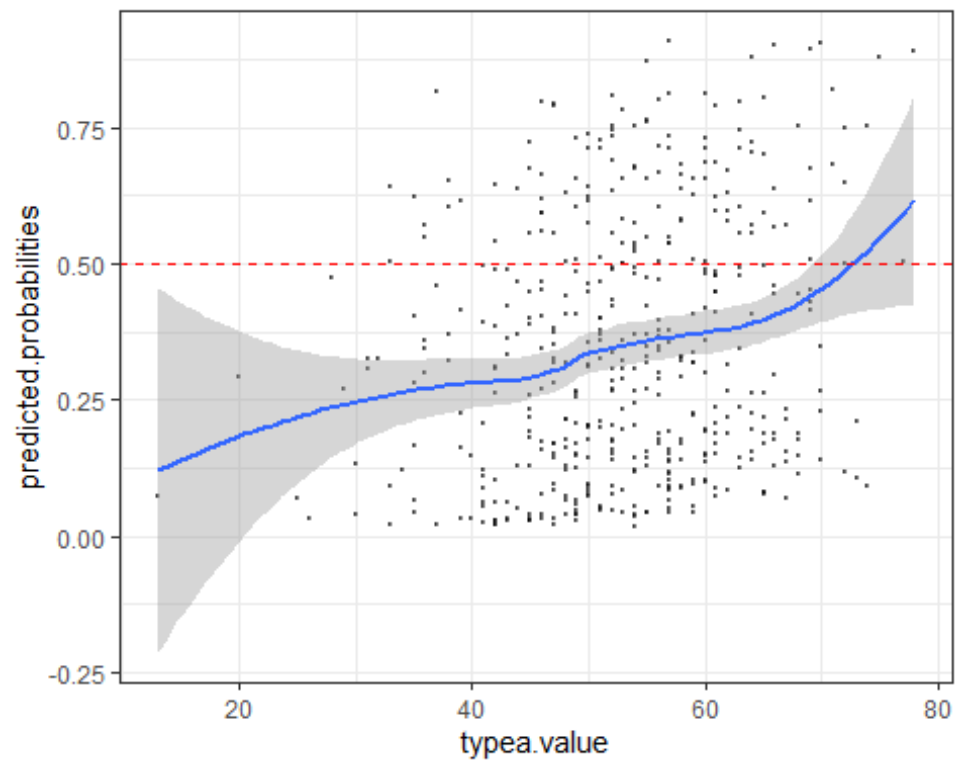


```
mydata_typea <- mydata %>%
  filter(predictors == "typea")

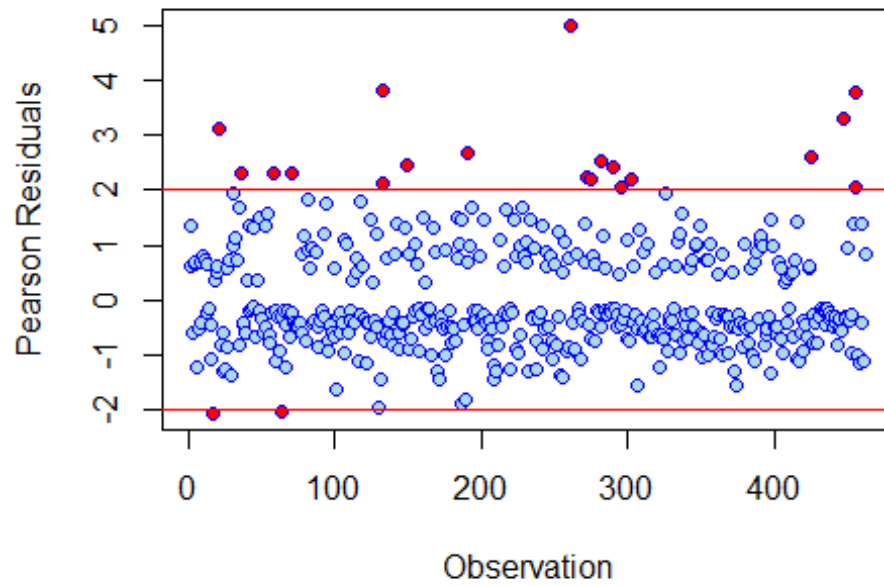
# Plot the graph for "typea" predictor
ggplot(mydata_typea, aes(predictor.value, logit)) +
  geom_point(size = 0.5, alpha = 0.5) +
  labs(x="typea.value", y="predicted.probabilities") +
  geom_smooth(method = "loess") +
```

```
theme_bw() + geom_hline(yintercept = 0.5, linetype = "dashed", color =  
"red")
```

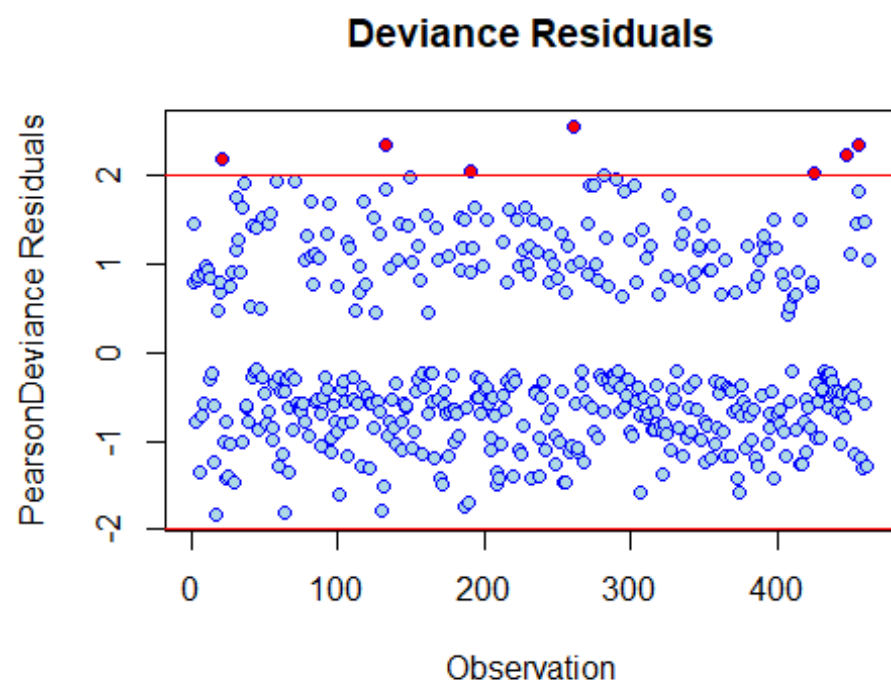
```
## `geom_smooth()` using formula = 'y ~ x'
```



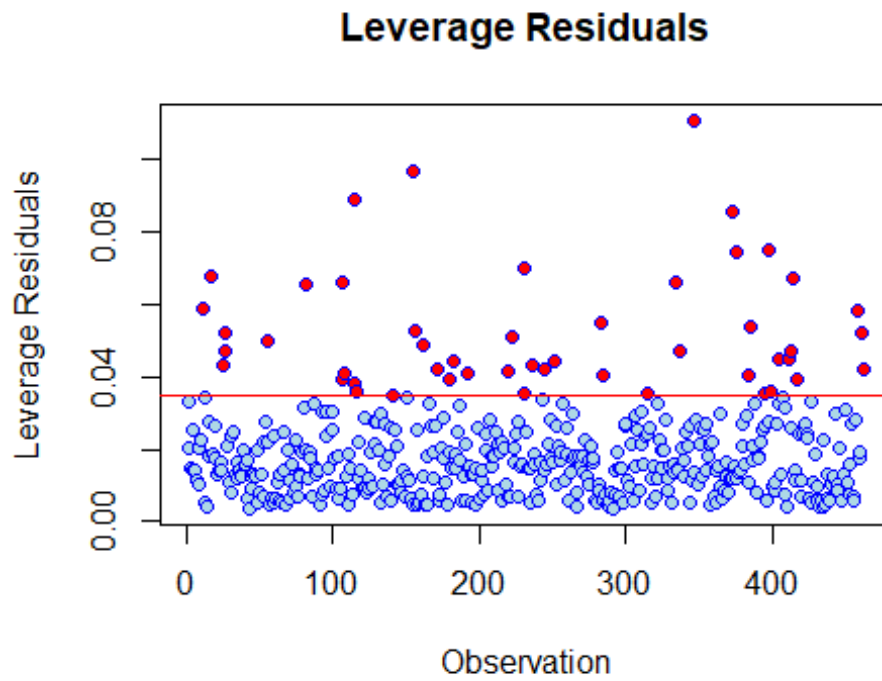
Pearson Residuals



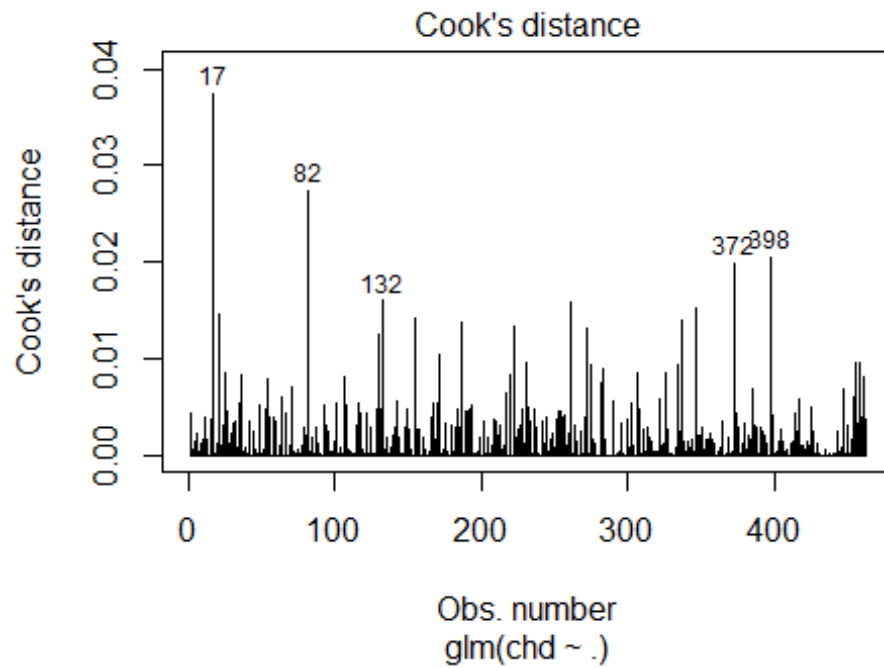
```
## 17 21 36 58 63 70 132 133 149 191 261 272 275 281 290 296 302 425
447 455
## 17 21 36 58 63 70 132 133 149 191 261 272 275 281 290 296 302 425
447 455
## 456
## 456
```



```
## 21 133 191 261 425 447 456
## 21 133 191 261 425 447 456
## [1] 0.1106625
```

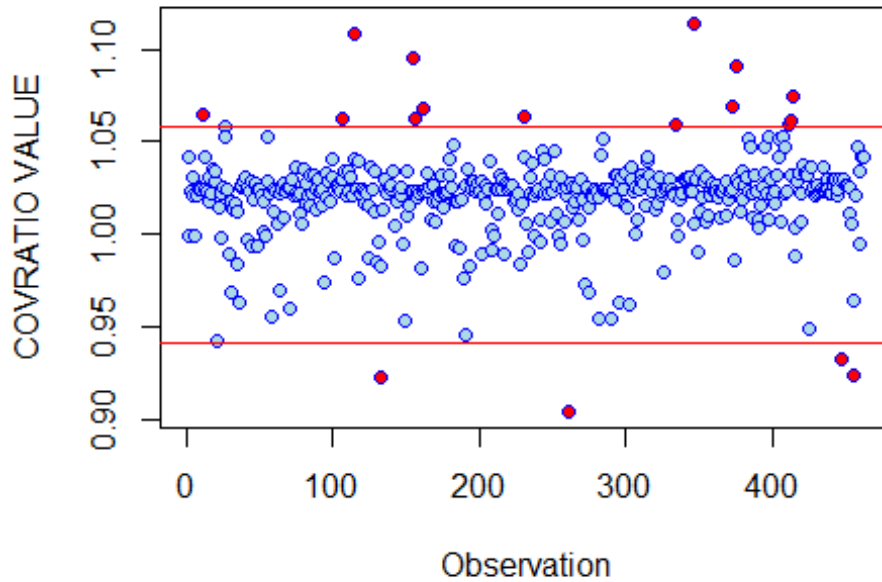


```
## 11 17 25 26 27 56 82 106 107 108 114 115 116 141 155 156 162 171
180 182
## 11 17 25 26 27 56 82 106 107 108 114 115 116 141 155 156 162 171
180 182
## 192 220 222 230 231 236 244 251 283 285 315 334 337 346 372 375 383 385
395 398
## 192 220 222 230 231 236 244 251 283 285 315 334 337 346 372 375 383 385
395 398
## 399 405 411 413 414 417 458 461 462
## 399 405 411 413 414 417 458 461 462
```



```
## named integer(0)
## Warning: package 'olsrr' was built under R version 4.3.1
##
## Attaching package: 'olsrr'
## The following object is masked from 'package:datasets':
##
##     rivers
```

COVRATIO Observations



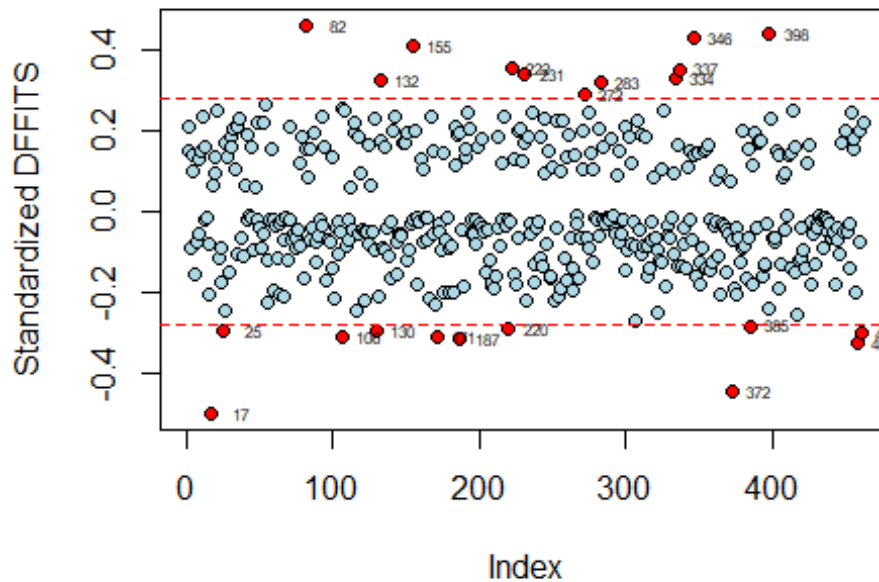
##	11	106	115	133	155	156	162	231	261	334	346	372	375	411	413	414	447	456
##	11	106	115	133	155	156	162	231	261	334	346	372	375	411	413	414	447	456
##		1				2			3			4			5			6
##		0.14631829				0.20668733			-0.09430808			0.09789284			0.13867155			-0.15876456
##		7				8			9			10			11			12
##		-0.07459322				0.12700958			-0.05762864			0.14815796			0.23429299			0.15733995
##		13				14			15			16			17			18
##		-0.02200407				-0.01532028			-0.20835283			-0.08045729			-0.50082821			0.06543484
##		19				20			21			22			23			24
##		0.09072837				0.13083809			0.24844711			-0.11869250			-0.17863937			-0.09372365
##		25				26			27			28			29			30
##		-0.29615988				0.16799339			-0.24534598			0.13239633			-0.15132770			0.18176259
##		31				32			33			34			35			36
##		0.15606250				0.20179298			0.10380793			0.20928631			-0.10868563			0.22534347
##		37				38			39			40			41			42
##		-0.06889027				-0.06562408			-0.11160699			0.06351756			0.18547816			-0.02266116
##		43				44			45			46			47			48
##		-0.01280775				0.15666967			-0.01681539			-0.09773989			0.05613471			0.21586111
##		49				50			51			52			53			54
##		-0.01959018				-0.04206899			-0.09003201			-0.05482582			0.21632957			0.26443734
##		55				56			57			58			59			60
##		-0.12418954				-0.22815829			-0.02278092			0.15210530			-0.02056740			-0.19660573
##		61				62			63			64			65			66
##		-0.03558132				-0.11635036			-0.20529923			-0.03268247			-0.02147978			-0.21286792
##		67				68			69			70			71			72
##		-0.07388830				-0.01746062			-0.12399058			0.20597109			-0.02421277			-0.07109328

##	73	74	75	76	77	78
##	-0.05594267	-0.09412222	-0.06477749	-0.04852909	-0.08648021	0.11646651
##	79	80	81	82	83	84
##	0.18084066	-0.16884383	0.15422633	0.45797592	0.08330627	0.15501793
##	85	86	87	88	89	90
##	-0.07379944	-0.04342902	0.19474179	-0.07724310	-0.12550014	-0.06006455
##	91	92	93	94	95	96
##	-0.01991563	0.23524544	-0.03607890	0.15682773	-0.17043934	-0.17402371
##	97	98	99	100	101	102
##	-0.07103957	-0.07756778	0.13125881	-0.14091791	-0.21420793	-0.04038978
##	103	104	105	106	107	108
##	-0.10719813	-0.02588937	-0.05239886	-0.31376184	0.25113733	0.24693000
##	109	110	111	112	113	114
##	-0.08720974	-0.01965693	-0.07108648	0.05691650	-0.04788762	0.19321145
##	115	116	117	118	119	120
##	0.21802885	-0.24768394	0.18467104	-0.05741308	0.09224339	-0.04685844
##	121	122	123	124	125	126
##	-0.22146359	-0.05351269	-0.05175178	0.16341011	-0.08011781	0.06090677
##	127	128	129	130	131	132
##	-0.05333361	-0.09761353	0.22647219	-0.29818731	-0.21261367	0.32296546
##	133	134	135	136	137	138
##	0.17121879	-0.09310966	-0.09804127	0.15667239	-0.04370002	-0.11320396
##	139	140	141	142	143	144
##	-0.02567884	-0.16516785	0.19765361	0.23425204	-0.07841234	-0.15705589
##	145	146	147	148	149	150
##	-0.05753827	-0.05515638	-0.06260947	0.16761805	0.16539534	0.19175210
##	151	152	153	154	155	156
##	-0.12022390	-0.09489137	-0.03359151	-0.02013981	0.40780625	0.19614020
##	157	158	159	160	161	162
##	-0.01603727	-0.18133937	-0.02899340	0.13015026	-0.07661265	0.10442643
##	163	164	165	166	167	168
##	-0.01664955	-0.07409039	-0.01601235	-0.21688675	0.23189704	0.14635604
##	169	170	171	172	173	174
##	-0.05598501	-0.23230517	-0.31378609	-0.03414631	-0.05201926	-0.09904209
##	175	176	177	178	179	180
##	-0.20067142	0.14350265	-0.06995962	-0.01747946	-0.08952923	-0.20120886
##	181	182	183	184	185	186
##	-0.08494139	-0.20244270	0.11287752	0.20838113	0.18663461	0.19489269
##	187	188	189	190	191	192
##	-0.31827527	-0.04536607	-0.18767354	0.13191407	0.15211174	0.24224612
##	193	194	195	196	197	198
##	0.20443476	-0.05660445	-0.02025569	-0.05695236	-0.01976145	-0.08372722
##	199	200	201	202	203	204
##	0.15903616	-0.03360205	-0.04198138	0.17833156	-0.04941268	-0.14932703
##	205	206	207	208	209	210
##	-0.15209924	-0.04612460	-0.12106573	-0.18577794	-0.17917280	-0.18998799
##	211	212	213	214	215	216
##	-0.16142794	0.18514777	-0.04366483	-0.08867640	-0.02408900	0.11929268
##	217	218	219	220	221	222
##	0.23234204	-0.03650813	-0.02061635	-0.29095624	-0.02722128	0.35216546

##	223	224	225	226	227	228
##	-0.15686047	0.12638075	-0.18079203	-0.06701028	0.19556406	0.20054155
##	229	230	231	232	233	234
##	0.12481160	0.16739221	0.33678411	-0.22252543	0.18112531	-0.03131657
##	235	236	237	238	239	240
##	-0.05542395	0.24235549	-0.13040307	-0.17543725	-0.04249275	-0.02516715
##	241	242	243	244	245	246
##	0.18487057	-0.11455678	-0.13510023	0.22634904	0.09292084	-0.08175188
##	247	248	249	250	251	252
##	0.14802305	-0.16488418	-0.16060705	0.10672257	-0.21931603	-0.04333132
##	253	254	255	256	257	258
##	0.22186218	-0.21099830	-0.19211581	0.09590730	0.22151434	-0.11825577
##	259	260	261	262	263	264
##	0.13185567	-0.16926568	0.18991531	-0.06663164	-0.19522277	-0.14317608
##	265	266	267	268	269	270
##	0.13666280	-0.01378624	-0.03291085	-0.16639540	-0.06689263	0.10114297
##	271	272	273	274	275	276
##	0.19622472	0.28652357	-0.06764891	-0.08947190	0.24362259	0.14469661
##	277	278	279	280	281	282
##	0.10537604	-0.12699739	-0.01895283	-0.02158368	0.20283129	-0.06203322
##	283	284	285	286	287	288
##	0.31548043	-0.02229854	0.15627248	-0.03266782	-0.01727729	-0.01620961
##	289	290	291	292	293	294
##	-0.02391930	0.17766334	-0.01214925	-0.04746638	-0.05744261	0.08681772
##	295	296	297	298	299	300
##	-0.07501191	0.14885622	-0.05826131	-0.02202102	0.20676805	-0.14793846
##	301	302	303	304	305	306
##	-0.09868714	0.18510266	-0.08159552	0.11664237	-0.02934577	-0.08757234
##	307	308	309	310	311	312
##	-0.27363919	0.22078826	-0.04751729	-0.10676837	0.18602229	-0.08543235
##	313	314	315	316	317	318
##	0.18515694	-0.08588965	-0.16405609	-0.14261390	-0.07242913	-0.02673628
##	319	320	321	322	323	324
##	0.08225516	-0.06601269	-0.24919299	-0.09634494	-0.12264756	0.10423986
##	325	326	327	328	329	330
##	-0.12715840	0.24826380	-0.18701512	-0.03804831	-0.06236462	-0.06559334
##	331	332	333	334	335	336
##	-0.01459099	-0.10579204	0.09423402	0.32907259	0.16299580	-0.13779414
##	337	338	339	340	341	342
##	0.34970028	-0.13678113	-0.08058107	-0.04332717	-0.14367252	0.10910873
##	343	344	345	346	347	348
##	0.14780830	-0.02309397	-0.06943401	0.42888132	0.13907048	-0.16416825
##	349	350	351	352	353	354
##	0.14355168	-0.12809332	-0.18047406	-0.09160177	0.14956299	0.15159040
##	355	356	357	358	359	360
##	-0.14284523	0.16157610	-0.02177354	-0.14879436	-0.12724156	-0.03641169
##	361	362	363	364	365	366
##	0.07684051	-0.02556336	-0.11185364	-0.20464630	0.10024191	-0.03121649
##	367	368	369	370	371	372
##	-0.03761196	-0.04790435	-0.15029964	-0.05887418	0.07313327	-0.44589973

[illegible]

Standardized DFFITS, critical value = $2 \cdot \sqrt{p/n} = \pm 0.2791$



NEW DATASET FILTERED

```
Assignment_filtered = Assignment[-
c(outlier_variables_DR,outlier_variables_PR,leverage_obs,COVRATIO_cuoffvalues
,influence_point_dffits,influence_point_dffits),]
```

Calculate pseudo R-squared and p-value

nagelkerke function also reports the McFadden, Cox and Snell, and Nagelkerke pseudo R-squared value for the model

```
library(rcompanion)
```

```
## Warning: package 'rcompanion' was built under R version 4.3.1
```

```
varyingmodel <- glm(chd ~., family = binomial(link = 'logit'), data =
Assignment)
```

```
x <- varyingmodel
```

```
nagelkerke(x)
```

```
## $Models
```

```
##
```

```
## Model: "glm, chd ~ ., binomial(link = \"logit\"), Assignment"
```

```
## Null: "glm, chd ~ 1, binomial(link = \"logit\"), Assignment"
```

```
##
```

```
## $Pseudo.R.squared.for.model.vs.null
```

```
## Pseudo.R.squared
```

```
## McFadden 0.207283
```

```

## Cox and Snell (ML)                0.234674
## Nagelkerke (Cragg and Uhler)      0.323775
##
## $Likelihood.ratio.test
##   Df.diff LogLik.diff Chisq    p.value
##      -8      -61.782 123.56 6.0846e-23
##
## $Number.of.observations
##
## Model: 462
## Null:  462
##
## $Messages
## [1] "Note: For models fit with REML, these statistics are based on
refitting with ML"
##
## $Warnings
## [1] "None"

##### Efron's pseudo R-squared

efronRSquared(x)

## EfronRSquared
##      0.245

varyingmodel <- glm(chd ~.-alcohol, family = binomial(link = 'logit'), data =
Assignment_filtered)
x <- varyingmodel
nagelkerke(x)

## $Models
##
## Model: "glm, chd ~ . - alcohol, binomial(link = \"logit\"),
Assignment_filtered"
## Null:  "glm, chd ~ 1, binomial(link = \"logit\"), Assignment_filtered"
##
## $Pseudo.R.squared.for.model.vs.null
##                                Pseudo.R.squared
## McFadden                      0.360943
## Cox and Snell (ML)             0.351028
## Nagelkerke (Cragg and Uhler)   0.502786
##
## $Likelihood.ratio.test
##   Df.diff LogLik.diff Chisq    p.value
##      -7      -84.527 169.05 3.9721e-33
##
## $Number.of.observations
##
## Model: 391
## Null:  391

```

```
##
## $Messages
## [1] "Note: For models fit with REML, these statistics are based on
refitting with ML"
##
## $Warnings
## [1] "None"

##### Efron's pseudo R-squared

efronRSquared(x)

## EfronRSquared
##          0.372

#####AUC METHOD
(**)#####

### AUC ROC for Initial
library(prediction)

## Warning: package 'prediction' was built under R version 4.3.1

library(caret)

## Warning: package 'caret' was built under R version 4.3.1

## Loading required package: lattice

library(ROCR)

## Warning: package 'ROCR' was built under R version 4.3.1

##
## Attaching package: 'ROCR'

## The following object is masked from 'package:prediction':
##
##      prediction

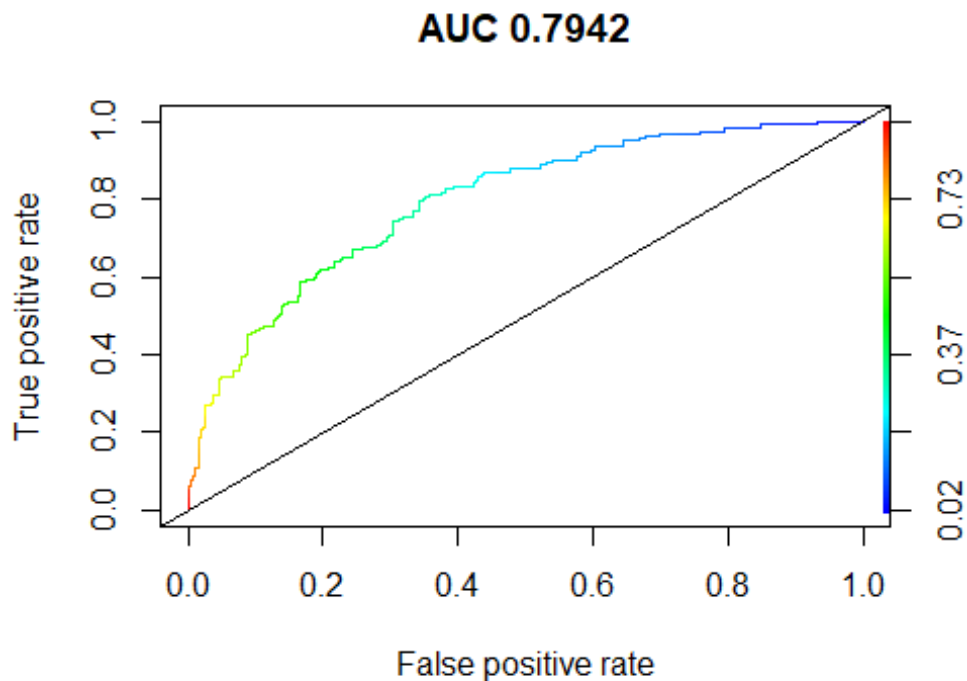
modell1 <- glm(chd ~. , family = binomial(link = 'logit'), data = Assignment)
pred <- predict(modell1, Assignment, type = "response")
pred.rocr <- prediction(pred, Assignment$chd)

perf.rocr <- performance(pred.rocr, measure = "auc", x.measure = "cutoff")

perf.rocr@y.values[[1]] <- round(perf.rocr@y.values[[1]], digits = 4)

perf.tpr.fpr.rocr <- performance(pred.rocr, "tpr", "fpr")
```

```
plot(perf.tpr.fpr.rocr, colorize = T, main =
paste("AUC", (perf.rocr@y.values)))
abline(a=0, b=1)
```



AUC ROC for Adjusted

```
model2 <- glm(chd ~ ., family = binomial(link = 'logit'), data =
Assignment_filtered)
pred <- predict(model2, Assignment_filtered, type = "response")
pred.rocr <- prediction(pred, Assignment_filtered$chd)
summary(model2)
```

```
##
## Call:
## glm(formula = chd ~ ., family = binomial(link = "logit"), data =
Assignment_filtered)
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept) -1.001e+01  1.966e+00  -5.091 3.56e-07 ***
## sbp          1.134e-02  8.744e-03   1.296 0.194838
## tobacco      1.406e-01  4.114e-02   3.417 0.000633 ***
## ldl          3.809e-01  9.141e-02   4.167 3.09e-05 ***
## famhist      1.396e+00  2.969e-01   4.703 2.57e-06 ***
## typea        6.208e-02  1.807e-02   3.436 0.000591 ***
```

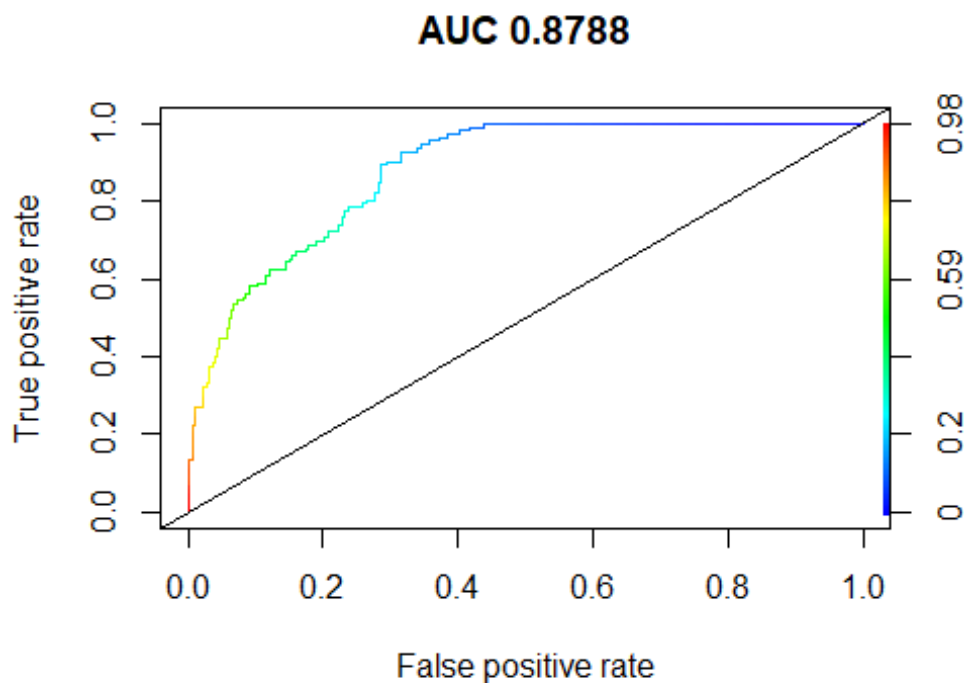
```
## obesity      -8.427e-02  4.377e-02  -1.925  0.054213 .
## alcohol      -6.428e-04  7.824e-03  -0.082  0.934528
## age          7.051e-02  1.471e-02   4.794  1.63e-06 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 468.37  on 390  degrees of freedom
## Residual deviance: 299.31  on 382  degrees of freedom
## AIC: 317.31
##
## Number of Fisher Scoring iterations: 6

perf.rocr <- performance(pred.rocr, measure = "auc", x.measure = "cutoff")

perf.rocr@y.values[[1]] <- round(perf.rocr@y.values[[1]], digits = 4)

perf.tpr.fpr.rocr <- performance(pred.rocr, "tpr", "fpr")

plot(perf.tpr.fpr.rocr, colorize = T, main =
paste("AUC", (perf.rocr@y.values)))
abline(a=0, b=1)
```



```
#####TESTING WHETHER BINOMIAL LOGISTICS IS
APPROPRIATE#####
```


#####USING LOGISTICS REGRESSION DIAGNOSTICS#####

```
library(tidyverse)
```

```
## Warning: package 'tidyverse' was built under R version 4.3.1
```

```
## Warning: package 'readr' was built under R version 4.3.1
```

```
## Warning: package 'purrr' was built under R version 4.3.1
```

```
## Warning: package 'stringr' was built under R version 4.3.1
```

```
## Warning: package 'forcats' was built under R version 4.3.1
```

```
## Warning: package 'lubridate' was built under R version 4.3.1
```

```
## — Attaching core tidyverse packages ————— tidyverse  
2.0.0 —
```

```
## ✓ forcats 1.0.0 ✓ readr 2.1.4
```

```
## ✓ lubridate 1.9.2 ✓ stringr 1.5.0
```

```
## ✓ purrr 1.0.1 ✓ tibble 3.2.1
```

```
## — Conflicts —————
```

```
tidyverse_conflicts() —
```

```
## ✗ dplyr::filter() masks stats::filter()
```

```
## ✗ dplyr::lag() masks stats::lag()
```

```
## ✗ purrr::lift() masks caret::lift()
```

```
## i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all  
conflicts to become errors
```

```
library(broom)
```

```
## Warning: package 'broom' was built under R version 4.3.1
```

```
new_model <- glm(chd ~., family = binomial(link = 'logit'), data =  
Assignment_filtered)
```

```
probabilities <- predict(new_model, type = "response")
```

```
predicted.classes <- ifelse(probabilities > 0.5, "pos", "neg")
```

```
#Select only numeric predictors
```

```
mydata <- Assignment_filtered %>%
```

```
  dplyr::select_if(is.numeric)
```

```
predictors <- colnames(mydata)
```

```
#Bind the Logit and tidying the data for plot
```

```
mydata <- mydata %>%
```

```
  mutate(logit = log(probabilities/(1-probabilities))) %>%
```

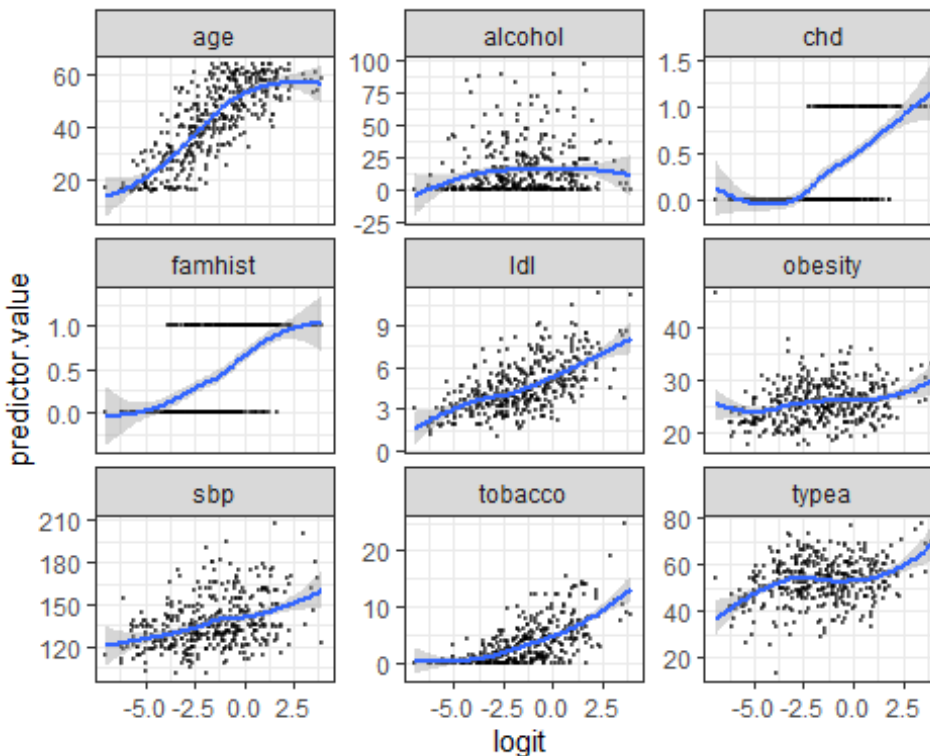
```
  gather(key = "predictors", value = "predictor.value", -logit)
```

```
#Graph visualization
```

```
ggplot(mydata, aes(logit, predictor.value))+
```

```
geom_point(size = 0.5, alpha = 0.5) +
geom_smooth(method = "loess") +
theme_bw() +
facet_wrap(~predictors, scales = "free_y")
```

```
## `geom_smooth()` using formula = 'y ~ x'
```



```
summary(new_model)
```

```
##
## Call:
## glm(formula = chd ~ ., family = binomial(link = "logit"), data =
Assignment_filtered)
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept) -1.001e+01  1.966e+00  -5.091 3.56e-07 ***
## sbp          1.134e-02  8.744e-03   1.296 0.194838
## tobacco      1.406e-01  4.114e-02   3.417 0.000633 ***
## ldl          3.809e-01  9.141e-02   4.167 3.09e-05 ***
## famhist      1.396e+00  2.969e-01   4.703 2.57e-06 ***
## typea        6.208e-02  1.807e-02   3.436 0.000591 ***
## obesity     -8.427e-02  4.377e-02  -1.925 0.054213 .
## alcohol     -6.428e-04  7.824e-03  -0.082 0.934528
## age          7.051e-02  1.471e-02   4.794 1.63e-06 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
##
## (Dispersion parameter for binomial family taken to be 1)
##
## Null deviance: 468.37 on 390 degrees of freedom
## Residual deviance: 299.31 on 382 degrees of freedom
## AIC: 317.31
##
## Number of Fisher Scoring iterations: 6

stepwise_newmodel <- step(new_model, direction = "both", trace = 0, k = 2)

# Print the final selected model
summary(stepwise_newmodel)

##
## Call:
## glm(formula = chd ~ tobacco + ldl + famhist + typea + obesity +
## age, family = binomial(link = "logit"), data = Assignment_filtered)
##
## Coefficients:
## Estimate Std. Error z value Pr(>|z|)
## (Intercept) -8.66248 1.64093 -5.279 1.30e-07 ***
## tobacco 0.14075 0.03959 3.556 0.000377 ***
## ldl 0.38155 0.09013 4.233 2.30e-05 ***
## famhist 1.37933 0.29506 4.675 2.94e-06 ***
## typea 0.05897 0.01780 3.313 0.000924 ***
## obesity -0.07732 0.04301 -1.798 0.072210 .
## age 0.07489 0.01425 5.255 1.48e-07 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
## Null deviance: 468.37 on 390 degrees of freedom
## Residual deviance: 301.04 on 384 degrees of freedom
## AIC: 315.04
##
## Number of Fisher Scoring iterations: 6

car::vif(new_model)

## sbp tobacco ldl famhist typea obesity alcohol age
## 1.177445 1.116131 1.186976 1.029565 1.174389 1.209209 1.138752 1.280999

#####
##### Calculate pseudo R-squared and p-value (CAN ANALYZE REMOVAL OF
RESIDUALS ****)
##### alternative test for p-value for a fitted model -> nagelkerke function
will report the p-value for a model using the LRT

##### nagelkerke function also reports the McFadden, Cox and Snell, and
```

Nagelkerke pseudo R-squared value for the model

```
library(rcompanion)

varyingmodel <- glm(chd ~., family = binomial(link = 'logit'), data =
Assignment)
x <- varyingmodel
nagelkerke(x)

## $Models
##
## Model: "glm, chd ~ ., binomial(link = \"logit\"), Assignment"
## Null:  "glm, chd ~ 1, binomial(link = \"logit\"), Assignment"
##
## $Pseudo.R.squared.for.model.vs.null
##                               Pseudo.R.squared
## McFadden                      0.207283
## Cox and Snell (ML)             0.234674
## Nagelkerke (Cragg and Uhler)   0.323775
##
## $Likelihood.ratio.test
## Df.diff LogLik.diff Chisq    p.value
##      -8      -61.782 123.56 6.0846e-23
##
## $Number.of.observations
##
## Model: 462
## Null:  462
##
## $Messages
## [1] "Note: For models fit with REML, these statistics are based on
refitting with ML"
##
## $Warnings
## [1] "None"

##### Efron's pseudo R-squared

efronRSquared(x)

## EfronRSquared
##           0.245

varyingmodel <- glm(chd ~.-alcohol, family = binomial(link = 'logit'), data =
Assignment_filtered)
x <- varyingmodel
nagelkerke(x)

## $Models
##
## Model: "glm, chd ~ . - alcohol, binomial(link = \"logit\"),
```

```

Assignment_filtered"
## Null:  "glm, chd ~ 1, binomial(link = \"logit\"), Assignment_filtered"
##
## $Pseudo.R.squared.for.model.vs.null
##                                Pseudo.R.squared
## McFadden                      0.360943
## Cox and Snell (ML)             0.351028
## Nagelkerke (Cragg and Uhler)   0.502786
##
## $Likelihood.ratio.test
## Df.diff LogLik.diff Chisq    p.value
##      -7      -84.527 169.05 3.9721e-33
##
## $Number.of.observations
##
## Model: 391
## Null:  391
##
## $Messages
## [1] "Note: For models fit with REML, these statistics are based on
refitting with ML"
##
## $Warnings
## [1] "None"

##### Efron's pseudo R-squared

efronRSquared(x)

## EfronRSquared
##      0.372

```