

Red Wines - upper

Katie, Rita, and Chang

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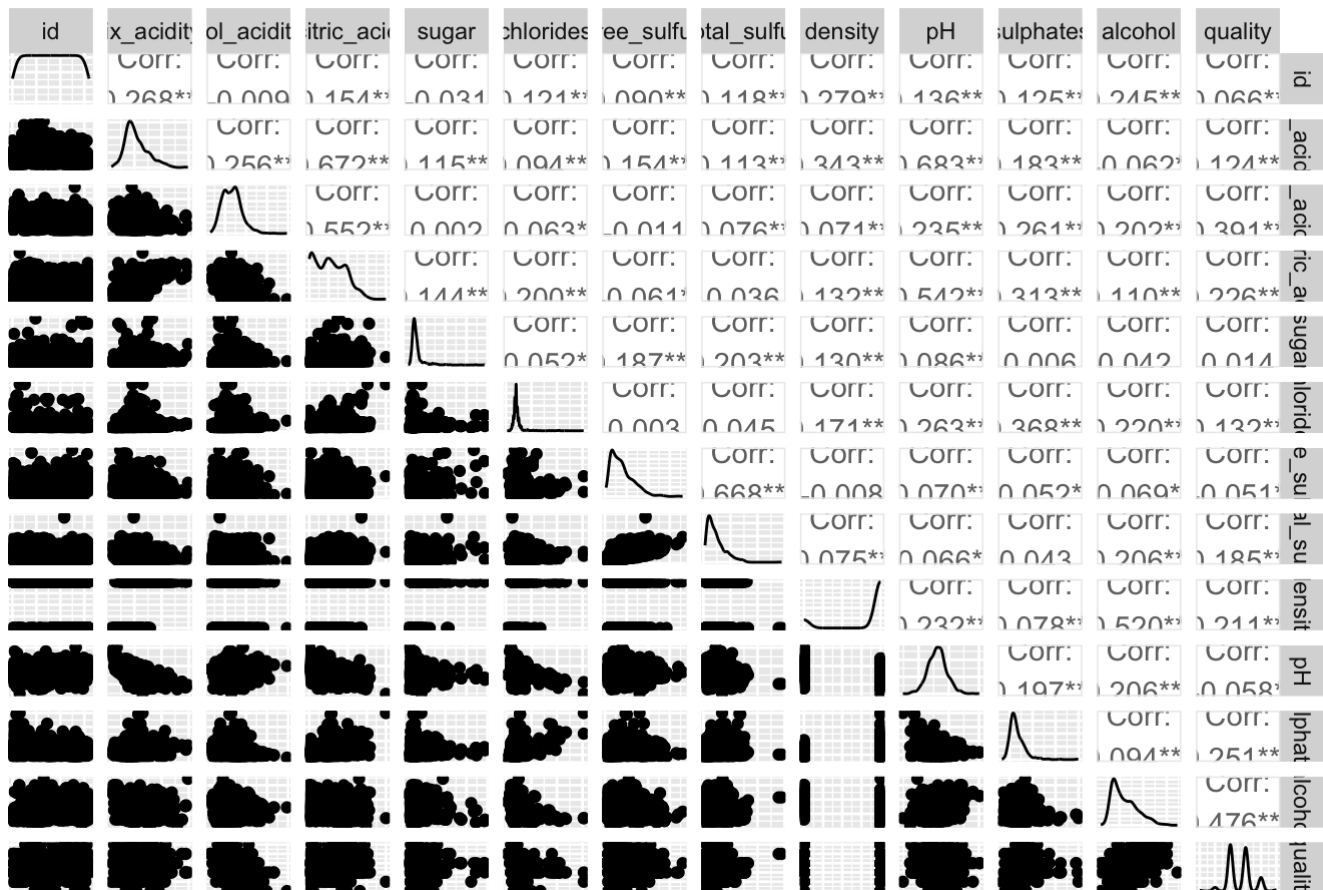
Scatterplot Matrix

```
library("GGally")
ggpairs(red, axisLabels = "none",
        title = "Scatterplot Matrix of Red Wines")
```

```
# corr codes
```

Scatterplot Matrix

Scatterplot Matrix of Red Wines



Create Binary Dependent Variable

```
red$highquality = factor((red$quality >= 6))  
red$highquality <- as.integer(as.logical(red$highquality))
```

Create Test and Training Data

```
library("caTools")  
set.seed = 100  
split = sample.split(red$highquality, SplitRatio = 0.6)  
train = subset(red, split == TRUE)  
test = subset(red, split == FALSE)  
print(dim(train)); print(dim(test))
```

```
## [1] 959  14
```

```
## [1] 640  14
```

Descriptive Statistics

```
library("Rmisc")
```

```
## Loading required package: lattice
```

```
## Loading required package: plyr
```

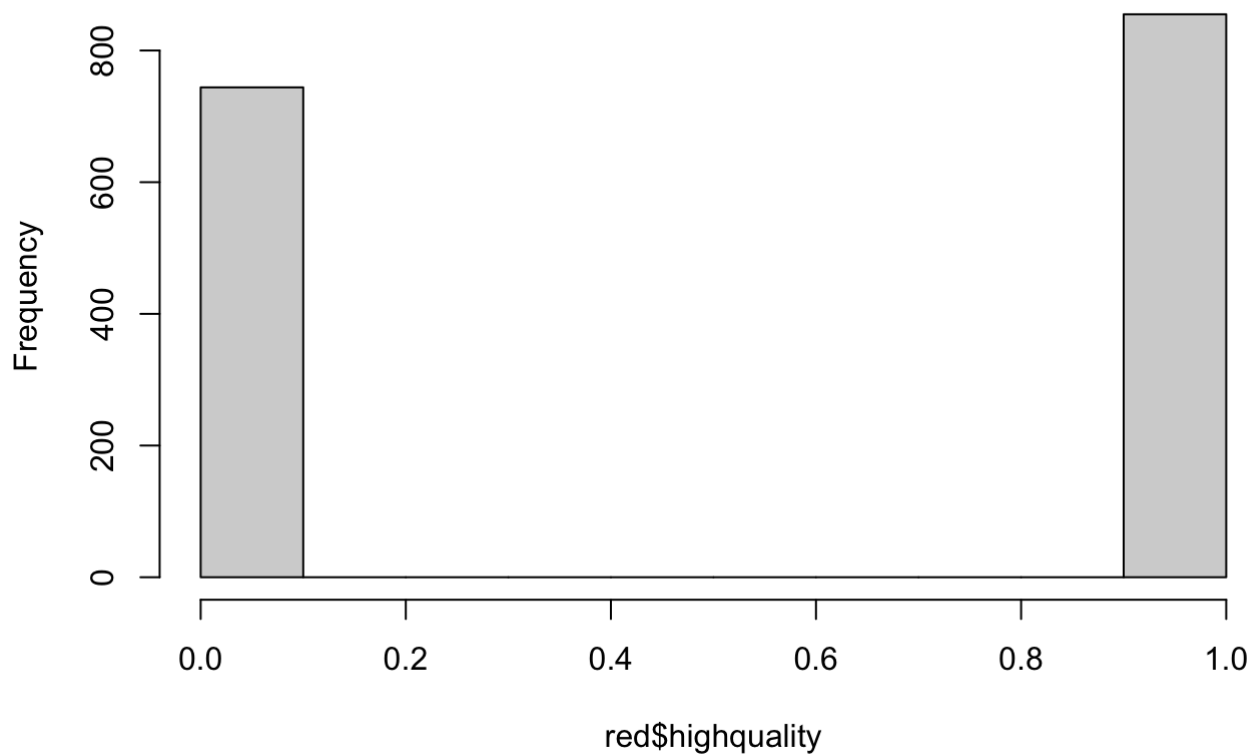
```
sum = summary(red)  
sum
```

```
##          id          fix_acidity    vol_acidity    citric_acid
## Min.      : 1.0    Min.      : 4.60    Min.      :0.1200    Min.      :0.000
## 1st Qu.: 400.5    1st Qu.: 7.10    1st Qu.:0.3900    1st Qu.:0.090
## Median : 800.0    Median : 7.90    Median :0.5200    Median :0.260
## Mean      : 800.0    Mean      : 8.32    Mean      :0.5284    Mean      :0.271
## 3rd Qu.:1199.5    3rd Qu.: 9.20    3rd Qu.:0.6400    3rd Qu.:0.420
## Max.      :1599.0    Max.      :15.90    Max.      :1.5800    Max.      :1.000
##          sugar          chlorides          free_sulfur          total_sulfur
## Min.      : 0.900    Min.      :0.01000    Min.      : 1.00    Min.      : 6.00
## 1st Qu.: 1.900    1st Qu.:0.07000    1st Qu.: 7.00    1st Qu.: 22.00
## Median : 2.200    Median :0.08000    Median :14.00    Median : 38.00
## Mean      : 2.539    Mean      :0.08787    Mean      :15.87    Mean      : 46.47
## 3rd Qu.: 2.600    3rd Qu.:0.09000    3rd Qu.:21.00    3rd Qu.: 62.00
## Max.      :15.500    Max.      :0.61000    Max.      :72.00    Max.      :289.00
##          density          pH          sulphates          alcohol
## Min.      :0.9900    Min.      :2.740    Min.      :0.3300    Min.      : 8.40
## 1st Qu.:1.0000    1st Qu.:3.210    1st Qu.:0.5500    1st Qu.: 9.50
## Median :1.0000    Median :3.310    Median :0.6200    Median :10.20
## Mean      :0.9985    Mean      :3.311    Mean      :0.6581    Mean      :10.42
## 3rd Qu.:1.0000    3rd Qu.:3.400    3rd Qu.:0.7300    3rd Qu.:11.10
## Max.      :1.0000    Max.      :4.010    Max.      :2.0000    Max.      :14.90
##          quality          highquality
## Min.      :3.000    Min.      :0.0000
## 1st Qu.:5.000    1st Qu.:0.0000
## Median :6.000    Median :1.0000
## Mean      :5.636    Mean      :0.5347
## 3rd Qu.:6.000    3rd Qu.:1.0000
## Max.      :8.000    Max.      :1.0000
```

Plot high quality vs low quality distribution

```
hist (red$highquality)
```

Histogram of red\$highquality



Random Forest

```
library("randomForest")
```

```
## randomForest 4.7-1.1
```

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

```
##  
## Attaching package: 'randomForest'
```

```
## The following object is masked from 'package:ggplot2':  
##  
## margin
```

```
library("caret")
library("e1071")
library("rpart")
```

```
rf <- randomForest(highquality ~ . - quality, data = train, mtry = 4, importance = TRUE,
ntree = 50, na.action = na.omit)
```

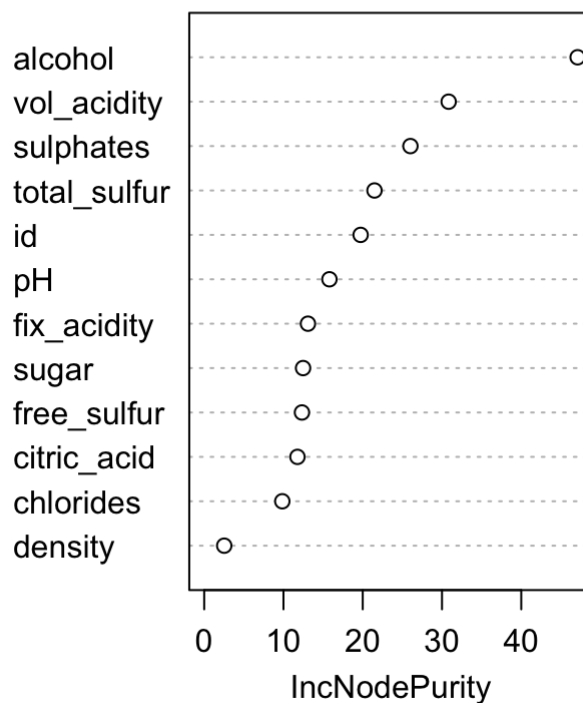
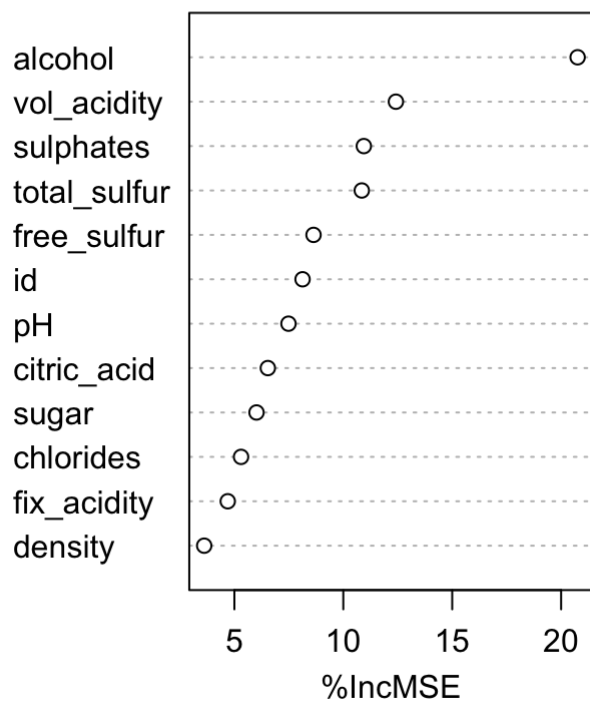
```
## Warning in randomForest.default(m, y, ...): The response has five or fewer
## unique values. Are you sure you want to do regression?
```

```
print(rf)
```

```
##
## Call:
## randomForest(formula = highquality ~ . - quality, data = train,      mtry = 4, impor
tance = TRUE, ntree = 50, na.action = na.omit)
##              Type of random forest: regression
##              Number of trees: 50
## No. of variables tried at each split: 4
##
##              Mean of squared residuals: 0.1549242
##              % Var explained: 37.73
```

```
varImpPlot(rf)
```

rf



```
# predictions on test set
set.seed(100)
predictTest = predict(rf, newdata = test, type = "response")

# confusion matrix on test set
table(test$highquality, predictTest >= 0.5)
```

```
##
##      FALSE TRUE
##    0    236   62
##    1     64  278
```

Random Forest Model

```
# Logit
randomforestmodlogit <- glm(highquality ~ alcohol + sulphates + total_sulfur + vol_acidity, data = red, family = "binomial"(link = "logit"))
summary(randomforestmodlogit)
```

```
##
## Call:
## glm(formula = highquality ~ alcohol + sulphates + total_sulfur +
##       vol_acidity, family = binomial(link = "logit"), data = red)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -3.1638  -0.8675   0.3076   0.8629   2.3262
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  -8.588813   0.795118 -10.802  < 2e-16 ***
## alcohol       0.927362   0.069268  13.388  < 2e-16 ***
## sulphates     2.059047   0.365976   5.626 1.84e-08 ***
## total_sulfur -0.011976   0.001924  -6.225 4.83e-10 ***
## vol_acidity  -3.083277   0.364832  -8.451  < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 2209.0  on 1598  degrees of freedom
## Residual deviance: 1684.2  on 1594  degrees of freedom
## AIC: 1694.2
##
## Number of Fisher Scoring iterations: 4
```

```
# Cloglog
randomforestmodcloglog <- glm(highquality ~ alcohol + sulphates + total_sulfur + vol_aci
dity, data = red, family = "binomial"(link = "cloglog"))
summary(randomforestmodcloglog)
```

```
##
## Call:
## glm(formula = highquality ~ alcohol + sulphates + total_sulfur +
##       vol_acidity, family = binomial(link = "cloglog"), data = red)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -4.5006  -0.9020   0.2185   0.9295   2.0506
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  -4.958517   0.478252 -10.368 < 2e-16 ***
## alcohol       0.505807   0.038543  13.123 < 2e-16 ***
## sulphates     1.324184   0.221318   5.983 2.19e-09 ***
## total_sulfur -0.009109   0.001364  -6.679 2.41e-11 ***
## vol_acidity  -2.022997   0.238813  -8.471 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 2209  on 1598  degrees of freedom
## Residual deviance: 1701  on 1594  degrees of freedom
## AIC: 1711
##
## Number of Fisher Scoring iterations: 7
```

```
# The logit model performed better with a lower AIC value
```

Cart

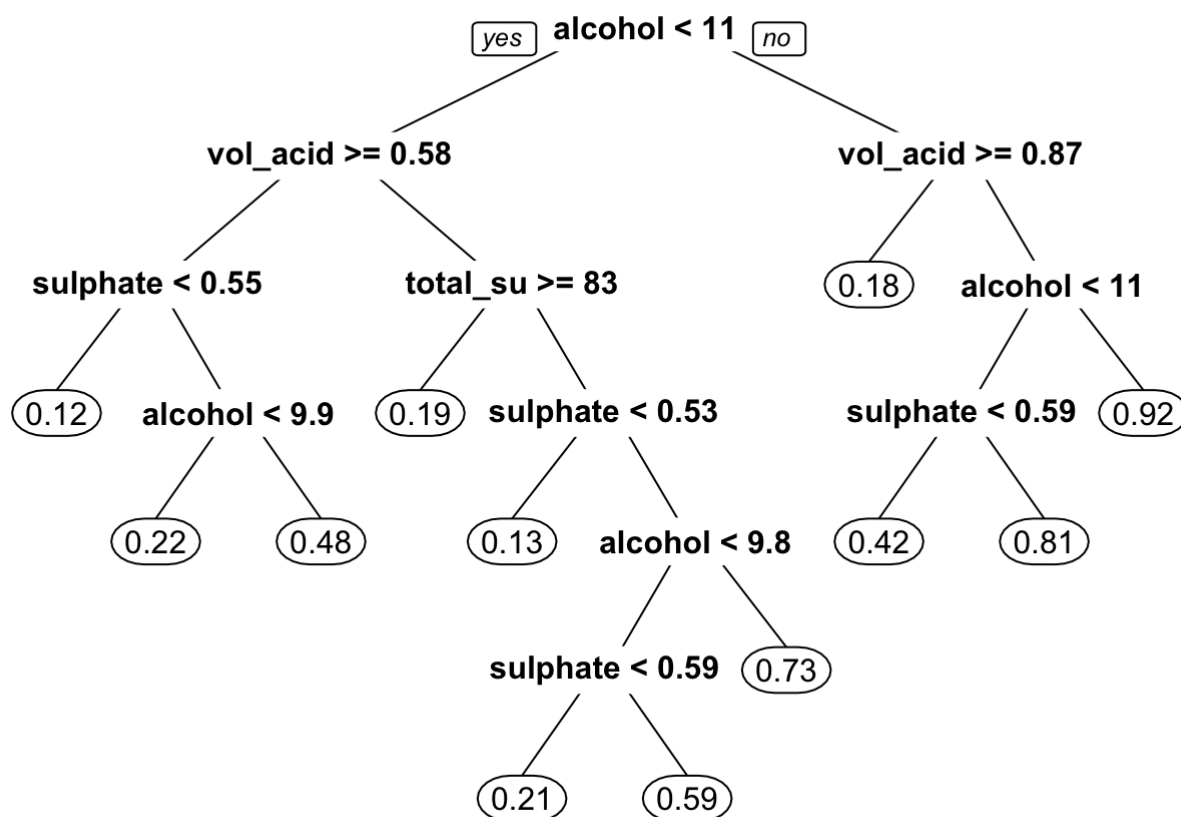
```
library("caret")
library("e1071")
library("rpart")
library("rpart.plot")

cartmodel = rpart(highquality ~ . - quality, data = train)
print(cartmodel)
```



```
## n= 959
##
## node), split, n, deviance, yval
##      * denotes terminal node
##
## 1) root 959 238.579800 0.5349322
##    2) alcohol< 10.525 589 137.826800 0.3735144
##      4) vol_acidity>=0.575 280 50.442860 0.2357143
##        8) sulphates< 0.545 109 11.449540 0.1192661 *
##        9) sulphates>=0.545 171 36.573100 0.3099415
##       18) alcohol< 9.85 111 18.810810 0.2162162 *
##       19) alcohol>=9.85 60 14.983330 0.4833333 *
##    5) vol_acidity< 0.575 309 77.249190 0.4983819
##      10) total_sulfur>=82.5 53 8.113208 0.1886792 *
##      11) total_sulfur< 82.5 256 63.000000 0.5625000
##        22) sulphates< 0.525 30 3.466667 0.1333333 *
##        23) sulphates>=0.525 226 53.274340 0.6194690
##       46) alcohol< 9.75 111 27.747750 0.5045045
##         92) sulphates< 0.585 24 3.958333 0.2083333 *
##         93) sulphates>=0.585 87 21.103450 0.5862069 *
##       47) alcohol>=9.75 115 22.643480 0.7304348 *
##    3) alcohol>=10.525 370 60.975680 0.7918919
##      6) vol_acidity>=0.87 17 2.470588 0.1764706 *
##      7) vol_acidity< 0.87 353 51.756370 0.8215297
##        14) alcohol< 11.45 182 36.263740 0.7252747
##          28) sulphates< 0.585 40 9.775000 0.4250000 *
##          29) sulphates>=0.585 142 21.866200 0.8098592 *
##        15) alcohol>=11.45 171 12.011700 0.9239766 *
```

```
prp(cartmodel)
```



```

# predictions on test set
set.seed(100)
predictTest = predict(cartmodel, newdata = test, type = "matrix")

# confusion matrix on test set
table(test$highquality, predictTest >= 0.5)

```

```

##
##      FALSE TRUE
##    0    216   82
##    1     86  256

```

Cart Model

```

# Logit
cartmodlogit <- glm(highquality ~ alcohol + sulphates + total_sulfur + vol_acidity, data
= red, family = "binomial"(link = "logit"))
summary(cartmodlogit)

```

```
##
## Call:
## glm(formula = highquality ~ alcohol + sulphates + total_sulfur +
##       vol_acidity, family = binomial(link = "logit"), data = red)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -3.1638  -0.8675   0.3076   0.8629   2.3262
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  -8.588813   0.795118 -10.802  < 2e-16 ***
## alcohol       0.927362   0.069268  13.388  < 2e-16 ***
## sulphates     2.059047   0.365976   5.626 1.84e-08 ***
## total_sulfur -0.011976   0.001924  -6.225 4.83e-10 ***
## vol_acidity  -3.083277   0.364832  -8.451  < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 2209.0  on 1598  degrees of freedom
## Residual deviance: 1684.2  on 1594  degrees of freedom
## AIC: 1694.2
##
## Number of Fisher Scoring iterations: 4
```

```
# Cloglog
cartmodcloglog <- glm(highquality ~ alcohol + sulphates + total_sulfur + fix_acidity, data = red, family = "binomial"(link = "cloglog"))
summary(cartmodcloglog)
```

```
##
## Call:
## glm(formula = highquality ~ alcohol + sulphates + total_sulfur +
##       fix_acidity, family = binomial(link = "cloglog"), data = red)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -4.7058  -0.9408   0.3075   0.9490   1.9387
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  -6.835907   0.481268 -14.204 < 2e-16 ***
## alcohol       0.542953   0.037720  14.394 < 2e-16 ***
## sulphates     1.639060   0.217233   7.545 4.52e-14 ***
## total_sulfur -0.009315   0.001384  -6.732 1.67e-11 ***
## fix_acidity   0.027351   0.021284   1.285  0.199
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 2209.0  on 1598  degrees of freedom
## Residual deviance: 1777.5  on 1594  degrees of freedom
## AIC: 1787.5
##
## Number of Fisher Scoring iterations: 18
```

```
# The logit model performed better with the lower AIC value
```

Compare best logit model with AIC

```
library("AICcmodavg")
```

```
##
## Attaching package: 'AICcmodavg'
```

```
## The following object is masked from 'package:randomForest':
##
##      importance
```

```
models <- list(randomforestmodlogit, cartmodlogit)
mod.names <- c('RandomForest', 'Cart')
aictab(cand.set = models, modnames = mod.names)
```

```
## Warning in aictab.AICglm.lm(cand.set = models, modnames = mod.names):
## Check model structure carefully as some models may be redundant
```

```
##
## Model selection based on AICc:
##
##           K      AICc Delta_AICc AICcWt Cum.Wt      LL
## RandomForest 5 1694.21          0    0.5    0.5 -842.09
## Cart         5 1694.21          0    0.5    1.0 -842.09
```

```
# The random forest logit model performed the best
```

Compare best model with BIC

```
library("flexmix")
BIC(randomforestmodlogit)
```

```
## [1] 1721.058
```

```
BIC(randomforestmodcloglog)
```

```
## [1] 1737.845
```

```
BIC(cartmodlogit)
```

```
## [1] 1721.058
```

```
BIC(cartmodcloglog)
```

```
## [1] 1814.418
```

```
# The random forest logit model performed the best
```

Confusion matrix for random forest logit model

```
confusionred = predict(randomforestmodlogit, newdata = red, type = "response")

# confusion matrix on test set
table(red$highquality, confusionred >= 0.5)
```

```
##  
##      FALSE TRUE  
##    0    548  196  
##    1    216  639
```

Predictions for random forest logit model

```
pred_test <- predict(randomforestmodlogit, test, type = "response")  
  
pred_test
```

##	1	2	8	10	12	14	15
##	0.21686414	0.16572704	0.35364079	0.50812152	0.50812152	0.69723806	0.13086405
##	16	17	18	19	20	25	27
##	0.13867752	0.64471340	0.56805457	0.20109254	0.62465624	0.49785778	0.51001918
##	29	30	33	35	37	41	42
##	0.20629006	0.38187954	0.17799414	0.37517558	0.65305951	0.60541307	0.20653385
##	45	46	47	52	54	56	60
##	0.28787990	0.91140600	0.05638181	0.41159910	0.21158666	0.24266029	0.37446288
##	61	67	69	70	73	74	76
##	0.35540196	0.36302140	0.76772166	0.66550098	0.11468249	0.17116376	0.67706290
##	77	78	79	81	82	83	84
##	0.67706290	0.41857071	0.19429727	0.38434763	0.65378965	0.19907576	0.44994570
##	86	90	91	95	97	98	102
##	0.45450681	0.20581734	0.12488112	0.11648871	0.45180389	0.36239110	0.59957642
##	104	106	107	108	110	113	115
##	0.17168420	0.17168420	0.65531540	0.25986779	0.05669027	0.21064410	0.46170336
##	116	119	122	127	129	131	143
##	0.61534547	0.59589245	0.59589245	0.15237332	0.73731458	0.08091771	0.98548676
##	144	145	146	147	148	150	152
##	0.47019352	0.98548676	0.08710726	0.18741376	0.29279616	0.65840863	0.86018228
##	154	162	164	168	172	173	175
##	0.24118269	0.45773435	0.10323416	0.18953659	0.42272086	0.42272086	0.40084611
##	178	180	181	182	183	185	186
##	0.58567146	0.27931853	0.27931853	0.25686482	0.16145478	0.19399307	0.38321589
##	187	190	191	193	199	200	208
##	0.28302491	0.13218057	0.16007768	0.12192200	0.81955290	0.29660130	0.11227661
##	209	212	215	219	224	229	230
##	0.27161312	0.23949363	0.30759231	0.37301967	0.39670029	0.56522890	0.64457164
##	231	232	234	238	243	244	245
##	0.77766942	0.47608964	0.64457164	0.20849103	0.17002266	0.67646997	0.67646997
##	248	249	250	252	260	267	268
##	0.14653731	0.35125228	0.56422747	0.44429231	0.80917975	0.19291723	0.97152743
##	269	272	273	283	285	287	291
##	0.50344768	0.87483178	0.61403829	0.22895580	0.32999609	0.63496399	0.69463023
##	295	301	305	308	315	320	322
##	0.55129126	0.59572880	0.11220744	0.51647504	0.73885733	0.29176280	0.22451563
##	326	327	330	333	335	338	343
##	0.32170068	0.85787373	0.47496687	0.12860589	0.79014250	0.56472573	0.66228068
##	345	346	352	353	356	359	360
##	0.68982583	0.34930907	0.28077747	0.31982246	0.86315416	0.76935573	0.55892705
##	363	368	369	371	372	374	377
##	0.40286206	0.28955508	0.37745429	0.28282101	0.55754674	0.21115302	0.87515687
##	379	381	384	389	390	391	392
##	0.97217666	0.65714868	0.65714868	0.35745073	0.59089106	0.80025092	0.62763464
##	393	394	396	397	399	400	402
##	0.45488358	0.09608587	0.95715624	0.11072664	0.63287534	0.16572036	0.75725600
##	406	409	413	414	419	427	428
##	0.74785302	0.89010131	0.13582974	0.92335864	0.77020269	0.58695311	0.40277653
##	431	434	436	437	441	442	443
##	0.91967155	0.45313982	0.45313982	0.31324574	0.70777919	0.84990517	0.63537526
##	444	447	448	452	456	458	461
##	0.86779419	0.64968242	0.83321168	0.64119919	0.95768591	0.22920186	0.82420951

##	465	467	468	469	473	477	478
##	0.55427468	0.80312295	0.98771509	0.46457042	0.68760803	0.69504790	0.94349416
##	480	482	484	486	489	490	491
##	0.43048988	0.95378202	0.72818356	0.35050201	0.82229147	0.71109362	0.38820824
##	492	493	494	495	497	499	501
##	0.97829349	0.98423867	0.64444930	0.81342325	0.27636155	0.84287574	0.27636155
##	502	506	507	508	514	515	520
##	0.88317746	0.94197356	0.92658894	0.36855117	0.87663195	0.87663195	0.69166380
##	521	523	526	528	542	544	547
##	0.79768845	0.43596448	0.34605728	0.79882025	0.81658345	0.60467189	0.44918649
##	551	553	556	563	565	567	568
##	0.37808419	0.56543070	0.72091694	0.14387840	0.92935383	0.20667837	0.20667837
##	573	574	575	576	577	579	580
##	0.77494478	0.28595807	0.53585414	0.75659792	0.51154935	0.26273769	0.52861287
##	581	583	584	587	588	595	598
##	0.38002007	0.29689198	0.69766881	0.86089516	0.09780108	0.20381128	0.49453337
##	599	600	602	605	610	614	620
##	0.33386682	0.35305750	0.28353983	0.19528536	0.93597592	0.61614952	0.58672865
##	622	623	627	628	630	632	638
##	0.17261769	0.30620045	0.23676532	0.23676532	0.13683730	0.60526652	0.03693294
##	641	644	645	646	647	648	651
##	0.32321697	0.24690166	0.32321697	0.52329572	0.44260928	0.50449082	0.40958829
##	653	657	658	661	662	665	668
##	0.99329501	0.40958829	0.62339714	0.53465925	0.32678756	0.58642668	0.51847411
##	671	672	674	676	678	679	680
##	0.52667590	0.24700448	0.24700448	0.63951248	0.22140871	0.15831687	0.50931169
##	681	684	685	686	687	688	692
##	0.26144196	0.61274001	0.02353841	0.61274001	0.19820088	0.30853776	0.08792767
##	695	697	704	705	706	711	713
##	0.17339004	0.33118031	0.39592542	0.26438374	0.09514858	0.09703634	0.18728926
##	716	717	720	722	724	727	728
##	0.27155586	0.30467669	0.22428109	0.15248206	0.74648551	0.59402154	0.43341582
##	730	732	734	735	736	738	741
##	0.69434981	0.56601333	0.31147678	0.29287694	0.11882456	0.24236950	0.71932009
##	742	744	748	752	754	760	761
##	0.12964712	0.33233049	0.32846257	0.24423781	0.24423781	0.20828698	0.18944864
##	765	766	769	774	775	782	786
##	0.21925919	0.21287996	0.19045961	0.58787221	0.57996000	0.49407054	0.36376503
##	793	797	798	800	802	804	806
##	0.21742711	0.41405130	0.81010666	0.66715805	0.42345371	0.35326525	0.97678970
##	809	810	812	813	818	819	822
##	0.48396705	0.60727563	0.74847855	0.74001422	0.93891093	0.15085766	0.98401461
##	824	825	828	832	834	835	836
##	0.43070016	0.59452832	0.61809251	0.75438553	0.57515459	0.21302698	0.16263773
##	838	839	840	842	843	844	846
##	0.73172430	0.90163251	0.45005102	0.46356092	0.65608836	0.13891921	0.38077515
##	848	853	854	860	862	867	868
##	0.37044591	0.46348365	0.77796635	0.81336010	0.29801703	0.85828874	0.84439382
##	872	873	877	878	884	886	887
##	0.53427892	0.53560900	0.74967371	0.68842602	0.17504411	0.44881228	0.39943319
##	890	893	894	896	898	901	902
##	0.07413691	0.44299541	0.18472125	0.54837049	0.54837049	0.91308562	0.61749313

##	903	907	911	917	920	922	923
##	0.61749313	0.61655717	0.97132513	0.47643991	0.84865159	0.73857291	0.84865159
##	924	927	930	931	933	935	941
##	0.57254719	0.75197156	0.94254625	0.61571145	0.35714165	0.61571145	0.95924086
##	943	944	946	950	951	952	953
##	0.51350024	0.39208642	0.85009985	0.96739607	0.96739607	0.96007673	0.89304382
##	956	958	959	962	963	964	966
##	0.81941210	0.75906185	0.76200136	0.34752279	0.42379344	0.88720837	0.85318054
##	970	972	975	977	978	981	983
##	0.53399341	0.91601469	0.93910091	0.28346202	0.08591010	0.70374096	0.93219598
##	986	987	988	989	992	993	994
##	0.76216515	0.87904782	0.31207909	0.40407434	0.32169795	0.33637960	0.32169795
##	995	997	998	999	1003	1004	1005
##	0.24614218	0.91773179	0.91773179	0.13340402	0.94122194	0.96113831	0.55234587
##	1007	1008	1017	1021	1022	1023	1024
##	0.94122194	0.94084017	0.95824366	0.91151720	0.91151720	0.64109441	0.94561683
##	1025	1028	1032	1039	1040	1043	1044
##	0.57387540	0.53740392	0.71727212	0.94854700	0.80300719	0.80300719	0.82020006
##	1045	1054	1055	1058	1059	1060	1061
##	0.89700536	0.97117164	0.14570490	0.20603080	0.81405117	0.90442354	0.77354568
##	1064	1077	1078	1079	1089	1092	1095
##	0.92780908	0.91711368	0.82775448	0.82775448	0.59911292	0.87464611	0.49531036
##	1100	1111	1114	1116	1120	1121	1123
##	0.34983074	0.55003024	0.64019617	0.67849027	0.91323567	0.95982776	0.87671921
##	1126	1128	1130	1131	1133	1134	1136
##	0.88512330	0.58545658	0.59990536	0.59465043	0.98621359	0.81790944	0.90991385
##	1140	1142	1146	1147	1152	1154	1155
##	0.25663548	0.79255354	0.81699614	0.62188669	0.84747620	0.78814787	0.69856005
##	1156	1162	1163	1164	1165	1166	1168
##	0.32861855	0.68472828	0.93097631	0.34911691	0.34911691	0.64143459	0.95811695
##	1169	1170	1171	1173	1176	1178	1180
##	0.89257541	0.83796361	0.78148743	0.94556825	0.71702966	0.91536349	0.83725702
##	1182	1184	1187	1190	1193	1194	1195
##	0.83146700	0.18325571	0.75844840	0.12264121	0.96428659	0.39866755	0.16205917
##	1196	1198	1201	1203	1204	1205	1209
##	0.36820273	0.43683111	0.43683111	0.90407751	0.12664335	0.84882533	0.84882533
##	1210	1211	1212	1213	1222	1224	1227
##	0.89081590	0.53707859	0.33645556	0.53707859	0.89545422	0.93813689	0.19332528
##	1228	1229	1230	1236	1244	1247	1248
##	0.43944184	0.95997919	0.35954537	0.70226268	0.22116768	0.23445159	0.60960225
##	1249	1250	1251	1256	1258	1259	1262
##	0.82547588	0.61830476	0.61830476	0.50363201	0.60797958	0.69403671	0.37701038
##	1263	1265	1266	1268	1269	1270	1271
##	0.38122840	0.88943306	0.50245646	0.90213067	0.37253912	0.97152715	0.98284751
##	1273	1274	1275	1279	1282	1286	1288
##	0.71983985	0.19873507	0.66138806	0.13504841	0.57652097	0.64986391	0.90845738
##	1291	1292	1293	1294	1301	1306	1307
##	0.59908827	0.59190236	0.93827703	0.25512295	0.87211203	0.20683713	0.23809254
##	1308	1310	1311	1312	1313	1315	1318
##	0.58216493	0.18736687	0.20683713	0.93088901	0.11783860	0.48312749	0.88286689
##	1319	1321	1324	1325	1326	1327	1328
##	0.20771741	0.27840724	0.82228483	0.65726913	0.65726913	0.65726913	0.65726913

```
##      1330      1331      1333      1334      1339      1343      1346
## 0.21628945 0.21628945 0.44771727 0.16577829 0.33063230 0.51740015 0.56699331
##      1348      1350      1358      1365      1367      1376      1379
## 0.23966391 0.56190596 0.68815611 0.76732852 0.21556429 0.22071490 0.47985390
##      1394      1395      1396      1397      1399      1406      1409
## 0.38905380 0.15440151 0.33954980 0.34921494 0.49039723 0.92222761 0.96956379
##      1411      1416      1417      1419      1420      1421      1424
## 0.55013345 0.46359329 0.67996830 0.39766821 0.11430274 0.39766821 0.72836169
##      1433      1437      1438      1441      1442      1443      1447
## 0.92574770 0.16556629 0.38062331 0.87184058 0.09511377 0.43582198 0.43582198
##      1448      1450      1453      1455      1456      1461      1467
## 0.35375316 0.87394441 0.78758198 0.78198047 0.45960983 0.59735007 0.47414501
##      1468      1472      1473      1474      1475      1482      1486
## 0.30916495 0.86017441 0.84882869 0.58301194 0.19663663 0.81973890 0.31103235
##      1488      1490      1493      1496      1501      1506      1508
## 0.66158978 0.75070329 0.79984815 0.77693701 0.26400037 0.36554744 0.85817699
##      1510      1511      1515      1517      1523      1525      1526
## 0.91307103 0.57620576 0.12840124 0.82595385 0.82595385 0.60658881 0.45939419
##      1529      1530      1532      1533      1534      1538      1541
## 0.63560644 0.34422384 0.42672777 0.52363402 0.23235943 0.50712482 0.81445748
##      1542      1543      1545      1546      1547      1548      1553
## 0.84898279 0.36046967 0.89062172 0.45450879 0.51435731 0.82069018 0.62386686
##      1556      1557      1558      1562      1563      1568      1570
## 0.64128386 0.23011305 0.48014133 0.14727854 0.37557065 0.37557065 0.78071517
##      1574      1575      1582      1584      1586      1592      1593
## 0.88033989 0.51978506 0.73417847 0.30660149 0.88371638 0.67450739 0.75118218
##      1594      1598      1599
## 0.38765251 0.45044094 0.81940411
```

AUC and ROC

```
library("pROC")
```

```
## Type 'citation("pROC")' for a citation.
```

```
##
## Attaching package: 'pROC'
```

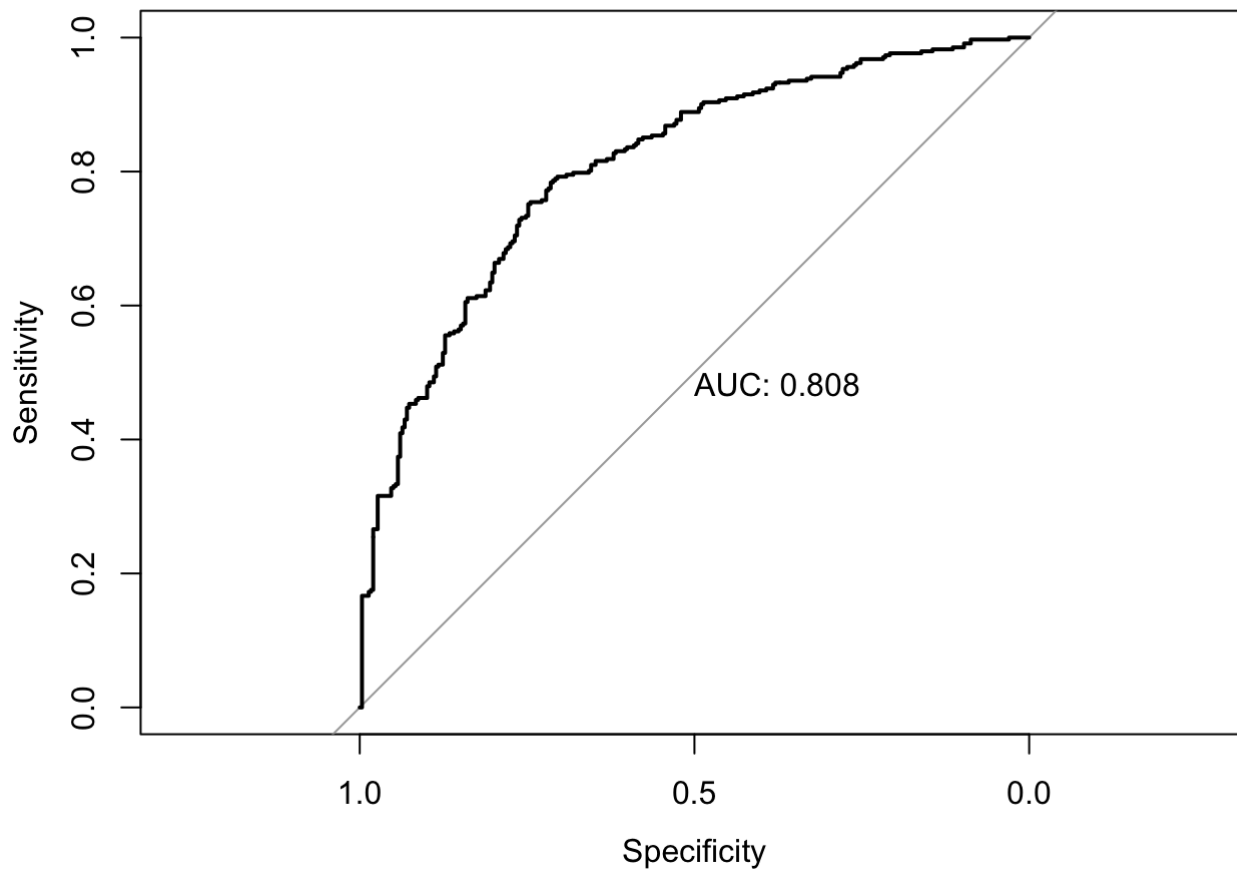
```
## The following objects are masked from 'package:stats':
##
##      cov, smooth, var
```

```
test_prob = predict(randomforestmodlogit, test, type = "response")

test_roc = roc(test$highquality ~ test_prob, plot = TRUE, print.auc = TRUE)
```

```
## Setting levels: control = 0, case = 1
```

```
## Setting direction: controls < cases
```



```
as.numeric(test_roc$auc)
```

```
## [1] 0.8083324
```

AUC and ROC with just one variable

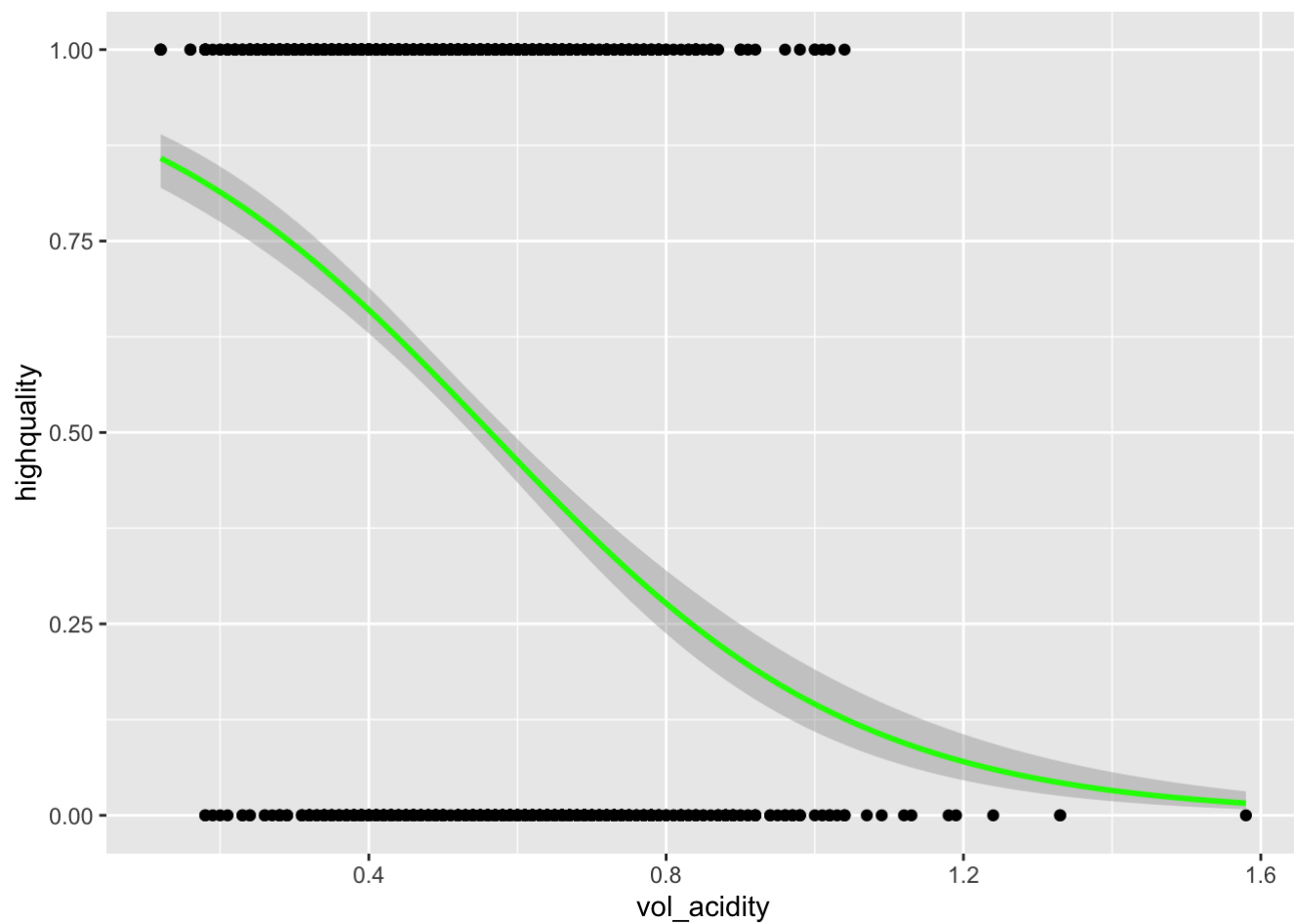
```
library("ggplot2")

simple <- glm(highquality ~ vol_acidity, data = red, family = "binomial"(link = "logit"))
summary(simple)
```

```
##
## Call:
## glm(formula = highquality ~ vol_acidity, family = binomial(link = "logit"),
##      data = red)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -1.8697  -1.1148   0.7156   1.0375   2.0349
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    2.2874     0.1838   12.45  <2e-16 ***
## vol_acidity   -4.0607     0.3334  -12.18  <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 2209.0  on 1598  degrees of freedom
## Residual deviance: 2033.4  on 1597  degrees of freedom
## AIC: 2037.4
##
## Number of Fisher Scoring iterations: 4
```

```
ggplot(red, aes(x = vol_acidity, y = highquality)) +geom_point()+stat_smooth(method="glm", color="green", se=TRUE, method.args = list(family=binomial))
```

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

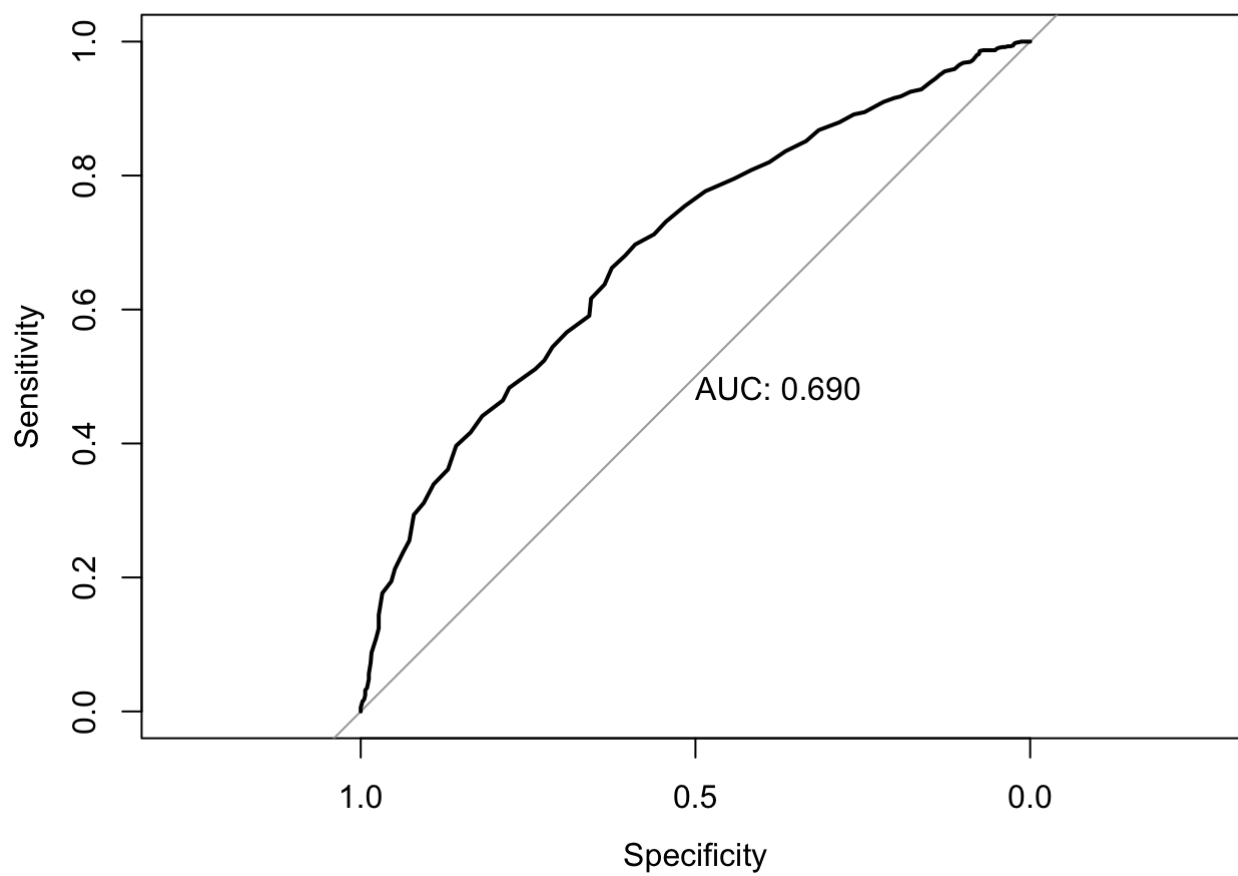


```
test_prop1 = predict(simple, red, type = "response")
```

```
test_roc1 = roc(red$highquality ~ test_prop1, plot = TRUE, print.auc = TRUE)
```

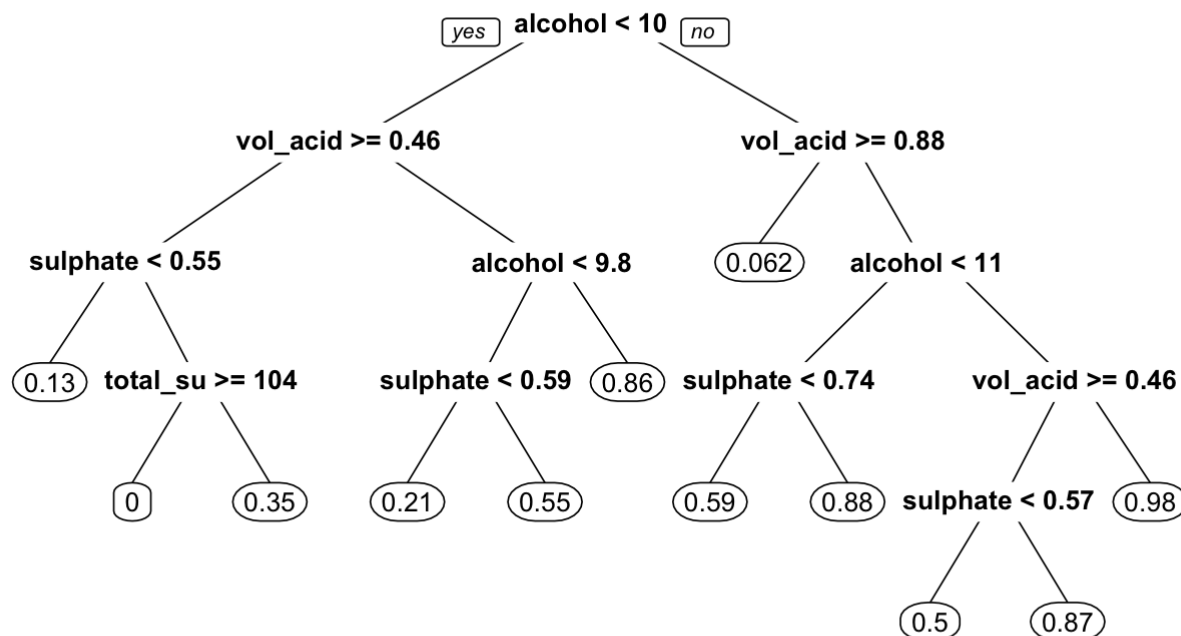
```
## Setting levels: control = 0, case = 1
```

```
## Setting direction: controls < cases
```



```
as.numeric(test_roc1$auc)
```

```
## [1] 0.6900011
```



```

# predictions on test set
set.seed(100)
predictTest = predict(cartmodel, newdata = test, type = "matrix")

# confusion matrix on test set
table(test$highquality, predictTest >= 0.5)

```

```

##
##      FALSE TRUE
##    0    195  103
##    1     82  260

```

Cart Model

```

# Logit
cartmodlogit <- glm(highquality ~ alcohol + sulphates + total_sulfur + fix_acidity, data
= red, family = "binomial"(link = "logit"))
summary(cartmodlogit)

```

```
##
## Call:
## glm(formula = highquality ~ alcohol + sulphates + total_sulfur +
##      fix_acidity, family = binomial(link = "logit"), data = red)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -3.3737  -0.9154   0.3562   0.8762   2.0206
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  -12.172146   0.828396  -14.694 < 2e-16 ***
## alcohol       0.989178   0.068276   14.488 < 2e-16 ***
## sulphates     2.587844   0.370028    6.994 2.68e-12 ***
## total_sulfur  -0.011171   0.001895   -5.895 3.75e-09 ***
## fix_acidity   0.109461   0.035511    3.082 0.00205 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 2209.0  on 1598  degrees of freedom
## Residual deviance: 1752.5  on 1594  degrees of freedom
## AIC: 1762.5
##
## Number of Fisher Scoring iterations: 4
```

```
# Cloglog
cartmodcloglog <- glm(highquality ~ alcohol + sulphates + total_sulfur + fix_acidity, data = red, family = "binomial"(link = "cloglog"))
summary(cartmodcloglog)
```



```
##
## Call:
## glm(formula = highquality ~ alcohol + sulphates + total_sulfur +
##       fix_acidity, family = binomial(link = "cloglog"), data = red)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -4.7058  -0.9408   0.3075   0.9490   1.9387
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  -6.835907   0.481268 -14.204 < 2e-16 ***
## alcohol       0.542953   0.037720  14.394 < 2e-16 ***
## sulphates     1.639060   0.217233   7.545 4.52e-14 ***
## total_sulfur -0.009315   0.001384  -6.732 1.67e-11 ***
## fix_acidity   0.027351   0.021284   1.285  0.199
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 2209.0  on 1598  degrees of freedom
## Residual deviance: 1777.5  on 1594  degrees of freedom
## AIC: 1787.5
##
## Number of Fisher Scoring iterations: 18
```

```
# The logit model performed better with the lower AIC value
```

Compare best logit model with AIC

```
library("AICcmodavg")
```

```
##
## Attaching package: 'AICcmodavg'
```

```
## The following object is masked from 'package:randomForest':
##
##      importance
```

```
models <- list(randomforestmodlogit, cartmodlogit)
mod.names <- c('RandomForest', 'Cart')
aictab(cand.set = models, modnames = mod.names)
```

```
##
## Model selection based on AICc:
##
##           K      AICc Delta_AICc AICcWt Cum.Wt      LL
## RandomForest 5 1694.21      0.00      1      1 -842.09
## Cart         5 1762.56     68.35      0      1 -876.26
```

```
# The random forest logit model performed the best
```

Compare best model with BIC

```
library("flexmix")
BIC(randomforestmodlogit)
```

```
## [1] 1721.058
```

```
BIC(randomforestmodcloglog)
```

```
## [1] 1737.845
```

```
BIC(cartmodlogit)
```

```
## [1] 1789.404
```

```
BIC(cartmodcloglog)
```

```
## [1] 1814.418
```

```
# The random forest logit model performed the best
```

Confusion matrix for random forest logit model

```
confusionred = predict(randomforestmodlogit, newdata = red, type = "response")

# confusion matrix on test set
table(red$highquality, confusionred >= 0.5)
```

```
##  
##      FALSE TRUE  
##    0    548  196  
##    1    216  639
```

Predictions for random forest logit model

```
pred_test <- predict(randomforestmodlogit, test, type = "response")  
  
pred_test
```

##	3	4	5	10	16	18	19
##	0.24006880	0.52789064	0.21686414	0.50812152	0.13867752	0.56805457	0.20109254
##	22	28	30	38	41	47	48
##	0.42364540	0.58076699	0.38187954	0.59357547	0.60541307	0.05638181	0.51951414
##	49	51	53	56	59	60	65
##	0.43019840	0.30073364	0.43459537	0.24266029	0.36211052	0.37446288	0.48498736
##	66	70	71	73	74	77	80
##	0.48498736	0.66550098	0.28235396	0.11468249	0.17116376	0.67706290	0.22946232
##	81	84	85	87	88	91	92
##	0.38434763	0.44994570	0.71372134	0.81269343	0.43771981	0.12488112	0.81269343
##	97	100	102	104	105	106	110
##	0.45180389	0.24182616	0.59957642	0.17168420	0.30465596	0.17168420	0.05669027
##	111	112	114	117	119	120	122
##	0.46170336	0.20639675	0.56940622	0.44937409	0.59589245	0.10540486	0.59589245
##	125	130	133	136	141	144	145
##	0.17454268	0.47573192	0.88460563	0.25641373	0.25641373	0.47019352	0.98548676
##	147	151	153	155	159	161	163
##	0.18741376	0.71199465	0.24118269	0.44034290	0.21501940	0.15424192	0.45821490
##	167	168	172	177	185	188	189
##	0.22409934	0.18953659	0.42272086	0.40084611	0.19399307	0.25256465	0.13003255
##	190	192	193	201	203	208	211
##	0.13218057	0.51206935	0.12192200	0.79856559	0.43027971	0.11227661	0.94372425
##	212	214	215	216	217	218	220
##	0.23949363	0.29462991	0.30759231	0.43866216	0.56377995	0.14689457	0.12986042
##	222	224	228	230	231	232	235
##	0.23247399	0.39670029	0.26752645	0.64457164	0.77766942	0.47608964	0.06682702
##	237	238	242	243	244	245	247
##	0.19063712	0.20849103	0.82087382	0.17002266	0.67646997	0.67646997	0.19535300
##	248	249	251	254	256	264	265
##	0.14653731	0.35125228	0.70288800	0.11931143	0.12202885	0.42251877	0.77916500
##	266	269	272	276	279	285	286
##	0.80711622	0.50344768	0.87483178	0.57130009	0.97446971	0.32999609	0.32999609
##	287	289	290	295	300	301	302
##	0.63496399	0.69463023	0.36131597	0.55129126	0.33509483	0.59572880	0.79587338
##	303	304	307	313	320	321	324
##	0.41165107	0.27204195	0.20435986	0.25470497	0.29176280	0.68916140	0.26289903
##	326	327	328	331	333	338	340
##	0.32170068	0.85787373	0.90657600	0.94054119	0.12860589	0.56472573	0.88983001
##	344	345	347	354	363	364	366
##	0.66228068	0.68982583	0.84745884	0.92386355	0.40286206	0.74339678	0.93834097
##	367	370	373	375	376	377	378
##	0.44355920	0.97850481	0.87743922	0.78031275	0.90189455	0.87515687	0.97850481
##	380	381	382	384	385	392	396
##	0.70150349	0.65714868	0.62763464	0.65714868	0.32025802	0.62763464	0.95715624
##	397	405	413	419	420	422	423
##	0.11072664	0.20675079	0.13582974	0.77020269	0.26802335	0.69125350	0.22216089
##	425	428	430	435	441	445	446
##	0.22216089	0.40277653	0.32784541	0.62568563	0.70777919	0.93413308	0.32689043
##	451	460	462	463	465	467	472
##	0.74289381	0.19464737	0.29995555	0.92477154	0.55427468	0.80312295	0.76084070
##	474	478	480	481	484	485	488
##	0.83494087	0.94349416	0.43048988	0.54068853	0.72818356	0.97215159	0.34928339

##	489	490	491	494	497	498	501
##	0.82229147	0.71109362	0.38820824	0.64444930	0.27636155	0.71705477	0.27636155
##	506	508	510	512	515	519	522
##	0.94197356	0.36855117	0.87805407	0.40270830	0.87663195	0.93281411	0.35179506
##	523	525	526	527	539	540	541
##	0.43596448	0.23574293	0.34605728	0.69166380	0.91684119	0.79067776	0.33434428
##	545	546	549	550	551	553	555
##	0.42365538	0.16061952	0.66697078	0.36606459	0.37808419	0.56543070	0.72091694
##	558	561	563	565	567	569	571
##	0.72091694	0.74635299	0.14387840	0.92935383	0.20667837	0.76484762	0.87692510
##	574	575	578	580	582	583	587
##	0.28595807	0.53585414	0.22627342	0.52861287	0.38002007	0.29689198	0.86089516
##	590	593	595	597	598	600	602
##	0.83511782	0.33060442	0.20381128	0.43790682	0.49453337	0.35305750	0.28353983
##	607	611	613	616	621	624	626
##	0.94521424	0.40906300	0.40122182	0.24600419	0.16493964	0.92396940	0.38478177
##	632	633	638	639	640	641	644
##	0.60526652	0.63251372	0.03693294	0.28835011	0.89903179	0.32321697	0.24690166
##	649	652	653	655	657	658	660
##	0.79606704	0.19901970	0.99329501	0.34792620	0.40958829	0.62339714	0.51004428
##	664	665	671	672	673	677	683
##	0.81425791	0.58642668	0.52667590	0.24700448	0.01313453	0.49457701	0.37713648
##	685	688	691	692	693	695	696
##	0.02353841	0.30853776	0.19926076	0.08792767	0.47830984	0.17339004	0.92968944
##	697	699	703	704	706	708	711
##	0.33118031	0.21822469	0.32282879	0.39592542	0.09514858	0.56086414	0.09703634
##	713	715	716	719	720	723	731
##	0.18728926	0.16052021	0.27155586	0.31642036	0.22428109	0.50744957	0.39042992
##	733	734	737	740	743	744	746
##	0.17713379	0.31147678	0.11882456	0.21223885	0.30029207	0.33233049	0.45423361
##	749	750	751	753	754	756	759
##	0.47090034	0.45423361	0.24423781	0.25521181	0.24423781	0.24061283	0.17874689
##	762	764	766	770	771	772	773
##	0.21832154	0.21832154	0.21287996	0.23550990	0.19045961	0.07048717	0.09786165
##	775	776	779	781	783	789	790
##	0.57996000	0.16206695	0.66183909	0.28239984	0.12989081	0.37973849	0.09437875
##	791	801	805	810	811	814	815
##	0.42161900	0.11964960	0.48671568	0.60727563	0.60772687	0.86887538	0.84554487
##	819	820	824	825	827	836	837
##	0.15085766	0.21151380	0.43070016	0.59452832	0.88097191	0.16263773	0.73172430
##	838	839	840	841	842	847	850
##	0.73172430	0.90163251	0.45005102	0.92522550	0.46356092	0.38077515	0.38523107
##	851	852	853	854	855	856	860
##	0.44134957	0.44134957	0.46348365	0.77796635	0.77796635	0.78141884	0.81336010
##	862	863	866	868	870	873	877
##	0.29801703	0.59081792	0.18470531	0.84439382	0.59381839	0.53560900	0.74967371
##	878	882	883	885	886	887	888
##	0.68842602	0.66426561	0.94676764	0.27035098	0.44881228	0.39943319	0.87920584
##	890	891	892	895	898	899	900
##	0.07413691	0.57461423	0.18472125	0.21369246	0.54837049	0.95762419	0.34173363
##	903	904	908	909	912	917	922
##	0.61749313	0.64721685	0.79467120	0.57631602	0.85616038	0.47643991	0.73857291

##	923	924	925	927	928	930	935
##	0.84865159	0.57254719	0.84029182	0.75197156	0.14108907	0.94254625	0.61571145
##	939	940	943	945	948	950	952
##	0.96015192	0.82418379	0.51350024	0.93921165	0.96007673	0.96739607	0.96007673
##	953	955	958	964	965	966	976
##	0.89304382	0.91523844	0.75906185	0.88720837	0.82503678	0.85318054	0.28346202
##	977	981	982	983	984	988	991
##	0.28346202	0.70374096	0.26394962	0.93219598	0.70374096	0.31207909	0.40407434
##	994	995	1000	1002	1004	1007	1008
##	0.32169795	0.24614218	0.89973442	0.78374144	0.96113831	0.94122194	0.94084017
##	1012	1019	1021	1023	1024	1027	1029
##	0.72592191	0.90328264	0.91151720	0.64109441	0.94561683	0.94593820	0.50667880
##	1031	1032	1036	1042	1048	1051	1052
##	0.77233191	0.71727212	0.77472663	0.49873341	0.60752858	0.60752858	0.67804325
##	1054	1059	1064	1067	1068	1072	1078
##	0.97117164	0.81405117	0.92780908	0.92897417	0.89593617	0.08796075	0.82775448
##	1079	1080	1082	1086	1087	1089	1091
##	0.82775448	0.40453879	0.37324283	0.28421079	0.91462483	0.59911292	0.87689507
##	1092	1094	1096	1098	1099	1102	1103
##	0.87464611	0.95181067	0.66933405	0.34983074	0.97135198	0.88355318	0.74464695
##	1108	1110	1111	1115	1116	1118	1120
##	0.94262703	0.67961164	0.55003024	0.96048872	0.67849027	0.67849027	0.91323567
##	1121	1124	1126	1129	1131	1133	1134
##	0.95982776	0.82569184	0.88512330	0.42390723	0.59465043	0.98621359	0.81790944
##	1137	1139	1141	1142	1145	1147	1150
##	0.85680402	0.20660676	0.34979431	0.79255354	0.55703889	0.62188669	0.91677782
##	1153	1157	1159	1160	1164	1169	1182
##	0.32861855	0.90970330	0.84798042	0.75007048	0.34911691	0.89257541	0.83146700
##	1184	1185	1186	1187	1191	1195	1199
##	0.18325571	0.39228204	0.83216767	0.75844840	0.93726418	0.16205917	0.80568967
##	1200	1201	1203	1205	1208	1209	1211
##	0.18261742	0.43683111	0.90407751	0.84882533	0.44462253	0.84882533	0.53707859
##	1212	1214	1215	1217	1218	1221	1222
##	0.33645556	0.79664197	0.71480329	0.25697732	0.90659821	0.89545422	0.89545422
##	1225	1228	1230	1231	1233	1234	1235
##	0.77019805	0.43944184	0.35954537	0.93764299	0.35954537	0.47608892	0.82878055
##	1236	1242	1246	1247	1248	1252	1257
##	0.70226268	0.44890191	0.60960225	0.23445159	0.60960225	0.31158618	0.15058729
##	1258	1260	1263	1264	1265	1267	1270
##	0.60797958	0.69403671	0.38122840	0.18918454	0.88943306	0.50245646	0.97152715
##	1272	1277	1278	1281	1284	1285	1288
##	0.78782193	0.91209807	0.35315581	0.57652097	0.35403270	0.72038518	0.90845738
##	1289	1292	1293	1296	1298	1299	1300
##	0.52796774	0.59190236	0.93827703	0.15149263	0.87066317	0.85966749	0.06683777
##	1301	1303	1304	1309	1310	1312	1315
##	0.87211203	0.89325440	0.81440601	0.23809254	0.18736687	0.93088901	0.48312749
##	1317	1320	1322	1323	1325	1326	1328
##	0.79987457	0.37260225	0.79987457	0.86432689	0.65726913	0.65726913	0.65726913
##	1330	1331	1339	1341	1343	1350	1351
##	0.21628945	0.21628945	0.33063230	0.57850387	0.51740015	0.56190596	0.29613263
##	1354	1355	1356	1357	1366	1369	1371
##	0.31505119	0.40899359	0.57777135	0.57441114	0.36658102	0.15039302	0.30081941

##	1372	1374	1376	1377	1378	1379	1380
##	0.91160867	0.08195041	0.22071490	0.18491208	0.83265348	0.47985390	0.62870483
##	1384	1386	1387	1392	1396	1399	1400
##	0.18145955	0.06251845	0.32286340	0.60775822	0.33954980	0.49039723	0.69724047
##	1401	1403	1404	1407	1408	1409	1411
##	0.10211360	0.94896201	0.85527482	0.92066898	0.73685309	0.96956379	0.55013345
##	1422	1423	1426	1428	1430	1433	1435
##	0.35313054	0.80043906	0.66331164	0.81035058	0.92040632	0.92574770	0.18849461
##	1437	1438	1439	1443	1444	1445	1449
##	0.16556629	0.38062331	0.50395641	0.43582198	0.77179526	0.48436224	0.37160792
##	1450	1451	1453	1455	1456	1458	1466
##	0.87394441	0.87184058	0.78758198	0.78198047	0.45960983	0.17120492	0.33900665
##	1469	1470	1474	1476	1477	1479	1483
##	0.47414501	0.11210856	0.58301194	0.96416743	0.19663663	0.28090696	0.55985828
##	1484	1485	1486	1487	1488	1491	1492
##	0.80671170	0.47494678	0.31103235	0.45669868	0.66158978	0.95967599	0.79522301
##	1494	1497	1499	1500	1501	1505	1516
##	0.14135590	0.14135590	0.42318892	0.71090938	0.26400037	0.85817699	0.13049129
##	1517	1527	1528	1529	1530	1531	1532
##	0.82595385	0.43733113	0.70351630	0.63560644	0.34422384	0.85054796	0.42672777
##	1533	1534	1535	1536	1543	1552	1553
##	0.52363402	0.23235943	0.81214100	0.37906001	0.36046967	0.24069329	0.62386686
##	1558	1559	1560	1561	1567	1569	1571
##	0.48014133	0.12503776	0.14727854	0.14727854	0.89505671	0.25504396	0.96429949
##	1572	1576	1579	1581	1582	1585	1587
##	0.82810158	0.86164481	0.72404537	0.90818659	0.73417847	0.91848799	0.88684408
##	1594	1597	1598				
##	0.38765251	0.75118218	0.45044094				

Model Diagnostics

```
accuracy = (548+639)/(548+196+216+639)
accuracy
```

```
## [1] 0.742339
```

```
sensitivity = 639/(639+196)
sensitivity
```

```
## [1] 0.7652695
```

```
specificity = 548/(548+216)
specificity
```

```
## [1] 0.7172775
```

AUC and ROC

```
library("pROC")
```

```
## Type 'citation("pROC")' for a citation.
```

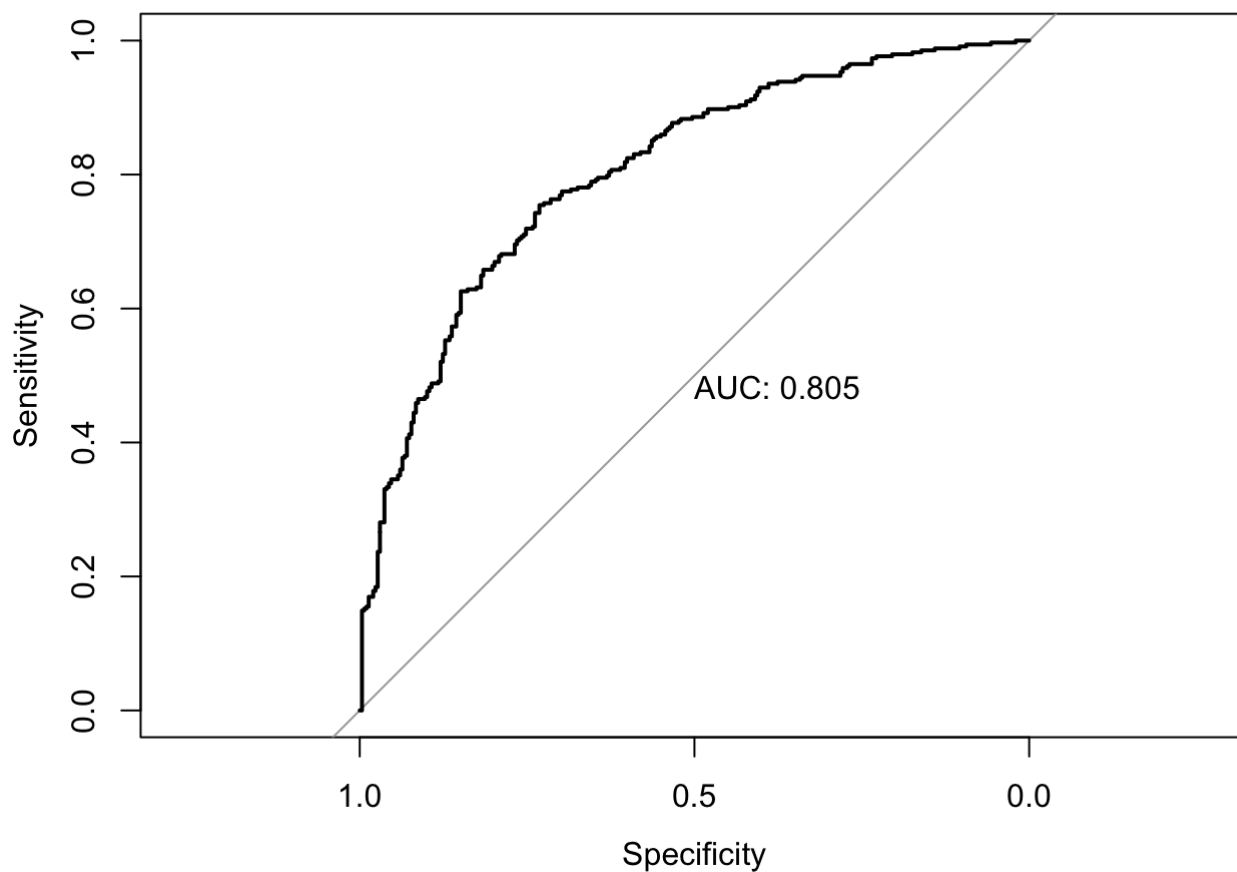
```
##  
## Attaching package: 'pROC'
```

```
## The following objects are masked from 'package:stats':  
##  
## cov, smooth, var
```

```
test_prob = predict(randomforestmodlogit, test, type = "response")  
  
test_roc = roc(test$highquality ~ test_prob, plot = TRUE, print.auc = TRUE)
```

```
## Setting levels: control = 0, case = 1
```

```
## Setting direction: controls < cases
```




```
as.numeric(test_roc$auc)
```

```
## [1] 0.8052023
```

AUC and ROC with just one variable

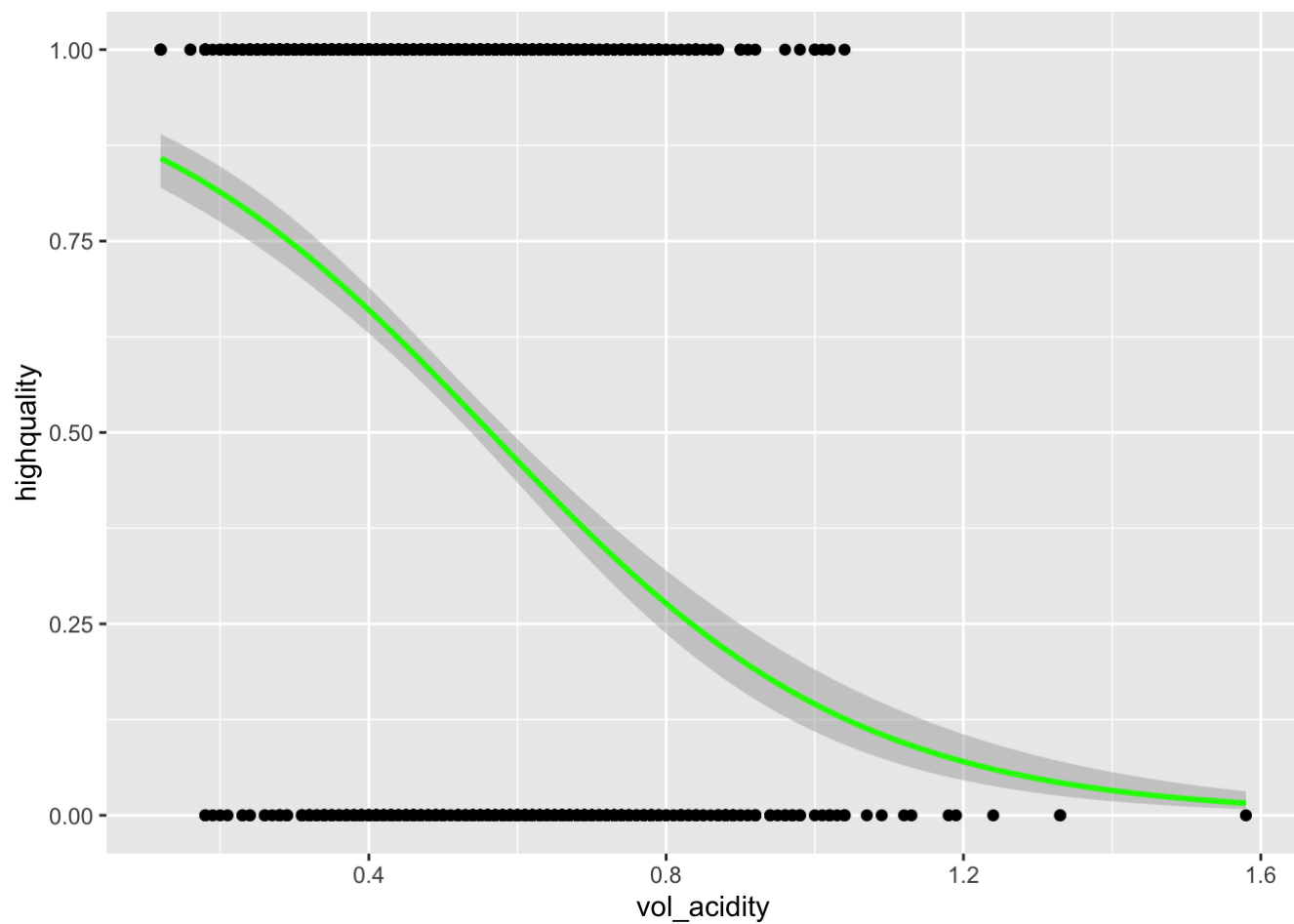
```
library("ggplot2")
```

```
simple <- glm(highquality ~ vol_acidity, data = red, family = "binomial"(link = "logit"))
summary(simple)
```

```
##
## Call:
## glm(formula = highquality ~ vol_acidity, family = binomial(link = "logit"),
##      data = red)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -1.8697  -1.1148   0.7156   1.0375   2.0349
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)   2.2874     0.1838  12.45  <2e-16 ***
## vol_acidity  -4.0607     0.3334  -12.18  <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 2209.0  on 1598  degrees of freedom
## Residual deviance: 2033.4  on 1597  degrees of freedom
## AIC: 2037.4
##
## Number of Fisher Scoring iterations: 4
```

```
ggplot(red, aes(x = vol_acidity, y = highquality)) +geom_point()+stat_smooth(method="glm", color="green", se=TRUE, method.args = list(family=binomial))
```

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

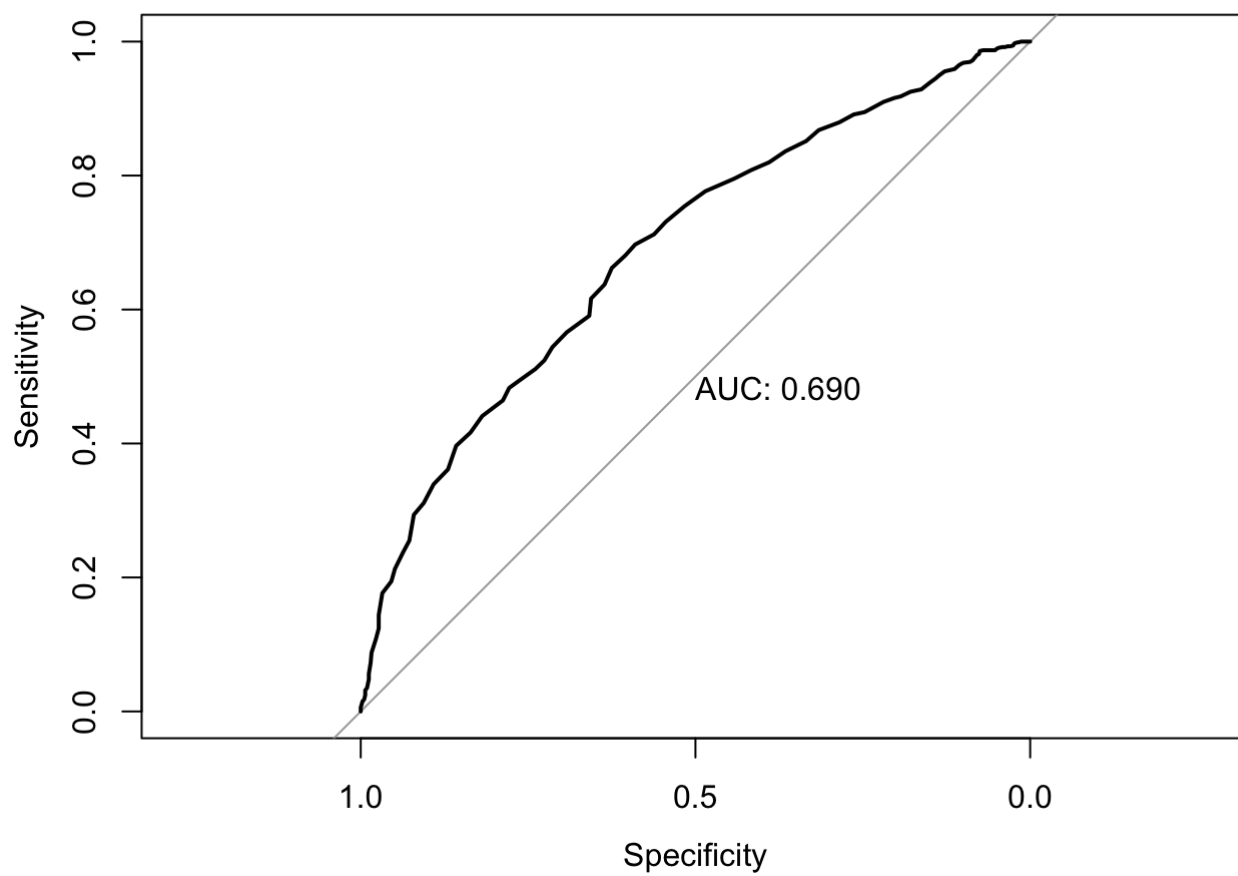


```
test_prop1 = predict(simple, red, type = "response")
```

```
test_roc1 = roc(red$highquality ~ test_prop1, plot = TRUE, print.auc = TRUE)
```

```
## Setting levels: control = 0, case = 1
```

```
## Setting direction: controls < cases
```



```
as.numeric(test_roc1$auc)
```

```
## [1] 0.6900011
```