

# STAT467 PROJECT

## GROUP 6

Mustafa Ugur Yalcin - 2502458

Melih Can Kanmaz - 2502169

Tevfik Oguz - 2442051

# CONTENT

- About the dataset
- Inference of Mean Vector
- Comparisons of Several Multivariate Means
- Principal Component Analysis and PCR
- Factor Analysis and Factor Rotation
- Discrimination and Classification
- Clustering
- Canonical Correlation Analysis

# PROJECT AIM

- Does the quality differ across wine colours?
- What factors influence the quality of wine?
- What factors affect the alcohol content?



# ABOUT THE DATA

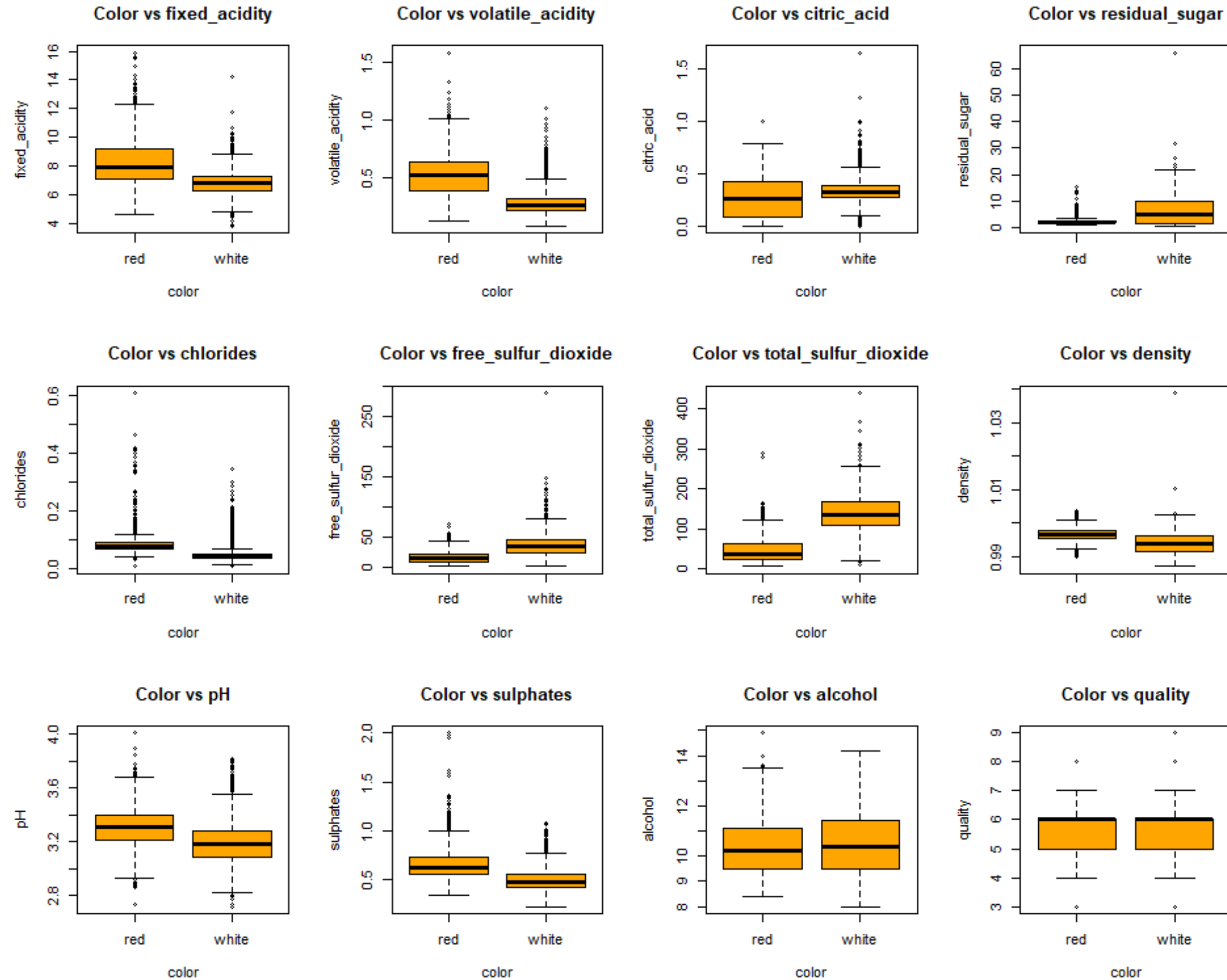




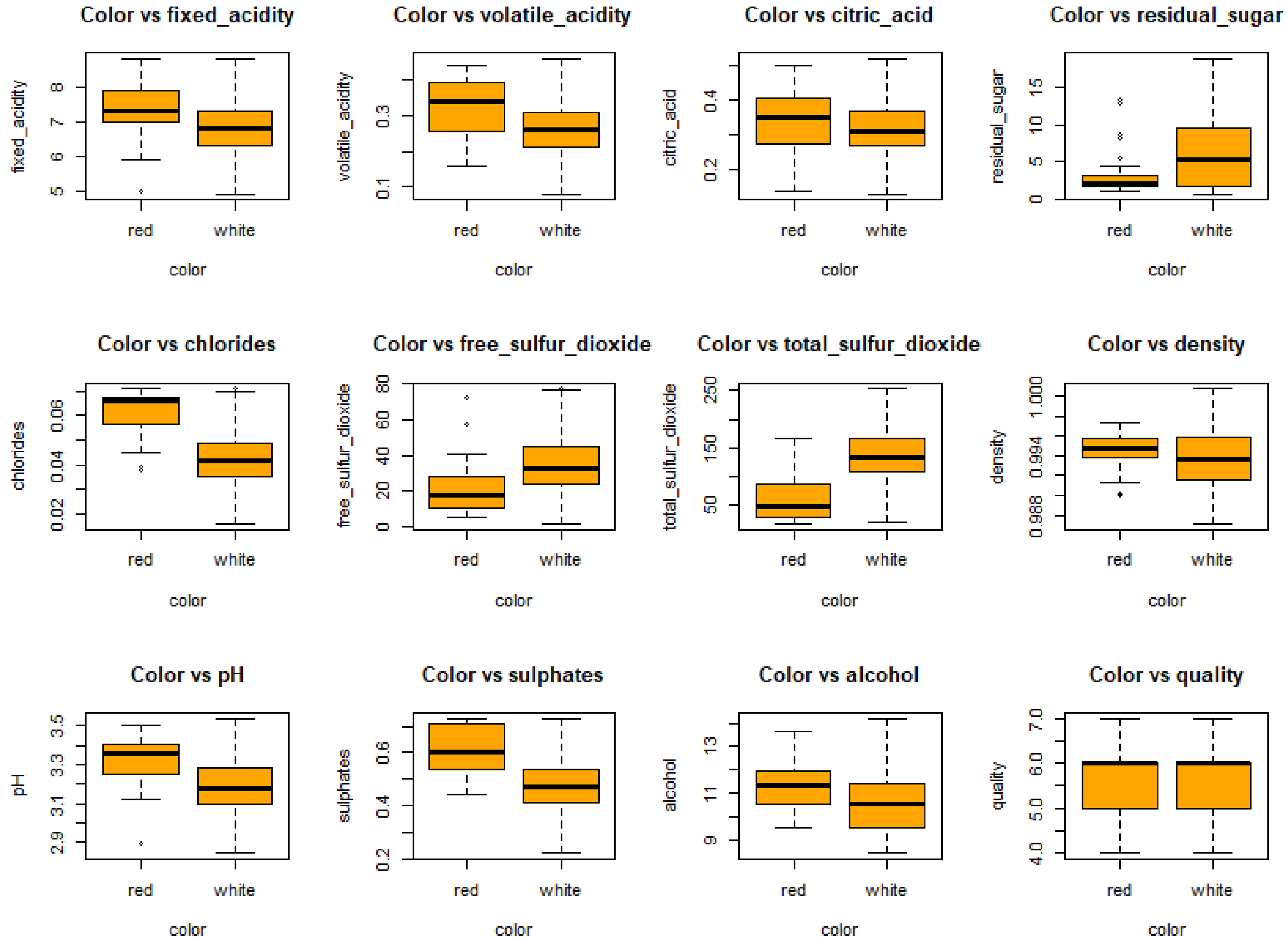
# SUMMARY OF THE DATA

```
> summary(data)
fixed_acidity    volatile_acidity    citric_acid    residual_sugar    chlorides    free_sulfur_dioxide
Min.   : 3.800    Min.   :0.0800    Min.   :0.0000    Min.   : 0.600    Min.   :0.00900    Min.   : 1.00
1st Qu.: 6.400    1st Qu.:0.2300    1st Qu.:0.2500    1st Qu.: 1.800    1st Qu.:0.03800    1st Qu.: 17.00
Median : 7.000    Median :0.2900    Median :0.3100    Median : 3.000    Median :0.04700    Median : 29.00
Mean   : 7.215    Mean   :0.3397    Mean   :0.3186    Mean   : 5.443    Mean   :0.05603    Mean   : 30.53
3rd Qu.: 7.700    3rd Qu.:0.4000    3rd Qu.:0.3900    3rd Qu.: 8.100    3rd Qu.:0.06500    3rd Qu.: 41.00
Max.   :15.900    Max.   :1.5800    Max.   :1.6600    Max.   :65.800    Max.   :0.61100    Max.   :289.00
total_sulfur_dioxide    density    pH    sulphates    alcohol    quality
Min.   : 6.0    Min.   :0.9871    Min.   :2.720    Min.   :0.2200    Min.   : 8.00    Min.   :3.000
1st Qu.: 77.0    1st Qu.:0.9923    1st Qu.:3.110    1st Qu.:0.4300    1st Qu.: 9.50    1st Qu.:5.000
Median :118.0    Median :0.9949    Median :3.210    Median :0.5100    Median :10.30    Median :6.000
Mean   :115.7    Mean   :0.9947    Mean   :3.219    Mean   :0.5313    Mean   :10.49    Mean   :5.818
3rd Qu.:156.0    3rd Qu.:0.9970    3rd Qu.:3.320    3rd Qu.:0.6000    3rd Qu.:11.30    3rd Qu.:6.000
Max.   :440.0    Max.   :1.0390    Max.   :4.010    Max.   :2.0000    Max.   :14.90    Max.   :9.000
color
Length:6497
Class :character
Mode  :character
```

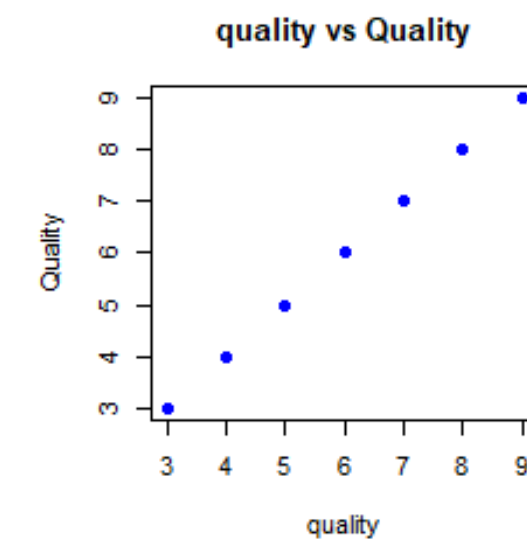
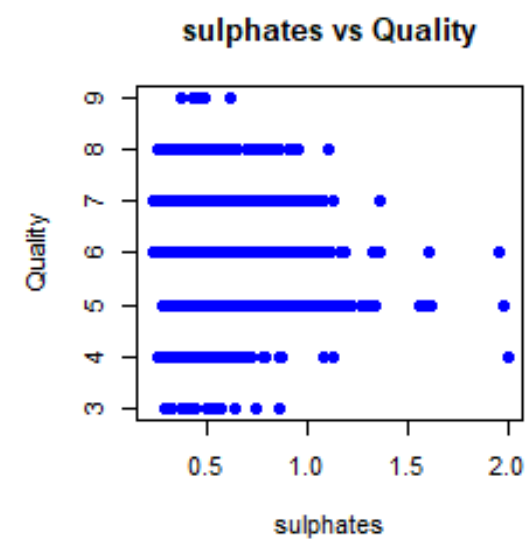
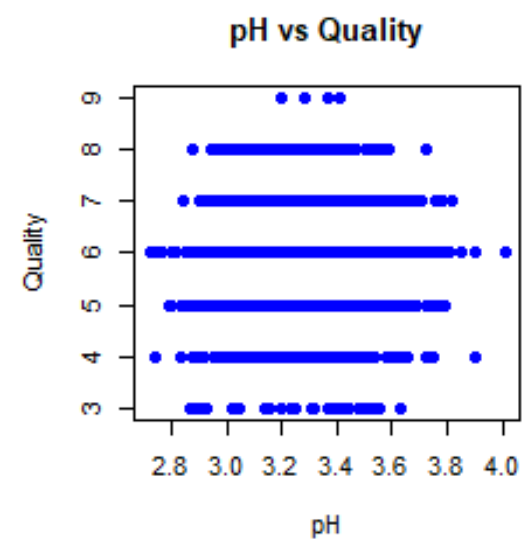
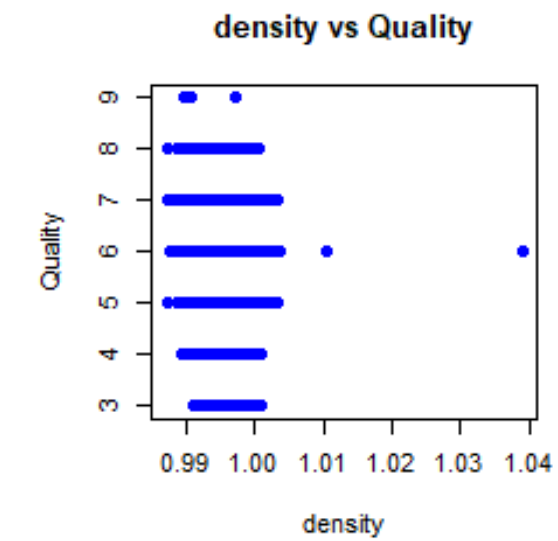
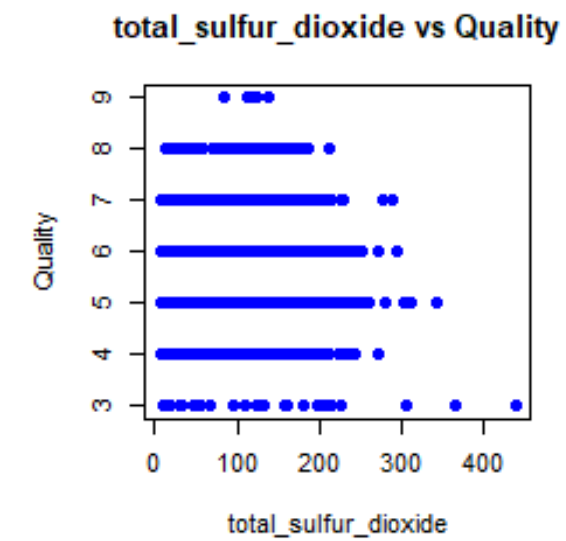
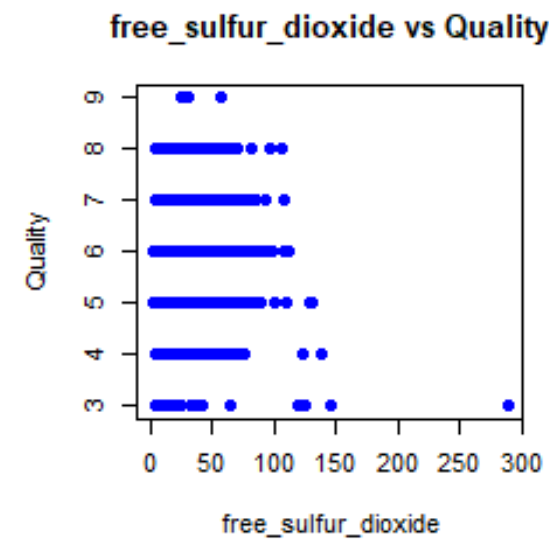
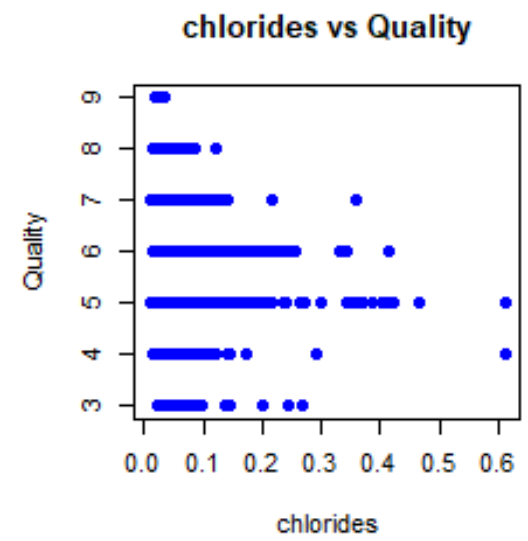
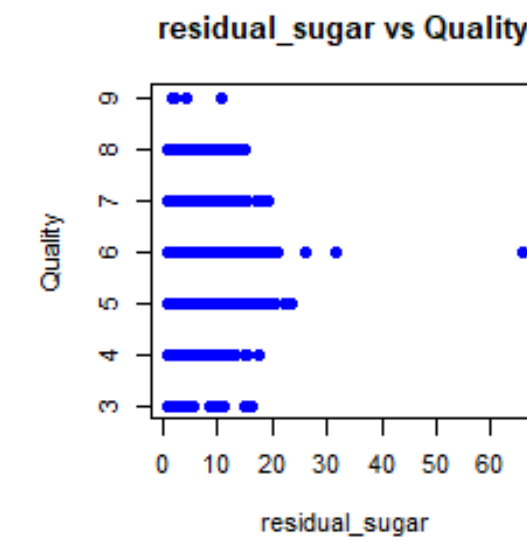
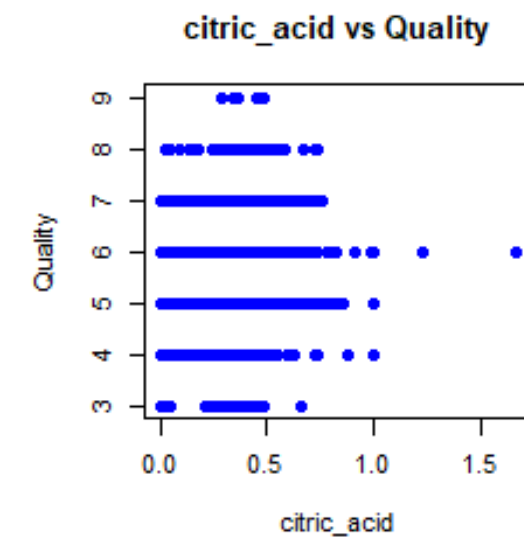
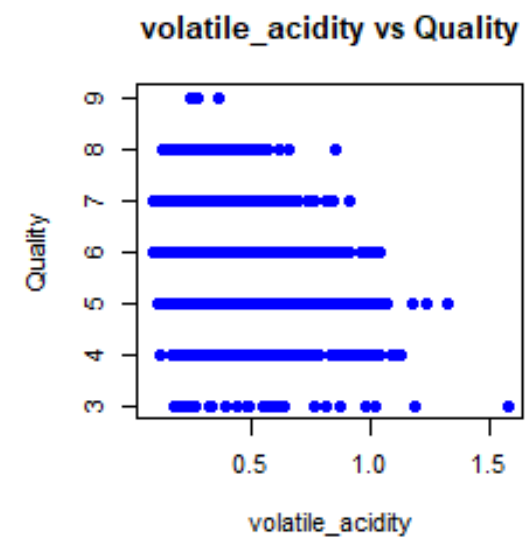
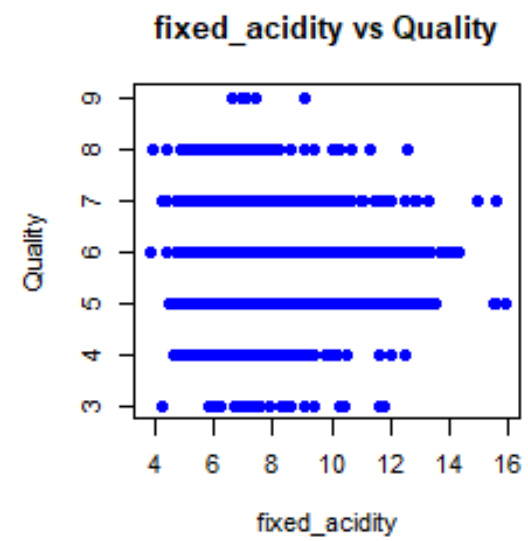
# DISTRIBUTIONS OF COLOR



# REMOVING OUTLIERS

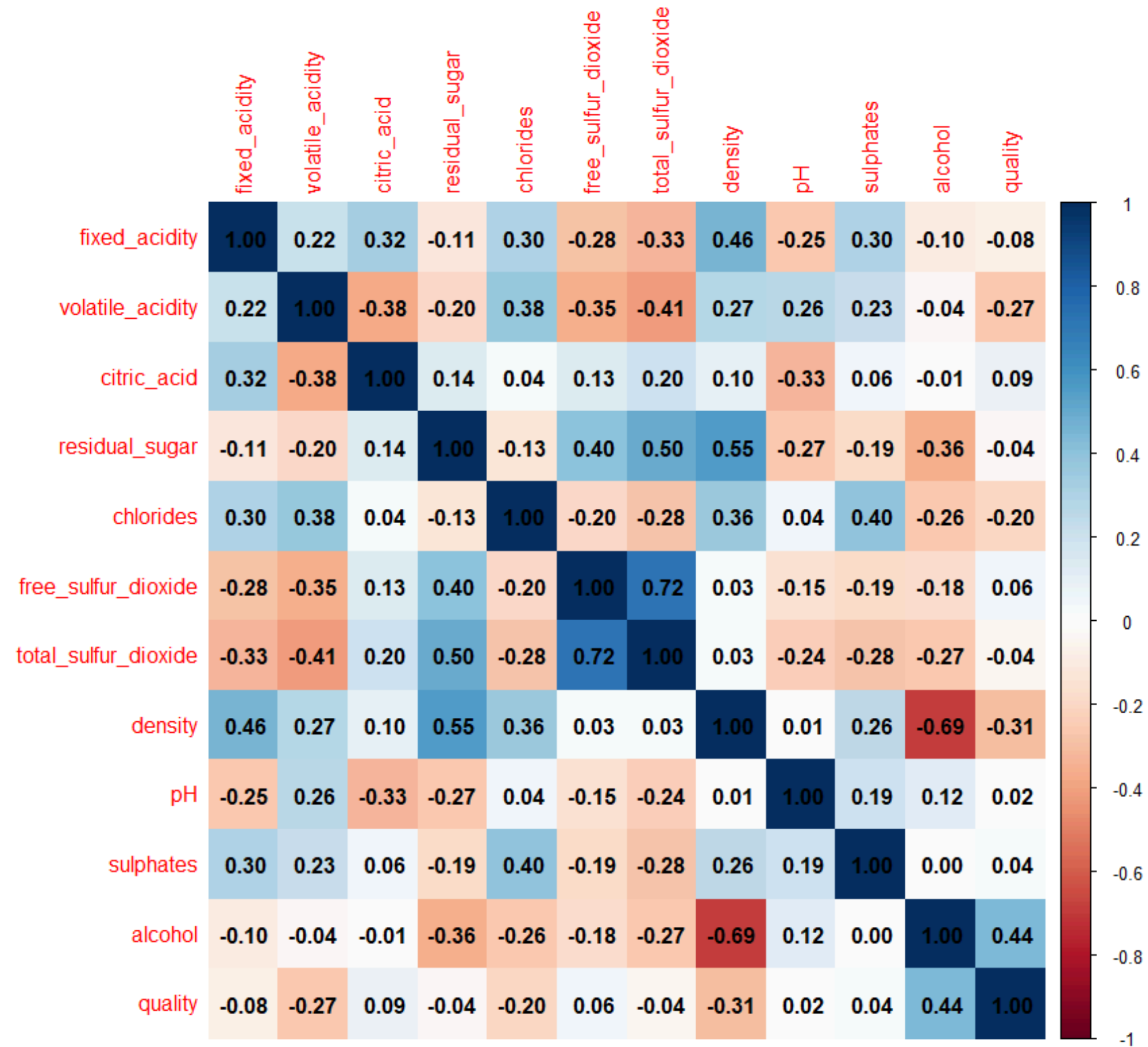


# DISTRIBUTIONS OF QUALITY

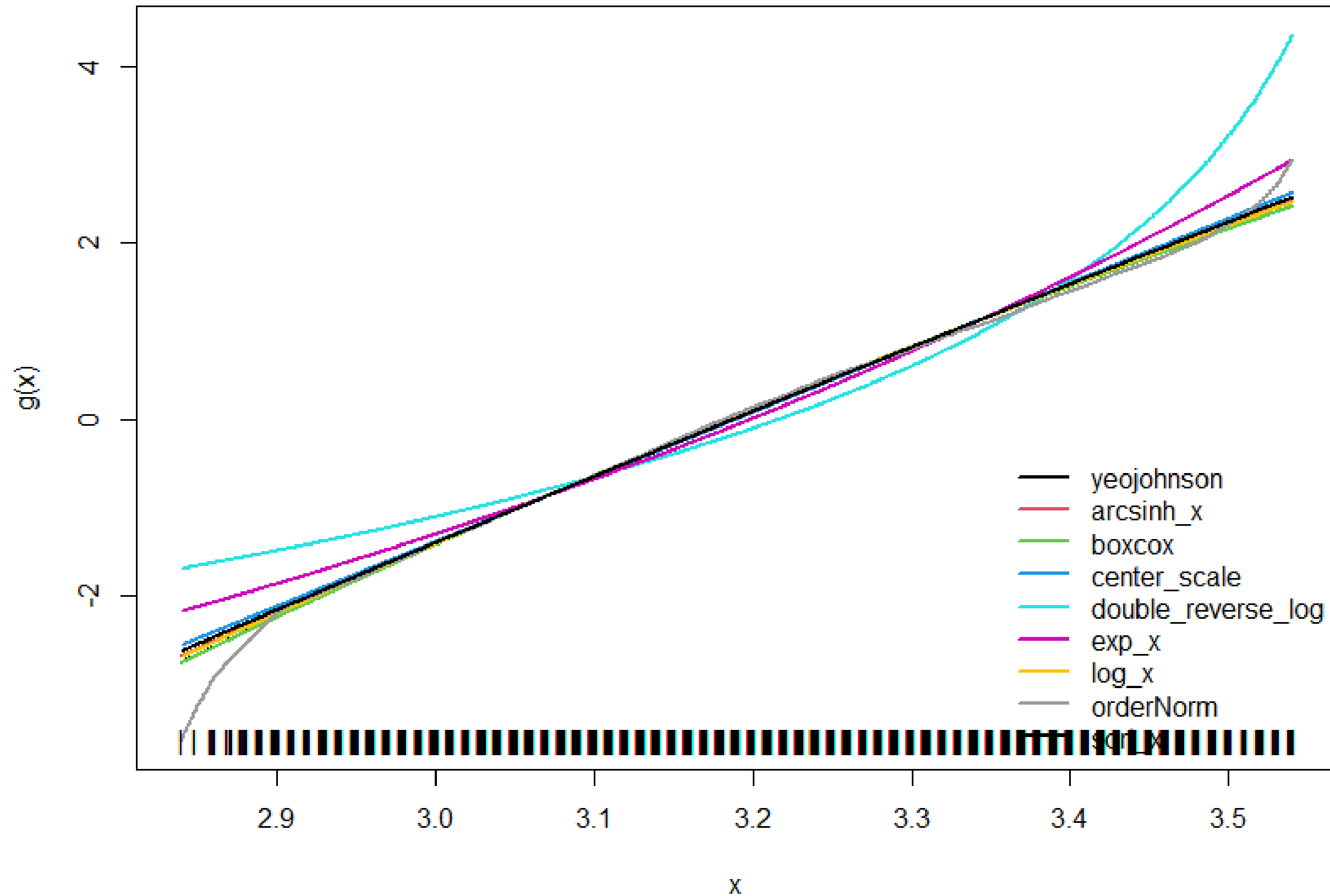




# CORRELATION OF VARIABLES



# INFERENCE OF MEAN VECTOR



# INFERENCE OF MEAN VECTOR

## NORMALITY ASSUMPTION

```
> test$multivariateNormality
```

	Test	Statistic	p value	Result
1	Mardia Skewness	138.822630619252	5.04302634059072e-29	NO
2	Mardia Kurtosis	-2.91040498967062	0.00360960693973933	NO
3	MVN	<NA>	<NA>	NO

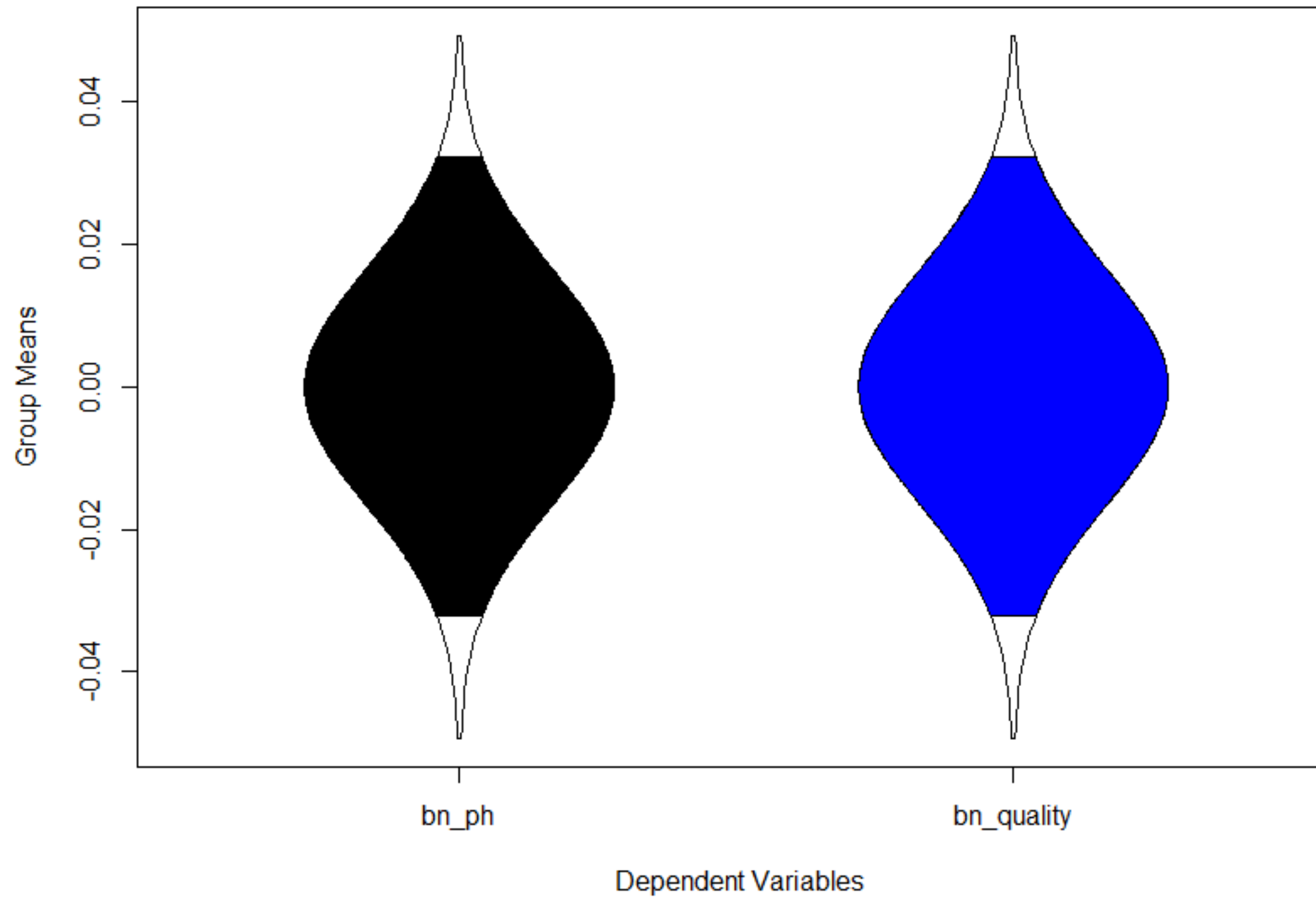
```
> test<-mvn(y,mvnTest = "mardia")
> test$univariateNormality
```

	Test	Variable	Statistic	p value	Normality
1	Anderson-Darling	bn_ph	2.1183	<0.001	NO
2	Anderson-Darling	bn_quality	271.7797	<0.001	NO

```
>
```

# VIOLIN GRAPHS

95% confidence limits



# INFERENCE OF MEAN VECTOR

## HOTELLING'S T TEST

```
> #Hotelling T test
> HotellingsT2(y,mu=mu0)

Hotelling's one sample T2-test

data:  y
T.2 = 3532.6, df1 = 2, df2 = 3703,
p-value < 2.2e-16
alternative hypothesis: true location is not equal to c(1,1)
```

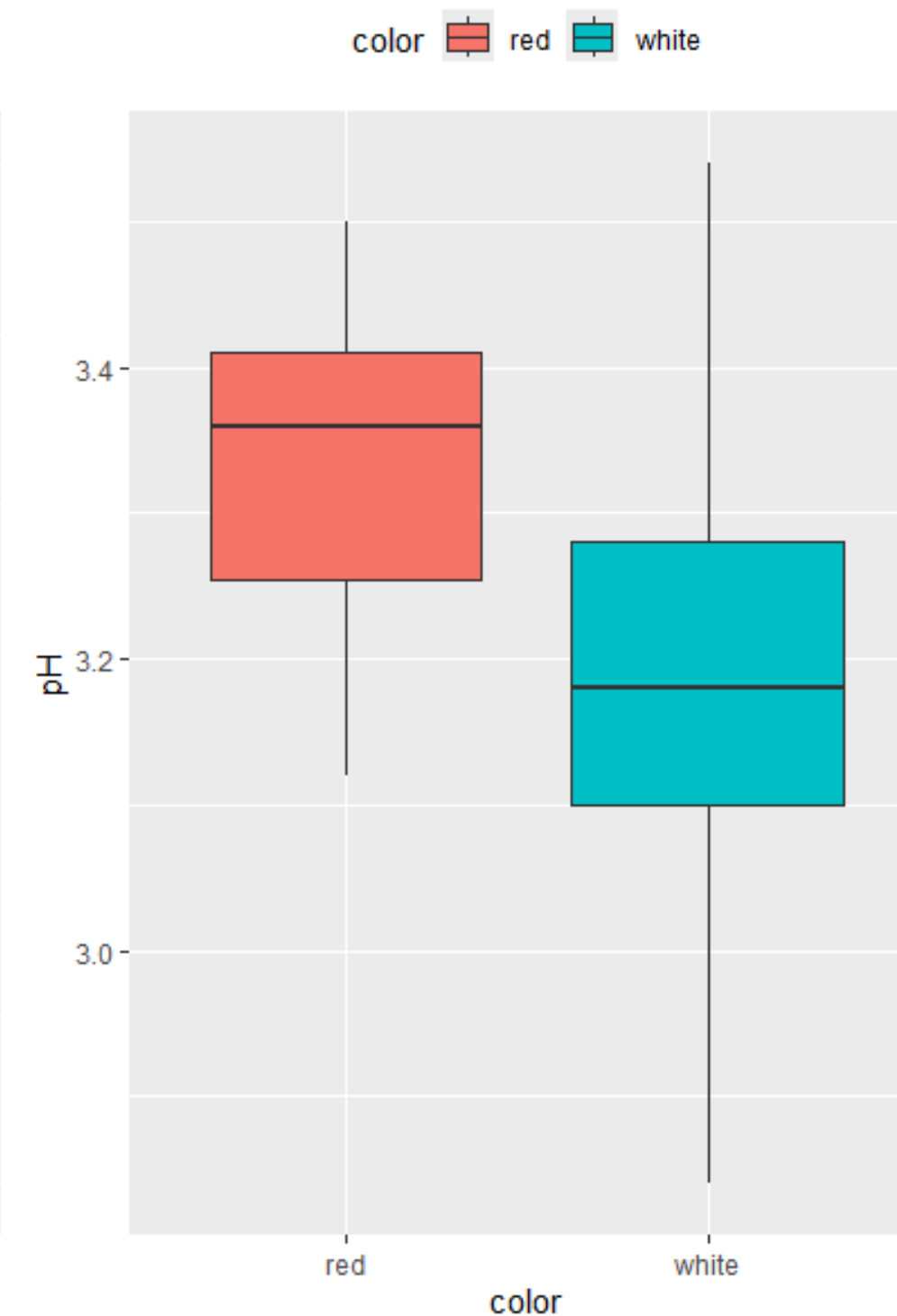


# COMPARISONS OF SEVERAL MULTIVARIATE MEANS

The Box Plot of quality by color.



The Box Plot of pH by color.



# NORMALITY TESTS

```
> subset_data1 %>% group_by(color) %>% shapiro_test(log_pH, log_quality)
# A tibble: 4 × 4
  color variable    statistic      p
  <chr> <chr>         <dbl>    <dbl>
1 red   log_pH         0.837 5.83e- 6
2 red   log_quality     0.815 1.61e- 6
3 white log_pH         0.996 1.73e- 8
4 white log_quality     0.835 6.35e-52

> library(heplots)
> boxM(Y = cbind(subset_data1$log_pH, subset_data1$log_quality), group = factor(subset_data1$color))

      Box's M-test for Homogeneity of Covariance
      Matrices

data:  cbind(subset_data1$log_pH, subset_data1$log_quality)
Chi-Sq (approx.) = 7.9875, df = 3, p-value
= 0.04627

> leveneTest(log_pH ~ color, data = subset_data1)
Levene's Test for Homogeneity of Variance (center = median)
      Df F value  Pr(>F)
group   1  3.2002 0.07371 .
      3703

---
signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

# MANOVA

```
> m1 <- manova(cbind(log_pH, log_quality) ~ color, data = subset_data1)
> summary(m1)
```

	Df	Pillai	approx F	num Df	den Df	Pr(>F)
color	1	0.01269	23.791	2	3702	5.412e-11 ***
Residuals	3703					

---

signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

```
> summary.aov(m1)
```

Response log\_pH :

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
color	1	0.0837	0.083700	46.401	1.12e-11 ***
Residuals	3703	6.6797	0.001804		

---

signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Response log\_quality :

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
color	1	0.009	0.0094775	0.5515	0.4577
Residuals	3703	63.633	0.0171842		

# PRINCIPAL COMPONENTS ANALYSIS

```
> cor(scaled_data)
```

	fixed_acidity	volatile_acidity	citric_acid	residual_sugar
fixed_acidity	1.000000000	-0.038485191	0.26043580	0.04671636
volatile_acidity	-0.038485191	1.000000000	-0.12265398	0.07314759
citric_acid	0.260435797	-0.122653976	1.000000000	-0.01897515
residual_sugar	0.046716357	0.073147589	-0.01897515	1.000000000
chlorides	0.106754560	0.012152361	0.02005761	0.25504336
free_sulfur_dioxide	-0.042415793	-0.039255844	0.05114958	0.35113924
total_sulfur_dioxide	0.050937391	0.120747029	0.05999404	0.41218759
density	0.219015982	0.008326949	0.03351907	0.83427178
pH	-0.360461196	-0.013102201	-0.09090630	-0.18255852
sulphates	0.002514512	0.035174451	0.06378779	-0.04183721
alcohol	-0.069948466	0.107407499	0.01991982	-0.49110790
quality	-0.053708662	-0.123667412	0.03199637	-0.11020532

```
> cov(scaled_data)
```

	fixed_acidity	volatile_acidity	citric_acid	residual_sugar
fixed_acidity	1.000000000	-0.038485191	0.26043580	0.04671636
volatile_acidity	-0.038485191	1.000000000	-0.12265398	0.07314759
citric_acid	0.260435797	-0.122653976	1.000000000	-0.01897515
residual_sugar	0.046716357	0.073147589	-0.01897515	1.000000000
chlorides	0.106754560	0.012152361	0.02005761	0.25504336
free_sulfur_dioxide	-0.042415793	-0.039255844	0.05114958	0.35113924
total_sulfur_dioxide	0.050937391	0.120747029	0.05999404	0.41218759
density	0.219015982	0.008326949	0.03351907	0.83427178
pH	-0.360461196	-0.013102201	-0.09090630	-0.18255852
sulphates	0.002514512	0.035174451	0.06378779	-0.04183721
alcohol	-0.069948466	0.107407499	0.01991982	-0.49110790
quality	-0.053708662	-0.123667412	0.03199637	-0.11020532

# PRINCIPAL COMPONENTS ANALYSIS

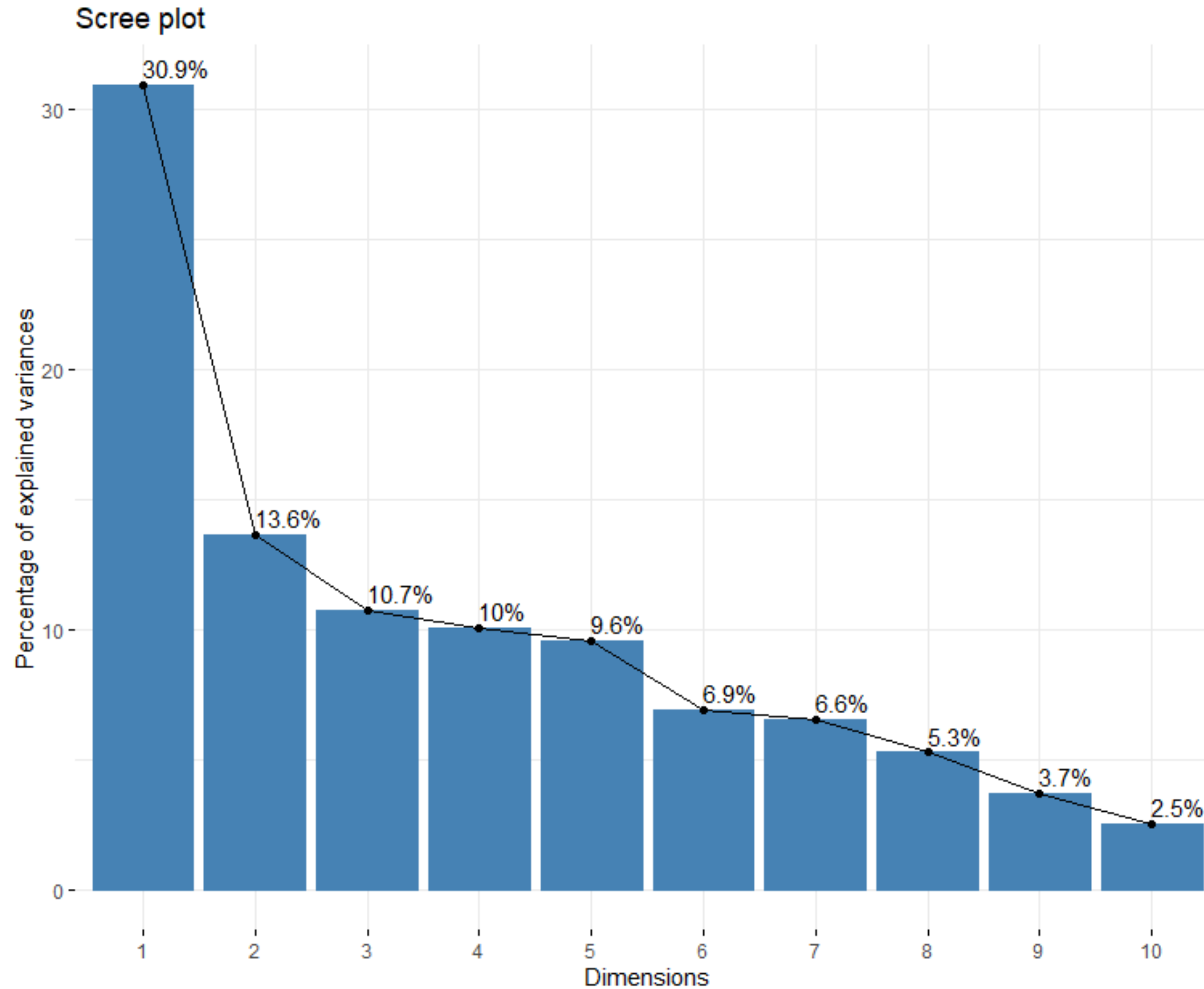
```
> summary(pca1)
Importance of components:

               PC1      PC2      PC3      PC4      PC5      PC6      PC7      PC8
standard deviation  1.8434  1.2240  1.0865  1.0514  1.02498  0.87300  0.84942  0.76205
Proportion of Variance 0.3089 0.1362 0.1073 0.1005 0.09551 0.06928 0.06559 0.05279
Cumulative Proportion 0.3089 0.4451 0.5524 0.6529 0.74842 0.81770 0.88329 0.93608

               PC9      PC10      PC11
standard deviation  0.63802  0.52881  0.12790
Proportion of Variance 0.03701 0.02542 0.00149
Cumulative Proportion 0.97309 0.99851 1.00000
```



# PRINCIPAL COMPONENTS ANALYSIS



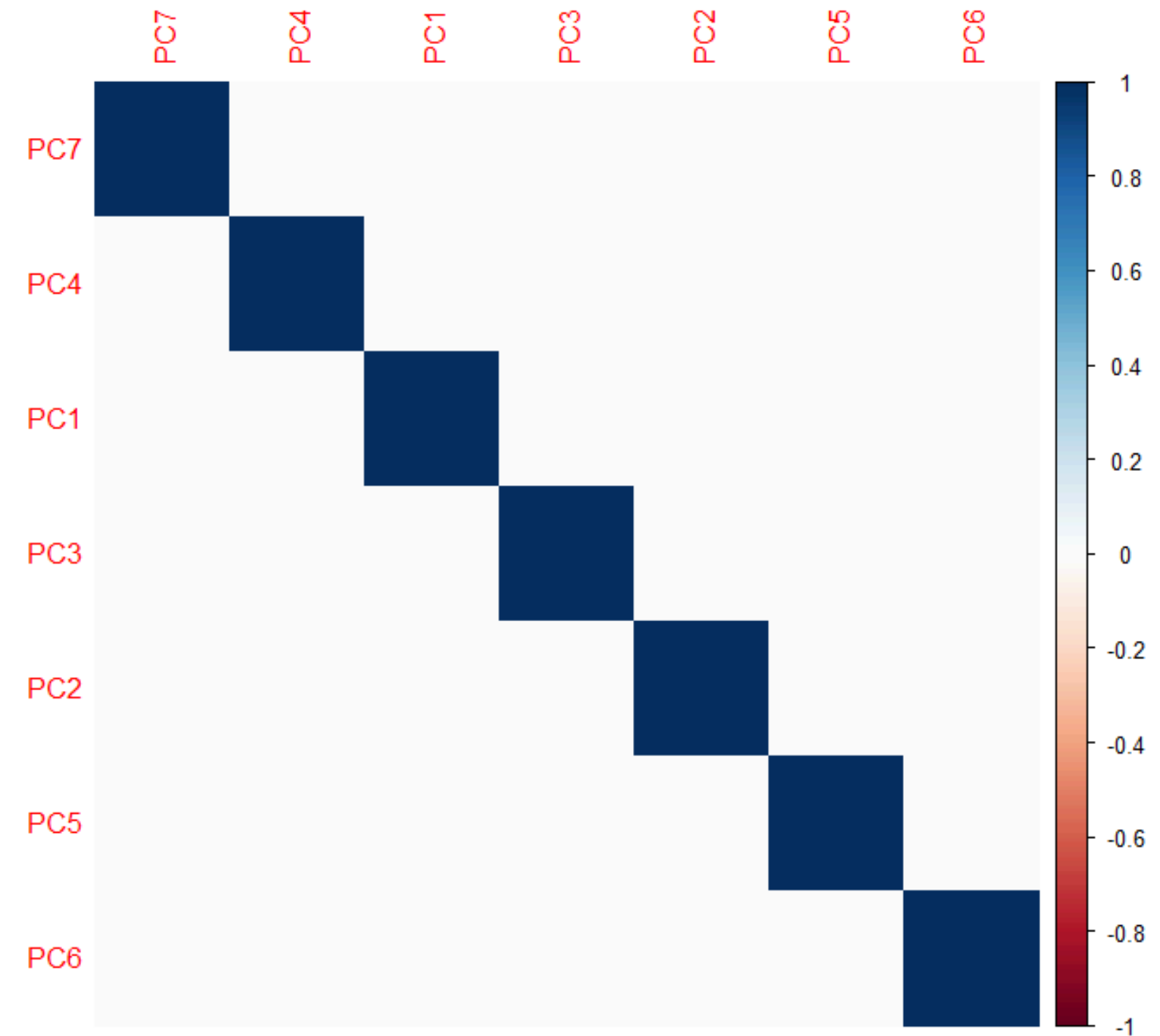
# PRINCIPAL COMPONENTS ANALYSIS

```
> head(pca)
```

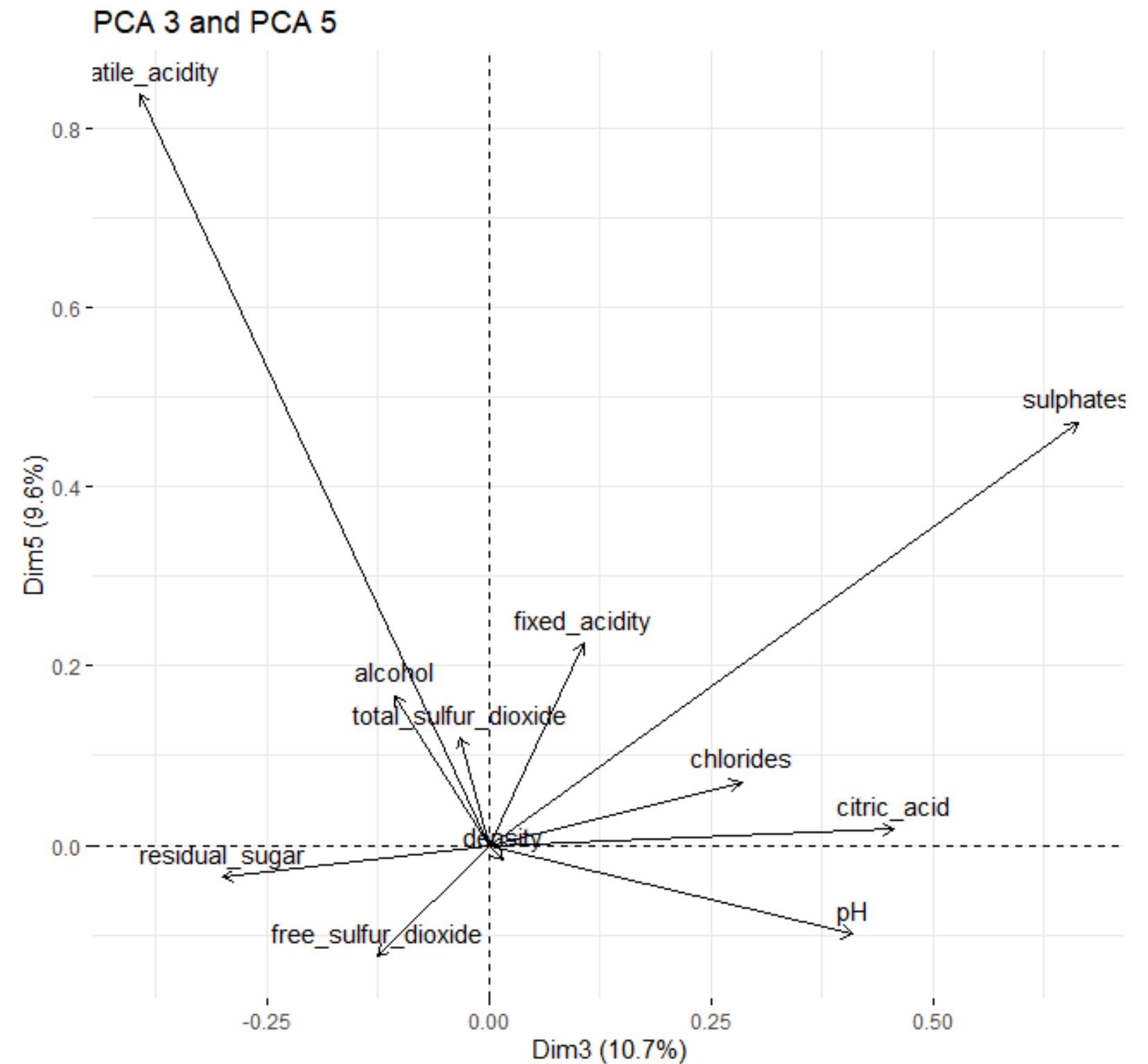
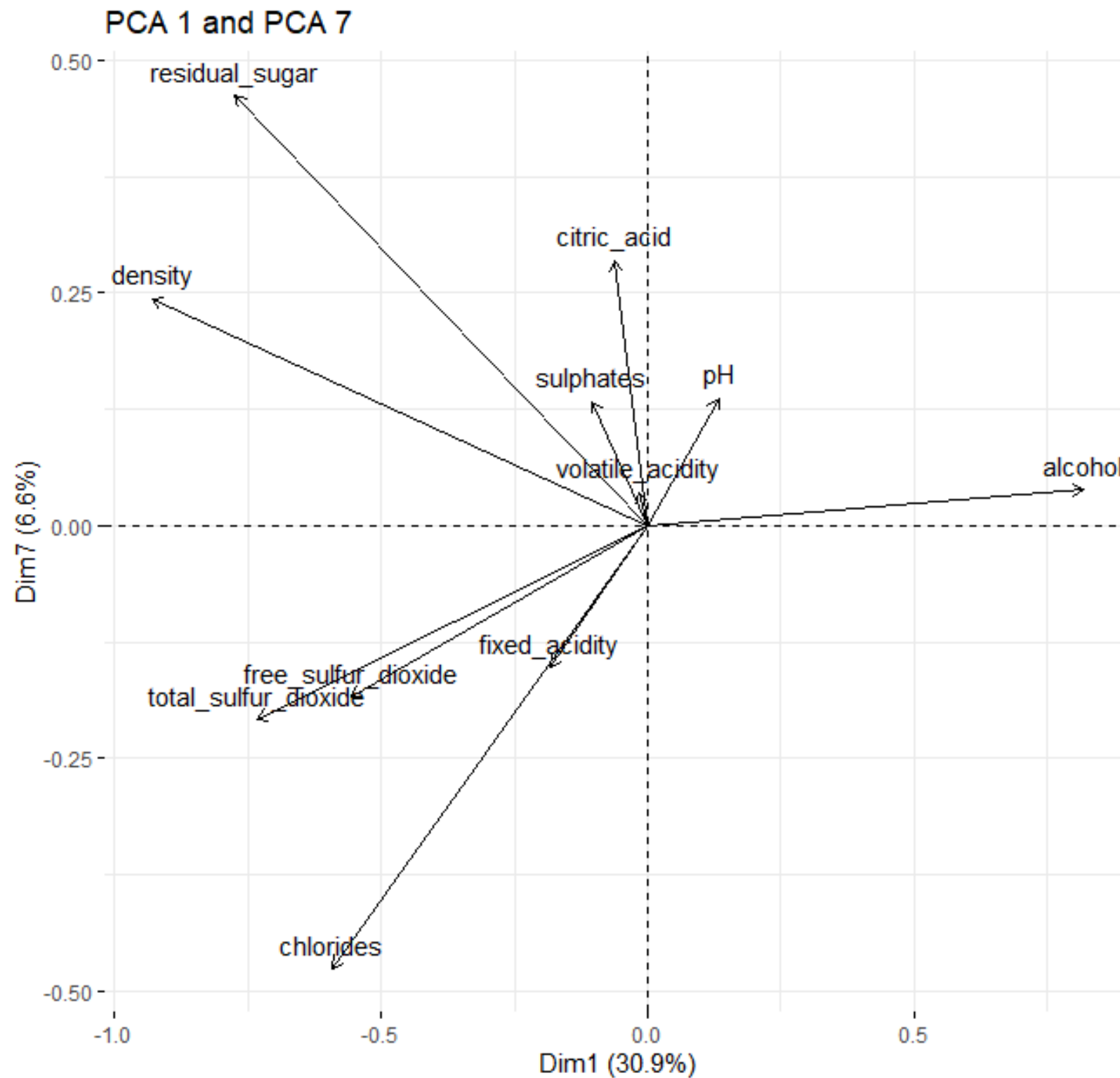
	PC1	PC2	PC3
38	-0.09207777	0.5036059	2.277253
155	-1.39337917	-0.9525258	2.736982
156	-1.38954992	-0.9271605	2.707217
157	-1.39337917	-0.9525258	2.736982
158	-1.38954992	-0.9271605	2.707217
310	0.35424132	0.2469293	2.315242

	PC4	PC5	PC6
38	-3.356240	2.7571187	-0.4209366
155	-1.004639	3.1083479	1.4162984
156	-1.097483	3.0720185	1.5147305
157	-1.004639	3.1083479	1.4162984
158	-1.097483	3.0720185	1.5147305
310	-2.991603	0.4658695	1.5681166

	PC7
38	-0.52178595
155	-0.03462939
156	-0.08623480
157	-0.03462939
158	-0.08623480
310	0.12036952



# PRINCIPAL COMPONENTS ANALYSIS



# PRINCIPAL COMPONENTS REGRESSION

```
> summary(lmodel)

call:
lm(formula = quality ~ ., data = ols.data)

Residuals:
    Min       1Q   Median       3Q      Max
-3.11588 -0.66977  0.02529  0.62446  2.33707

Coefficients:
              Estimate Std. Error t value Pr(>|t|)
(Intercept) -5.096e-16  1.505e-02   0.000    1.000
PC1          1.600e-01  8.167e-03  19.587 < 2e-16 ***
PC2          4.218e-03  1.230e-02   0.343   0.732
PC3          9.750e-03  1.386e-02   0.704   0.482
PC4          1.993e-01  1.432e-02  13.916 < 2e-16 ***
PC5         -6.405e-02  1.469e-02  -4.361 1.33e-05 ***
PC6         -1.295e-01  1.725e-02  -7.508 7.46e-14 ***
PC7          1.393e-01  1.772e-02   7.861 4.94e-15 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.9162 on 3697 degrees of freedom
Multiple R-squared:  0.1621,    Adjusted R-squared:  0.1605
F-statistic: 102.2 on 7 and 3697 DF,  p-value: < 2.2e-16
```

# FACTOR ANALYSIS AND ROTATION METHODS

There are a lot of rotation methods that can be used for factor analysis but we prefer Varimax (most preferred) and Quartimax since the estimated weights for the factor scores are unmatched for the other methods.



# ANALYSIS METHODS

For our dataset, we've been researching for the best analysis method related to factor analysis such as principal axis, maximum likelihood, generalized weighted least squares methods.

	ML3	ML2	ML1
SS loadings	3.44	2.28	1.31
Proportion Var	0.26	0.18	0.10
Cumulative Var	0.26	0.44	0.54
Proportion Explained	0.49	0.32	0.19

**BIC = 12703.87** Model fits the data.

## Maximum Likelihood Method

	[,1]	[,2]	[,3]
SS loadings	3.29	2.22	1.42
Proportion Var	0.25	0.17	0.11
Cumulative Var	0.25	0.42	0.53
Proportion Explained	0.47	0.32	0.20

**BIC = 14696.07** Model fits the data.

## Ordinary Least Squares Method

# VARIABLES AND PATTERNS

	ML3	ML2	ML1	h2	u2	com
fixed_acidity	-0.61	0.15	0.78	1.00	0.0050	2.0
volatile_acidity	-0.63	0.14	-0.23	0.47	0.5298	1.4
citric_acid	0.14	0.05	0.52	0.29	0.7078	1.2
residual_sugar	0.46	0.72	0.08	0.74	0.2582	1.7
chlorides	-0.49	0.24	-0.04	0.30	0.6978	1.5
free_sulfur_dioxide	0.57	0.20	0.04	0.36	0.6359	1.3
total_sulfur_dioxide	0.76	0.25	0.12	0.66	0.3427	1.3
density	-0.30	0.93	0.18	1.00	0.0049	1.3
pH	-0.28	0.03	-0.54	0.37	0.6277	1.5
sulphates	-0.48	0.13	-0.01	0.25	0.7511	1.1
alcohol	-0.05	-0.74	-0.02	0.56	0.4446	1.0
quality	0.10	-0.30	0.04	0.10	0.8969	1.3
color*	0.94	-0.14	0.13	0.93	0.0741	1.1

ML1 estimator is related to Fixed Acidity, Citric Acid, pH. So, we can correspond this factor loading as a Tendency to Acidity.

ML2 is related to Residual Sugar, Density, Alcohol, Quality. So, we can correspond this factor loading as a Sweetness Profile.

ML3 is related to Volatile Acidity, Chlorides, Free Sulfur Dioxide, Total Sulfur Dioxide, Sulphates, Color. So, we can correspond this factor loading as a Wine Stability.

# BEST AND WORST FEATURE THAT IS EXPLAINED BY FACTOR ANALYSIS

	ML3	ML2	ML1	h2	u2	com
fixed_acidity	-0.61	0.15	0.78	1.00	0.0050	2.0
volatile_acidity	-0.63	0.14	-0.23	0.47	0.5298	1.4
citric_acid	0.14	0.05	0.52	0.29	0.7078	1.2
residual_sugar	0.46	0.72	0.08	0.74	0.2582	1.7
chlorides	-0.49	0.24	-0.04	0.30	0.6978	1.5
free_sulfur_dioxide	0.57	0.20	0.04	0.36	0.6359	1.3
total_sulfur_dioxide	0.76	0.25	0.12	0.66	0.3427	1.3
density	-0.30	0.93	0.18	1.00	0.0049	1.3
pH	-0.28	0.03	-0.54	0.37	0.6277	1.5
sulphates	-0.48	0.13	-0.01	0.25	0.7511	1.1
alcohol	-0.05	-0.74	-0.02	0.56	0.4446	1.0
quality	0.10	-0.30	0.04	0.10	0.8969	1.3
color*	0.94	-0.14	0.13	0.93	0.0741	1.1

Since  $h^2$  value of Color of Wine is equal to 0.93, it says 93% of the total variance in the variable is explained by the factors.

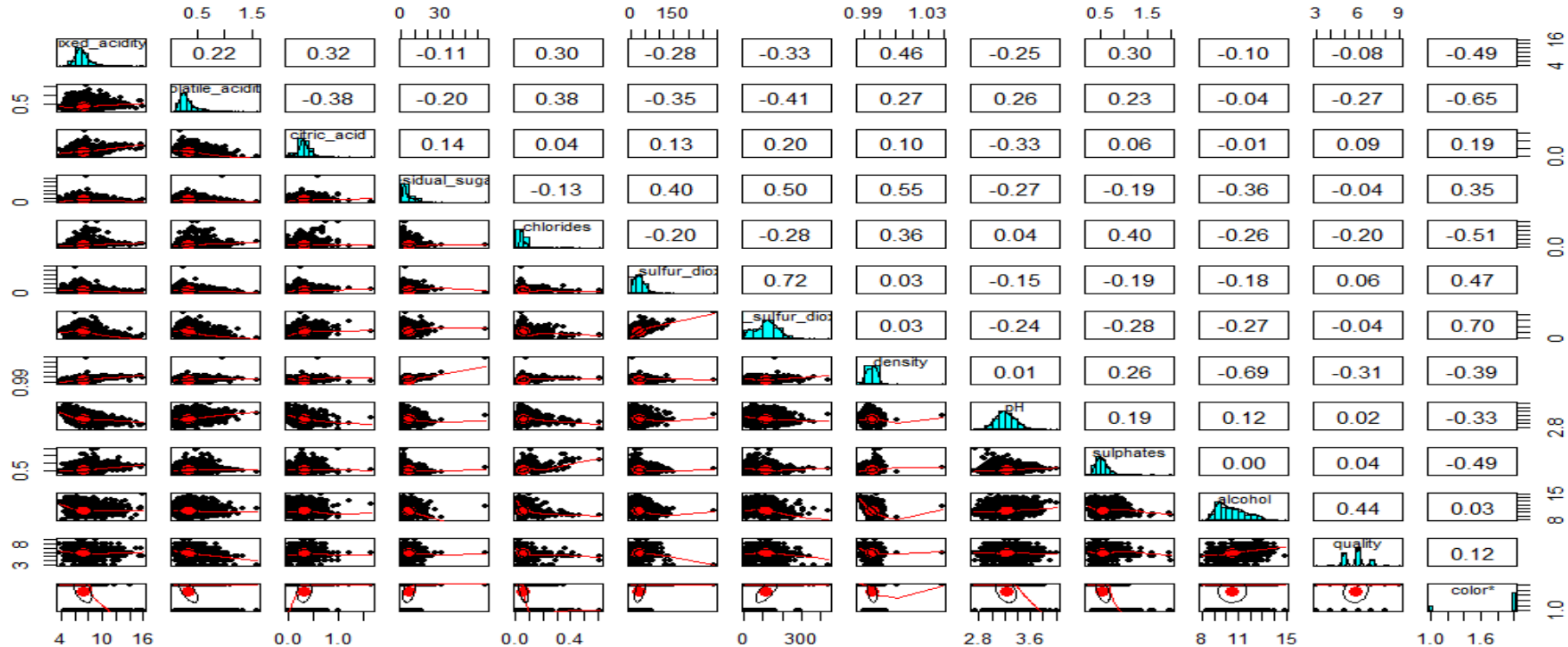
As for the worst, Quality variable has a 0.89 value of  $u^2$ , it says 89% of the total variance in the variable is failed to explained by the factors.

# DISCRIMINATION AND CLASSIFICATION

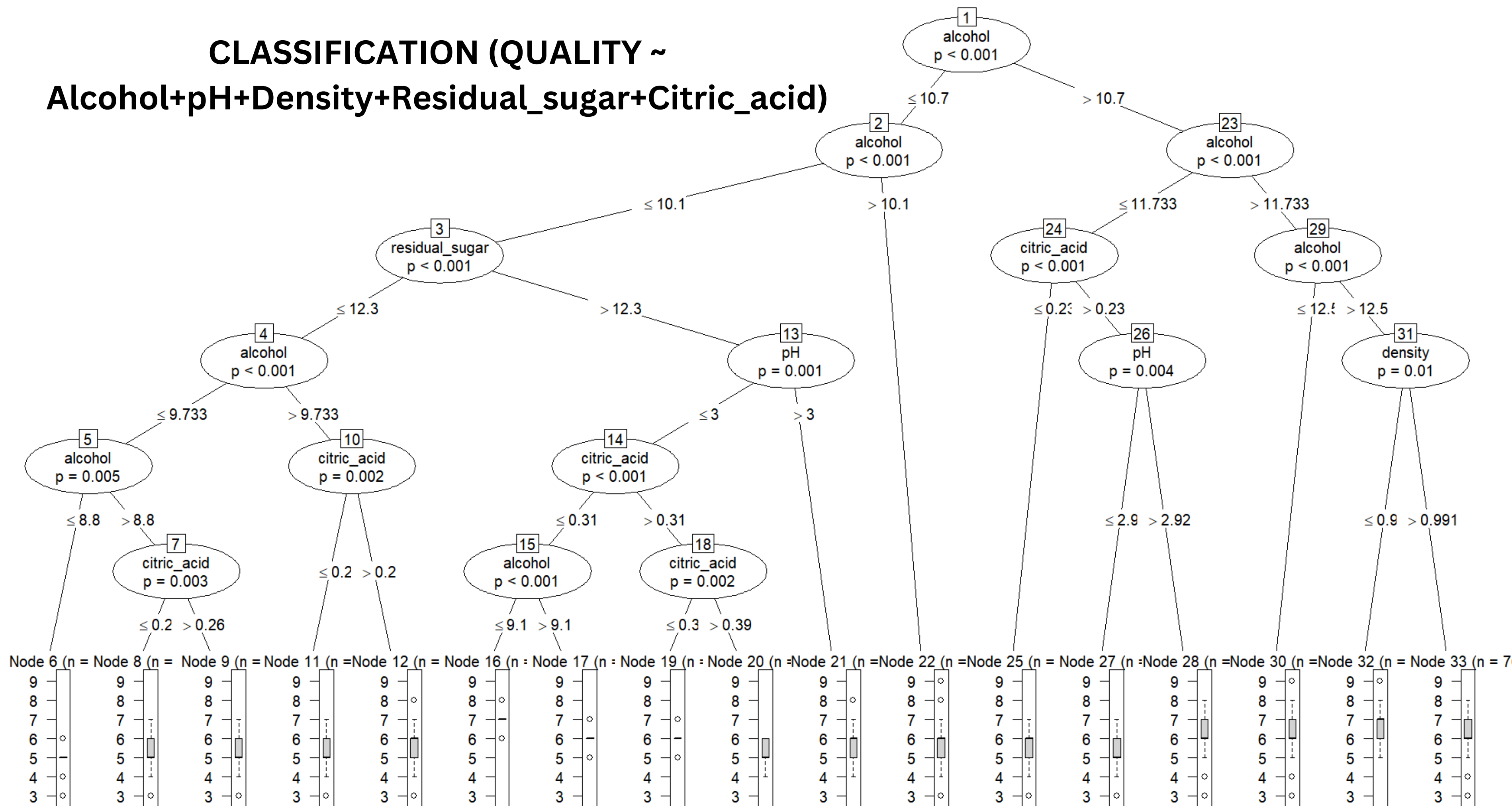
We aim to visualize these variables and their relationships through a graphical representation, choosing a pair plot for clarity.



# VARIABLES AND THEIR RELATIONS



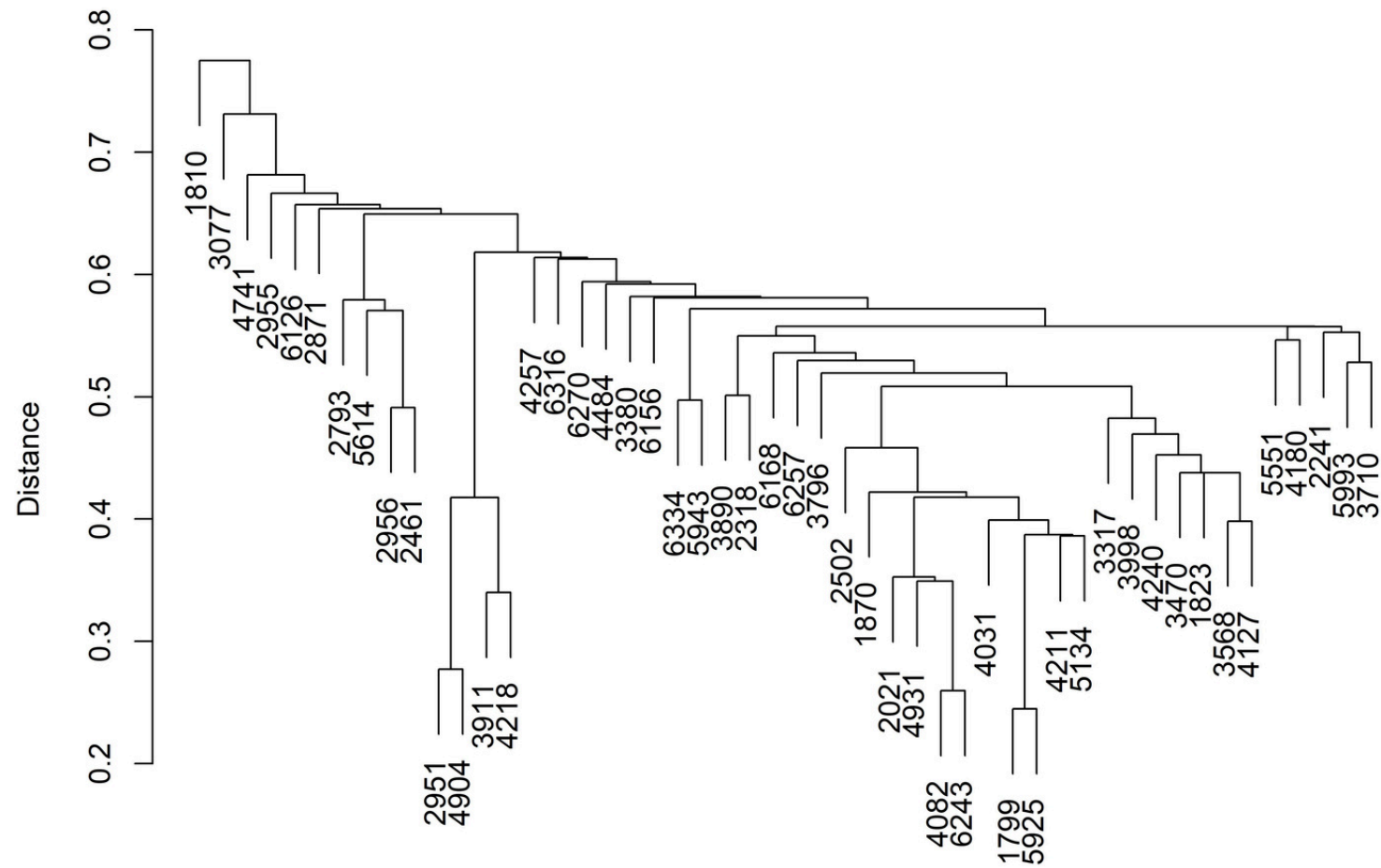
# CLASSIFICATION (QUALITY ~ Alcohol+pH+Density+Residual\_sugar+Citric\_acid)



# CLUSTERING

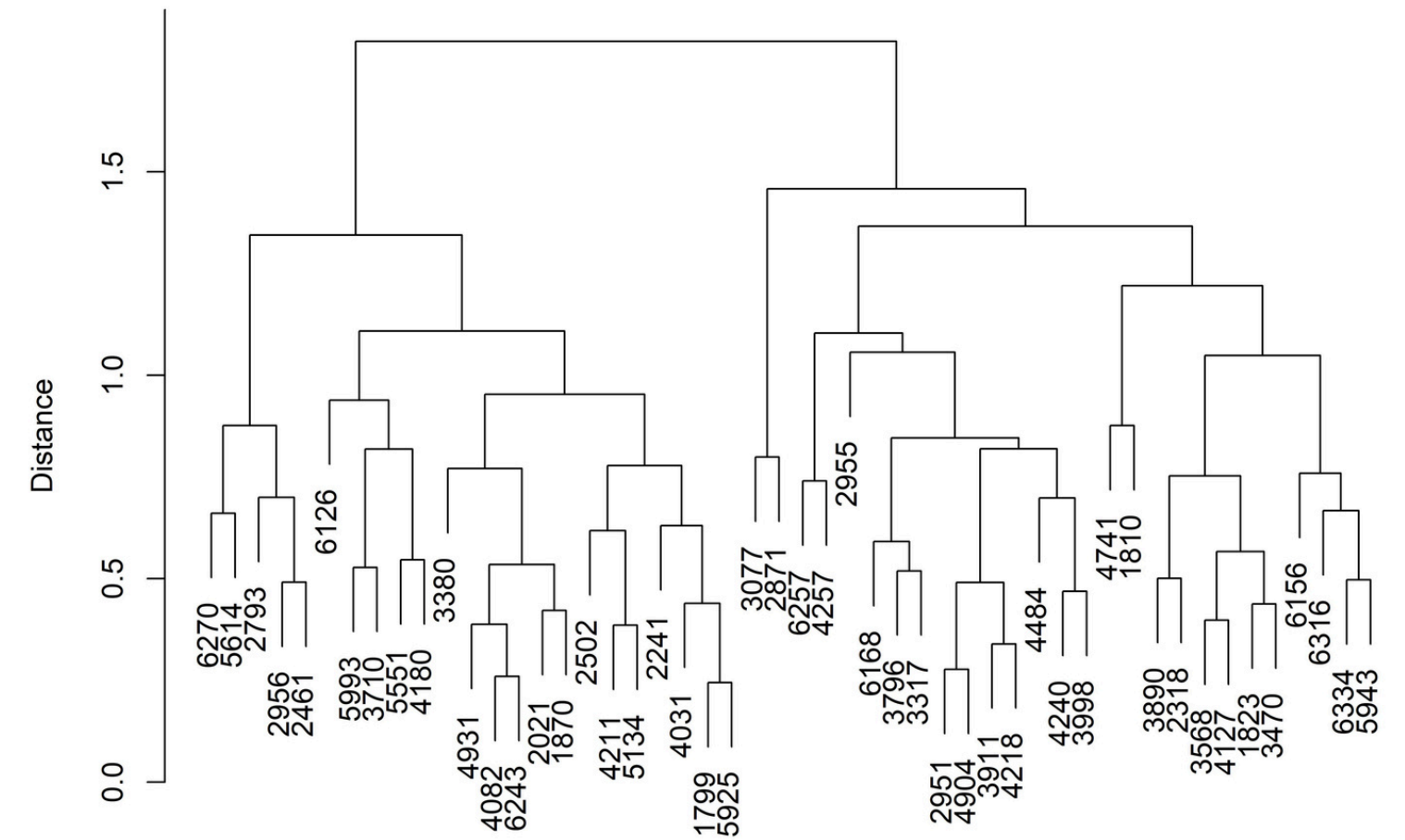
## Hierarchical

Single Linkage Dendrogram



Sample Index  
Random sample of 50 observations

Complete Linkage Dendrogram

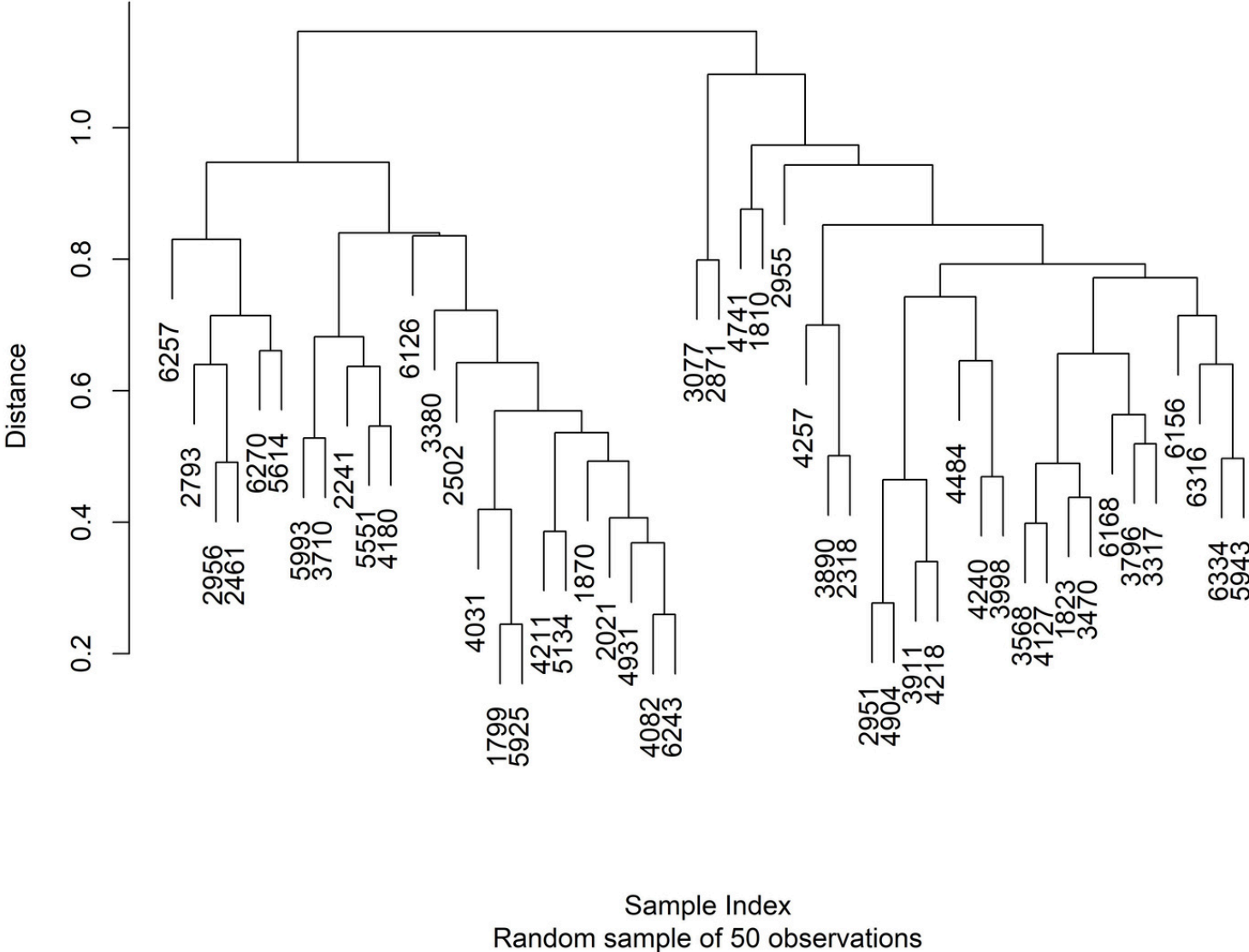


Sample Index  
Random sample of 50 observations

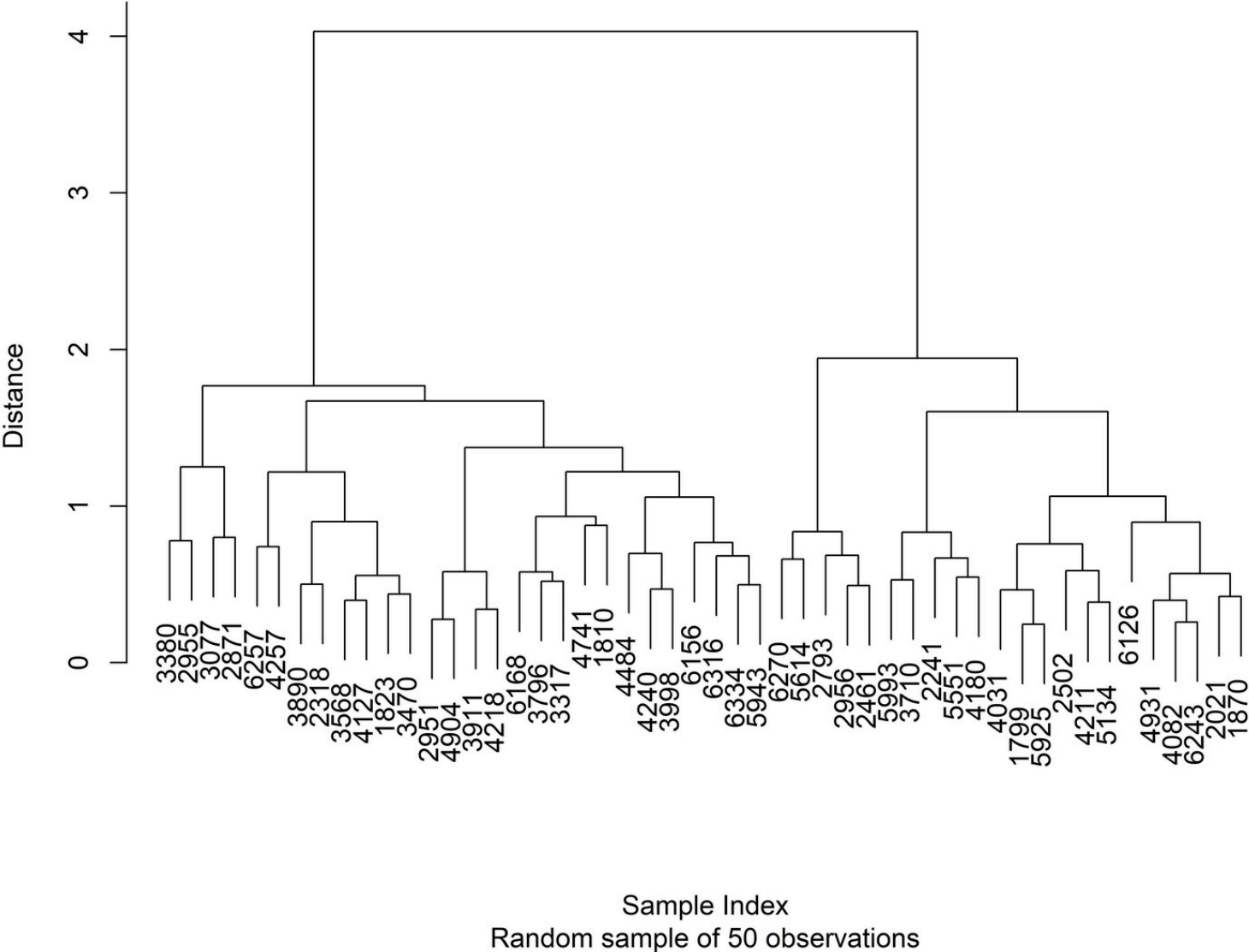
# CLUSTERING

## Hierarchical

Average Linkage Dendrogram



Ward Method Dendrogram

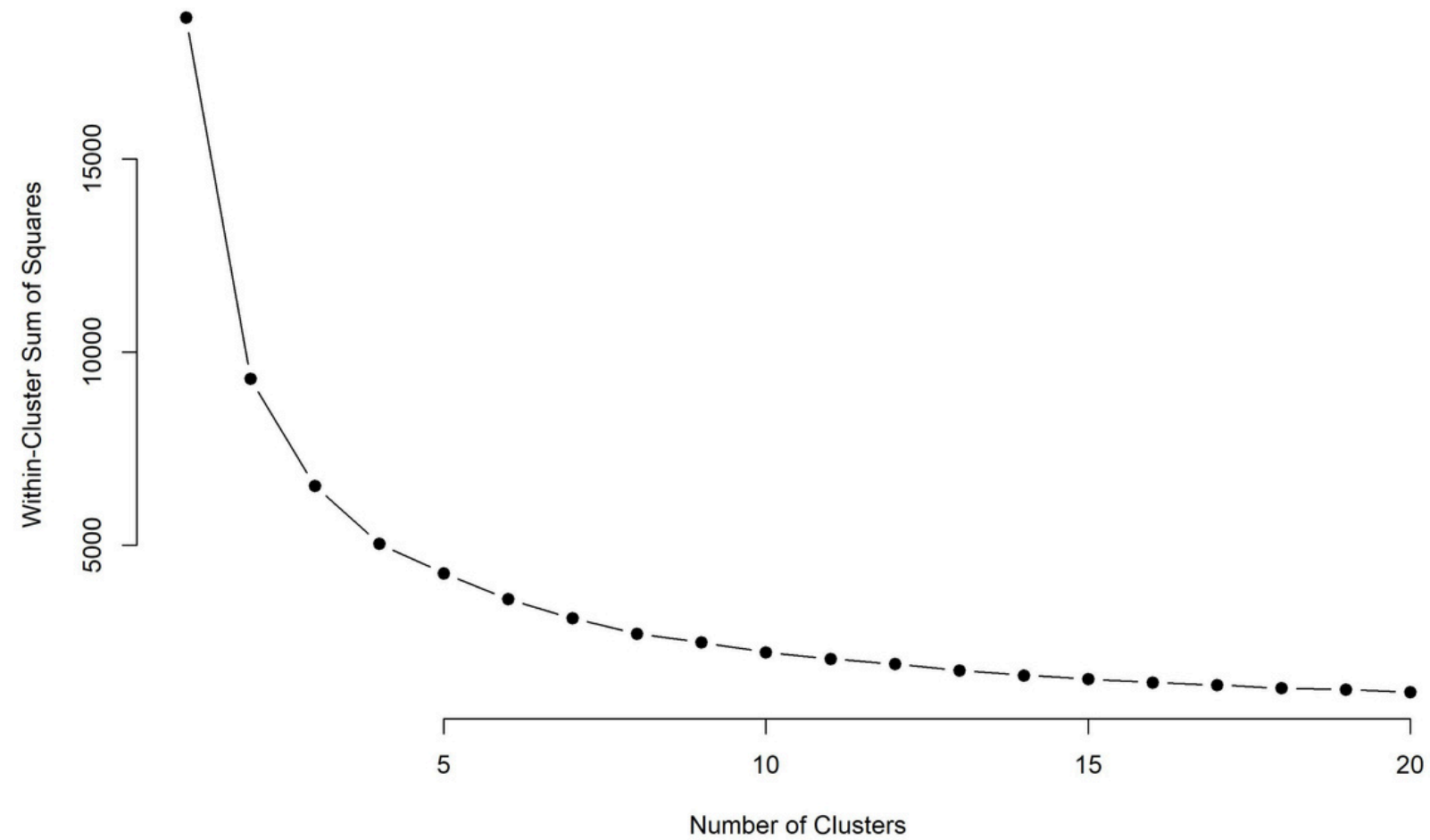




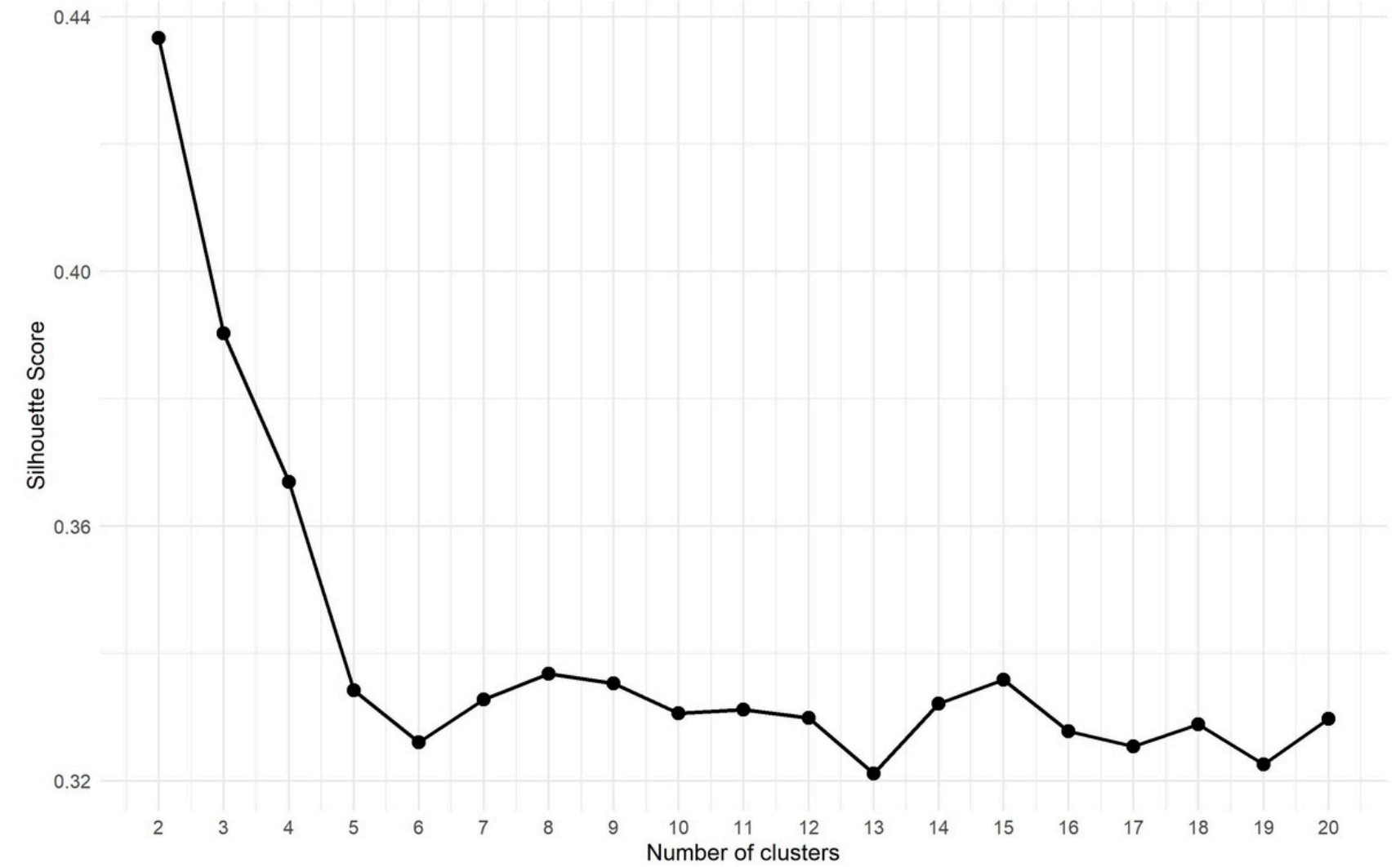
# CLUSTERING

## K-Means

Elbow Plot for Optimal k

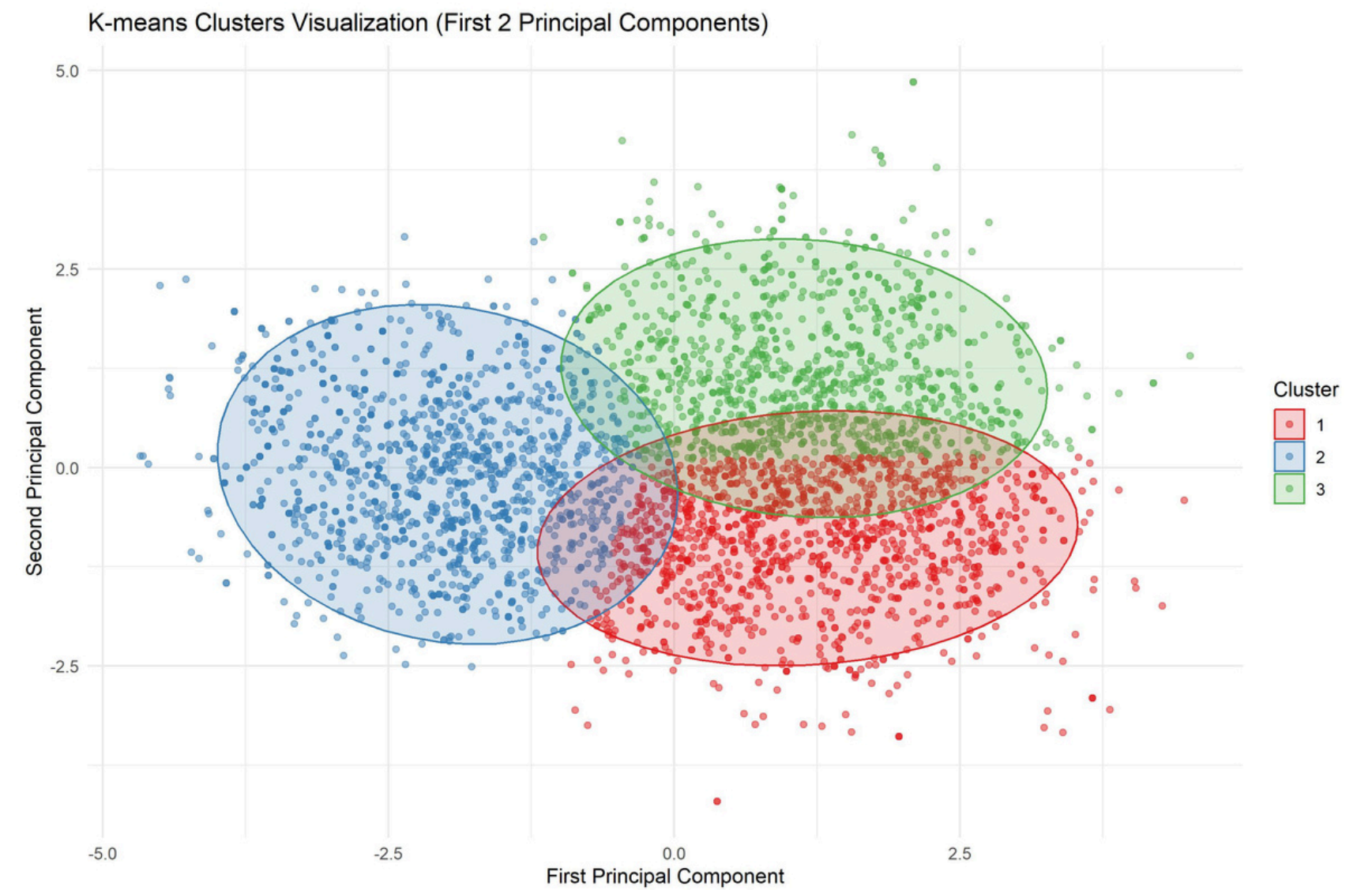
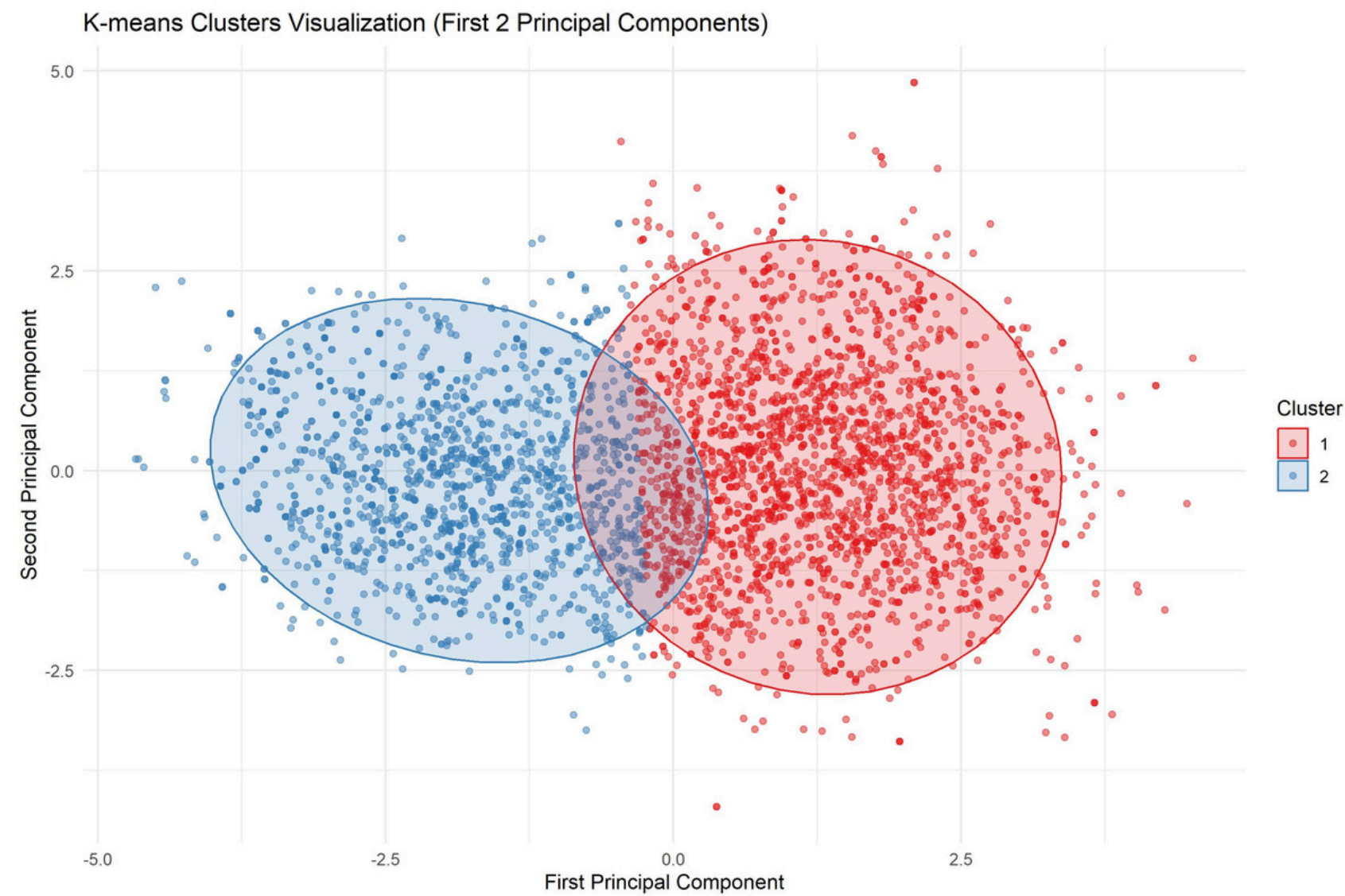


Silhouette Score for different cluster amount



# CLUSTERING

## K-Means



# CCA

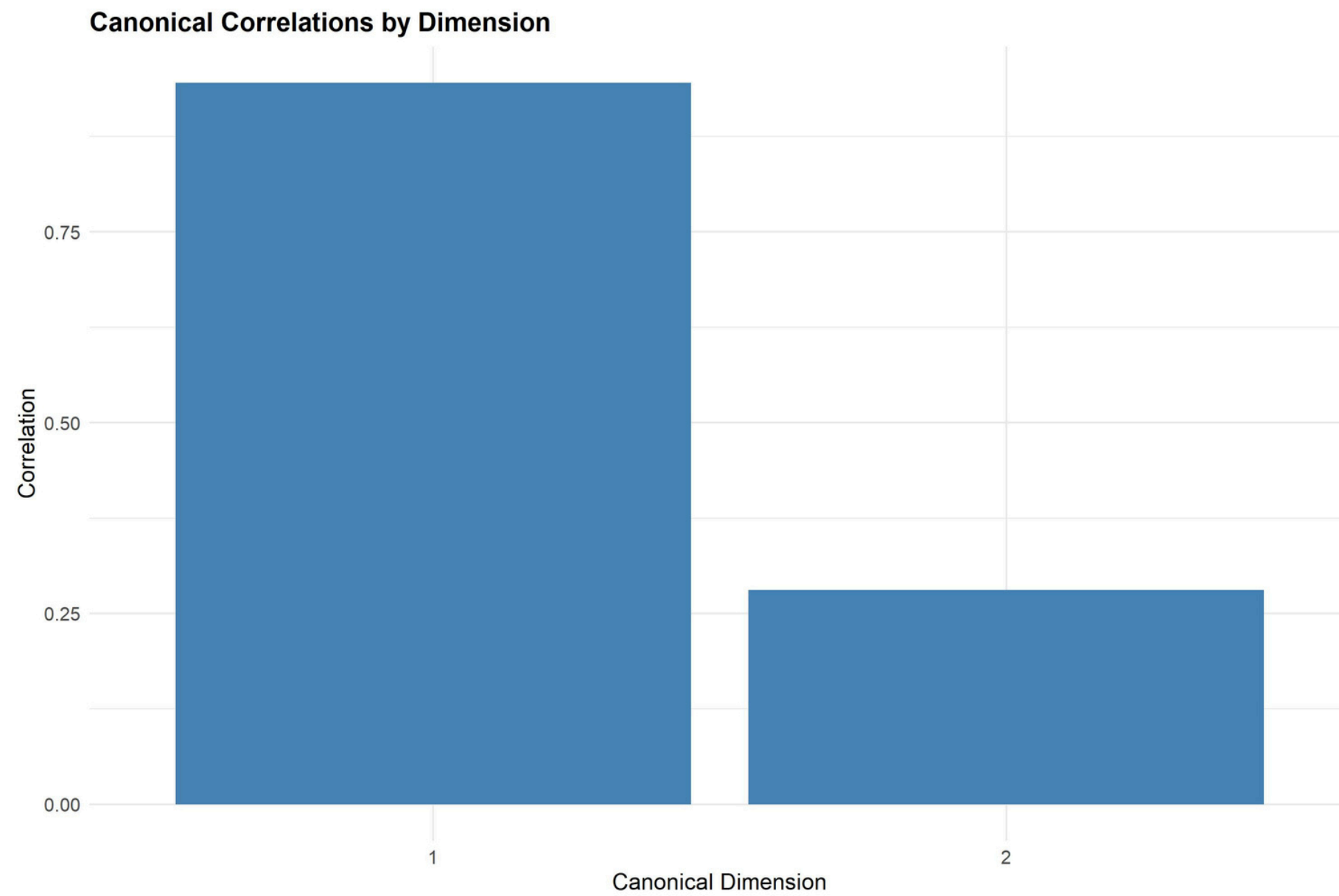
## Features of the Wine

- fixed\_acidity
- volatile\_acidity
- citric\_acid
- residual\_sugar
- chlorides
- free\_sulfur\_dioxide
- total\_sulfur\_dioxide
- density
- pH
- sulphates

## Responses

- alcohol
- quality

# CCA



```
> p.asym(cca_result$cor, n, p, q, tstat = "wilks")
wilks' Lambda, using F-approximation (Rao's F):
      stat   approx df1  df2 p.value
1 to 2:  0.09821539 843.9263  20 7704      0
2 to 2:  0.92121920  36.6112   9 3853      0
```



# CCA

\$xcoef										
	[,1]	[,2]	[,3]	[,4]	[,5]	[,6]	[,7]	[,8]	[,9]	[,10]
fixed_acidity	-8.255103e-03	-1.613168e-03	-4.991130e-03	-7.220327e-03	7.415668e-03	1.406704e-03	-0.0040074578	0.0156053883	-1.592585e-02	-2.698613e-03
volatile_acidity	-1.771809e-02	1.333880e-01	3.150766e-02	9.546204e-02	3.886881e-02	7.734632e-02	-0.0539888400	-0.0374460624	-2.660643e-02	-5.373661e-02
citric_acid	-7.183468e-03	7.115340e-04	2.045824e-01	7.785046e-03	-4.112579e-03	-1.603061e-02	-0.0086390207	-0.0092121358	4.127932e-03	1.596504e-02
residual_sugar	-3.390414e-03	-3.025775e-03	-2.310143e-04	4.001277e-03	-9.464951e-04	-1.631095e-03	0.0001805842	0.0036279364	-1.831122e-03	-4.856156e-04
chlorides	-7.998196e-02	2.626506e-01	-4.098336e-04	1.124304e-01	-1.628610e+00	-2.726232e-02	-0.3334550568	0.9679406469	-2.292681e-01	-4.497120e-02
free_sulfur_dioxide	1.835910e-05	-4.263284e-04	-5.428822e-06	-1.325776e-04	-4.203248e-05	1.087511e-03	-0.0007888072	-0.0001652701	6.645632e-05	1.115670e-04
total_sulfur_dioxide	1.648582e-05	6.881169e-05	1.939561e-06	1.268899e-05	1.477944e-05	1.064496e-05	0.0005112672	0.0001618843	-4.701803e-05	-1.737873e-05
density	1.016806e+01	3.151752e+00	6.034626e-01	-3.634155e+00	4.546440e+00	-2.452156e-02	-2.5466400443	-7.5487995238	2.431802e+00	9.190540e-01
pH	-3.726593e-02	-2.993412e-02	-2.489056e-03	7.957038e-03	-1.880053e-02	-2.993513e-03	0.0108116414	-0.0211720623	-1.247475e-01	1.953263e-02
sulphates	-1.792747e-02	-5.960948e-02	-1.879533e-03	-9.359193e-03	-1.430191e-02	-9.023040e-03	0.0088364935	-0.0095647395	1.195664e-02	-1.542182e-01
\$ycoef										
	[,1]	[,2]								
alcohol	-0.0134215501	0.006316683								
quality	-0.0004153857	-0.023409717								
\$xcenter										
fixed_acidity	volatile_acidity	citric_acid	residual_sugar	chlorides	free_sulfur_dioxide	total_sulfur_dioxide	density			
6.83976449	0.26691382	0.32189182	6.16900880	0.04281004	34.10623706	135.25582298	0.99387597			
pH	sulphates									
3.19189700	0.48480072									
\$ycenter										
alcohol	quality									
10.578972	5.881988									

# CCA

```
$corr.X.xscores
      [,1]      [,2]      [,3]      [,4]      [,5]      [,6]      [,7]      [,8]      [,9]     [,10]
fixed_acidity    0.07338023  0.14498328  0.060791109 -0.43992468  0.45748974 -0.034959352 -0.253310835  0.61973202 -0.305596996 -0.155638052
volatile_acidity -0.10414682  0.66749145  0.019175134  0.53540429  0.14892270  0.285641730 -0.109399488 -0.08627773 -0.170039463 -0.323246789
citric_acid      -0.02241931 -0.08173228  0.967321500 -0.16663961  0.07110988 -0.048441487 -0.035472343  0.13803126 -0.037283475  0.015277399
residual_sugar   0.50823649 -0.37915069  0.003624151  0.65781724  0.18593128 -0.084762099 -0.096550036  0.32028349 -0.107438630  0.001495136
chlorides        0.53549616  0.19594209  0.027994471  0.01199054 -0.67445539  0.005375436 -0.164148490  0.36860724 -0.201577699 -0.124717822
free_sulfur_dioxide 0.27614093 -0.46041238  0.069624392  0.11467862 -0.01868219  0.818080986 -0.004239598  0.13036353  0.007788292  0.084196635
total_sulfur_dioxide 0.48508772 -0.09544151  0.095567342  0.17795312  0.04116799  0.479396905  0.607918114  0.30218374 -0.137943475 -0.057933992
density          0.85507690 -0.17850153  0.035005624  0.28436558  0.13091466 -0.063645111 -0.104665759  0.19976490 -0.271363817 -0.095931846
pH               -0.06750050 -0.12280381 -0.019533275 -0.01150834 -0.30201761 -0.003752170  0.152320260 -0.59225290 -0.713503439  0.074619631
sulphates        0.06280753 -0.25720792  0.072728135 -0.15505425 -0.13849144 -0.011551518  0.059669555 -0.17571764 -0.056060641 -0.918517742

$corr.Y.xscores
      [,1]      [,2]      [,3]      [,4]      [,5]      [,6]      [,7]      [,8]      [,9]     [,10]
alcohol -0.9450419  0.004979632 -5.165005e-16  1.957909e-15 -1.843174e-15 -6.103785e-16  1.283771e-15  3.504828e-15 -2.466439e-15  3.563178e-16
quality -0.4024938 -0.253958967 -2.471846e-16  8.254339e-16 -4.955094e-16 -1.017174e-16  1.963456e-16  8.853033e-16 -9.753260e-16  3.117138e-16

$corr.X.yscores
      [,1]      [,2]
fixed_acidity    0.06935831  0.04069379
volatile_acidity -0.09843860  0.18735095
citric_acid      -0.02119052 -0.02294055
residual_sugar   0.48038040 -0.10641970
chlorides        0.50614599  0.05499687
free_sulfur_dioxide 0.26100584 -0.12922817
total_sulfur_dioxide 0.45850040 -0.02678845
density          0.80821072 -0.05010166
pH               -0.06380085 -0.03446847
sulphates        0.05936509 -0.07219291

$corr.Y.yscores
      [,1]      [,2]
alcohol -0.9998426  0.01774137
quality -0.4258334 -0.90480158
```

# CONCLUSION



**THANK YOU**



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- Lecture Notes