

Red Wine Association: Onima Shah

2025-05-01

Setting up R file

```
#Downloading necessary packages
```

```
#install.packages("tidyverse")
```

```
#install.packages("corrplot")
```

```
#install.packages("regclass")
```

```
#Loading packages
```

```
library(tidyverse)
```

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

```
tidyverse 2.0.0 —
```

```
## ✓ dplyr      1.1.4      ✓ readr      2.1.4
```

```
## ✓ forcats   1.0.0      ✓ stringr   1.5.1
```

```
## ✓ ggplot2   3.5.1      ✓ tibble    3.2.1
```

```
## ✓ lubridate 1.9.3      ✓ tidyr     1.3.0
```

```
## ✓ purrr     1.0.2
```

```
## — Conflicts —————
```

```
tidyverse_conflicts() —
```

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

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

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

```
library(corrplot)
```

```
## corrplot 0.95 loaded
```

```
library(regclass)
```

```
## Loading required package: bestglm
```

```
## Loading required package: leaps
```

```
## Loading required package: VGAM
```

```
## Loading required package: stats4
```

```
## Loading required package: splines
```

```
## Loading required package: rpart
```

```
## Loading required package: 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:dplyr':
```

```
##
```

```
##      combine
```

```
##
```

```
## The following object is masked from 'package:ggplot2':
##
##     margin
##
## Important regclass change from 1.3:
## All functions that had a . in the name now have an _
## all.correlations -> all_correlations, cor.demo -> cor_demo, etc.
```

Loading file and checking dataset

#Loading csv file

```
wine <- read.csv("winequality-red.csv")
```

#Checking dataset

```
head(wine)
```

```
##   fixed.acidity volatile.acidity citric.acid residual.sugar chlorides
## 1           7.4           0.70         0.00           1.9      0.076
## 2           7.8           0.88         0.00           2.6      0.098
## 3           7.8           0.76         0.04           2.3      0.092
## 4          11.2           0.28         0.56           1.9      0.075
## 5           7.4           0.70         0.00           1.9      0.076
## 6           7.4           0.66         0.00           1.8      0.075
##   free.sulfur.dioxide total.sulfur.dioxide density    pH sulphates alcohol
## 1                   11                   34 0.9978 3.51      0.56      9.4
## 2                   25                   67 0.9968 3.20      0.68      9.8
## 3                   15                   54 0.9970 3.26      0.65      9.8
## 4                   17                   60 0.9980 3.16      0.58      9.8
## 5                   11                   34 0.9978 3.51      0.56      9.4
## 6                   13                   40 0.9978 3.51      0.56      9.4
##   quality
## 1        5
## 2        5
## 3        5
## 4        6
## 5        5
## 6        5
```

```
str(wine)
```

```
## 'data.frame':   1599 obs. of  12 variables:
## $ fixed.acidity      : num  7.4 7.8 7.8 11.2 7.4 7.4 7.9 7.3 7.8 7.5 ...
## $ volatile.acidity   : num  0.7 0.88 0.76 0.28 0.7 0.66 0.6 0.65 0.58
0.5 ...
## $ citric.acid        : num  0 0 0.04 0.56 0 0 0.06 0 0.02 0.36 ...
## $ residual.sugar     : num  1.9 2.6 2.3 1.9 1.9 1.8 1.6 1.2 2 6.1 ...
## $ chlorides          : num  0.076 0.098 0.092 0.075 0.076 0.075 0.069
0.065 0.073 0.071 ...
## $ free.sulfur.dioxide : num  11 25 15 17 11 13 15 15 9 17 ...
## $ total.sulfur.dioxide: num  34 67 54 60 34 40 59 21 18 102 ...
## $ density            : num  0.998 0.997 0.997 0.998 0.998 ...
```

```
## $ pH : num 3.51 3.2 3.26 3.16 3.51 3.51 3.3 3.39 3.36
3.35 ...
## $ sulphates : num 0.56 0.68 0.65 0.58 0.56 0.56 0.46 0.47 0.57
0.8 ...
## $ alcohol : num 9.4 9.8 9.8 9.8 9.4 9.4 9.4 10 9.5 10.5 ...
## $ quality : int 5 5 5 6 5 5 5 7 7 5 ...
```

summary(wine)

```
## fixed.acidity volatile.acidity citric.acid residual.sugar
## Min. : 4.60 Min. :0.1200 Min. :0.000 Min. : 0.900
## 1st Qu.: 7.10 1st Qu.:0.3900 1st Qu.:0.090 1st Qu.: 1.900
## Median : 7.90 Median :0.5200 Median :0.260 Median : 2.200
## Mean : 8.32 Mean :0.5278 Mean :0.271 Mean : 2.539
## 3rd Qu.: 9.20 3rd Qu.:0.6400 3rd Qu.:0.420 3rd Qu.: 2.600
## Max. :15.90 Max. :1.5800 Max. :1.000 Max. :15.500
## chlorides free.sulfur.dioxide total.sulfur.dioxide density
## Min. :0.01200 Min. : 1.00 Min. : 6.00 Min. :0.9901
## 1st Qu.:0.07000 1st Qu.: 7.00 1st Qu.: 22.00 1st Qu.:0.9956
## Median :0.07900 Median :14.00 Median : 38.00 Median :0.9968
## Mean :0.08747 Mean :15.87 Mean : 46.47 Mean :0.9967
## 3rd Qu.:0.09000 3rd Qu.:21.00 3rd Qu.: 62.00 3rd Qu.:0.9978
## Max. :0.61100 Max. :72.00 Max. :289.00 Max. :1.0037
## pH sulphates alcohol quality
## Min. :2.740 Min. :0.3300 Min. : 8.40 Min. :3.000
## 1st Qu.:3.210 1st Qu.:0.5500 1st Qu.: 9.50 1st Qu.:5.000
## Median :3.310 Median :0.6200 Median :10.20 Median :6.000
## Mean :3.311 Mean :0.6581 Mean :10.42 Mean :5.636
## 3rd Qu.:3.400 3rd Qu.:0.7300 3rd Qu.:11.10 3rd Qu.:6.000
## Max. :4.010 Max. :2.0000 Max. :14.90 Max. :8.000
```

names(wine)

```
## [1] "fixed.acidity" "volatile.acidity" "citric.acid"
## [4] "residual.sugar" "chlorides" "free.sulfur.dioxide"
## [7] "total.sulfur.dioxide" "density" "pH"
## [10] "sulphates" "alcohol" "quality"
```

#Checking for missing values

sum(is.na(wine))

```
## [1] 0
```

colSums(is.na(wine))

```
## fixed.acidity volatile.acidity citric.acid
## 0 0 0
## residual.sugar chlorides free.sulfur.dioxide
## 0 0 0
## total.sulfur.dioxide density pH
## 0 0 0
```

```
##          sulphates          alcohol          quality
##              0              0              0

#Checking for duplicates
sum(duplicated(wine))

## [1] 240

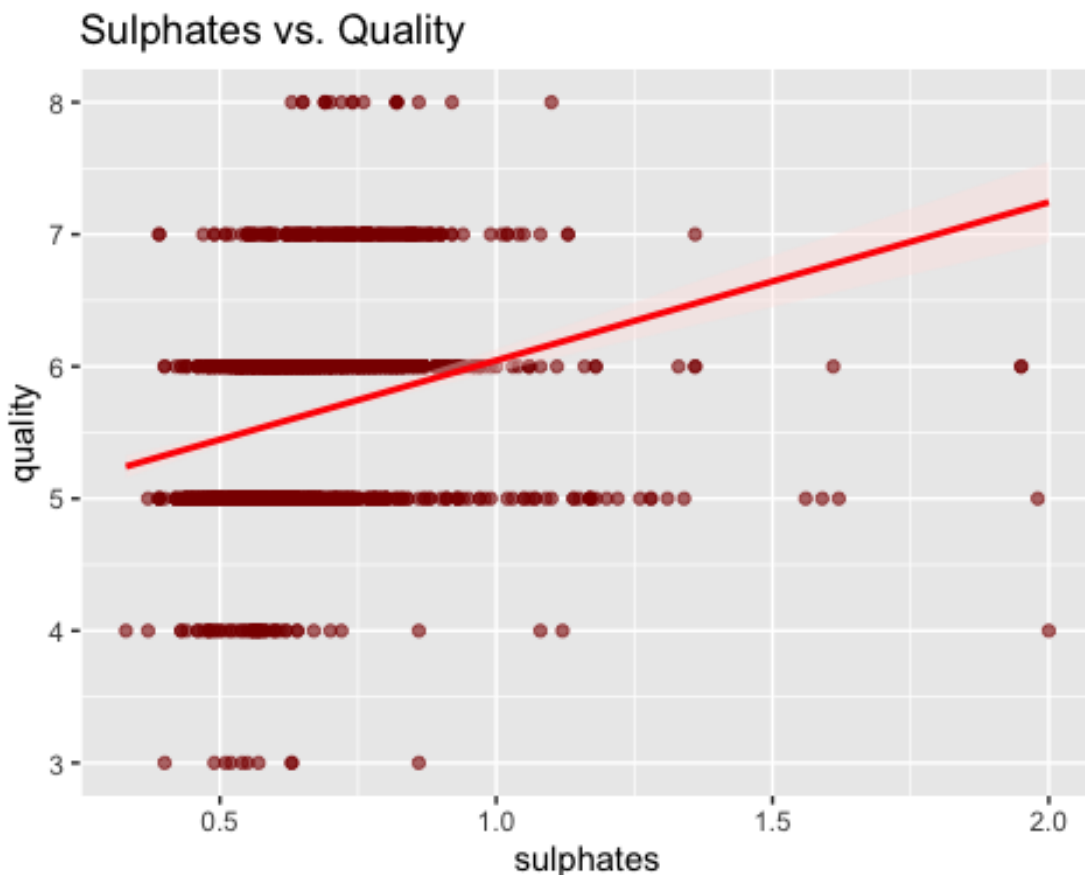
#Should we remove the duplicates??
```

Variables for association analysis: $y = \text{quality}$; $x = \text{sulphates, residual.sugar}$

Sulphates

```
#Graph
ggplot(wine, aes(x = sulphates, y = quality)) +
  geom_point(color = "darkred", alpha = 0.6) +
  geom_smooth(method = "lm", color = "red", fill = "mistyrose", se = TRUE) +
  ggtitle("Sulphates vs. Quality")

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

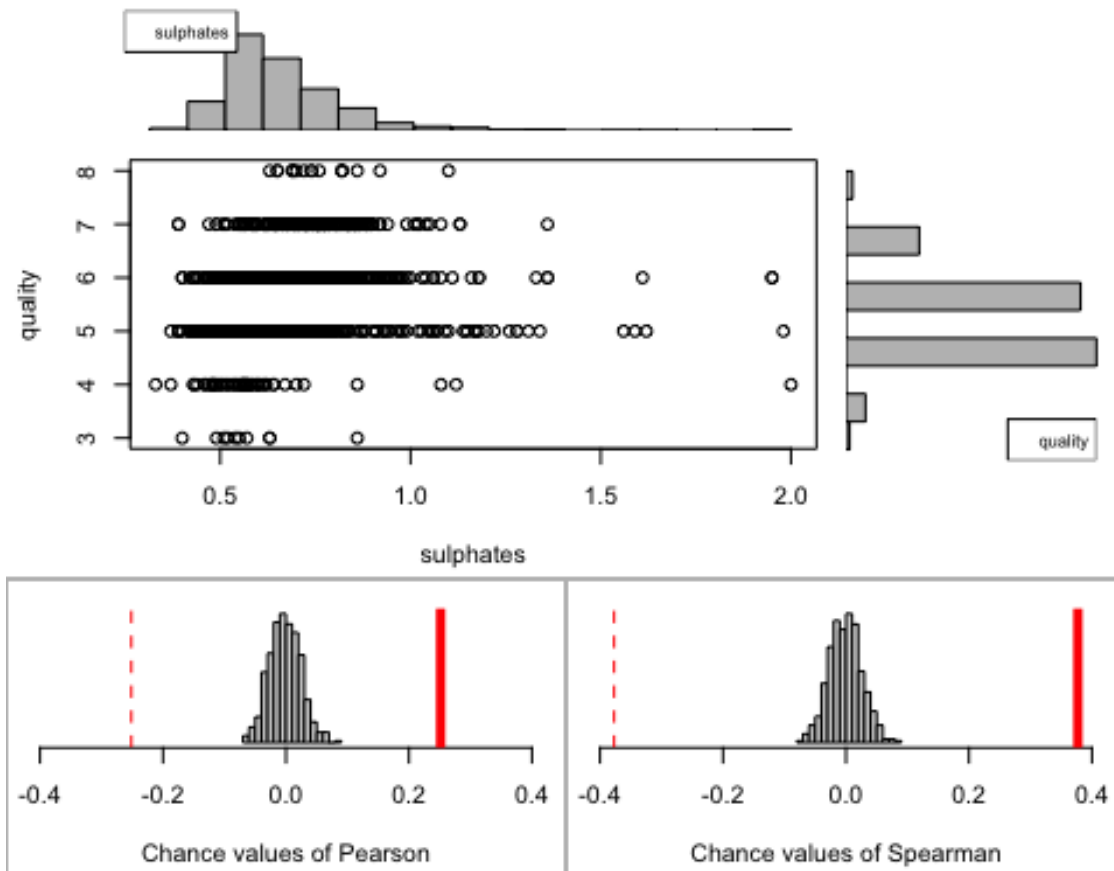


#Checking for association

```
associate(quality~sulphates, data = wine)
```

```
## Association between sulphates (numerical) and quality (numerical)
```

```
## using 1599 complete cases
```



```
## Permutation procedure:
```

```
##                                     Value Estimated p-value
```

```
## Pearson's r                        0.2513971                0
```

```
## Spearman's rank correlation 0.3770602                0
```

```
## With 500 permutations, we are 95% confident that:
```

```
## the p-value of Pearson's correlation (r) is between 0 and 0.007
```

```
## the p-value of Spearman's rank correlation is between 0 and 0.007
```

```
## Note: If 0.05 is in this range, increase the permutations= argument.
```

```
##
```

```
##
```

```
##
```

```
## Advice: If stream of points is well described by an ellipse, use Pearson's  
## r.
```

```
## Otherwise, as long as stream is monotonic, use Spearman's rank correlation
```

```
## or try logs, e.g. associate( log10(y)~log10(x) )
```

#Checking for correlation against all variables

```
all_correlations(wine, interest = "sulphates", sorted = "magnitude")
```

##		var1	var2	correlation	pval
## 1		chlorides	sulphates	0.371260481	1.986310e-53
## 2		citric.acid	sulphates	0.312770044	1.265262e-37
## 3		volatile.acidity	sulphates	-0.260986685	2.606926e-26
## 4		sulphates	quality	0.251397079	1.802088e-24
## 5		pH	sulphates	-0.196647602	2.106734e-15
## 6		fixed.acidity	sulphates	0.183005664	1.648652e-13
## 7		density	sulphates	0.148506412	2.418474e-09
## 8		sulphates	alcohol	0.093594750	1.783053e-04
## 9		free.sulfur.dioxide	sulphates	0.051657572	3.888321e-02
## 10		total.sulfur.dioxide	sulphates	0.042946836	8.601835e-02
## 11		residual.sugar	sulphates	0.005527121	8.252134e-01

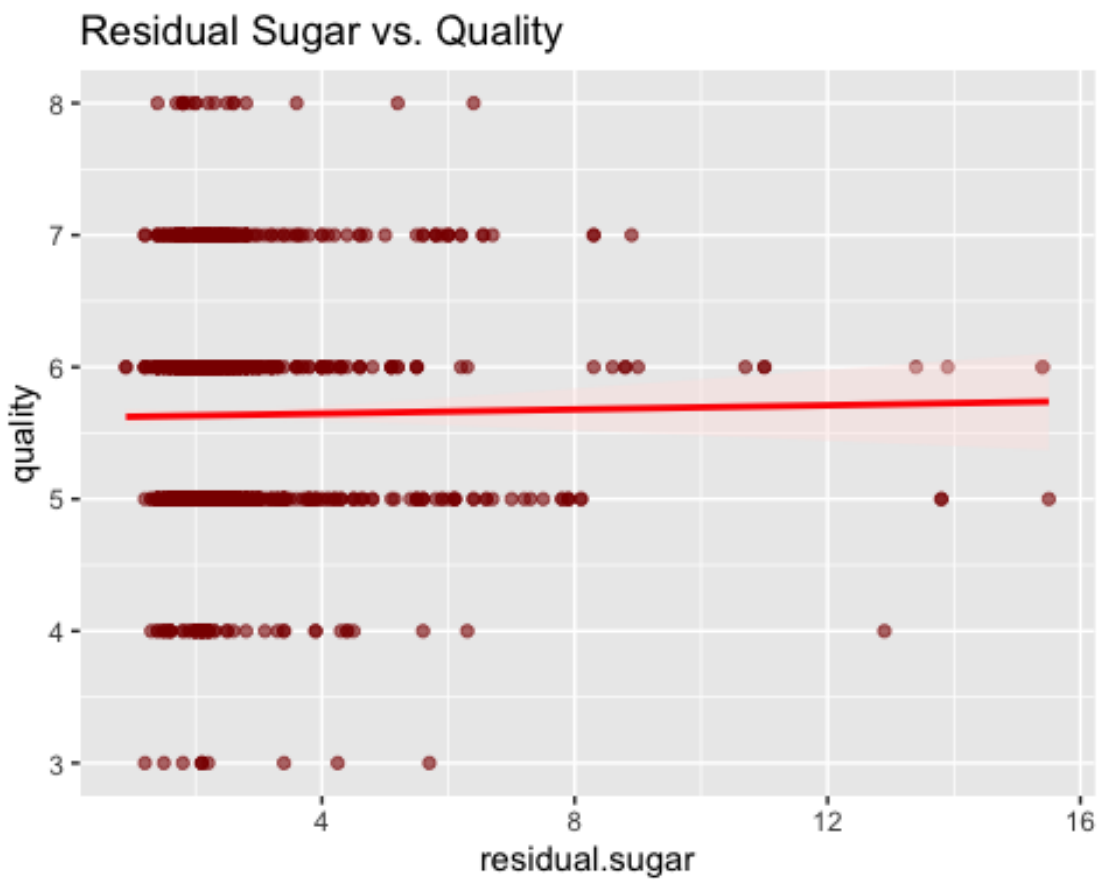
Analysis: According to the Pearson correlation ($r = 0.25$), there is a moderate positive linear relationship between the quality of wine and sulphates. As sulphates increase, the quality of wine tends to increase slightly, but it is not a very strong correlation. P-values are lower than 0.007 for both quality and sulphates, so the correlations are significant between the two and sulphates are meaningfully associated with the quality of wine, even if the linear relationship is weaker. Sulphates appear to correlate strongly with other variables such as citric acid, suggesting that sulphates is more present in the chemical composition of the wine rather than the quality.

Residual Sugar

#Graph

```
ggplot(wine, aes(x = residual.sugar, y = quality)) +  
  geom_point(color = "darkred", alpha = 0.6) +  
  geom_smooth(method = "lm", color = "red", fill = "mistyrose", se = TRUE) +  
  ggtitle("Residual Sugar vs. Quality")
```

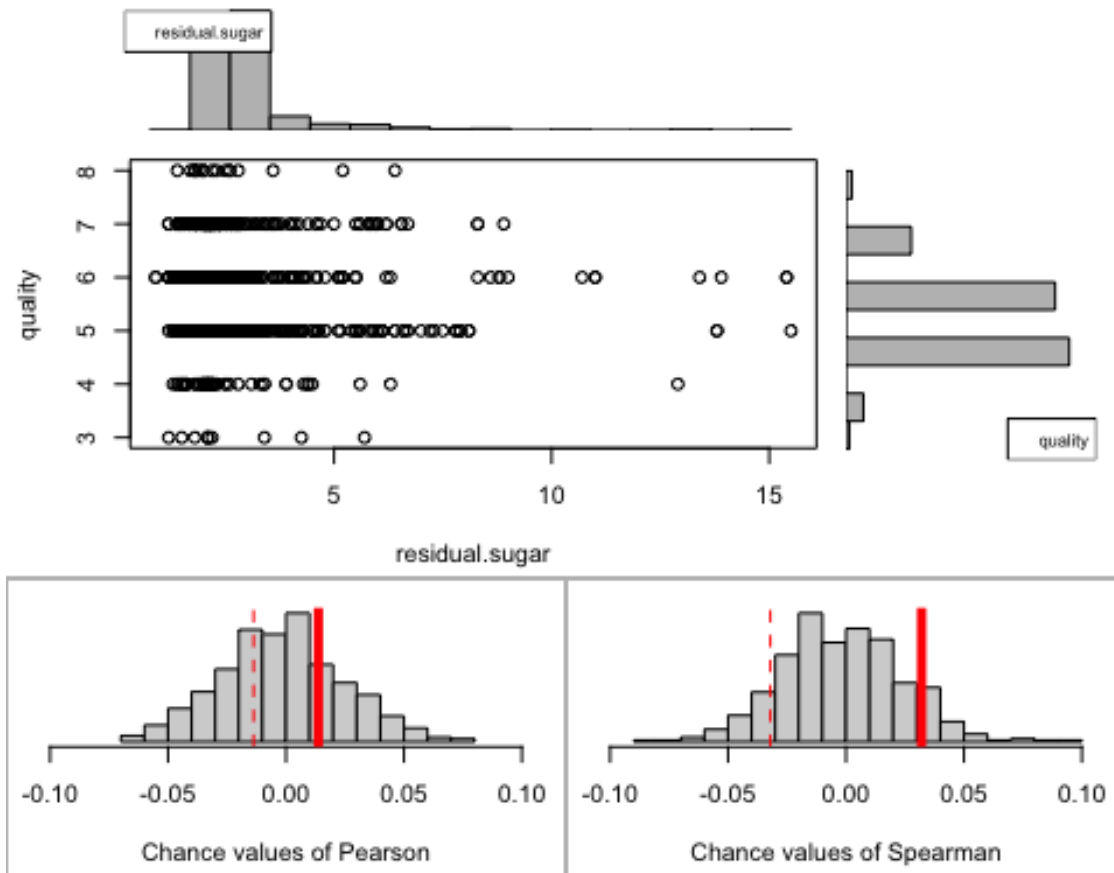
```
## `geom_smooth()` using formula = 'y ~ x'
```



#Checking for association

```
associate(quality~residual.sugar, data = wine)
```

```
## Association between residual.sugar (numerical) and  quality (numerical)  
## using 1599 complete cases
```



```
## Permutation procedure:
##                               Value Estimated p-value
## Pearson's r                  0.01373164          0.600
## Spearman's rank correlation 0.03204817          0.202
## With 500 permutations, we are 95% confident that:
## the p-value of Pearson's correlation (r) is between 0.556 and 0.643
## the p-value of Spearman's rank correlation is between 0.168 and 0.24
## Note: If 0.05 is in this range, increase the permutations= argument.
##
##
## Advice: If stream of points is well described by an ellipse, use Pearson's
## r.
## Otherwise, as long as stream is monotonic, use Spearman's rank correlation
## or try logs, e.g. associate( log10(y)~log10(x) )
```

#Checking for correlation against all variables

```
all_correlations(wine, interest = "residual.sugar", sorted = "magnitude")
```

	var1	var2	correlation	pval
## 1	residual.sugar	density	0.355283371	9.013042e-49
## 2	residual.sugar	total.sulfur.dioxide	0.203027882	2.449285e-16
## 3	residual.sugar	free.sulfur.dioxide	0.187048995	4.684735e-14

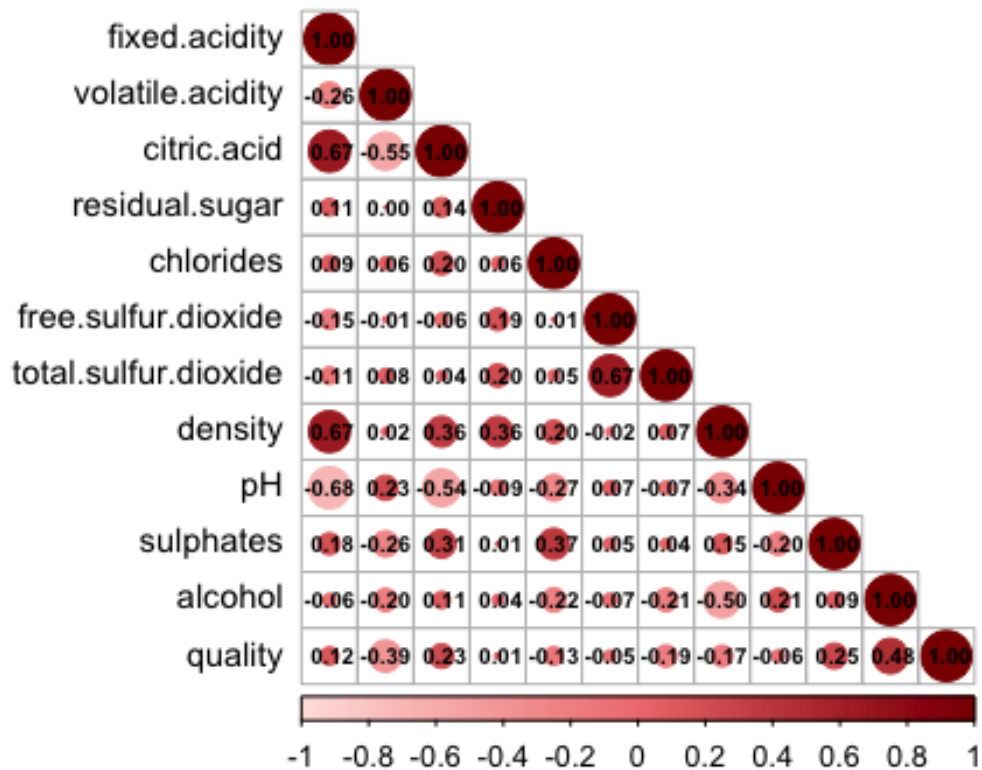

```
## 4      citric.acid      residual.sugar  0.143577162 8.083723e-09
## 5      fixed.acidity    residual.sugar  0.114776724 4.199465e-06
## 6      residual.sugar    pH -0.085652422 6.065915e-04
## 7      residual.sugar    chlorides  0.055609535 2.617079e-02
## 8      residual.sugar    alcohol  0.042075437 9.258425e-02
## 9      residual.sugar    quality  0.013731637 5.832180e-01
## 10     residual.sugar    sulphates  0.005527121 8.252134e-01
## 11 volatile.acidity     residual.sugar  0.001917882 9.389168e-01
```

Analysis: According to the Pearson correlation ($r = 0.0137$), there is nearly no linear relationship between the quality of wine and residual sugar. P-values are higher than 0.05, so the correlations are not statistically significant. We fail to reject the null hypothesis, meaning residual sugar is not associated with the quality of wine. In the graph, there is no visible pattern and the data points are concentrated at low sugar levels, with no clear slope. Residual sugar has some correlation with density, as sugar adds weight to the wine. Overall, there is no meaningful correlation between the quality of wine and residual sugar.

Correlations Visualization

```
cor_matrix <- cor(wine)
wine_colors <- colorRampPalette(c("mistyrose", "lightcoral", "darkred"))(200)
corrplot(cor_matrix, method = "circle", type = "lower", col = wine_colors,
tl.col = "black", tl.cex = 0.9, tl.pos = "l", cl.pos = "b", cl.cex = 0.8,
number.cex = 0.6, addCoef.col = "black", title = "Red Wine Quality
Correlation Matrix", mar = c(0, 0, 2, 0))
```

Red Wine Quality Correlation Matrix



Analysis: This correlation matrix creates a visualization of the linear relationships between the quality of wine and other variables in the chemical properties. The darker red shows a stronger positive correlation, and the lighter colors show weaker correlations. Negative correlations are shown in lighter colors. The bigger circles show stronger correlation, whether it is positive or negative, and smaller circles show weak or no correlation.

Alcohol appears to have the strongest positive correlations with quality at 0.48, meanwhile volatile acidity shows a weak negative correlation with quality at -0.26. Residual sugar and chlorides show little or no correlation with quality. Citric acid and fixed acidity are strongly correlated with each other, and weakly correlated with pH. Residual sugar, chlorides, and sulphates have either little or no direct relationship with quality of wine.

Multi linear regression

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	2.8431068	0.2050732	13.864	< 2e-16	***
alcohol	0.2953419	0.0160375	18.416	< 2e-16	***
volatile.acidity	-1.2223102	0.1124774	-10.867	< 2e-16	***
sulphates	0.7207881	0.1027039	7.018	3.32e-12	***
citric.acid	-0.0427246	0.1035810	-0.412	0.68	
total.sulfur.dioxide	-0.0022182	0.0005126	-4.327	1.60e-05	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.6552 on 1593 degrees of freedom

Multiple R-squared: 0.3439, Adjusted R-squared: 0.3418

Analysis: Of the coefficients all are statistically significant except for citric acid using a standard 95% confidence interval ($p < .05$). From the significant coefficients two of them are positive and two are negative with alcohol level and sulfates both providing a positive impact though the model shows that sulphates have a higher impact on the wines quality. From this wine makers could conclude that focusing on a wine that emphasizes sulphates over one that does not would result in a better end product.

On the other hand, volatile acidity has a very negative impact on wine quality. The presence of excess citric acid causes an undesirable taste in the wine and because this coefficient is high it shows that it has a large impact on how wine is made and should be considered for wine makers Total sulfur dioxide value is low indicating that even though it is statistically significant the impact that it has is negligible. This shows that sulfur dioxide may be unnoticeable to people who are drinking the wine and is not something that may need to be considered for wine makers.

The R squared value for this model indicates that it explains an adequate amount of the variance but there may be other factors that contribute to a wines quality outside of what is in the dataset currently such as vineyard or vintage.