GPH 2338 Project

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2023-03-13

Data Upload

```
library(haven)
library(psych)
```

```
## Warning: package 'psych' was built under R version 4.2.2
```

```
library(ggplot2)
```

```
## Warning: package 'ggplot2' was built under R version 4.2.2
```

```
##
## Attaching package: 'ggplot2'
```

```
## The following objects are masked from 'package:psych':
##
##     %+%, alpha
```

```
library(psych)
library(pander)
```

```
## Warning: package 'pander' was built under R version 4.2.2
```

```
library(corrplot)
```

```
## Warning: package 'corrplot' was built under R version 4.2.2
```

```
## corrplot 0.92 loaded
```

```
require(readr)
```

```
## Loading required package: readr
```

```
## Warning: package 'readr' was built under R version 4.2.2
```

```
require(r02pro)
```

```
## Loading required package: r02pro
```

```
## Warning: package 'r02pro' was built under R version 4.2.2
```

```
wine <- read_csv("Wine_Quality_Data.csv")
```

```
## Rows: 6497 Columns: 13
```

```
## -- Column specification --------------------------------------------------
## Delimiter: ","
## chr  (1): color
## dbl (12): fixed_acidity, volatile_acidity, citric_acid, residual_sugar, chlo...
##
## i Use 'spec()' to retrieve the full column specification for this data.
## i Specify the column types or set 'show_col_types = FALSE' to quiet this message.
```

```
head(wine)
```

```
## # A tibble: 6 x 13
##    fixed_~1 volat~2 citri~3 resid~4 chlor~5 free_~6 total~7 density    pH sulph~8
##       <dbl>   <dbl>   <dbl>   <dbl>   <dbl>   <dbl>   <dbl>   <dbl> <dbl>   <dbl>
## 1      7.4    0.7    0         1.9   0.076      11      34   0.998  3.51    0.56
## 2      7.8    0.88   0         2.6   0.098      25      67   0.997  3.2     0.68
## 3      7.8    0.76   0.04      2.3   0.092      15      54   0.997  3.26    0.65
## 4     11.2    0.28   0.56      1.9   0.075      17      60   0.998  3.16    0.58
## 5      7.4    0.7    0         1.9   0.076      11      34   0.998  3.51    0.56
## 6      7.4    0.66   0         1.8   0.075      13      40   0.998  3.51    0.56
## # ... with 3 more variables: alcohol <dbl>, quality <dbl>, color <chr>, and
## #   abbreviated variable names 1: fixed_acidity, 2: volatile_acidity,
## #   3: citric_acid, 4: residual_sugar, 5: chlorides, 6: free_sulfur_dioxide,
## #   7: total_sulfur_dioxide, 8: sulphates
```

Data Preparation

```
sum(wine$color == "white")
```

```
## [1] 4898
```

```
range(wine$fixed_acidity)
```

```
## [1]  3.8 15.9
```

```
range(wine$volatile_acidity)
```

```
## [1] 0.08 1.58
```

```
range(wine$citric_acid)
```

```
## [1] 0.00 1.66
```

```
range(wine$residual_sugar)
```

```
## [1]  0.6 65.8
```

```
range(wine$chlorides)
```

```
## [1] 0.009 0.611
```

```
range(wine$free_sulfur_dioxide)
```

```
## [1]   1 289
```

```
range(wine$total_sulfur_dioxide)
```

```
## [1]   6 440
```

```
range(wine$density)
```

```
## [1] 0.98711 1.03898
```

```
range(wine$pH)
```

```
## [1] 2.72 4.01
```

```
range(wine$sulphates)
```

```
## [1] 0.22 2.00
```

```
range(wine$alcohol)
```

```
## [1]  8.0 14.9
```

```
range(wine$quality)
```

```
## [1] 3 9
```

```
range(wine$color)
```

```
## [1] "red"   "white"
```

```r
sum(is.na(wine$fixed_acidity))
```

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

```r
sum(is.na(wine$fixed_acidity))
```

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

```r
sum(is.na(wine$volatile_acidity))
```

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

```r
sum(is.na(wine$citric_acid))
```

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

```r
sum(is.na(wine$residual_sugar))
```

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

```r
sum(is.na(wine$chlorides))
```

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

```r
sum(is.na(wine$free_sulfur_dioxide))
```

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

```r
sum(is.na(wine$total_sulfur_dioxide))
```

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

```r
sum(is.na(wine$density))
```

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

```r
sum(is.na(wine$pH))
```

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

```r
sum(is.na(wine$sulphates))
```

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

```
sum(is.na(wine$alcohol))
```

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

```
sum(is.na(wine$quality))
```

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

```
dim(wine)
```

```
## [1] 6497    13
```