

Veri Biliminde R Uygulamaları Odev

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Contents

1	Veri setinin detaylı incelenmesi ve özet halinde açıklanması	2
2	Veri Ön İşleme	3
2.1	Veri öz nitelikleri	3
2.2	Değişken seçimi ve dönüşüm işlemleri	4
2.3	dplyr paketi ile temel işlemler(veri seçme ve filtreleme)	5
3	Veri Manipülasyonu	5
3.1	Veri setinin özelliklerinin analize hazır hale getirilmesi(reshaping data)	5
3.2	Eksik veri ve aykırı değerlerin tespiti	6
3.3	Eksik verilerin tamamlanması ya da analiz dışı bırakılması	6
3.4	Veri normalizasyonu ya da standardizasyonu	7
3.5	Veri seçme ve filtreleme işlemlerinin gerçekleştirilmesi	8
3.6	Yeni hesaplamaların veri setine dâhil edilmesi	8
3.7	Temel istatistiklerin hesaplanması	9
4	Keşifçi ve Açıklayıcı Veri Analizi	13
4.1	ggplot2 paketi ile uygun özelliklere ait veri görselleştirmenin gerçekleştirilmesi	13

Veri setine erişim linki: <https://archive.ics.uci.edu/dataset/109/wine>

“Wine” veri seti, üç farklı sınıfa ait üzüm şaraplarından elde edilen kimyasal bileşenleri içerir. Bu veri setinin özellikleri şunlardır:

- **Wine:** Her bir şarap örneğinin sınıfını belirten bir değişkeni ifade eder. Bu değişken, şarap örneklerinin sınıflarını temsil eden kategorik bir değişkendir. Üç farklı sınıfa ait şarap örneklerini içerir
- **Alcohol:** Şaraptaki alkol oranını ölçen sayısal bir özellik.
- **Malic Acid:** Şaraptaki elma asidi miktarını ölçen sayısal bir özellik.
- **Ash:** Şaraptaki kül miktarını ölçen sayısal bir özellik.
- **Alcalinity(Acl):** Şaraptaki külün alkalinitesini ölçen sayısal bir özellik.
- **Magnesium(Mg):** Şaraptaki magnezyum miktarını ölçen sayısal bir özellik.
- **Phenols:** Şaraptaki toplam fenol miktarını ölçen sayısal bir özellik.
- **Flavanoids:** Şaraptaki flavanoid miktarını ölçen sayısal bir özellik.

- **Nonflavanoid Phenols:** Şaraptaki nonflavanoid fenol miktarını ölçen sayısal bir özellik.
- **Proanthocyanins:** Şaraptaki proantosiyandin miktarını ölçen sayısal bir özellik.
- **Color Intensity:** Şaraptaki renk yoğunluğunu ölçen sayısal bir özellik.
- **Hue:** Şaraptaki renk tonunu ölçen sayısal bir özellik.
- **OD:** Şarabın 280/315 oranındaki optik yoğunluğunu ölçen sayısal bir özellik.
- **Proline:** Şaraptaki prolin miktarını ölçen sayısal bir özellik.

1 Veri setinin detaylı incelenmesi ve özet halinde açıklanması

```
library(dplyr)
library(tidyverse)
wine_data = read.csv("wine.csv",header = T, sep=",")
wine_data = as_tibble(wine_data)
head(wine_data)
```

```
## # A tibble: 6 x 14
##   Wine Alcohol Malic.acid  Ash  Acl  Mg Phenols Flavanoids
##   <int>   <dbl>    <dbl> <dbl> <dbl> <int>   <dbl>    <dbl>
## 1     1    14.2      1.71  2.43  15.6  127     2.8      3.06
## 2     1    13.2      1.78  2.14  11.2  100     2.65      2.76
## 3     1    13.2      2.36  2.67  18.6  101     2.8      3.24
## 4     1    14.4      1.95  2.5   16.8  113     3.85      3.49
## 5     1    13.2      2.59  2.87  21    118     2.8      2.69
## 6     1    14.2      1.76  2.45  15.2  112     3.27      3.39
## # i 6 more variables: Nonflavanoid.phenols <dbl>, Proanth <dbl>,
## #   Color.int <dbl>, Hue <dbl>, OD <dbl>, Proline <int>
```

```
glimpse(wine_data)
```

```
## Rows: 178
## Columns: 14
## $ Wine           <int> 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1~
## $ Alcohol        <dbl> 14.23, 13.20, 13.16, 14.37, 13.24, 14.20, 14.39, ~
## $ Malic.acid     <dbl> 1.71, 1.78, 2.36, 1.95, 2.59, 1.76, 1.87, 2.15, 1~
## $ Ash            <dbl> 2.43, 2.14, 2.67, 2.50, 2.87, 2.45, 2.45, 2.61, 2~
## $ Acl            <dbl> 15.6, 11.2, 18.6, 16.8, 21.0, 15.2, 14.6, 17.6, 1~
## $ Mg             <int> 127, 100, 101, 113, 118, 112, 96, 121, 97, 98, 10~
## $ Phenols        <dbl> 2.80, 2.65, 2.80, 3.85, 2.80, 3.27, 2.50, 2.60, 2~
## $ Flavanoids     <dbl> 3.06, 2.76, 3.24, 3.49, 2.69, 3.39, 2.52, 2.51, 2~
## $ Nonflavanoid.phenols <dbl> 0.28, 0.26, 0.30, 0.24, 0.39, 0.34, 0.30, 0.31, 0~
## $ Proanth        <dbl> 2.29, 1.28, 2.81, 2.18, 1.82, 1.97, 1.98, 1.25, 1~
## $ Color.int      <dbl> 5.64, 4.38, 5.68, 7.80, 4.32, 6.75, 5.25, 5.05, 5~
## $ Hue            <dbl> 1.04, 1.05, 1.03, 0.86, 1.04, 1.05, 1.02, 1.06, 1~
## $ OD             <dbl> 3.92, 3.40, 3.17, 3.45, 2.93, 2.85, 3.58, 3.58, 2~
## $ Proline        <int> 1065, 1050, 1185, 1480, 735, 1450, 1290, 1295, 10~
```

```
class(wine_data)
```

```
## [1] "tbl_df"      "tbl"        "data.frame"
```

2 Veri Ön İşleme

2.1 Veri öz nitelikleri

2.1.1 Seçilen veri setinde analiz için kullanılacak özelliklerin belirlenmesi

```
features = select(wine_data, Alcohol:Proline)
features
```

```
## # A tibble: 178 x 13
##   Alcohol Malic.acid Ash   Acl   Mg Phenols Flavanoids Nonflavanoid.phenols
##   <dbl>    <dbl> <dbl> <dbl> <int> <dbl>    <dbl>          <dbl>
## 1    14.2      1.71  2.43  15.6  127   2.8      3.06          0.28
## 2    13.2      1.78  2.14  11.2  100   2.65     2.76          0.26
## 3    13.2      2.36  2.67  18.6  101   2.8      3.24          0.3
## 4    14.4      1.95  2.5   16.8  113   3.85     3.49          0.24
## 5    13.2      2.59  2.87  21    118   2.8      2.69          0.39
## 6    14.2      1.76  2.45  15.2  112   3.27     3.39          0.34
## 7    14.4      1.87  2.45  14.6   96   2.5      2.52          0.3
## 8    14.1      2.15  2.61  17.6  121   2.6      2.51          0.31
## 9    14.8      1.64  2.17  14     97   2.8      2.98          0.29
## 10   13.9      1.35  2.27  16     98   2.98     3.15          0.22
## # i 168 more rows
## # i 5 more variables: Proanth <dbl>, Color.int <dbl>, Hue <dbl>, OD <dbl>,
## #   Proline <int>
```

```
summary(features)
```

```
##      Alcohol      Malic.acid      Ash      Acl
## Min.   :11.03 Min.   :0.740 Min.   :1.360 Min.   :10.60
## 1st Qu.:12.36 1st Qu.:1.603 1st Qu.:2.210 1st Qu.:17.20
## Median :13.05 Median :1.865 Median :2.360 Median :19.50
## Mean   :13.00 Mean   :2.336 Mean   :2.367 Mean   :19.49
## 3rd Qu.:13.68 3rd Qu.:3.083 3rd Qu.:2.558 3rd Qu.:21.50
## Max.   :14.83 Max.   :5.800 Max.   :3.230 Max.   :30.00
##      Mg      Phenols      Flavanoids      Nonflavanoid.phenols
## Min.   : 70.00 Min.   :0.980 Min.   :0.340 Min.   :0.1300
## 1st Qu.: 88.00 1st Qu.:1.742 1st Qu.:1.205 1st Qu.:0.2700
## Median : 98.00 Median :2.355 Median :2.135 Median :0.3400
## Mean   : 99.74 Mean   :2.295 Mean   :2.029 Mean   :0.3619
## 3rd Qu.:107.00 3rd Qu.:2.800 3rd Qu.:2.875 3rd Qu.:0.4375
## Max.   :162.00 Max.   :3.880 Max.   :5.080 Max.   :0.6600
##      Proanth      Color.int      Hue      OD
## Min.   :0.410 Min.   : 1.280 Min.   :0.4800 Min.   :1.270
## 1st Qu.:1.250 1st Qu.: 3.220 1st Qu.:0.7825 1st Qu.:1.938
## Median :1.555 Median : 4.690 Median :0.9650 Median :2.780
## Mean   :1.591 Mean   : 5.058 Mean   :0.9574 Mean   :2.612
## 3rd Qu.:1.950 3rd Qu.: 6.200 3rd Qu.:1.1200 3rd Qu.:3.170
## Max.   :3.580 Max.   :13.000 Max.   :1.7100 Max.   :4.000
##      Proline
## Min.   : 278.0
```

```
## 1st Qu.: 500.5
## Median : 673.5
## Mean : 746.9
## 3rd Qu.: 985.0
## Max. :1680.0
```

```
correlation_matrix = cor(features)
head(correlation_matrix)
```

```
##           Alcohol Malic.acid      Ash      Acl      Mg      Phenols
## Alcohol      1.00000000 0.09439694 0.2115446 -0.31023514 0.27079823 0.2891011
## Malic.acid   0.09439694 1.00000000 0.1640455 0.28850040 -0.05457510 -0.3351670
## Ash          0.21154460 0.16404547 1.00000000 0.44336719 0.28658669 0.1289795
## Acl          -0.31023514 0.28850040 0.4433672 1.00000000 -0.08333309 -0.3211133
## Mg           0.27079823 -0.05457510 0.2865867 -0.08333309 1.00000000 0.2144012
## Phenols      0.28910112 -0.33516700 0.1289795 -0.32111332 0.21440123 1.0000000
##           Flavanoids Nonflavanoid.phenols      Proanth      Color.int      Hue
## Alcohol      0.2368149          -0.1559295 0.136697912 0.54636420 -0.07174720
## Malic.acid   -0.4110066          0.2929771 -0.220746187 0.24898534 -0.56129569
## Ash          0.1150773          0.1862304 0.009651935 0.25888726 -0.07466689
## Acl          -0.3513699          0.3619217 -0.197326836 0.01873198 -0.27395522
## Mg           0.1957838          -0.2562940 0.236440610 0.19995001 0.05539820
## Phenols      0.8645635          -0.4499353 0.612413084 -0.05513642 0.43368134
##           OD      Proline
## Alcohol      0.072343187 0.6437200
## Malic.acid   -0.368710428 -0.1920106
## Ash          0.003911231 0.2236263
## Acl          -0.276768549 -0.4405969
## Mg           0.066003936 0.3933508
## Phenols      0.699949365 0.4981149
```

2.2 Değişken seçimi ve dönüşüm işlemleri

```
#Seçilen sayısal değişkenler gather fonksiyonu ile uzun formatlı hale getirildi.
(long_data = wine_data %>% keep(is.numeric) %>% gather())
```

```
## # A tibble: 2,492 x 2
##   key   value
##   <chr> <dbl>
## 1 Wine      1
## 2 Wine      1
## 3 Wine      1
## 4 Wine      1
## 5 Wine      1
## 6 Wine      1
## 7 Wine      1
## 8 Wine      1
## 9 Wine      1
## 10 Wine     1
## # i 2,482 more rows
```

2.3 dplyr paketi ile temel işlemler(veri seçme ve filtreleme)

```
filter(wine_data, Alcohol > 13 & Phenols > 2)
```

```
## # A tibble: 66 x 14
##   Wine Alcohol Malic.acid Ash Acl Mg Phenols Flavanoids
##   <int>   <dbl>   <dbl> <dbl> <dbl> <int>   <dbl>   <dbl>
## 1     1     14.2     1.71  2.43  15.6  127     2.8     3.06
## 2     1     13.2     1.78  2.14  11.2  100     2.65     2.76
## 3     1     13.2     2.36  2.67  18.6  101     2.8     3.24
## 4     1     14.4     1.95  2.5   16.8  113     3.85     3.49
## 5     1     13.2     2.59  2.87  21    118     2.8     2.69
## 6     1     14.2     1.76  2.45  15.2  112     3.27     3.39
## 7     1     14.4     1.87  2.45  14.6   96     2.5     2.52
## 8     1     14.1     2.15  2.61  17.6  121     2.6     2.51
## 9     1     14.8     1.64  2.17  14     97     2.8     2.98
## 10    1     13.9     1.35  2.27  16     98     2.98     3.15
## # i 56 more rows
## # i 6 more variables: Nonflavanoid.phenols <dbl>, Proanth <dbl>,
## #   Color.int <dbl>, Hue <dbl>, OD <dbl>, Proline <int>
```

```
wine_data %>%
  group_by(Wine) %>%
  summarise(count = n())
```

```
## # A tibble: 3 x 2
##   Wine count
##   <int> <int>
## 1     1     59
## 2     2     71
## 3     3     48
```

```
grouped_data <- wine_data %>%
  group_by(Wine) %>%
  summarise(mean_Alcohol = mean(Alcohol), mean_Color_Int = mean(Color.int))
print(grouped_data)
```

```
## # A tibble: 3 x 3
##   Wine mean_Alcohol mean_Color_Int
##   <int>   <dbl>   <dbl>
## 1     1     13.7     5.53
## 2     2     12.3     3.09
## 3     3     13.2     7.40
```

3 Veri Manipülasyonu

3.1 Veri setinin özelliklerinin analize hazır hale getirilmesi(reshaping data)

```
normalized_data = scale(wine_data[, 2:ncol(wine_data)])
head(normalized_data)
```

```
##           Alcohol Malic.acid      Ash      Acl      Mg      Phenols
## [1,] 1.5143408 -0.56066822  0.2313998 -1.1663032 1.90852151 0.8067217
## [2,] 0.2455968 -0.49800856 -0.8256672 -2.4838405 0.01809398 0.5670481
## [3,] 0.1963252  0.02117152  1.1062139 -0.2679823 0.08810981 0.8067217
## [4,] 1.6867914 -0.34583508  0.4865539 -0.8069748 0.92829983 2.4844372
## [5,] 0.2948684  0.22705328  1.8352256  0.4506745 1.27837900 0.8067217
## [6,] 1.4773871 -0.51591132  0.3043010 -1.2860793 0.85828399 1.5576991
##           Flavanoids Nonflavanoid.phenols      Proanth      Color.int      Hue      OD
## [1,] 1.0319081          -0.6577078  1.2214385  0.2510088  0.3611585 1.8427215
## [2,] 0.7315653          -0.8184106 -0.5431887 -0.2924962  0.4049085 1.1103172
## [3,] 1.2121137          -0.4970050  2.1299594  0.2682629  0.3174085 0.7863692
## [4,] 1.4623994          -0.9791134  1.0292513  1.1827317 -0.4263410 1.1807407
## [5,] 0.6614853           0.2261576  0.4002753 -0.3183774  0.3611585 0.4483365
## [6,] 1.3622851          -0.1755994  0.6623487  0.7298108  0.4049085 0.3356589
##           Proline
## [1,] 1.01015939
## [2,] 0.96252635
## [3,] 1.39122370
## [4,] 2.32800680
## [5,] -0.03776747
## [6,] 2.23274072
```

3.2 Eksik veri ve aykırı değerlerin tespiti

```
missing_values = wine_data %>%
  summarise_all(~ sum(is.na(.)))
missing_values
```

```
## # A tibble: 1 x 14
##   Wine Alcohol Malic.acid      Ash      Acl      Mg Phenols Flavanoids
##   <int>   <int>      <int> <int> <int> <int>   <int>      <int>
## 1     0     0          0     0     0     0     0          0
## # i 6 more variables: Nonflavanoid.phenols <int>, Proanth <int>,
## #   Color.int <int>, Hue <int>, OD <int>, Proline <int>
```

```
outliers = wine_data %>%
  filter_all(all_vars(!is.na(.) & (. < quantile(., 0.25) - 1.5 * IQR(.) | . > quantile(., 0.75) + 1.5 *
outliers
```

```
## # A tibble: 0 x 14
## # i 14 variables: Wine <int>, Alcohol <dbl>, Malic.acid <dbl>, Ash <dbl>,
## #   Acl <dbl>, Mg <int>, Phenols <dbl>, Flavanoids <dbl>,
## #   Nonflavanoid.phenols <dbl>, Proanth <dbl>, Color.int <dbl>, Hue <dbl>,
## #   OD <dbl>, Proline <int>
```

3.3 Eksik verilerin tamamlanması ya da analiz dışı bırakılması

Eksik veri bulunmamıştır.

3.4 Veri normalizasyonu ya da standardizasyonu

```
normalize_et = function(x) {
  return((x - min(x)) / (max(x) - min(x)))
}

veri_setini_normalize_et = function(veri_seti) {
  normalize_edilmis_set = as.data.frame(lapply(veri_seti, function(col) {
    if (is.numeric(col)) {
      return(normalize_et(col))
    } else {
      return(col)
    }
  })))
  return(normalize_edilmis_set)
}

normalize_data = veri_setini_normalize_et(wine_data)
head(normalize_data)
```

```
##   Wine   Alcohol Malic.acid      Ash      Acl      Mg   Phenols Flavanoids
## 1    0 0.8421053 0.1916996 0.5721925 0.25773196 0.6195652 0.6275862 0.5738397
## 2    0 0.5710526 0.2055336 0.4171123 0.03092784 0.3260870 0.5758621 0.5105485
## 3    0 0.5605263 0.3201581 0.7005348 0.41237113 0.3369565 0.6275862 0.6118143
## 4    0 0.8789474 0.2391304 0.6096257 0.31958763 0.4673913 0.9896552 0.6645570
## 5    0 0.5815789 0.3656126 0.8074866 0.53608247 0.5217391 0.6275862 0.4957806
## 6    0 0.8342105 0.2015810 0.5828877 0.23711340 0.4565217 0.7896552 0.6434599
## Nonflavanoid.phenols  Proanth Color.int      Hue      OD   Proline
## 1                0.2830189 0.5930599 0.3720137 0.4552846 0.9706960 0.5613409
## 2                0.2452830 0.2744479 0.2645051 0.4634146 0.7802198 0.5506419
## 3                0.3207547 0.7570978 0.3754266 0.4471545 0.6959707 0.6469330
## 4                0.2075472 0.5583596 0.5563140 0.3089431 0.7985348 0.8573466
## 5                0.4905660 0.4447950 0.2593857 0.4552846 0.6080586 0.3259629
## 6                0.3962264 0.4921136 0.4667235 0.4634146 0.5787546 0.8359486
```

```
standardize_et = function(x) {
  return((x - mean(x)) / sd(x))
}

veri_setini_standardize_et = function(veri_seti) {
  standardize_edilmis_set = as.data.frame(lapply(veri_seti, function(col) {
    if (is.numeric(col)) {
      return(standardize_et(col))
    } else {
      return(col)
    }
  })))
  return(standardize_edilmis_set)
}

standardize_data = veri_setini_standardize_et(wine_data)
head(standardize_data)
```

```
##      Wine Alcohol Malic.acid      Ash      Acl      Mg Phenols
## 1 -1.210529 1.5143408 -0.56066822 0.2313998 -1.1663032 1.90852151 0.8067217
## 2 -1.210529 0.2455968 -0.49800856 -0.8256672 -2.4838405 0.01809398 0.5670481
## 3 -1.210529 0.1963252 0.02117152 1.1062139 -0.2679823 0.08810981 0.8067217
## 4 -1.210529 1.6867914 -0.34583508 0.4865539 -0.8069748 0.92829983 2.4844372
## 5 -1.210529 0.2948684 0.22705328 1.8352256 0.4506745 1.27837900 0.8067217
## 6 -1.210529 1.4773871 -0.51591132 0.3043010 -1.2860793 0.85828399 1.5576991
##      Flavanoids Nonflavanoid.phenols      Proanth      Color.int      Hue      OD
## 1 1.0319081 -0.6577078 1.2214385 0.2510088 0.3611585 1.8427215
## 2 0.7315653 -0.8184106 -0.5431887 -0.2924962 0.4049085 1.1103172
## 3 1.2121137 -0.4970050 2.1299594 0.2682629 0.3174085 0.7863692
## 4 1.4623994 -0.9791134 1.0292513 1.1827317 -0.4263410 1.1807407
## 5 0.6614853 0.2261576 0.4002753 -0.3183774 0.3611585 0.4483365
## 6 1.3622851 -0.1755994 0.6623487 0.7298108 0.4049085 0.3356589
##      Proline
## 1 1.01015939
## 2 0.96252635
## 3 1.39122370
## 4 2.32800680
## 5 -0.03776747
## 6 2.23274072
```

3.5 Veri seçme ve filtreleme işlemlerinin gerçekleştirilmesi

```
Alcohol_Category = cut(wine_data$Alcohol, breaks = c(0, 12, 14, 16), labels = c("Low", "Medium", "High"))
```

3.6 Yeni hesaplamaların veri setine dâhil edilmesi

```
wine_data = wine_data %>%
  mutate(Alcohol_Category = cut(Alcohol, breaks = c(0, 12, 14, 16), labels = c("Low", "Medium", "High")))
head(wine_data)
```

```
## # A tibble: 6 x 15
##      Wine Alcohol Malic.acid      Ash      Acl      Mg Phenols Flavanoids
##      <int>   <dbl>      <dbl> <dbl> <dbl> <int>   <dbl>      <dbl>
## 1      1    14.2      1.71  2.43  15.6  127    2.8      3.06
## 2      1    13.2      1.78  2.14  11.2  100    2.65     2.76
## 3      1    13.2      2.36  2.67  18.6  101    2.8      3.24
## 4      1    14.4      1.95  2.5   16.8  113    3.85     3.49
## 5      1    13.2      2.59  2.87  21    118    2.8      2.69
## 6      1    14.2      1.76  2.45  15.2  112    3.27     3.39
## # i 7 more variables: Nonflavanoid.phenols <dbl>, Proanth <dbl>,
## #      Color.int <dbl>, Hue <dbl>, OD <dbl>, Proline <int>, Alcohol_Category <fct>
```

```
wine_data %>%
  group_by(Alcohol_Category) %>%
  summarise(count = n())
```



```
## # A tibble: 3 x 2
##   Alcohol_Category count
##   <fct>             <int>
## 1 Low                22
## 2 Medium            134
## 3 High              22
```

3.7 Temel istatistiklerin hesaplanması

```
summary(wine_data)
```

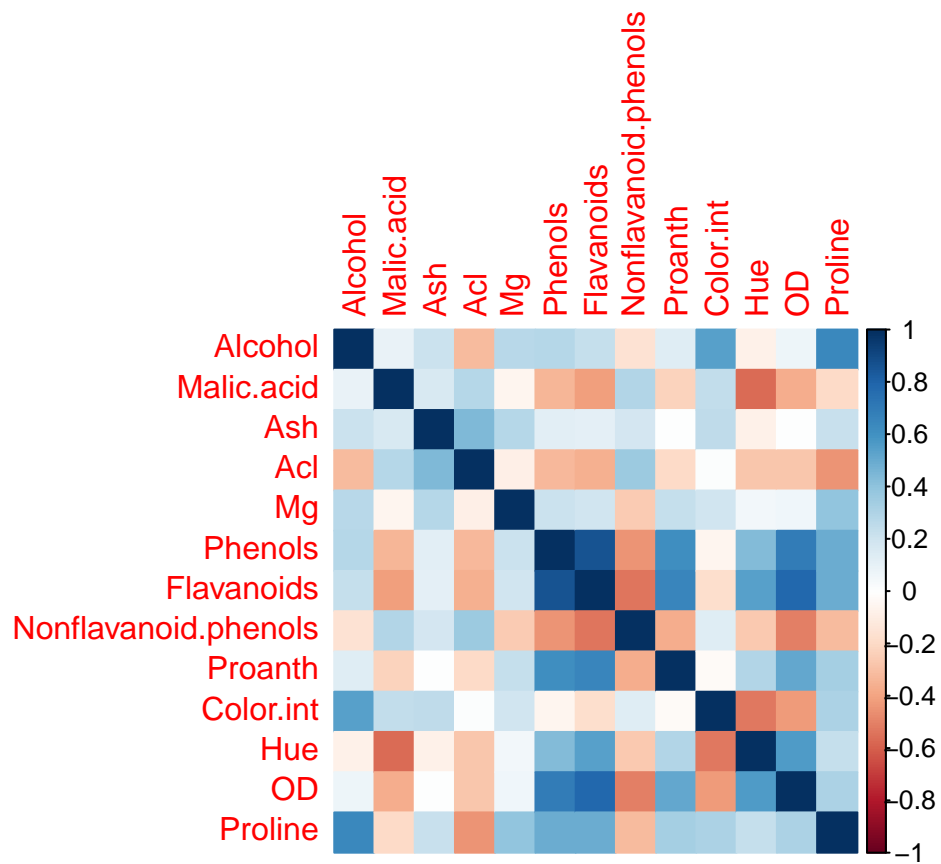
```
##      Wine      Alcohol      Malic.acid      Ash
##  Min.   :1.000   Min.   :11.03   Min.   :0.740   Min.   :1.360
## 1st Qu.:1.000   1st Qu.:12.36   1st Qu.:1.603   1st Qu.:2.210
## Median :2.000   Median :13.05   Median :1.865   Median :2.360
## Mean   :1.938   Mean   :13.00   Mean   :2.336   Mean   :2.367
## 3rd Qu.:3.000   3rd Qu.:13.68   3rd Qu.:3.083   3rd Qu.:2.558
## Max.   :3.000   Max.   :14.83   Max.   :5.800   Max.   :3.230
##      Acl      Mg      Phenols      Flavanoids
##  Min.   :10.60   Min.   : 70.00   Min.   :0.980   Min.   :0.340
## 1st Qu.:17.20   1st Qu.: 88.00   1st Qu.:1.742   1st Qu.:1.205
## Median :19.50   Median : 98.00   Median :2.355   Median :2.135
## Mean   :19.49   Mean   : 99.74   Mean   :2.295   Mean   :2.029
## 3rd Qu.:21.50   3rd Qu.:107.00   3rd Qu.:2.800   3rd Qu.:2.875
## Max.   :30.00   Max.   :162.00   Max.   :3.880   Max.   :5.080
## Nonflavanoid.phenols Proanth      Color.int      Hue
##  Min.   :0.1300   Min.   :0.410   Min.   : 1.280   Min.   :0.4800
## 1st Qu.:0.2700   1st Qu.:1.250   1st Qu.: 3.220   1st Qu.:0.7825
## Median :0.3400   Median :1.555   Median : 4.690   Median :0.9650
## Mean   :0.3619   Mean   :1.591   Mean   : 5.058   Mean   :0.9574
## 3rd Qu.:0.4375   3rd Qu.:1.950   3rd Qu.: 6.200   3rd Qu.:1.1200
## Max.   :0.6600   Max.   :3.580   Max.   :13.000   Max.   :1.7100
##      OD      Proline      Alcohol_Category
##  Min.   :1.270   Min.   : 278.0   Low   : 22
## 1st Qu.:1.938   1st Qu.: 500.5   Medium:134
## Median :2.780   Median : 673.5   High  : 22
## Mean   :2.612   Mean   : 746.9
## 3rd Qu.:3.170   3rd Qu.: 985.0
## Max.   :4.000   Max.   :1680.0
```

```
library(psych)
describe(wine_data)
```

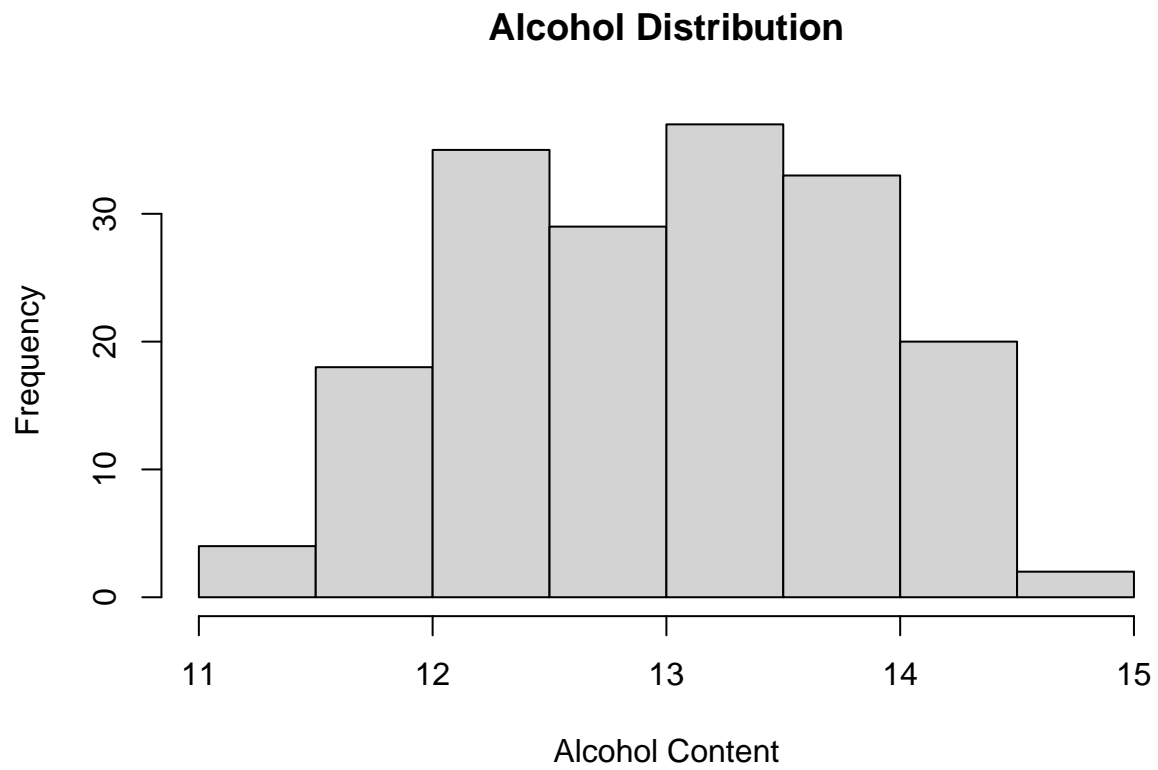
```
##      vars    n  mean    sd median trimmed    mad    min
## Wine      1 178  1.94  0.78   2.00   1.92   1.48   1.00
## Alcohol   2 178 13.00  0.81  13.05  13.01   1.01  11.03
## Malic.acid 3 178  2.34  1.12   1.87   2.21   0.77   0.74
## Ash       4 178  2.37  0.27   2.36   2.37   0.24   1.36
## Acl       5 178 19.49  3.34  19.50  19.42   3.04  10.60
## Mg       6 178 99.74 14.28  98.00  98.44  14.83  70.00
## Phenols   7 178  2.30  0.63   2.36   2.29   0.75   0.98
```

## Flavanoids	8	178	2.03	1.00	2.13	2.02	1.24	0.34
## Nonflavanoid.phenols	9	178	0.36	0.12	0.34	0.36	0.13	0.13
## Proanth	10	178	1.59	0.57	1.56	1.56	0.56	0.41
## Color.int	11	178	5.06	2.32	4.69	4.83	2.24	1.28
## Hue	12	178	0.96	0.23	0.96	0.96	0.24	0.48
## OD	13	178	2.61	0.71	2.78	2.63	0.77	1.27
## Proline	14	178	746.89	314.91	673.50	719.30	300.23	278.00
## Alcohol_Category*	15	178	2.00	0.50	2.00	2.00	0.00	1.00
##		max	range	skew	kurtosis	se		
## Wine		3.00	2.00	0.11	-1.34	0.06		
## Alcohol		14.83	3.80	-0.05	-0.89	0.06		
## Malic.acid		5.80	5.06	1.02	0.22	0.08		
## Ash		3.23	1.87	-0.17	1.03	0.02		
## Acl		30.00	19.40	0.21	0.40	0.25		
## Mg		162.00	92.00	1.08	1.96	1.07		
## Phenols		3.88	2.90	0.09	-0.87	0.05		
## Flavanoids		5.08	4.74	0.02	-0.91	0.07		
## Nonflavanoid.phenols		0.66	0.53	0.44	-0.68	0.01		
## Proanth		3.58	3.17	0.51	0.47	0.04		
## Color.int		13.00	11.72	0.85	0.30	0.17		
## Hue		1.71	1.23	0.02	-0.40	0.02		
## OD		4.00	2.73	-0.30	-1.11	0.05		
## Proline		1680.00	1402.00	0.75	-0.31	23.60		
## Alcohol_Category*		3.00	2.00	0.00	1.00	0.04		

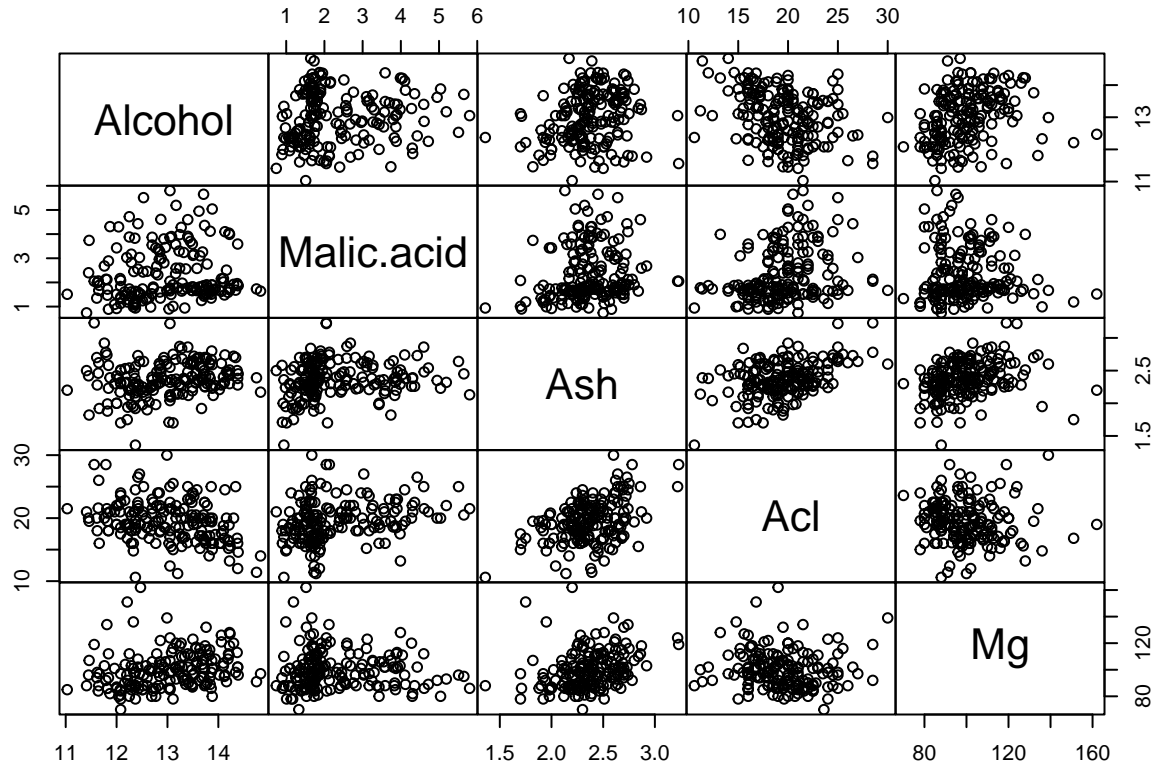
```
library(corrplot)
corrplot(correlation_matrix, method = "color")
```



```
# Histogram grafiği
hist(wine_data$Alcohol, main = "Alcohol Distribution", xlab = "Alcohol Content")
```



```
#Dağılım Grafiği  
pairs(wine_data[, 2:6], gap = 0.01)
```



4 Keşifçi ve Açıklayıcı Veri Analizi

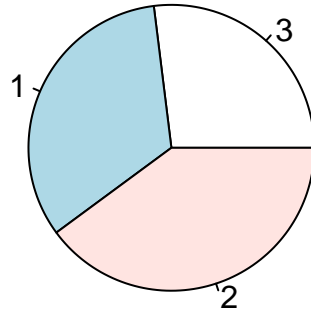
4.1 ggplot2 paketi ile uygun özelliklere ait veri görselleştirmenin gerçekleştirilmesi

```
par(wine_data, mfrow = c(1,2))
```

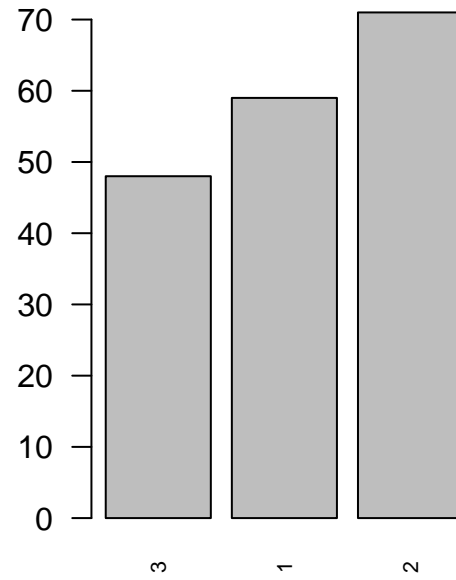
```
## Warning in par(wine_data, mfrow = c(1, 2)): argument 1 does not name a
## graphical parameter
```

```
tbl = sort(table(wine_data$Wine))
pie(tbl)
title("Wine Type Pie Chart")
barplot(tbl, las = 2, cex.names = 0.7)
title("Wine Type Bar Chart")
```

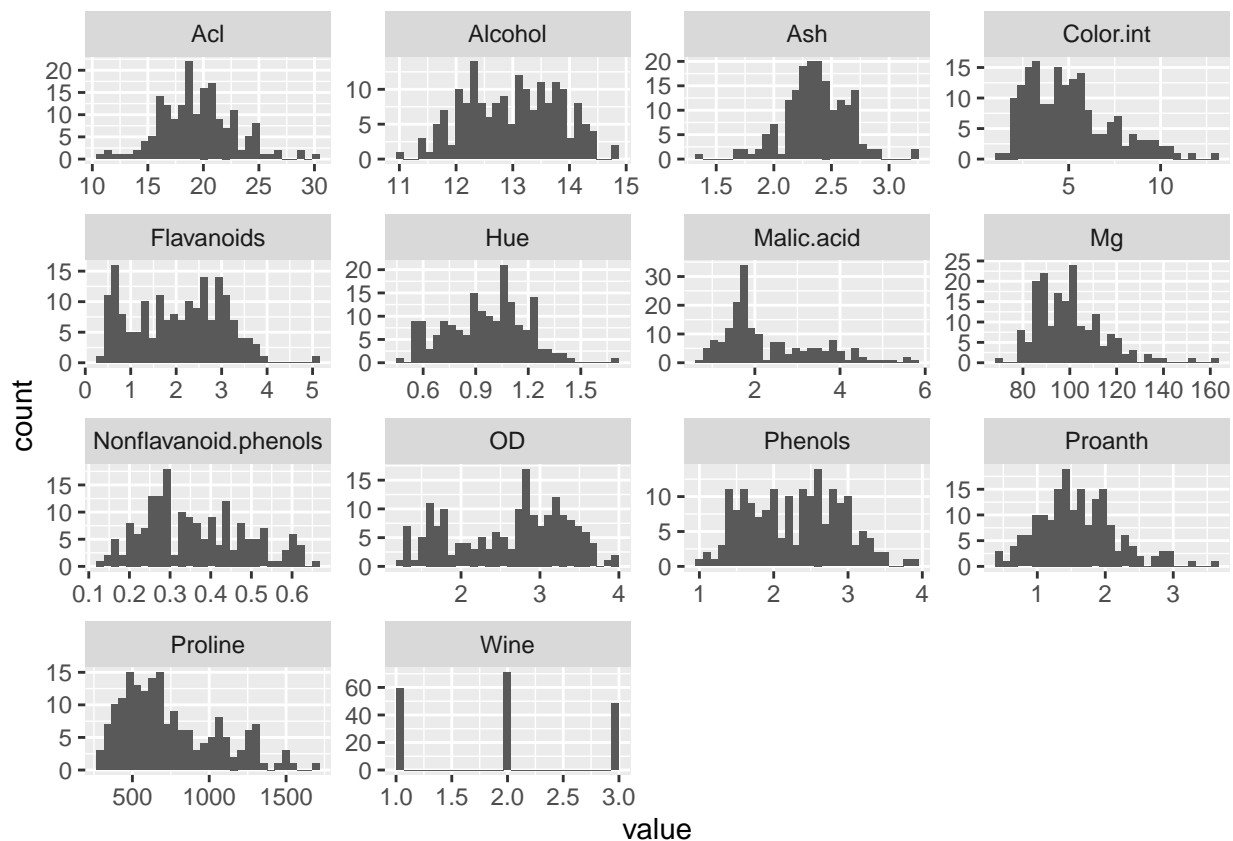
Wine Type Pie Chart



Wine Type Bar Chart



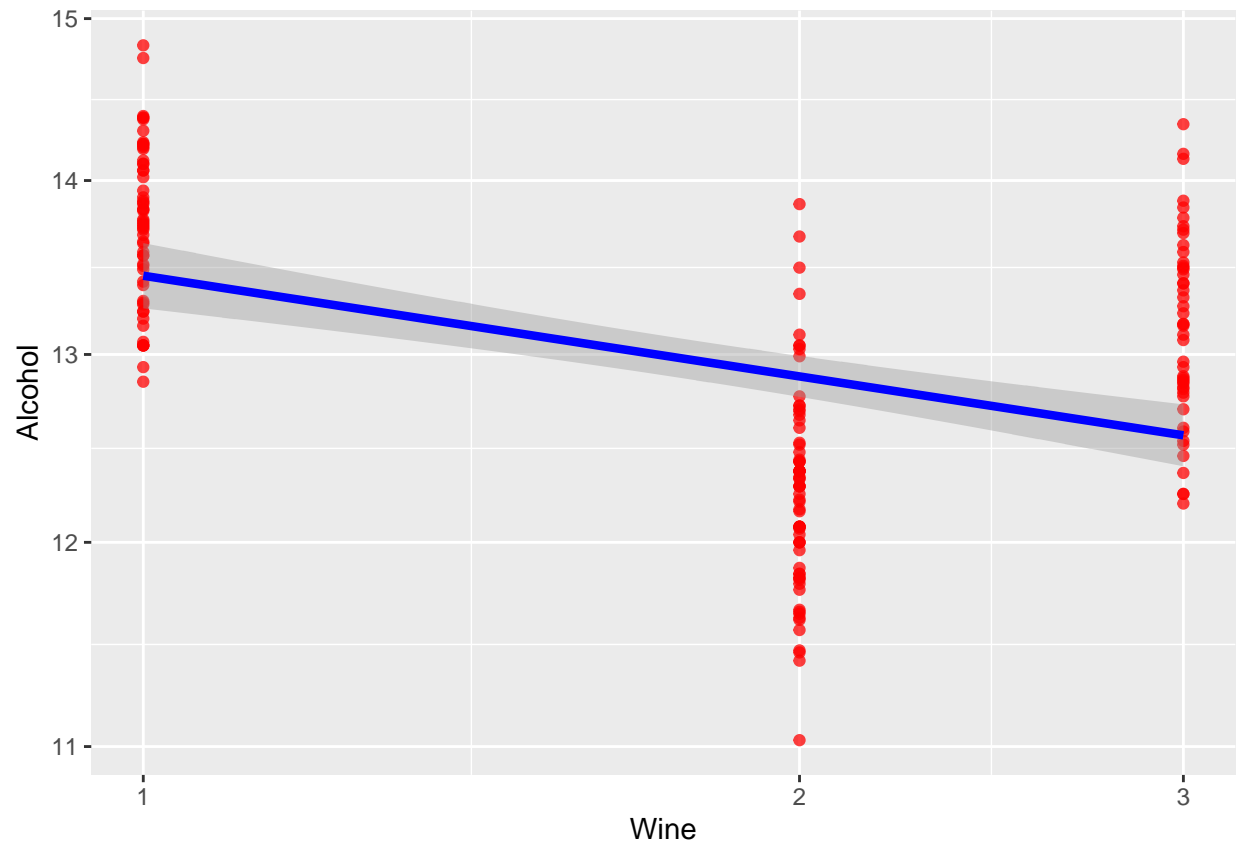
```
#Her bir sayısal değişkenin histogramı  
long_data %>% ggplot(aes(value)) +  
  facet_wrap(~ key, scales = "free") + geom_histogram(bins = 30)
```



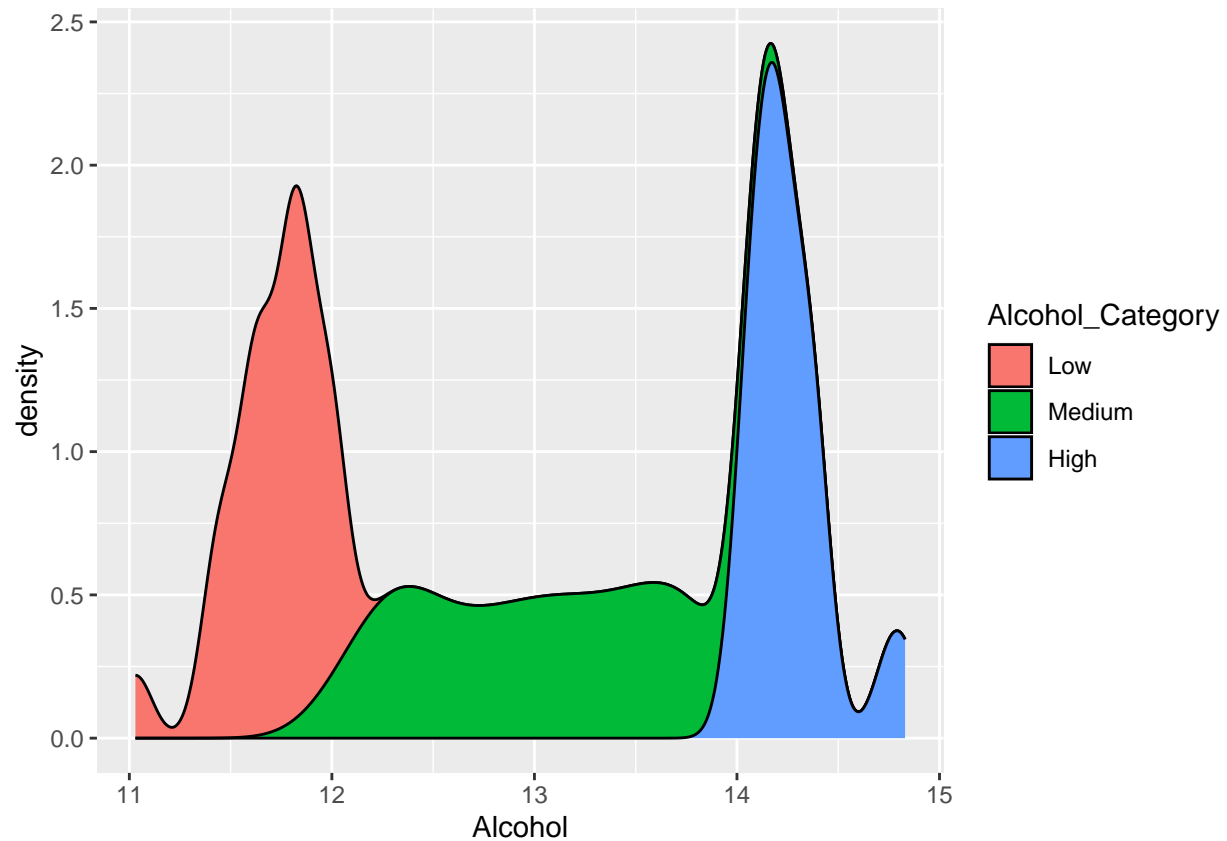
```
ggplot(wine_data, aes(x = Wine, y = Alcohol)) +
  geom_point(alpha = 0.75, col = "red") +
  scale_x_log10() +
  scale_y_log10() +
  stat_smooth(method = "lm", se = T, col = "blue", size = 1.5)
```

```
## Warning: Using 'size' aesthetic for lines was deprecated in ggplot2 3.4.0.
## i Please use 'linewidth' instead.
## This warning is displayed once every 8 hours.
## Call 'lifecycle::last_lifecycle_warnings()' to see where this warning was
## generated.
```

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



```
ggplot(data = wine_data, aes(Alcohol)) + geom_density(aes(fill = Alcohol_Category), position = "stack")
```

MACHINE LEARNING ALGORITHMS

KNN

```
library(class)
```

```
## Warning: package 'class' was built under R version 4.3.2
```

```
set.seed(123)
index = sample(1:nrow(wine_data), 0.7 * nrow(wine_data))
train_data = wine_data[index, ]
test_data = wine_data[-index, ]
```

```
k <- 3
knn_model = knn(train = train_data[, 2:ncol(normalize_data)],
                 test = test_data[, 2:ncol(normalize_data)],
                 cl = train_data$Wine,
                 k = k)
```

```
# Confusion matrix
conf_matrix = table(Actual = test_data$Wine, Predicted = knn_model)
conf_matrix
```

```
##          Predicted
## Actual  1  2  3
```

```
##      1 17  0  2
##      2  1 14  9
##      3  0  1 10
```

```
# Accuracy değeri
accuracy = sum(diag(conf_matrix)) / sum(conf_matrix)
cat("Accuracy:", accuracy, "\n")
```

```
## Accuracy: 0.7592593
```

Logistic Regression

```
set.seed(123)
index = sample(1:nrow(wine_data), 0.7 * nrow(wine_data))
train_data = wine_data[index, ]
test_data = wine_data[-index, ]
```

```
glm_model = glm(as.factor(Wine) ~ ., data = train_data, family = "binomial")
```

```
glm_predictions = predict(glm_model, test_data, type = "response")
```

```
glm_predictions = ifelse(glm_predictions > 0.5, "Class_2", "Class_1")
```

```
# Confusion matrix
conf_matrix_glm = table(Actual = test_data$Wine, Predicted = glm_predictions)
conf_matrix_glm
```

```
##      Predicted
## Actual Class_1 Class_2
##      1      19      0
##      2       0     24
##      3       0     11
```

```
# Accuracy değeri
accuracy_glm = sum(diag(conf_matrix_glm)) / sum(conf_matrix_glm)
cat("Logistic Regression Accuracy:", accuracy_glm, "\n")
```

```
## Logistic Regression Accuracy: 0.7962963
```

DECISION TREE

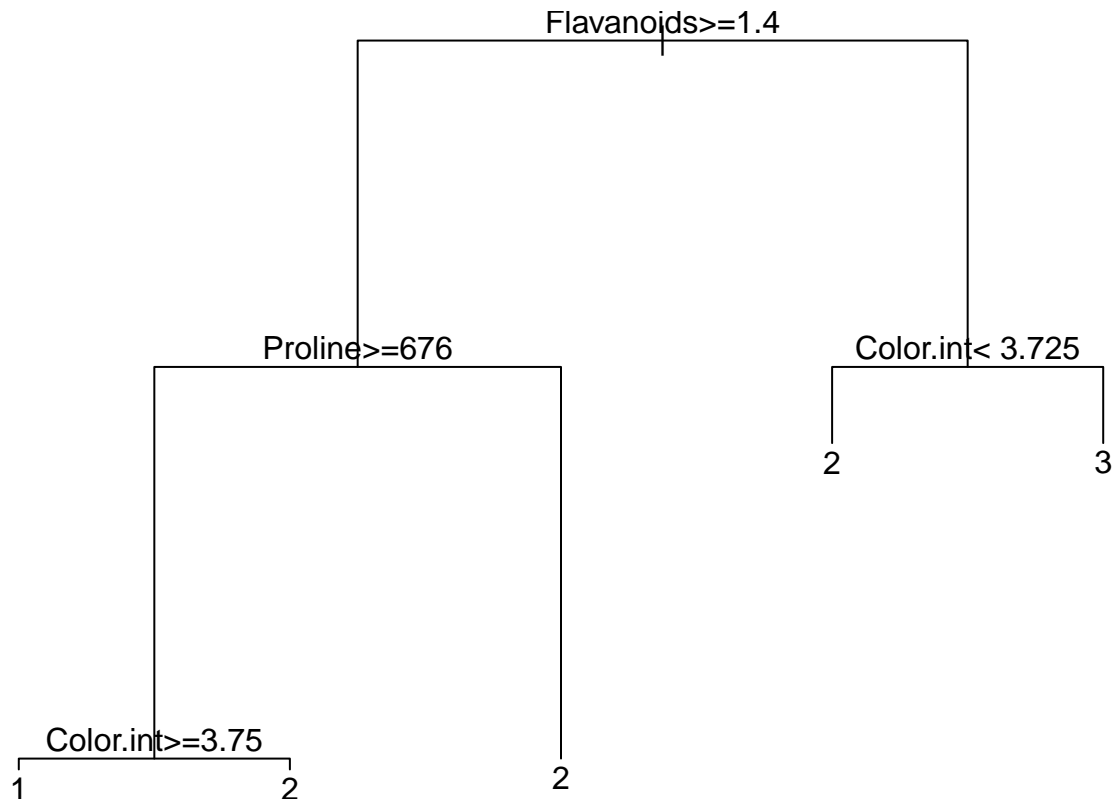
```
library(rpart)
```

```
## Warning: package 'rpart' was built under R version 4.3.2
```

```
set.seed(123)
index = sample(1:nrow(wine_data), 0.7 * nrow(wine_data))
train_data = wine_data[index, ]
test_data = wine_data[-index, ]
```

```
tree_model <- rpart(as.factor(Wine) ~ ., data = train_data, method = "class")
```

```
par(mar = c(1, 1, 1, 1))
plot(tree_model)
text(tree_model)
```



```
tree_predictions = predict(tree_model, test_data, type = "class")
```

```
# Confusion matrix
conf_matrix_tree = table(Actual = test_data$Wine, Predicted = tree_predictions)
conf_matrix_tree
```

```
##      Predicted
## Actual  1  2  3
##      1 19  0  0
##      2  2 22  0
##      3  0  1 10
```

```
# Accuracy değeri
accuracy_tree = sum(diag(conf_matrix_tree)) / sum(conf_matrix_tree)
cat("Decision Tree Accuracy:", accuracy_tree, "\n")
```

```
## Decision Tree Accuracy: 0.9444444
```

SVM

```
library(e1071)
```

```
## Warning: package 'e1071' was built under R version 4.3.2
```

```
wine_data$Wine = as.factor(wine_data$Wine)
```

```
set.seed(123)
indices = sample(1:nrow(wine_data), 0.7 * nrow(wine_data))
train_data = wine_data[indices, ]
test_data = wine_data[-indices, ]
```

```
svm_model = svm(Wine ~ ., data = train_data, kernel = "linear")
predictions = predict(svm_model, newdata = test_data)
```

```
# Confusion matrix
conf_matrix_svm = table(Actual = test_data$Wine, Predicted = predictions)
conf_matrix_svm
```

```
##      Predicted
## Actual  1  2  3
##      1 19  0  0
##      2  0 24  0
##      3  0  1 10
```

```
# Accuracy değeri
accuracy = sum(predictions == test_data$Wine) / nrow(test_data)
cat("SVM Accuracy:", accuracy, "\n")
```

```
## SVM Accuracy: 0.9814815
```