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Section: CPE22S3

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```
import numpy as np
import pandas as pd
import statistics as stats
from google.colab import drive
drive.mount('/content/drive')
df_wine = pd.read_csv('/content/drive/MyDrive/CSVs/winequality-red.csv')
```

Drive already mounted at /content/drive; to attempt to forcibly remount, call drive.m

```
#Column names of df_wine
df_wine.columns
```

```
Index(['fixed acidity', 'volatile acidity', 'citric acid', 'residual sugar',
       'chlorides', 'free sulfur dioxide', 'total sulfur dioxide', 'density',
       'pH', 'sulphates', 'alcohol', 'quality'],
      dtype='object')
```

```
#Data types of the data
df_wine.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1599 entries, 0 to 1598
Data columns (total 12 columns):
#   Column                Non-Null Count  Dtype
---  -
0   fixed acidity          1599 non-null   float64
1   volatile acidity       1599 non-null   float64
2   citric acid            1599 non-null   float64
3   residual sugar         1599 non-null   float64
4   chlorides              1599 non-null   float64
5   free sulfur dioxide    1599 non-null   float64
6   total sulfur dioxide   1599 non-null   float64
7   density                1599 non-null   float64
8   pH                    1599 non-null   float64
9   sulphates              1599 non-null   float64
10  alcohol                1599 non-null   float64
11  quality                1599 non-null   int64
dtypes: float64(11), int64(1)
memory usage: 150.0 KB
```

```
#first 10 entries  
df_wine.head(10)
```

Next steps:



[View recommended plots](#)

```
#last 10 entries  
df_wine.tail(10)
```

```
#total number of records  
print(df_wine.shape[0])
```

```
1599
```

```
#Create a new column high quality that display 'Yes' if the value of 'quality' is above 5, otherwise 'No'
df_wine['high quality'] = np.where(df_wine['quality'] > 5, 'Yes', 'No')
df_wine.head(10)
```

Next steps:

 [View recommended plots](#)

```
#Create a new Dataframe 'highQuality' that gathers data of high quality red wines
highQuality = df_wine[df_wine['high quality'] == 'Yes'].copy()
highQuality.head()
```

Next steps:

 [View recommended plots](#)

```
#Create a new Dataframe 'lowQuality' that gathers data of low quality red wines
lowQuality = df_wine[df_wine['high quality'] == 'No'].copy()
lowQuality.head()
```

Next steps:

 [View recommended plots](#)

#25th and 75th percentile of both dataframes, column total sulfur dioxide

```
print("highQuality dataframe 25th and 75th [total sulfur dioxide]: "+str(np.percentile((highQuality['total sulfur dioxide'], lowQuality['total sulfur dioxide']), 25))+" and "+str(np.percentile((highQuality['total sulfur dioxide'], lowQuality['total sulfur dioxide']), 75))")
print("lowQuality dataframe 25th and 75th [total sulfur dioxide]: "+str(np.percentile((lowQuality['total sulfur dioxide'], highQuality['total sulfur dioxide']), 25))+" and "+str(np.percentile((lowQuality['total sulfur dioxide'], highQuality['total sulfur dioxide']), 75))")
```

```
highQuality dataframe 25th and 75th [total sulfur dioxide]: 20.0      50.0
lowQuality dataframe 25th and 75th [total sulfur dioxide]: 23.75     78.0
```

#median of both dataframes, column total sulfur dioxide

```
print("Median of the column total sulfur dioxide of the dataframe highQuality: ",np.mean(highQuality['total sulfur dioxide']))
print("Median of the column total sulfur dioxide of the dataframe lowQuality: ",np.mean(lowQuality['total sulfur dioxide']))
```

```
Median of the column total sulfur dioxide of the dataframe highQuality: 39.352046783
Median of the column total sulfur dioxide of the dataframe lowQuality: 54.645161290
```

#standard deviation of both dataframes, column total sulfur dioxide

```
print("Standard deviation of the column total sulfur dioxide of the dataframe highQuality: ",np.std(highQuality['total sulfur dioxide']))
print("Standard deviation of the column total sulfur dioxide of the dataframe lowQuality: ",np.std(lowQuality['total sulfur dioxide']))
```

```
Standard deviation of the column total sulfur dioxide of the dataframe highQuality: 11.576455542
Standard deviation of the column total sulfur dioxide of the dataframe lowQuality: 12.846296117
```

#min and max of both dataframes, column total sulfur dioxide

```
print("highQuality dataframe min and max [total sulfur dioxide]: "+str(np.min(highQuality['total sulfur dioxide']))+" and "+str(np.max(highQuality['total sulfur dioxide']))+"")
print("lowQuality dataframe min and max [total sulfur dioxide]: "+str(np.min(lowQuality['total sulfur dioxide']))+" and "+str(np.max(lowQuality['total sulfur dioxide']))+"")
```

```
highQuality dataframe min and max [total sulfur dioxide]: 6.0      289.0
lowQuality dataframe min and max [total sulfur dioxide]: 6.0      155.0
```

#Characteristics of a high quality and low quality red wine using their mean(average)

```
print("\t\t\t\t\tHIGH\t\t\tLOW")
print('Average fixed acidity:      '+str(np.mean(highQuality['fixed acidity']))+"\t\t"+str(np.mean(lowQuality['fixed acidity'])))
print('Average volatile acidity:    '+str(np.mean(highQuality['volatile acidity']))+"\t"+str(np.mean(lowQuality['volatile acidity'])))
print('Average citric acid:          '+str(np.mean(highQuality['citric acid']))+"\t"+str(np.mean(lowQuality['citric acid'])))
print('Average residual sugar:        '+str(np.mean(highQuality['residual sugar']))+"\t"+str(np.mean(lowQuality['residual sugar'])))
print('Average chlorides:             '+str(np.mean(highQuality['chlorides']))+"\t"+str(np.mean(lowQuality['chlorides'])))
print('Average free sulfur dioxide:    '+str(np.mean(highQuality['free sulfur dioxide']))+"\t\t"+str(np.mean(lowQuality['free sulfur dioxide'])))
print('Average total sulfur dioxide:   '+str(np.mean(highQuality['total sulfur dioxide']))+"\t\t"+str(np.mean(lowQuality['total sulfur dioxide'])))
print('Average density:               '+str(np.mean(highQuality['density']))+"\t"+str(np.mean(lowQuality['density'])))
print('Average pH:                    '+str(np.mean(highQuality['pH']))+"\t"+str(np.mean(lowQuality['pH'])))
print('Average sulphates:             '+str(np.mean(highQuality['sulphates']))+"\t"+str(np.mean(lowQuality['sulphates'])))
print('Average alcohol:               '+str(np.mean(highQuality['alcohol']))+"\t\t"+str(np.mean(lowQuality['alcohol'])))
```

	HIGH	LOW
Average fixed acidity:	8.474035087719297	8.142204301075267
Average volatile acidity:	0.4741461988304093	0.589502688172043
Average citric acid:	0.29988304093567253	0.237755376344086
Average residual sugar:	2.5359649122807015	2.5420698924731187
Average chlorides:	0.08266081871345027	0.09298924731182795
Average free sulfur dioxide:	15.27251461988304	16.567204301075268
Average total sulfur dioxide:	39.35204678362573	54.645161290322584
Average density:	0.9964666432748538	0.997068494623656
Average pH:	3.3106432748538013	3.3116532258064515
Average sulphates:	0.6926198830409357	0.6185349462365591
Average alcohol:	10.85502923976608	9.926478494623655

Data Analysis: all columns except the total sulfur dioxide column results in a big difference between the values of the high quality and low quality wines. Based on this difference, I can conclude that the characteristic that determines a high quality red wine is a low total sulfur dioxide value.