

```
In [1]: import pandas as pd
import numpy as np
```

```
-----
-
NameError                                Traceback (most recent call las
t)
Cell In[1], line 3
      1 import pandas as pd
      2 import numpy as np
----> 3 cd

NameError: name 'cd' is not defined
```

```
In [2]: cd
```

```
C:\Users\cvr
```

```
In [6]: import pandas as pd
import numpy as np
hd=pd.read_csv('C:\coe 67b6\wine_data.csv')
```

```
hd
```

```
Out[6]:
```

|       | fixed_acidity | volatile_acidity | citric_acid | residual_sugar | chlorides | free_sulfur_dioxide |
|-------|---------------|------------------|-------------|----------------|-----------|---------------------|
| 0     | 11.6          | 0.580            | 0.66        | 2.20           | 0.074     | 10.0                |
| 1     | 10.4          | 0.610            | 0.49        | 2.10           | 0.200     | 5.0                 |
| 2     | 7.4           | 1.185            | 0.00        | 4.25           | 0.097     | 5.0                 |
| 3     | 10.4          | 0.440            | 0.42        | 1.50           | 0.145     | 34.0                |
| 4     | 8.3           | 1.020            | 0.02        | 3.40           | 0.084     | 6.0                 |
| ...   | ...           | ...              | ...         | ...            | ...       | ...                 |
| 20995 | 9.7           | 1.020            | 0.91        | 50.00          | 0.412     | 114.6               |
| 20996 | 10.2          | 0.610            | 0.88        | 53.80          | 0.250     | 62.4                |
| 20997 | 13.4          | 0.460            | 1.04        | 52.10          | 0.449     | 63.0                |
| 20998 | 6.6           | 1.030            | 1.09        | 25.30          | 0.138     | 179.8               |
| 20999 | 9.3           | 0.930            | 1.32        | 33.60          | 0.412     | 128.7               |

```
21000 rows × 7 columns
```

◀  ▶

In [7]:

```
import pandas as pd
import numpy as np
hd=pd.read_csv('C:\coe 67b6\wine_data.csv')
hd.head()
hd.tail()
```

Out[7]:

|       | fixed_acidity | volatile_acidity | citric_acid | residual_sugar | chlorides | free_sulfur_dioxide |
|-------|---------------|------------------|-------------|----------------|-----------|---------------------|
| 20995 | 9.7           | 1.02             | 0.91        | 50.0           | 0.412     | 114.6               |
| 20996 | 10.2          | 0.61             | 0.88        | 53.8           | 0.250     | 62.4                |
| 20997 | 13.4          | 0.46             | 1.04        | 52.1           | 0.449     | 63.0                |
| 20998 | 6.6           | 1.03             | 1.09        | 25.3           | 0.138     | 179.8               |
| 20999 | 9.3           | 0.93             | 1.32        | 33.6           | 0.412     | 128.7               |

In [8]:

```
import pandas as pd
import numpy as np
hd=pd.read_csv('C:\coe 67b6\wine_data.csv')
hd.shape
```

Out[8]: (21000, 12)

In [9]:

```
import pandas as pd
import numpy as np
hd=pd.read_csv('C:\coe 67b6\wine_data.csv')
hd.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 21000 entries, 0 to 20999
Data columns (total 12 columns):
#   Column                                Non-Null Count  Dtype
---  -
0   fixed_acidity                        21000 non-null  float64
1   volatile_acidity                    21000 non-null  float64
2   citric_acid                         21000 non-null  float64
3   residual_sugar                      21000 non-null  float64
4   chlorides                           21000 non-null  float64
5   free_sulfur_dioxide                 21000 non-null  float64
6   total_sulfur_dioxide                21000 non-null  float64
7   density                             21000 non-null  float64
8   pH                                  21000 non-null  float64
9   sulphates                           21000 non-null  float64
10  alcohol                             21000 non-null  float64
11  quality                             21000 non-null  int64
dtypes: float64(11), int64(1)
memory usage: 1.9 MB
```

In [10]: hd.columns

```
Out[10]: Index(['fixed_acidity', 'volatile_acidity', 'citric_acid', 'residual_suga
r',
               'chlorides', 'free_sulfur_dioxide', 'total_sulfur_dioxide', 'densit
y',
               'pH', 'sulphates', 'alcohol', 'quality'],
              dtype='object')
```

In [11]: `hd.isnull()`

Out[11]:

|       | fixed_acidity | volatile_acidity | citric_acid | residual_sugar | chlorides | free_sulfur_dioxide |
|-------|---------------|------------------|-------------|----------------|-----------|---------------------|
| 0     | False         | False            | False       | False          | False     | False               |
| 1     | False         | False            | False       | False          | False     | False               |
| 2     | False         | False            | False       | False          | False     | False               |
| 3     | False         | False            | False       | False          | False     | False               |
| 4     | False         | False            | False       | False          | False     | False               |
| ...   | ...           | ...              | ...         | ...            | ...       | ...                 |
| 20995 | False         | False            | False       | False          | False     | False               |
| 20996 | False         | False            | False       | False          | False     | False               |
| 20997 | False         | False            | False       | False          | False     | False               |
| 20998 | False         | False            | False       | False          | False     | False               |
| 20999 | False         | False            | False       | False          | False     | False               |

21000 rows × 12 columns



In [12]: `hd.isnull().sum()`

Out[12]:

|                      |   |
|----------------------|---|
| fixed_acidity        | 0 |
| volatile_acidity     | 0 |
| citric_acid          | 0 |
| residual_sugar       | 0 |
| chlorides            | 0 |
| free_sulfur_dioxide  | 0 |
| total_sulfur_dioxide | 0 |
| density              | 0 |
| pH                   | 0 |
| sulphates            | 0 |
| alcohol              | 0 |
| quality              | 0 |

dtype: int64

In [13]: `hd.iloc[0]`

Out[13]:

|                      |         |
|----------------------|---------|
| fixed_acidity        | 11.6000 |
| volatile_acidity     | 0.5800  |
| citric_acid          | 0.6600  |
| residual_sugar       | 2.2000  |
| chlorides            | 0.0740  |
| free_sulfur_dioxide  | 10.0000 |
| total_sulfur_dioxide | 47.0000 |
| density              | 1.0008  |
| pH                   | 3.2500  |
| sulphates            | 0.5700  |
| alcohol              | 9.0000  |
| quality              | 3.0000  |

Name: 0, dtype: float64

In [15]: `hd.pH.unique()`

Out[15]: array([3.25, 3.16, 3.63, 3.38, 3.48, 3.5 , 3.32, 3.31, 3.4 , 3.55, 3.02, 3.53, 3.23, 3.24, 2.89, 2.87, 3.2 , 3.04, 3.14, 2.93, 3.05, 3.42, 2.9 , 3.37, 3.44, 3.01, 3.03, 2.98, 3.08, 3.15, 3.1 , 3.58, 2.96, 2.91, 2.85, 2.99, 3.21, 3.39, 3.11, 3.34, 3.47, 2.78, 2.94, 3.19, 3.09, 3.18, 3.17, 2.95, 3.45, 3.29, 3.65, 3.22, 3.13, 3.51, 3.26, 3.56, 2.92, 3.07, 3.33, 2.86, 3.28, 3.12, 3.41, 2.97, 3. , 3.3 , 3.46, 3.27, 2.88, 2.84, 3.06, 3.35, 3.36, 3.62, 2.81, 3.6 , 3.57, 2.8 , 3.43, 3.52, 3.59, 2.83, 3.68, 2.79, 3.61, 3.54, 3.49, 2.82, 3.72, 3.9 , 3.75, 2.74, 3.66, 2.75, 3.64, 2.77, 3.71, 3.67, 3.74, 3.69, 3.77, 3.79, 3.81, 3.85, 3.7 , 3.78, 4.01, 2.72, 3.8 , 3.76, 3.82, 3.73])

In [18]: `hd.drop_duplicates()`

Out[18]:

|       | fixed_acidity | volatile_acidity | citric_acid | residual_sugar | chlorides | free_sulfur_dioxide |
|-------|---------------|------------------|-------------|----------------|-----------|---------------------|
| 0     | 11.6          | 0.580            | 0.66        | 2.20           | 0.074     | 10.0                |
| 1     | 10.4          | 0.610            | 0.49        | 2.10           | 0.200     | 5.0                 |
| 2     | 7.4           | 1.185            | 0.00        | 4.25           | 0.097     | 5.0                 |
| 3     | 10.4          | 0.440            | 0.42        | 1.50           | 0.145     | 34.0                |
| 4     | 8.3           | 1.020            | 0.02        | 3.40           | 0.084     | 6.0                 |
| ...   | ...           | ...              | ...         | ...            | ...       | ...                 |
| 20985 | 10.3          | 1.080            | 0.53        | 50.80          | 0.290     | 84.2                |
| 20987 | 11.4          | 0.900            | 0.64        | 51.20          | 0.417     | 197.1               |
| 20988 | 11.8          | 1.100            | 0.84        | 46.50          | 0.255     | 198.4               |
| 20993 | 13.0          | 0.580            | 1.22        | 52.80          | 0.247     | 93.0                |
| 20994 | 12.8          | 0.850            | 1.12        | 44.90          | 0.188     | 215.3               |

14940 rows × 12 columns

<  >

In [19]: `hd.describe()`

Out[19]:

|       | fixed_acidity | volatile_acidity | citric_acid  | residual_sugar | chlorides    | free_sulfur_dioxide |
|-------|---------------|------------------|--------------|----------------|--------------|---------------------|
| count | 21000.000000  | 21000.000000     | 21000.000000 | 21000.000000   | 21000.000000 | 21000.000000        |
| mean  | 9.797079      | 0.774796         | 0.793870     | 31.289348      | 0.200245     | 129.000000          |
| std   | 2.413919      | 0.365015         | 0.384833     | 19.015391      | 0.124933     | 77.000000           |
| min   | 3.800000      | 0.080000         | 0.000000     | 0.600000       | 0.009000     | 1.000000            |
| 25%   | 7.600000      | 0.430000         | 0.410000     | 9.800000       | 0.072000     | 45.000000           |
| 50%   | 10.000000     | 0.830000         | 0.870000     | 37.600000      | 0.205000     | 145.000000          |
| 75%   | 11.800000     | 1.080000         | 1.110000     | 46.800000      | 0.298000     | 194.000000          |
| max   | 15.900000     | 1.580000         | 1.660000     | 65.800000      | 0.611000     | 289.000000          |

<  >

```
In [22]: hd.rename(columns={'free_sulfur_dioxide':'free_so2'})
```

```
Out[22]:
```

|       | fixed_acidity | volatile_acidity | citric_acid | residual_sugar | chlorides | free_so2 | total_sulf |
|-------|---------------|------------------|-------------|----------------|-----------|----------|------------|
| 0     | 11.6          | 0.580            | 0.66        | 2.20           | 0.074     | 10.0     |            |
| 1     | 10.4          | 0.610            | 0.49        | 2.10           | 0.200     | 5.0      |            |
| 2     | 7.4           | 1.185            | 0.00        | 4.25           | 0.097     | 5.0      |            |
| 3     | 10.4          | 0.440            | 0.42        | 1.50           | 0.145     | 34.0     |            |
| 4     | 8.3           | 1.020            | 0.02        | 3.40           | 0.084     | 6.0      |            |
| ...   | ...           | ...              | ...         | ...            | ...       | ...      |            |
| 20995 | 9.7           | 1.020            | 0.91        | 50.00          | 0.412     | 114.6    |            |
| 20996 | 10.2          | 0.610            | 0.88        | 53.80          | 0.250     | 62.4     |            |
| 20997 | 13.4          | 0.460            | 1.04        | 52.10          | 0.449     | 63.0     |            |
| 20998 | 6.6           | 1.030            | 1.09        | 25.30          | 0.138     | 179.8    |            |
| 20999 | 9.3           | 0.930            | 1.32        | 33.60          | 0.412     | 128.7    |            |

21000 rows × 12 columns



```
In [23]: avg=hd['pH'].mean()  
print(avg)
```

3.1587119047619048

```
In [27]: avg=hd['pH'].median()  
print(avg)
```

3.15

```
In [29]: avg=hd['pH'].std()  
print(avg)
```

0.17137100174822836

In [30]: `hd.corr()`

Out[30]:

|                             | fixed_acidity | volatile_acidity | citric_acid | residual_sugar | chlorides | free_s |
|-----------------------------|---------------|------------------|-------------|----------------|-----------|--------|
| <b>fixed_acidity</b>        | 1.000000      | 0.562166         | 0.596388    | 0.653793       | 0.601889  |        |
| <b>volatile_acidity</b>     | 0.562166      | 1.000000         | 0.634551    | 0.730159       | 0.622915  |        |
| <b>citric_acid</b>          | 0.596388      | 0.634551         | 1.000000    | 0.765075       | 0.660209  |        |
| <b>residual_sugar</b>       | 0.653793      | 0.730159         | 0.765075    | 1.000000       | 0.747546  |        |
| <b>chlorides</b>            | 0.601889      | 0.622915         | 0.660209    | 0.747546       | 1.000000  |        |
| <b>free_sulfur_dioxide</b>  | 0.653679      | 0.746446         | 0.700625    | 0.781699       | 0.647345  |        |
| <b>total_sulfur_dioxide</b> | 0.506634      | 0.622015         | 0.636895    | 0.748704       | 0.606934  |        |
| <b>density</b>              | 0.625238      | 0.673260         | 0.740276    | 0.787859       | 0.715181  |        |
| <b>pH</b>                   | -0.254357     | -0.213949        | -0.185421   | -0.254473      | -0.113158 |        |
| <b>sulphates</b>            | 0.579008      | 0.660118         | 0.645684    | 0.679813       | 0.603356  |        |
| <b>alcohol</b>              | 0.167870      | 0.375957         | 0.358110    | 0.349538       | 0.278020  |        |
| <b>quality</b>              | 0.037545      | 0.020286         | 0.052341    | 0.049734       | 0.052905  |        |

In [35]: `hd.fillna(hd.mean())`

Out[35]:

|              | fixed_acidity | volatile_acidity | citric_acid | residual_sugar | chlorides | free_sulfur_dioxide |
|--------------|---------------|------------------|-------------|----------------|-----------|---------------------|
| <b>0</b>     | 11.6          | 0.580            | 0.66        | 2.20           | 0.074     | 10.0                |
| <b>1</b>     | 10.4          | 0.610            | 0.49        | 2.10           | 0.200     | 5.0                 |
| <b>2</b>     | 7.4           | 1.185            | 0.00        | 4.25           | 0.097     | 5.0                 |
| <b>3</b>     | 10.4          | 0.440            | 0.42        | 1.50           | 0.145     | 34.0                |
| <b>4</b>     | 8.3           | 1.020            | 0.02        | 3.40           | 0.084     | 6.0                 |
| ...          | ...           | ...              | ...         | ...            | ...       | ...                 |
| <b>20995</b> | 9.7           | 1.020            | 0.91        | 50.00          | 0.412     | 114.6               |
| <b>20996</b> | 10.2          | 0.610            | 0.88        | 53.80          | 0.250     | 62.4                |
| <b>20997</b> | 13.4          | 0.460            | 1.04        | 52.10          | 0.449     | 63.0                |
| <b>20998</b> | 6.6           | 1.030            | 1.09        | 25.30          | 0.138     | 179.8               |
| <b>20999</b> | 9.3           | 0.930            | 1.32        | 33.60          | 0.412     | 128.7               |

21000 rows × 12 columns

In [36]: `hd.index`

Out[36]: `RangeIndex(start=0, stop=21000, step=1)`

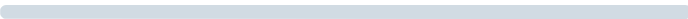
In [37]: `hd.to_numpy()`

```
Out[37]: array([[11.6 ,  0.58 ,  0.66 , ...,  0.57 ,  9.   ,  3.   ],
        [10.4 ,  0.61 ,  0.49 , ...,  0.63 ,  8.4  ,  3.   ],
        [ 7.4 ,  1.185,  0.   , ...,  0.54 , 10.7  ,  3.   ],
        ...,
        [13.4 ,  0.46 ,  1.04 , ...,  1.76 ,  9.3  ,  9.   ],
        [ 6.6 ,  1.03 ,  1.09 , ...,  1.54 , 12.9  ,  9.   ],
        [ 9.3 ,  0.93 ,  1.32 , ...,  1.42 , 13.   ,  9.   ]])
```

In [38]: `hd.iloc[0:2,:]`

```
Out[38]:
```

|   | fixed_acidity | volatile_acidity | citric_acid | residual_sugar | chlorides | free_sulfur_dioxide | total_sulfur_dioxide |
|---|---------------|------------------|-------------|----------------|-----------|---------------------|----------------------|
| 0 | 11.6          | 0.58             | 0.66        | 2.2            | 0.074     |                     | 10.0                 |
| 1 | 10.4          | 0.61             | 0.49        | 2.1            | 0.200     |                     | 5.0                  |

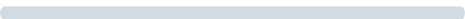
<  >

In [39]: `hd.T`

```
Out[39]:
```

|                      | 0       | 1       | 2       | 3        | 4        | 5        | 6        |        |
|----------------------|---------|---------|---------|----------|----------|----------|----------|--------|
| fixed_acidity        | 11.6000 | 10.4000 | 7.4000  | 10.40000 | 8.30000  | 7.60000  | 6.80000  | 7.300  |
| volatile_acidity     | 0.5800  | 0.6100  | 1.1850  | 0.44000  | 1.02000  | 1.58000  | 0.81500  | 0.980  |
| citric_acid          | 0.6600  | 0.4900  | 0.0000  | 0.42000  | 0.02000  | 0.00000  | 0.00000  | 0.050  |
| residual_sugar       | 2.2000  | 2.1000  | 4.2500  | 1.50000  | 3.40000  | 2.10000  | 1.20000  | 2.100  |
| chlorides            | 0.0740  | 0.2000  | 0.0970  | 0.14500  | 0.08400  | 0.13700  | 0.26700  | 0.061  |
| free_sulfur_dioxide  | 10.0000 | 5.0000  | 5.0000  | 34.00000 | 6.00000  | 5.00000  | 16.00000 | 20.000 |
| total_sulfur_dioxide | 47.0000 | 16.0000 | 14.0000 | 48.00000 | 11.00000 | 9.00000  | 29.00000 | 49.000 |
| density              | 1.0008  | 0.9994  | 0.9966  | 0.99832  | 0.99892  | 0.99476  | 0.99471  | 0.997  |
| pH                   | 3.2500  | 3.1600  | 3.6300  | 3.38000  | 3.48000  | 3.50000  | 3.32000  | 3.310  |
| sulphates            | 0.5700  | 0.6300  | 0.5400  | 0.86000  | 0.49000  | 0.40000  | 0.51000  | 0.550  |
| alcohol              | 9.0000  | 8.4000  | 10.7000 | 9.90000  | 11.00000 | 10.90000 | 9.80000  | 9.700  |
| quality              | 3.0000  | 3.0000  | 3.0000  | 3.00000  | 3.00000  | 3.00000  | 3.00000  | 3.000  |

12 rows × 21000 columns

<  >

In [40]: `hd.agg(lambda x: np.mean(x) * 5.6)`

```
Out[40]: fixed_acidity      54.863640
volatile_acidity      4.338859
citric_acid           4.445669
residual_sugar       175.220347
chlorides             1.121373
free_sulfur_dioxide   724.877067
total_sulfur_dioxide  1282.449067
density              5.655845
pH                   17.688787
sulphates            5.715592
alcohol              63.233608
quality              33.600000
dtype: float64
```

In [ ]: