

In []:

In [78]:

```
# IMPORT LIBRARIES
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
```

In [79]:

```
a=pd.read_csv(r"C:\Users\user\Downloads\11_winequality-red.csv")
a
```

Out[79]:

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	alcohol
0	7.4	0.700	0.00	1.9	0.076	11.0	34.0	0.99780	3.51	0.56	15.0
1	7.8	0.880	0.00	2.6	0.098	25.0	67.0	0.99680	3.20	0.68	15.0
2	7.8	0.760	0.04	2.3	0.092	15.0	54.0	0.99700	3.26	0.65	15.0
3	11.2	0.280	0.56	1.9	0.075	17.0	60.0	0.99800	3.16	0.58	15.0
4	7.4	0.700	0.00	1.9	0.076	11.0	34.0	0.99780	3.51	0.56	15.0
...
1594	6.2	0.600	0.08	2.0	0.090	32.0	44.0	0.99490	3.45	0.58	15.0
1595	5.9	0.550	0.10	2.2	0.062	39.0	51.0	0.99512	3.52	0.76	15.0
1596	6.3	0.510	0.13	2.3	0.076	29.0	40.0	0.99574	3.42	0.75	15.0
1597	5.9	0.645	0.12	2.0	0.075	32.0	44.0	0.99547	3.57	0.71	15.0
1598	6.0	0.310	0.47	3.6	0.067	18.0	42.0	0.99549	3.39	0.66	15.0

1599 rows × 12 columns



In [80]:

```
a=a.head(10)
a
```

Out[80]:

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	alco
0	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	0.56	
1	7.8	0.88	0.00	2.6	0.098	25.0	67.0	0.9968	3.20	0.68	
2	7.8	0.76	0.04	2.3	0.092	15.0	54.0	0.9970	3.26	0.65	
3	11.2	0.28	0.56	1.9	0.075	17.0	60.0	0.9980	3.16	0.58	
4	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	0.56	
5	7.4	0.66	0.00	1.8	0.075	13.0	40.0	0.9978	3.51	0.56	
6	7.9	0.60	0.06	1.6	0.069	15.0	59.0	0.9964	3.30	0.46	
7	7.3	0.65	0.00	1.2	0.065	15.0	21.0	0.9946	3.39	0.47	1
8	7.8	0.58	0.02	2.0	0.073	9.0	18.0	0.9968	3.36	0.57	
9	7.5	0.50	0.36	6.1	0.071	17.0	102.0	0.9978	3.35	0.80	1

In [81]:

```
# to find
a.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 10 entries, 0 to 9
Data columns (total 12 columns):
#   Column                Non-Null Count  Dtype
---  -
0   fixed acidity          10 non-null     float64
1   volatile acidity       10 non-null     float64
2   citric acid            10 non-null     float64
3   residual sugar         10 non-null     float64
4   chlorides              10 non-null     float64
5   free sulfur dioxide    10 non-null     float64
6   total sulfur dioxide   10 non-null     float64
7   density                10 non-null     float64
8   pH                    10 non-null     float64
9   sulphates              10 non-null     float64
10  alcohol                10 non-null     float64
11  quality                10 non-null     int64
dtypes: float64(11), int64(1)
memory usage: 1.1 KB
```

In [82]:

```
# to display summary of statistic  
a.describe()
```

Out[82]:

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density
count	10.000000	10.000	10.000000	10.000000	10.000000	10.000000	10.000000	10.000000
mean	7.950000	0.631	0.104000	2.330000	0.077000	14.800000	48.900000	0.997080
std	1.162612	0.161	0.194548	1.376025	0.010198	4.467164	25.066356	0.001038
min	7.300000	0.280	0.000000	1.200000	0.065000	9.000000	18.000000	0.994600
25%	7.400000	0.585	0.000000	1.825000	0.071500	11.500000	34.000000	0.996800
50%	7.650000	0.655	0.010000	1.900000	0.075000	15.000000	47.000000	0.997400
75%	7.800000	0.700	0.055000	2.225000	0.076000	16.500000	59.750000	0.997800
max	11.200000	0.880	0.560000	6.100000	0.098000	25.000000	102.000000	0.998000

In [85]:

```
# to display colum heading  
a.columns
```

Out[85]:

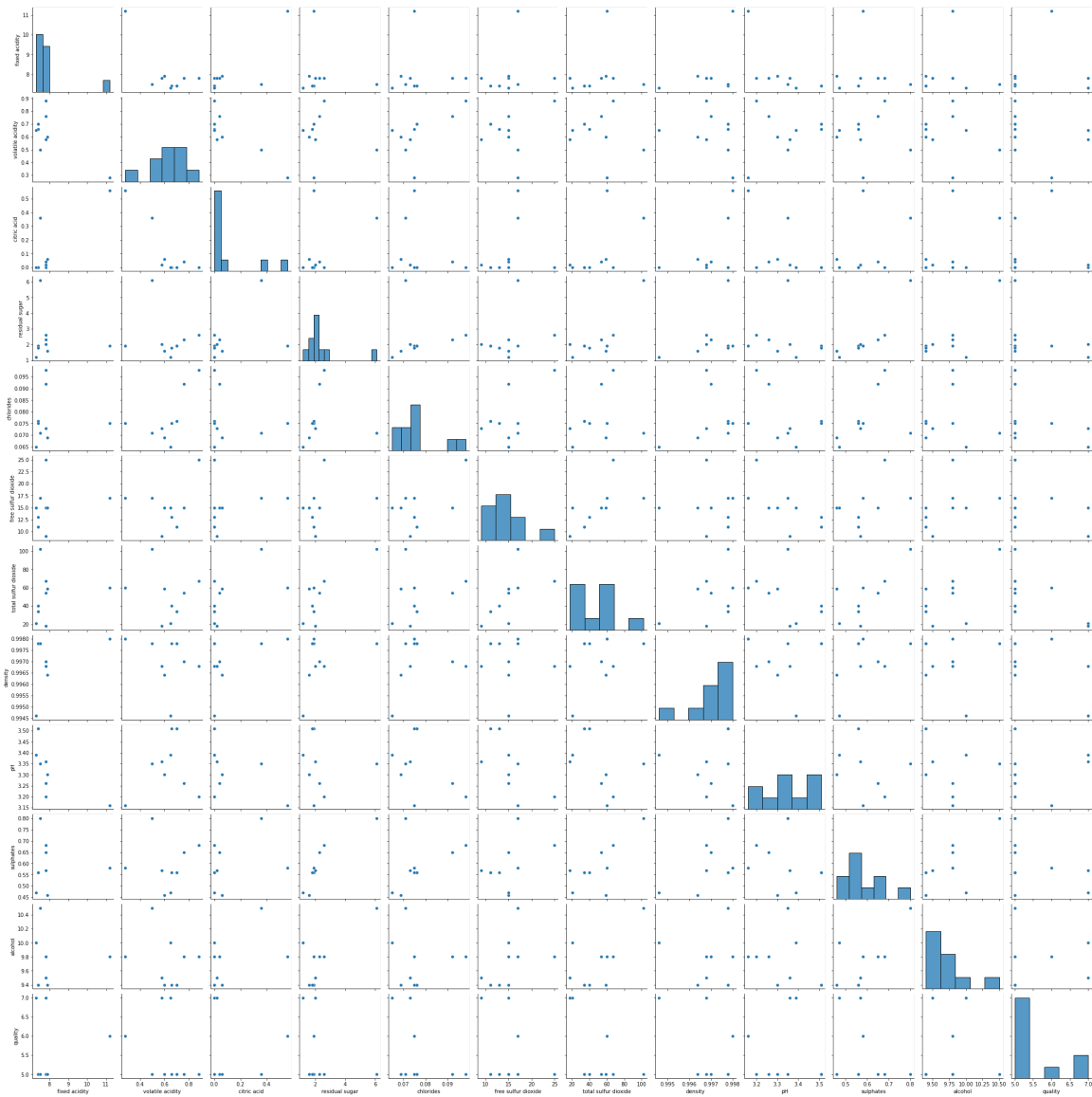
```
Index(['fixed acidity', 'volatile acidity', 'citric acid', 'residual suga  
r',  
      'chlorides', 'free sulfur dioxide', 'total sulfur dioxide', 'densit  
y',  
      'pH', 'sulphates', 'alcohol', 'quality'],  
      dtype='object')
```

In [84]:

```
sns.pairplot(a)
```

Out[84]:

<seaborn.axisgrid.PairGrid at 0x203ab501c40>

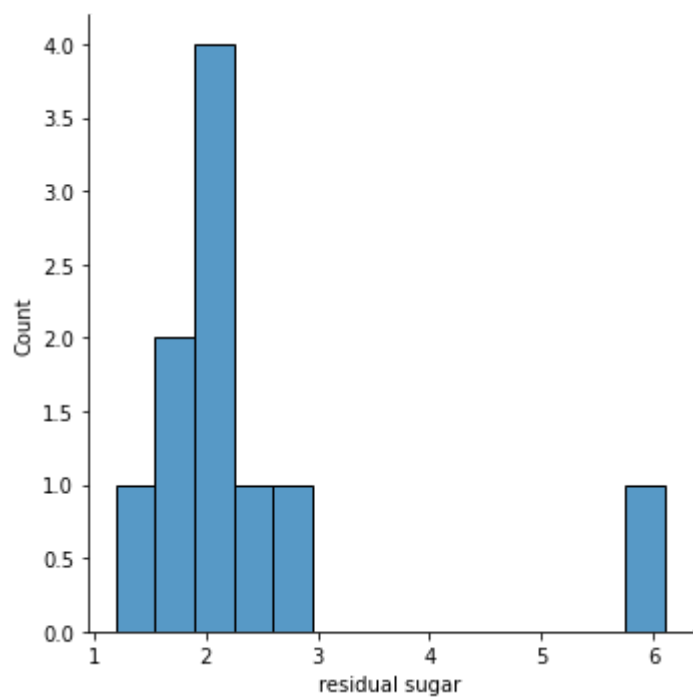


In [86]:

```
sns.displot(a["residual sugar"])
```

Out[86]:

<seaborn.axisgrid.FacetGrid at 0x203d6848ca0>



In [87]:

```
b=a[['fixed acidity', 'volatile acidity', 'citric acid', 'residual sugar',  
     'chlorides', 'free sulfur dioxide']]  
b
```

Out[87]:

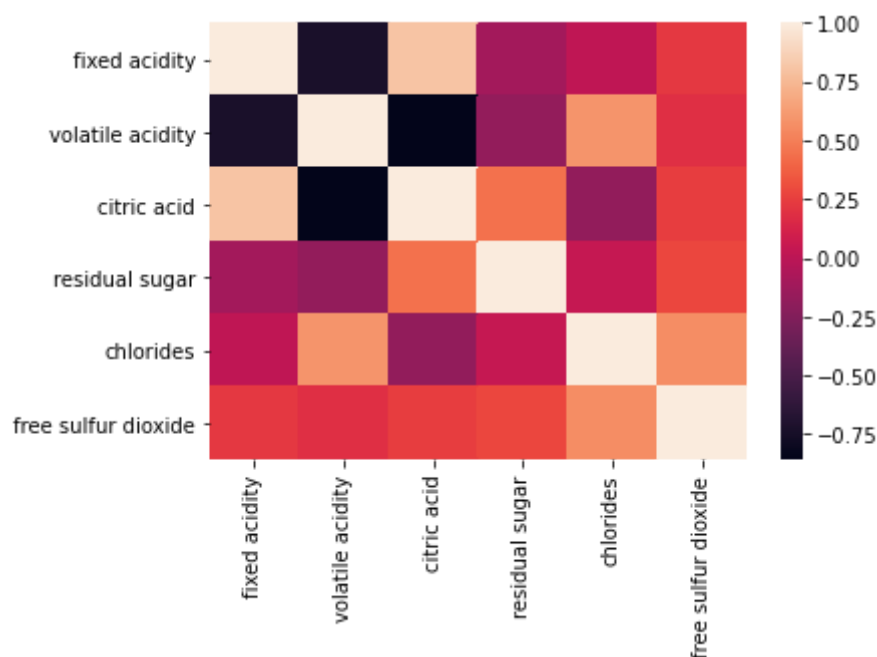
	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide
0	7.4	0.70	0.00	1.9	0.076	11.0
1	7.8	0.88	0.00	2.6	0.098	25.0
2	7.8	0.76	0.04	2.3	0.092	15.0
3	11.2	0.28	0.56	1.9	0.075	17.0
4	7.4	0.70	0.00	1.9	0.076	11.0
5	7.4	0.66	0.00	1.8	0.075	13.0
6	7.9	0.60	0.06	1.6	0.069	15.0
7	7.3	0.65	0.00	1.2	0.065	15.0
8	7.8	0.58	0.02	2.0	0.073	9.0
9	7.5	0.50	0.36	6.1	0.071	17.0

In [88]:

```
sns.heatmap(b.corr())
```

Out[88]:

<AxesSubplot:>



In [90]:

```
x=a[['fixed acidity', 'volatile acidity', 'citric acid', 'residual sugar',  
     'chlorides', 'free sulfur dioxide']]  
y=a['residual sugar']
```

In [91]:

```
from sklearn.model_selection import train_test_split  
x_train,x_test,y_train,y_test = train_test_split(x,y,test_size=0.3)
```

In [92]:

```
from sklearn.linear_model import LinearRegression  
lr=LinearRegression()  
lr.fit(x_train,y_train)
```

Out[92]:

LinearRegression()

In [93]:

```
lr.intercept_
```

Out[93]:

9.325873406851315e-15

In [94]:

```
coeff=pd.DataFrame(lr.coef_,x.columns,columns=['Co-efficient'])
coeff
```

Out[94]:

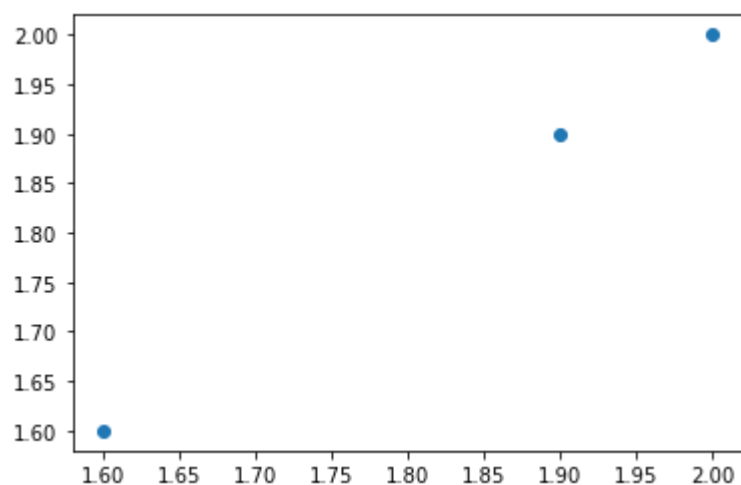
	Co-efficient
fixed acidity	-2.088683e-15
volatile acidity	1.137009e-14
citric acid	2.425779e-14
residual sugar	1.000000e+00
chlorides	2.007004e-14
free sulfur dioxide	1.509803e-17

In [95]:

```
prediction = lr.predict(x_test)
plt.scatter(y_test,prediction)
```

Out[95]:

<matplotlib.collections.PathCollection at 0x203e0bfe520>



In [96]:

```
lr.score(x_test,y_test)
```

Out[96]:

1.0

In [97]:

```
lr.score(x_train,y_train)
```

Out[97]:

1.0

In [98]:

```
from sklearn.linear_model import Ridge,Lasso
```

In [99]:

```
rr=Ridge(alpha=10)  
rr.fit(x_test,y_test)
```

Out[99]:

Ridge(alpha=10)

In [100]:

```
rr.score(x_test,y_test)
```

Out[100]:

0.8726431720424364

In [101]:

```
la=Lasso(alpha=10)  
la.fit(x_test,y_test)
```

Out[101]:

Lasso(alpha=10)

In [102]:

```
la.score(x_test,y_test)
```

Out[102]:

0.0

In []: