

```
In [1]: import numpy as np
import pandas as pd
import warnings
warnings.filterwarnings('ignore')
```

```
In [2]: red_wine_data = pd.read_csv('https://raw.githubusercontent.com/btkhimsar/DataSets')
```

```
In [3]: red_wine_data.head()
```

```
Out[3]:
```

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	alcohol
0	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	0.56	9.4
1	7.8	0.88	0.00	2.6	0.098	25.0	67.0	0.9968	3.20	0.68	9.8
2	7.8	0.76	0.04	2.3	0.092	15.0	54.0	0.9970	3.26	0.65	9.8
3	11.2	0.28	0.56	1.9	0.075	17.0	60.0	0.9980	3.16	0.58	9.8
4	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	0.56	9.4

```
In [4]: red_wine_data.shape
```

```
Out[4]: (1599, 12)
```

```
In [5]: red_wine_data.describe()
```

```
Out[5]:
```

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide
count	1599.000000	1599.000000	1599.000000	1599.000000	1599.000000	1599.000000	1599.000000
mean	8.319637	0.527821	0.270976	2.538806	0.087467	15.874922	46.467792
std	1.741096	0.179060	0.194801	1.409928	0.047065	10.460157	32.895324
min	4.600000	0.120000	0.000000	0.900000	0.012000	1.000000	6.000000
25%	7.100000	0.390000	0.090000	1.900000	0.070000	7.000000	22.000000
50%	7.900000	0.520000	0.260000	2.200000	0.079000	14.000000	38.000000
75%	9.200000	0.640000	0.420000	2.600000	0.090000	21.000000	62.000000
max	15.900000	1.580000	1.000000	15.500000	0.611000	72.000000	289.000000

```
In [6]: red_wine_data.columns
```

```
Out[6]: Index(['fixed acidity', 'volatile acidity', 'citric acid', 'residual sugar',
              'chlorides', 'free sulfur dioxide', 'total sulfur dioxide', 'density',
              'pH', 'sulphates', 'alcohol', 'quality'],
              dtype='object')
```

```
In [7]: red_wine_data['quality'].unique()
```

```
Out[7]: array([5, 6, 7, 4, 8, 3], dtype=int64)
```

```
In [8]: red_wine_data['quality'].nunique()
```

```
Out[8]: 6
```

```
In [9]: red_wine_data['quality'].value_counts()
```

```
Out[9]: 5    681
        6    638
        7    199
        4     53
        8     18
        3     10
        Name: quality, dtype: int64
```

```
In [10]: red_wine_data.rename(columns={'fixed acidity': 'fixed_acidity', 'volatile acidity':
```

```
Out[10]:
```

	fixed_acidity	volatile_acidity	citric_acid	residual_sugar	chlorides	free_sulfur_dioxide	total_sulfur_dioxide
0	7.4	0.700	0.00	1.9	0.076	11.0	
1	7.8	0.880	0.00	2.6	0.098	25.0	
2	7.8	0.760	0.04	2.3	0.092	15.0	
3	11.2	0.280	0.56	1.9	0.075	17.0	
4	7.4	0.700	0.00	1.9	0.076	11.0	
...	...	...	...	...	...	...	...
1594	6.2	0.600	0.08	2.0	0.090	32.0	
1595	5.9	0.550	0.10	2.2	0.062	39.0	
1596	6.3	0.510	0.13	2.3	0.076	29.0	
1597	5.9	0.645	0.12	2.0	0.075	32.0	
1598	6.0	0.310	0.47	3.6	0.067	18.0	

1599 rows × 12 columns

```
In [11]: red_wine_data.isna()
```

```
Out[11]:
```

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	alcol
0	False	False	False	False	False	False	False	False	False	False	Fa
1	False	False	False	False	False	False	False	False	False	False	Fa
2	False	False	False	False	False	False	False	False	False	False	Fa
3	False	False	False	False	False	False	False	False	False	False	Fa
4	False	False	False	False	False	False	False	False	False	False	Fa
...	...	...	...	...	...	...	...	...	...	...	...
1594	False	False	False	False	False	False	False	False	False	False	Fa
1595	False	False	False	False	False	False	False	False	False	False	Fa
1596	False	False	False	False	False	False	False	False	False	False	Fa
1597	False	False	False	False	False	False	False	False	False	False	Fa
1598	False	False	False	False	False	False	False	False	False	False	Fa

1599 rows × 12 columns

```
In [12]: red_wine_data.isna().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]: red_wine_data.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
```

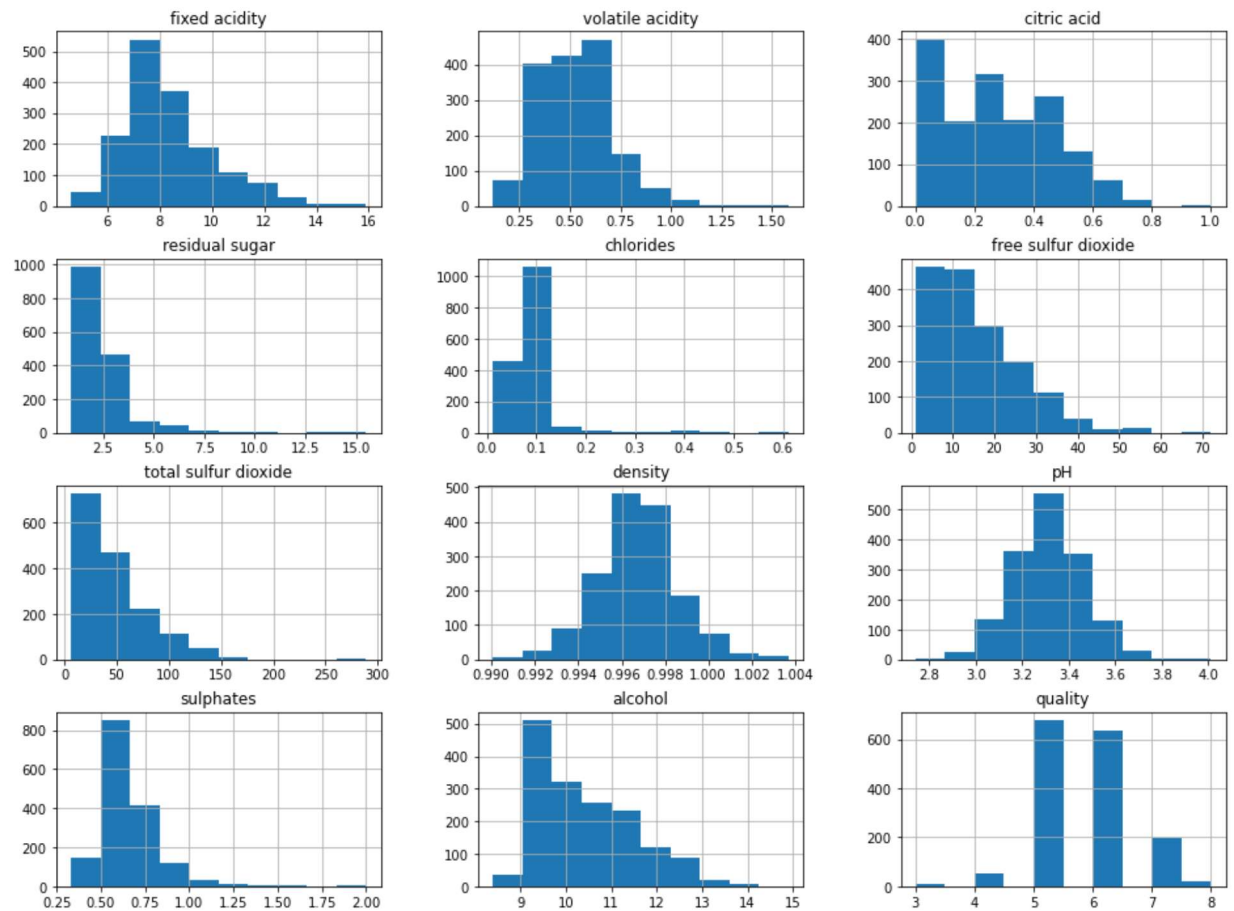
```
In [14]: duplicate=red_wine_data[red_wine_data.duplicated()]
duplicate.shape
```

```
Out[14]: (240, 12)
```

```
In [15]: import matplotlib.pyplot as plt
import seaborn as sns
```

```
In [16]: y=red_wine_data['quality']
X=red_wine_data.drop(red_wine_data['quality'],inplace=True)
```

```
In [17]: red_wine_data.hist(bins=10,figsize=(16,12))  
plt.show()
```



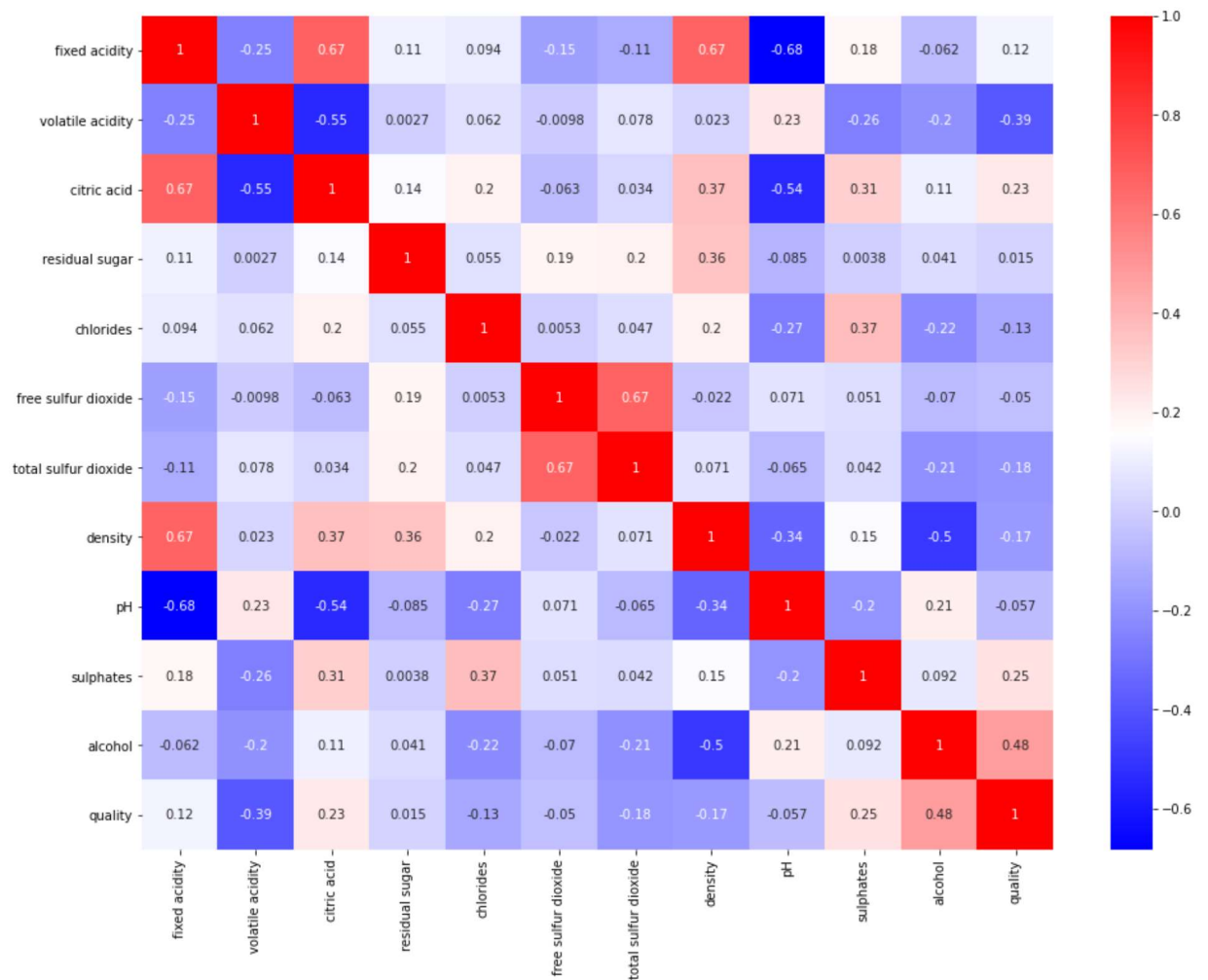
In [18]: `red_wine_data.corr()`

Out[18]:

	<b>fixed acidity</b>	<b>volatile acidity</b>	<b>citric acid</b>	<b>residual sugar</b>	<b>chlorides</b>	<b>free sulfur dioxide</b>	<b>total sulfur dioxide</b>	<b>density</b>	
<b>fixed acidity</b>	1.000000	-0.254398	0.671410	0.114616	0.093650	-0.154536	-0.114387	0.668582	-0.682678
<b>volatile acidity</b>	-0.254398	1.000000	-0.551206	0.002691	0.061790	-0.009841	0.077683	0.022580	0.233098
<b>citric acid</b>	0.671410	-0.551206	1.000000	0.142135	0.203428	-0.062613	0.033668	0.365729	-0.540720
<b>residual sugar</b>	0.114616	0.002691	0.142135	1.000000	0.054904	0.186785	0.202718	0.355577	-0.084971
<b>chlorides</b>	0.093650	0.061790	0.203428	0.054904	1.000000	0.005293	0.047110	0.200734	-0.265023
<b>free sulfur dioxide</b>	-0.154536	-0.009841	-0.062613	0.186785	0.005293	1.000000	0.667660	-0.021810	0.071316
<b>total sulfur dioxide</b>	-0.114387	0.077683	0.033668	0.202718	0.047110	0.667660	1.000000	0.070902	-0.065474
<b>density</b>	0.668582	0.022580	0.365729	0.355577	0.200734	-0.021810	0.070902	1.000000	-0.342598
<b>pH</b>	-0.682678	0.233098	-0.540720	-0.084971	-0.265023	0.071316	-0.065474	-0.342598	1.000000
<b>sulphates</b>	0.182851	-0.260545	0.311429	0.003772	0.370793	0.051157	0.042388	0.148474	-0.003772
<b>alcohol</b>	-0.062303	-0.201714	0.107572	0.040552	-0.222395	-0.070349	-0.206664	-0.496641	0.040552
<b>quality</b>	0.124330	-0.391616	0.228214	0.014504	-0.128786	-0.050443	-0.184133	-0.174029	-0.128786

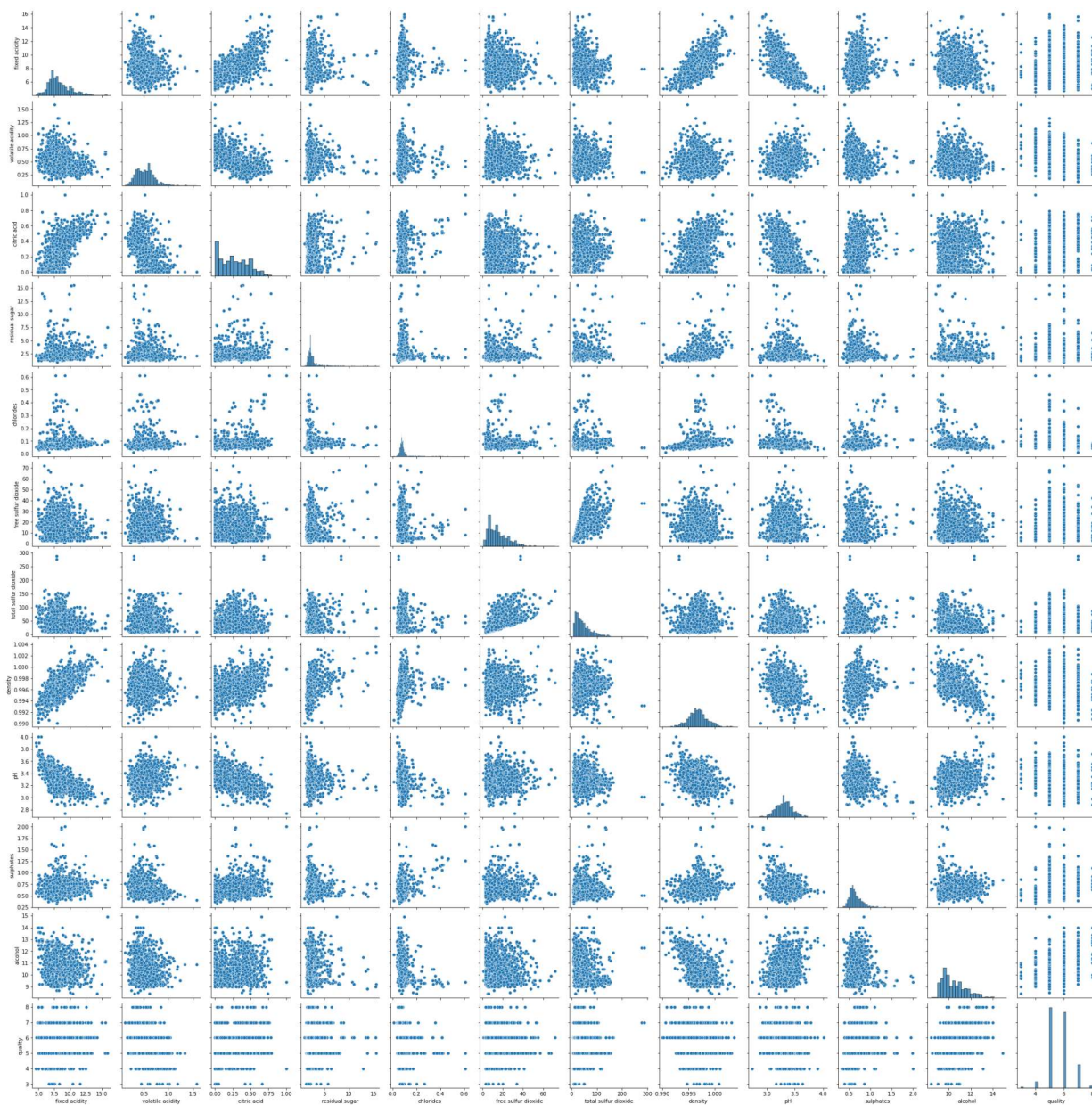
```
In [19]: plt.figure(figsize=(16,12))
sns.heatmap(red_wine_data.corr(),cmap='bwr',annot=True)
```

Out[19]: <AxesSubplot:>



```
In [20]: sns.pairplot(red_wine_data)
```

```
Out[20]: <seaborn.axisgrid.PairGrid at 0x22cb8753f70>
```



```
In [1]: #CUSTOMER CREDIT CARD FRAUD DETECTION
```

```
In [ ]:
```