

1llodbpid

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0.1 Data Preprocessing

```
[78]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
```

```
[79]: df = pd.read_csv('/content/winequality.csv')
df.head()
```

```
[79]:
```

	type	fixed acidity	volatile acidity	citric acid	residual sugar	\
0	white	7.0	0.27	0.36	20.7	
1	white	6.3	0.30	0.34	1.6	
2	white	8.1	0.28	0.40	6.9	
3	white	7.2	0.23	0.32	8.5	
4	white	7.2	0.23	0.32	8.5	

	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	\
0	0.045	45.0	170.0	1.0010	3.00	
1	0.049	14.0	132.0	0.9940	3.30	
2	0.050	30.0	97.0	0.9951	3.26	
3	0.058	47.0	186.0	0.9956	3.19	
4	0.058	47.0	186.0	0.9956	3.19	

	sulphates	alcohol	quality
0	0.45	8.8	6
1	0.49	9.5	6
2	0.44	10.1	6
3	0.40	9.9	6
4	0.40	9.9	6

```
[80]: df.shape
```

```
[80]: (6497, 13)
```

```
[81]: df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 6497 entries, 0 to 6496
Data columns (total 13 columns):
 #   Column                Non-Null Count  Dtype
---  -
 0   type                  6497 non-null   object
 1   fixed acidity         6487 non-null   float64
 2   volatile acidity      6489 non-null   float64
 3   citric acid           6494 non-null   float64
 4   residual sugar        6495 non-null   float64
 5   chlorides             6495 non-null   float64
 6   free sulfur dioxide    6497 non-null   float64
 7   total sulfur dioxide   6497 non-null   float64
 8   density               6497 non-null   float64
 9   pH                   6488 non-null   float64
10   sulphates             6493 non-null   float64
11   alcohol               6497 non-null   float64
12   quality               6497 non-null   int64
dtypes: float64(11), int64(1), object(1)
memory usage: 660.0+ KB
```

```
[82]: df.isnull().any()
```

```
[82]: type                False
fixed acidity           True
volatile acidity        True
citric acid             True
residual sugar          True
chlorides               True
free sulfur dioxide     False
total sulfur dioxide    False
density                False
pH                     True
sulphates              True
alcohol                False
quality                False
dtype: bool
```

```
[83]: df.isnull().sum()
```

```
[83]: type                0
fixed acidity          10
volatile acidity        8
citric acid            3
residual sugar         2
```

```

chlorides                2
free sulfur dioxide       0
total sulfur dioxide      0
density                  0
pH                       9
sulphates                4
alcohol                  0
quality                  0
dtype: int64

```

```
[84]: df.describe()
```

```

[84]:      fixed acidity  volatile acidity  citric acid  residual sugar  \
count      6487.000000      6489.000000   6494.000000      6495.000000
mean         7.216579         0.339691     0.318722         5.444326
std         1.296750         0.164649     0.145265         4.758125
min         3.800000         0.080000     0.000000         0.600000
25%         6.400000         0.230000     0.250000         1.800000
50%         7.000000         0.290000     0.310000         3.000000
75%         7.700000         0.400000     0.390000         8.100000
max        15.900000         1.580000     1.660000        65.800000

      chlorides  free sulfur dioxide  total sulfur dioxide      density  \
count      6495.000000      6497.000000      6497.000000      6497.000000
mean         0.056042      30.525319      115.744574         0.994697
std         0.035036      17.749400      56.521855         0.002999
min         0.009000         1.000000         6.000000         0.987110
25%         0.038000      17.000000      77.000000         0.992340
50%         0.047000      29.000000      118.000000         0.994890
75%         0.065000      41.000000      156.000000         0.996990
max         0.611000      289.000000      440.000000         1.038980

      pH  sulphates  alcohol  quality
count      6488.000000      6493.000000   6497.000000   6497.000000
mean         3.218395         0.531215     10.491801         5.818378
std         0.160748         0.148814         1.192712         0.873255
min         2.720000         0.220000         8.000000         3.000000
25%         3.110000         0.430000         9.500000         5.000000
50%         3.210000         0.510000        10.300000         6.000000
75%         3.320000         0.600000        11.300000         6.000000
max         4.010000         2.000000        14.900000         9.000000

```

```
[8]: df.quality.nunique()
```

```
[8]: 7
```

```
[9]: df.quality.unique()
```

```
[9]: array([6, 5, 7, 8, 4, 3, 9])
```

```
[10]: df.quality.value_counts()
```

```
[10]: 6    2836
      5    2138
      7    1079
      4     216
      8     193
      3      30
      9       5
      Name: quality, dtype: int64
```

```
[11]: df.alcohol.value_counts()
```

```
[11]: 9.500000    367
      9.400000    332
      9.200000    271
      10.000000   229
      10.500000   227
      ...
      10.533333     1
      11.366667     1
      12.333333     1
      14.050000     1
      10.750000     1
      Name: alcohol, Length: 111, dtype: int64
```

```
[12]: df.corr()
```

<ipython-input-12-2f6f6606aa2c>:1: FutureWarning: The default value of numeric_only in DataFrame.corr is deprecated. In a future version, it will default to False. Select only valid columns or specify the value of numeric_only to silence this warning.

```
df.corr()
```

```
[12]:
```

	fixed acidity	volatile acidity	citric acid	\
fixed acidity	1.000000	0.220172	0.323736	
volatile acidity	0.220172	1.000000	-0.378061	
citric acid	0.323736	-0.378061	1.000000	
residual sugar	-0.112319	-0.196702	0.142486	
chlorides	0.298421	0.377167	0.039315	
free sulfur dioxide	-0.283317	-0.353230	0.133437	
total sulfur dioxide	-0.329747	-0.414928	0.195218	
density	0.459204	0.271193	0.096320	
pH	-0.251814	0.260660	-0.328689	
sulphates	0.300380	0.225476	0.057613	

alcohol	-0.095603	-0.038248	-0.010433
quality	-0.077031	-0.265953	0.085706

	residual sugar	chlorides	free sulfur dioxide	\
fixed acidity	-0.112319	0.298421	-0.283317	
volatile acidity	-0.196702	0.377167	-0.353230	
citric acid	0.142486	0.039315	0.133437	
residual sugar	1.000000	-0.128902	0.403439	
chlorides	-0.128902	1.000000	-0.195042	
free sulfur dioxide	0.403439	-0.195042	1.000000	
total sulfur dioxide	0.495820	-0.279580	0.720934	
density	0.552498	0.362594	0.025717	
pH	-0.267050	0.044806	-0.145191	
sulphates	-0.185745	0.395332	-0.188489	
alcohol	-0.359706	-0.256861	-0.179838	
quality	-0.036825	-0.200886	0.055463	

	total sulfur dioxide	density	pH	sulphates	\
fixed acidity	-0.329747	0.459204	-0.251814	0.300380	
volatile acidity	-0.414928	0.271193	0.260660	0.225476	
citric acid	0.195218	0.096320	-0.328689	0.057613	
residual sugar	0.495820	0.552498	-0.267050	-0.185745	
chlorides	-0.279580	0.362594	0.044806	0.395332	
free sulfur dioxide	0.720934	0.025717	-0.145191	-0.188489	
total sulfur dioxide	1.000000	0.032395	-0.237687	-0.275381	
density	0.032395	1.000000	0.011920	0.259454	
pH	-0.237687	0.011920	1.000000	0.191248	
sulphates	-0.275381	0.259454	0.191248	1.000000	
alcohol	-0.265740	-0.686745	0.121002	-0.003261	
quality	-0.041385	-0.305858	0.019366	0.038729	

	alcohol	quality
fixed acidity	-0.095603	-0.077031
volatile acidity	-0.038248	-0.265953
citric acid	-0.010433	0.085706
residual sugar	-0.359706	-0.036825
chlorides	-0.256861	-0.200886
free sulfur dioxide	-0.179838	0.055463
total sulfur dioxide	-0.265740	-0.041385
density	-0.686745	-0.305858
pH	0.121002	0.019366
sulphates	-0.003261	0.038729
alcohol	1.000000	0.444319
quality	0.444319	1.000000

```
[13]: df.corr().quality.sort_values(ascending =False)
```

```
<ipython-input-13-faa9332a27f4>:1: FutureWarning: The default value of
numeric_only in DataFrame.corr is deprecated. In a future version, it will
default to False. Select only valid columns or specify the value of numeric_only
to silence this warning.
```

```
df.corr().quality.sort_values(ascending =False)
```

```
[13]: quality          1.000000
      alcohol         0.444319
      citric acid      0.085706
      free sulfur dioxide 0.055463
      sulphates        0.038729
      pH              0.019366
      residual sugar   -0.036825
      total sulfur dioxide -0.041385
      fixed acidity    -0.077031
      chlorides        -0.200886
      volatile acidity  -0.265953
      density          -0.305858
      Name: quality, dtype: float64
```

0.2 Visualization

0.2.1 Univariate Analysis

```
[14]: sns.distplot(df.density)
```

```
<ipython-input-14-d7414a6caaf8>:1: UserWarning:
```

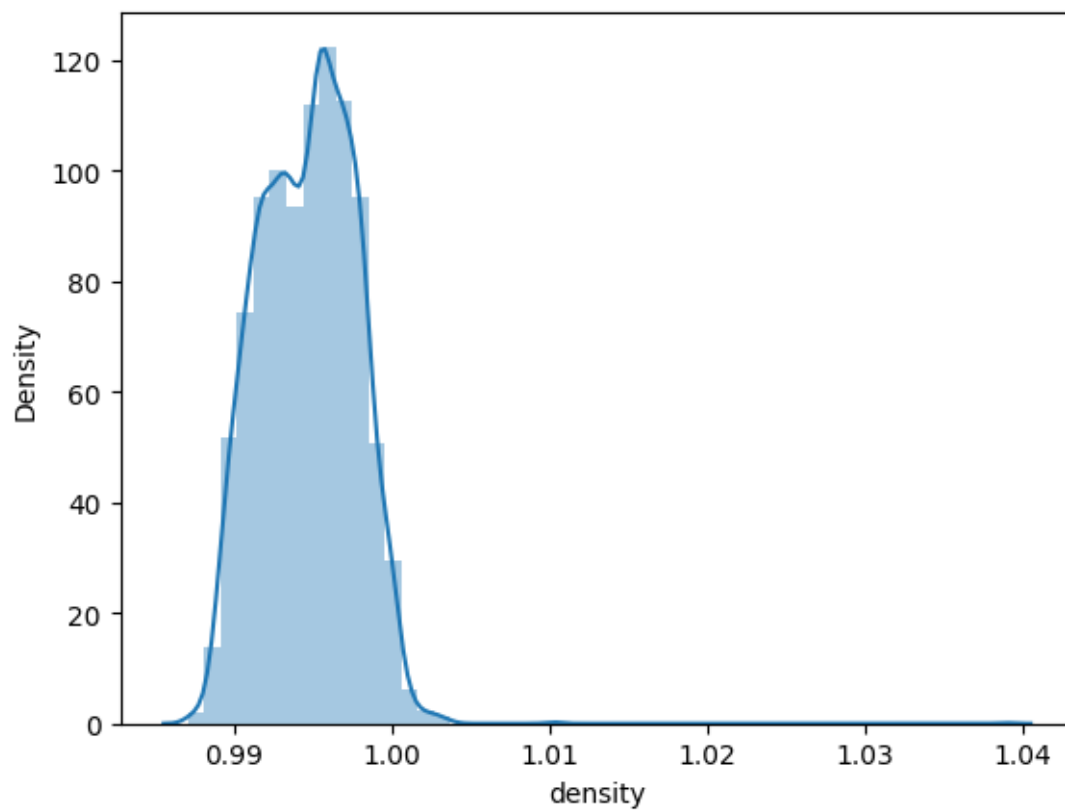
```
`distplot` is a deprecated function and will be removed in seaborn v0.14.0.
```

Please adapt your code to use either `displot` (a figure-level function with similar flexibility) or `histplot` (an axes-level function for histograms).

For a guide to updating your code to use the new functions, please see <https://gist.github.com/mwaskom/de44147ed2974457ad6372750bbe5751>

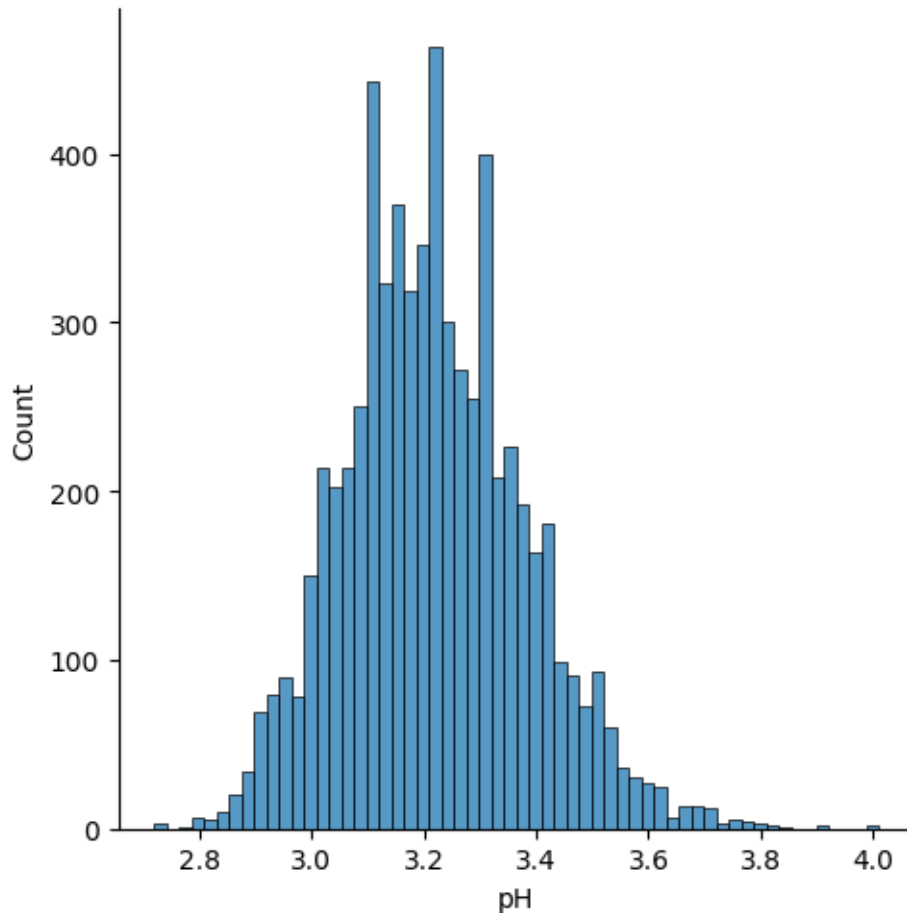
```
sns.distplot(df.density)
```

```
[14]: <Axes: xlabel='density', ylabel='Density'>
```



```
[15]: sns.displot(df.pH)
```

```
[15]: <seaborn.axisgrid.FacetGrid at 0x7ece8e5e7c40>
```



```
[68]: plt.pie(df.quality.value_counts(), [0,0,0,0.1,0.1,0.1], labels = _
↳ [5,6,7,4,8,3], autopct = '%1.1f%%', type = quality )
plt.title('QUALITY')
plt.show()
```

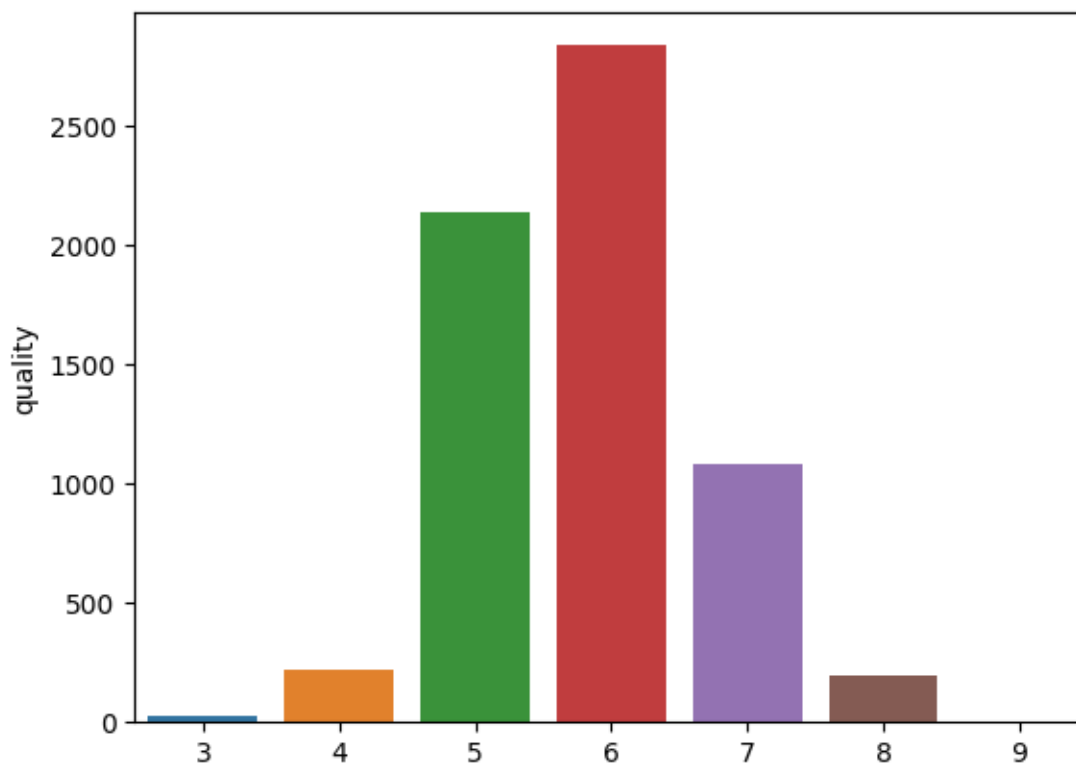
```
-----
NameError                                Traceback (most recent call last)
<ipython-input-68-c1163dbbc575> in <cell line: 1>()
----> 1 plt.pie(df.quality.value_counts(), [0,0,0,0.1,0.1,0.1], labels = _
↳ [5,6,7,4,8,3], autopct = '%1.1f%%', type = quality )
      2 plt.title('QUALITY')
      3 plt.show()

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

```
[17]: sns.barplot(x =df.quality.value_counts().index,y =df.quality.value_counts() )
```



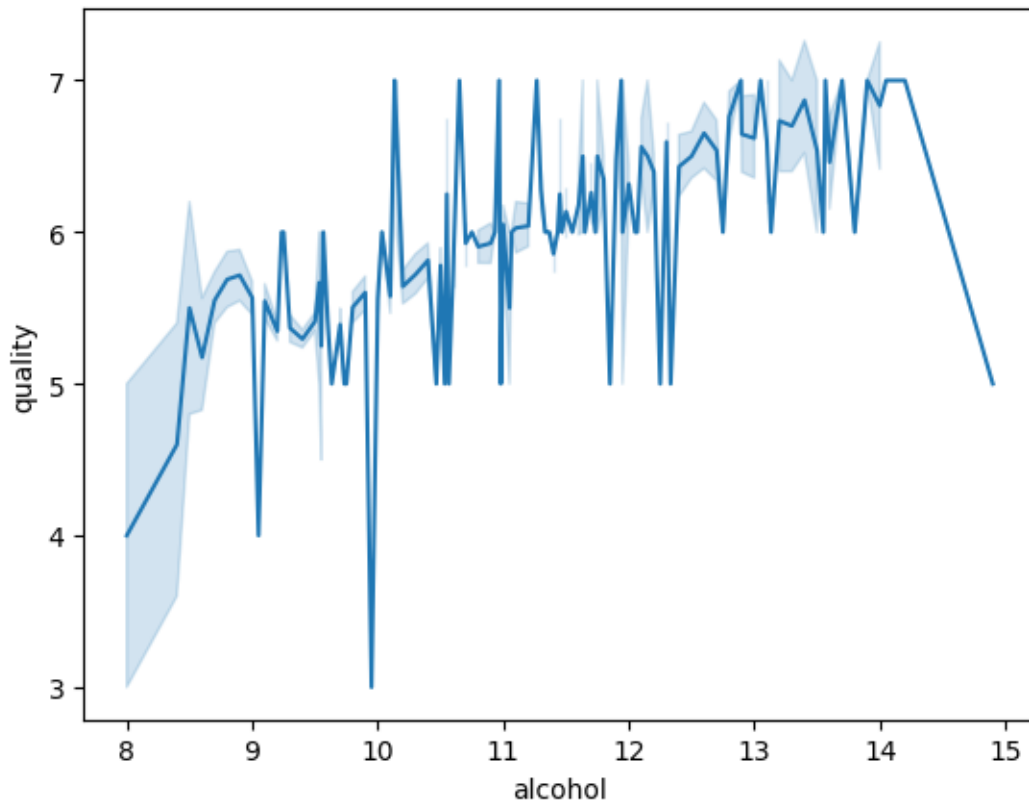
```
[17]: <Axes: ylabel='quality'>
```



0.2.2 Bivariate Analysis

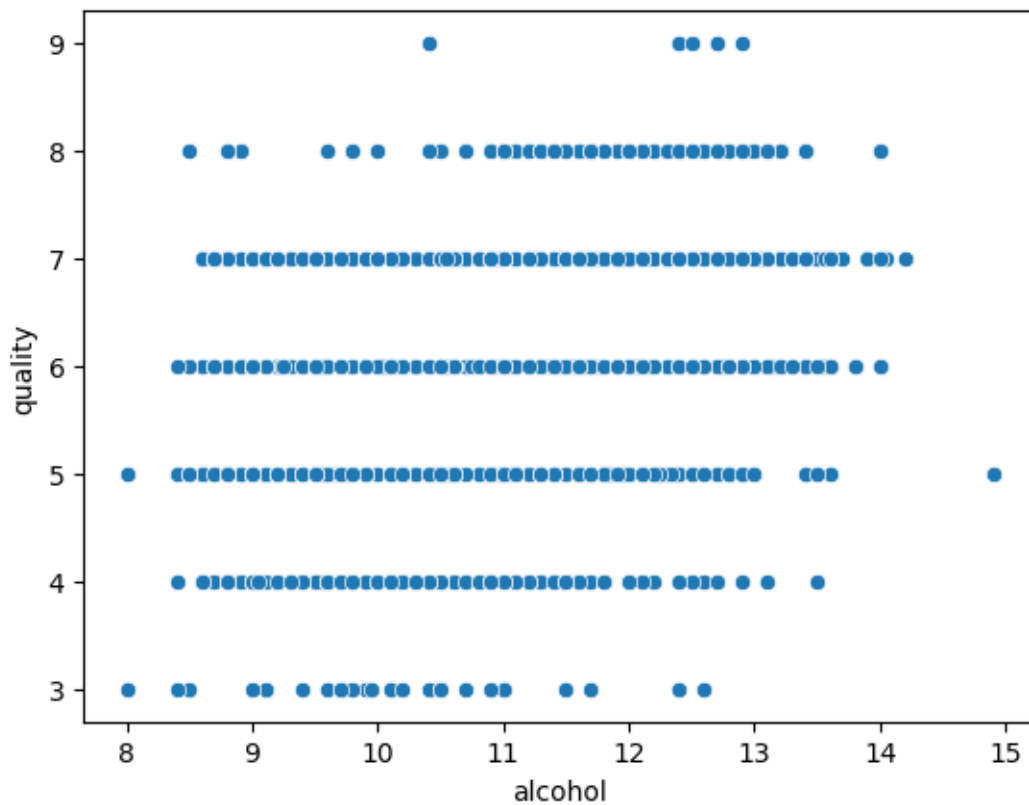
```
[18]: sns.lineplot(x = df.alcohol,y=df.quality)
```

```
[18]: <Axes: xlabel='alcohol', ylabel='quality'>
```



```
[19]: sns.scatterplot(x = df.alcohol,y=df.quality)
```

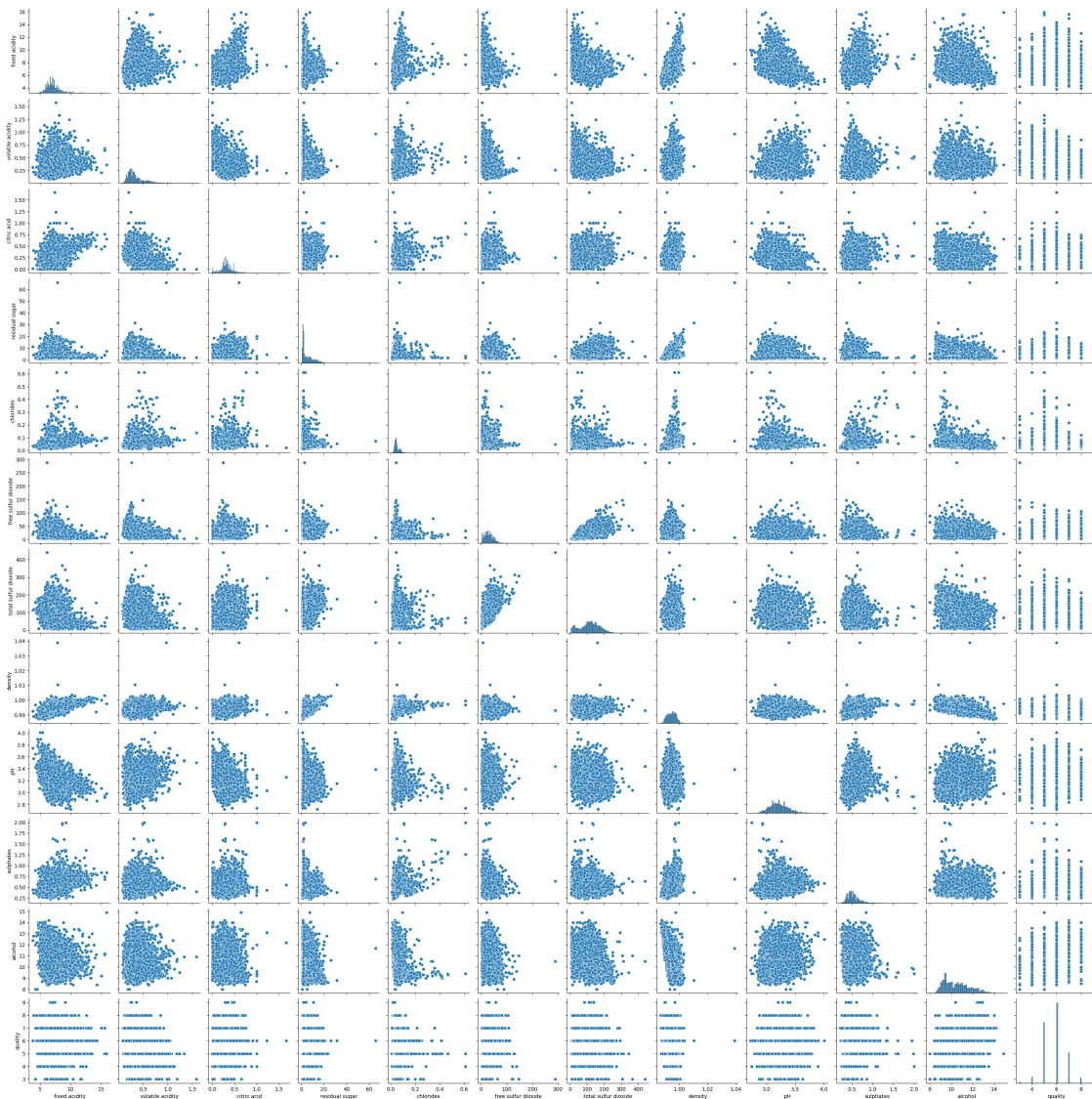
```
[19]: <Axes: xlabel='alcohol', ylabel='quality'>
```



0.2.3 Multivariate Analysis

```
[20]: sns.pairplot(df)
```

```
[20]: <seaborn.axisgrid.PairGrid at 0x7ece8e68d0c0>
```

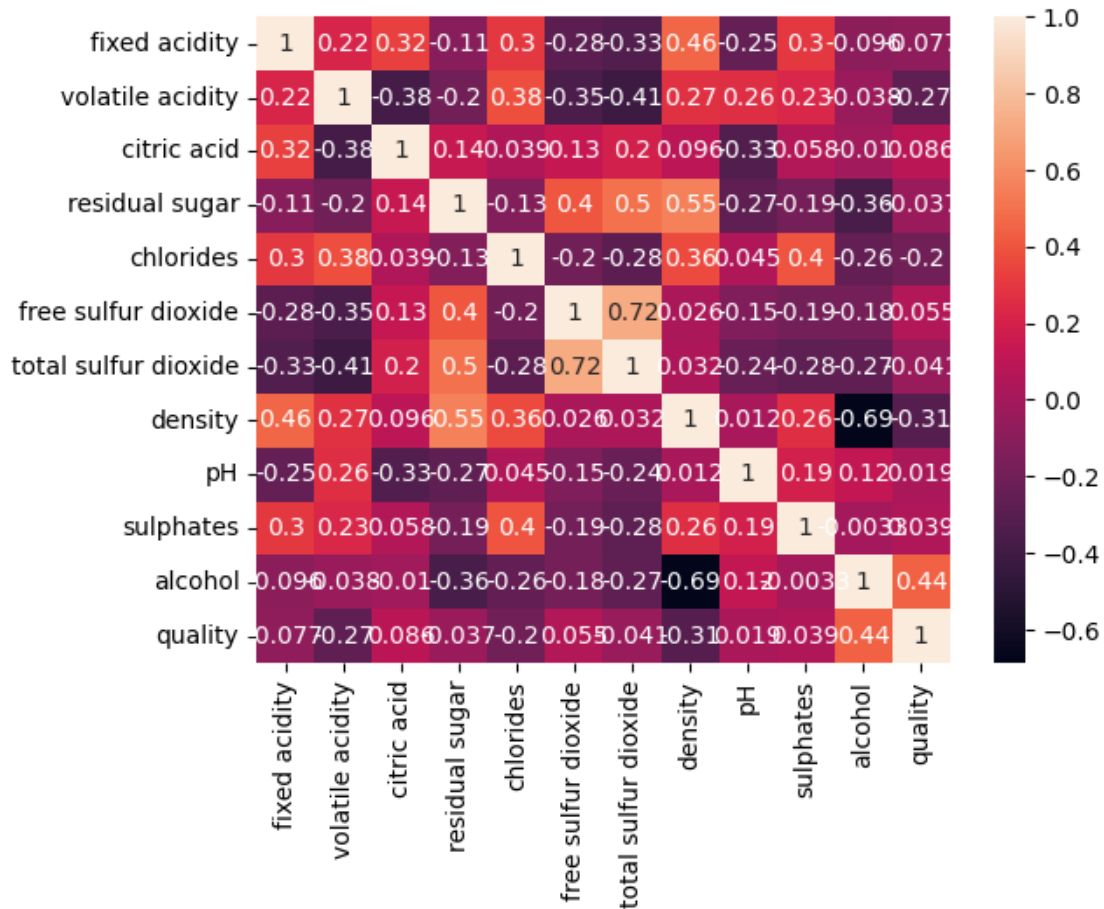


```
[21]: sns.heatmap(df.corr(),annot=True)
```

<ipython-input-21-8df7bcac526d>:1: FutureWarning: The default value of numeric_only in DataFrame.corr is deprecated. In a future version, it will default to False. Select only valid columns or specify the value of numeric_only to silence this warning.

```
sns.heatmap(df.corr(),annot=True)
```

```
[21]: <Axes: >
```



0.2.4 Outlier detection

```
[22]: df.head()
```

```
[22]:
```

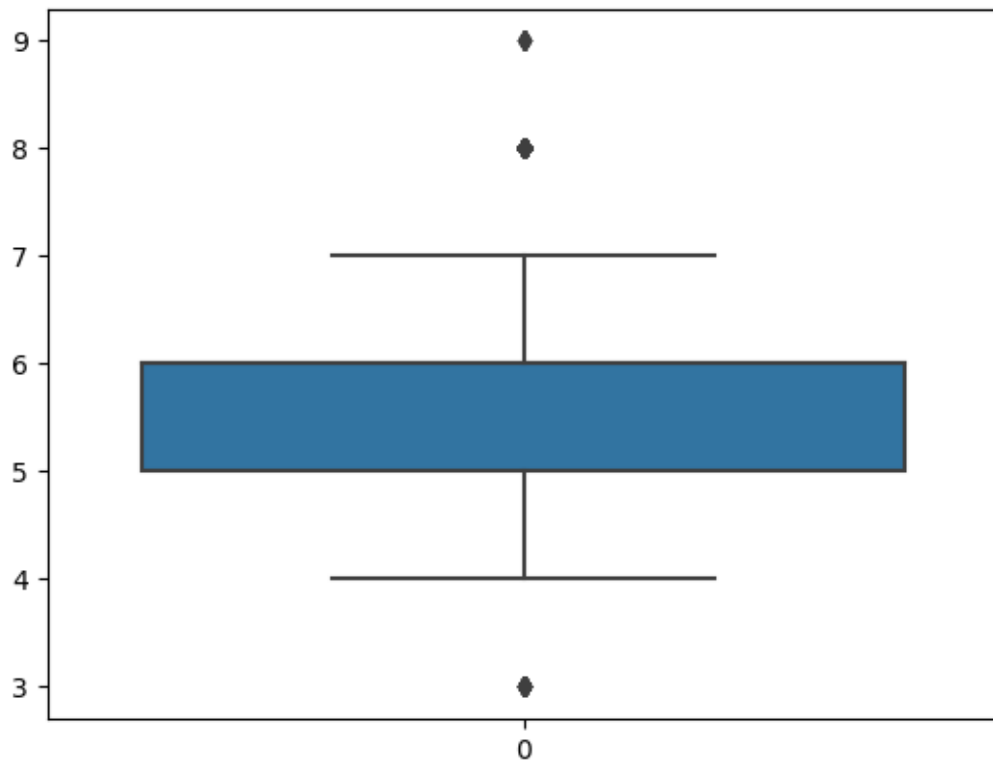
	type	fixed acidity	volatile acidity	citric acid	residual sugar	\
0	white	7.0	0.27	0.36	20.7	
1	white	6.3	0.30	0.34	1.6	
2	white	8.1	0.28	0.40	6.9	
3	white	7.2	0.23	0.32	8.5	
4	white	7.2	0.23	0.32	8.5	

	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	\
0	0.045	45.0	170.0	1.0010	3.00	
1	0.049	14.0	132.0	0.9940	3.30	
2	0.050	30.0	97.0	0.9951	3.26	
3	0.058	47.0	186.0	0.9956	3.19	
4	0.058	47.0	186.0	0.9956	3.19	

	sulphates	alcohol	quality
0	0.45	8.8	6
1	0.49	9.5	6
2	0.44	10.1	6
3	0.40	9.9	6
4	0.40	9.9	6

```
[23]: sns.boxplot(df.quality)
```

```
[23]: <Axes: >
```



0.2.5 Outlier removal using IQR

```
[24]: q1 = df.quality.quantile(0.25)
      q3 = df.quality.quantile(0.75)
```

```
[25]: IQR=q3-q1
```

```
[26]: upper_limit = q3+1.5*IQR
      upper_limit
```

```
[26]: 7.5
```

```
[27]: df.median()
```

```
<ipython-input-27-6d467abf240d>:1: FutureWarning: The default value of  
numeric_only in DataFrame.median is deprecated. In a future version, it will  
default to False. In addition, specifying 'numeric_only=None' is deprecated.  
Select only valid columns or specify the value of numeric_only to silence this  
warning.
```

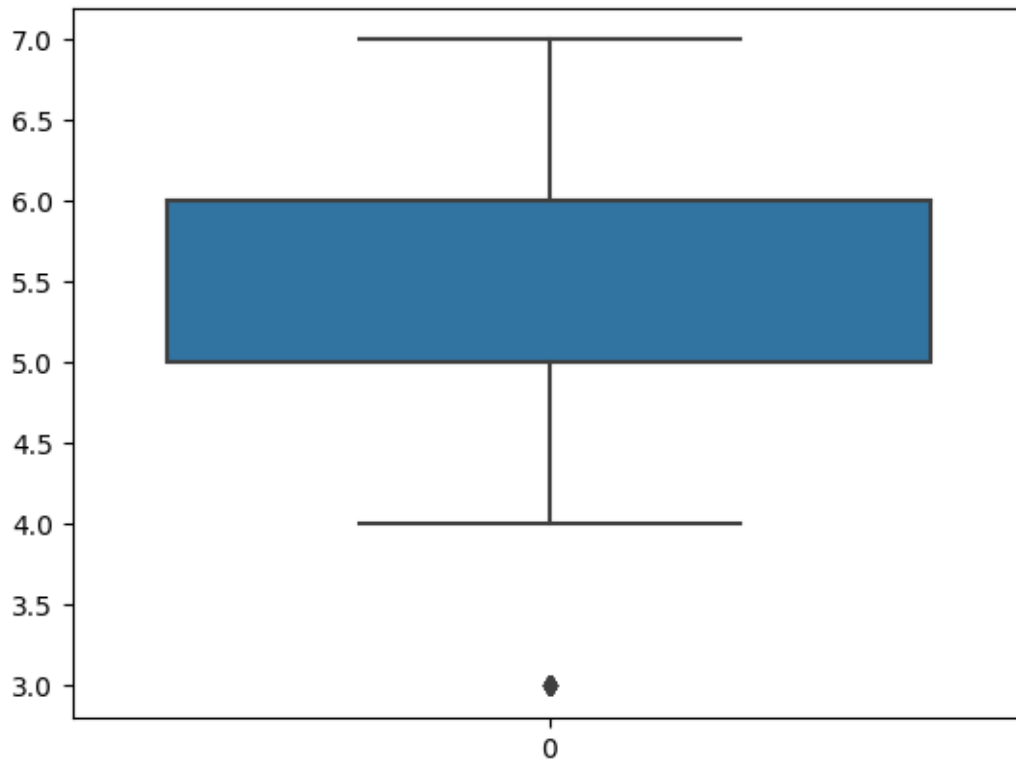
```
df.median()
```

```
[27]: fixed acidity          7.00000  
volatile acidity          0.29000  
citric acid               0.31000  
residual sugar            3.00000  
chlorides                 0.04700  
free sulfur dioxide       29.00000  
total sulfur dioxide      118.00000  
density                   0.99489  
pH                        3.21000  
sulphates                 0.51000  
alcohol                   10.30000  
quality                   6.00000  
dtype: float64
```

```
[28]: df['quality']=np.where(df['quality']>upper_limit,6,df['quality'])
```

```
[29]: sns.boxplot(df.quality)
```

```
[29]: <Axes: >
```



```
[30]: df.shape
```

```
[30]: (6497, 13)
```

0.2.6 Z-Score method

```
[31]: from scipy import stats
```

```
[32]: quality_zscore=stats.zscore(df.quality)
      quality_zscore
```

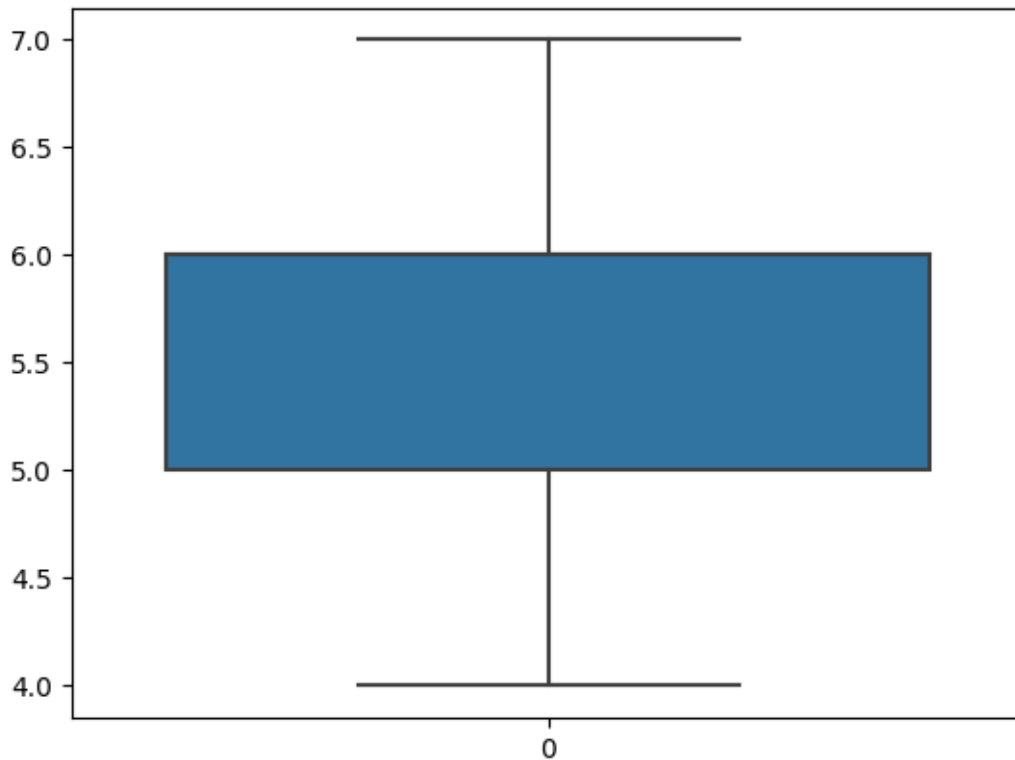
```
[32]: 0      0.311447
      1      0.311447
      2      0.311447
      3      0.311447
      4      0.311447
      ...
      6492  -0.968421
      6493   0.311447
      6494   0.311447
      6495  -0.968421
      6496   0.311447
```


Name: quality, Length: 6497, dtype: float64

```
[33]: df_z=df[np.abs(quality_zscore)<=3]
```

```
[34]: sns.boxplot(df_z.quality)
```

```
[34]: <Axes: >
```



0.2.7 Label Encoding

```
[35]: from sklearn.preprocessing import LabelEncoder
```

```
[36]: le = LabelEncoder()
```

```
[37]: df.alcohol = le.fit_transform(df.alcohol)
df.quality = le.fit_transform(df.quality)
```

```
[38]: df.head()
```

```
[38]:
```

	type	fixed acidity	volatile acidity	citric acid	residual sugar	\
0	white	7.0	0.27	0.36	20.7	
1	white	6.3	0.30	0.34	1.6	

2	white	8.1	0.28	0.40	6.9
3	white	7.2	0.23	0.32	8.5
4	white	7.2	0.23	0.32	8.5

	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH \
0	0.045	45.0	170.0	1.0010	3.00
1	0.049	14.0	132.0	0.9940	3.30
2	0.050	30.0	97.0	0.9951	3.26
3	0.058	47.0	186.0	0.9956	3.19
4	0.058	47.0	186.0	0.9956	3.19

	sulphates	alcohol	quality
0	0.45	5	3
1	0.49	15	3
2	0.44	29	3
3	0.40	25	3
4	0.40	25	3

0.2.8 One Hot encoding

```
[39]: df_main = pd.get_dummies(df, columns = ['quality'])
df_main.head()
```

```
[39]:
```

	type	fixed acidity	volatile acidity	citric acid	residual sugar \
0	white	7.0	0.27	0.36	20.7
1	white	6.3	0.30	0.34	1.6
2	white	8.1	0.28	0.40	6.9
3	white	7.2	0.23	0.32	8.5
4	white	7.2	0.23	0.32	8.5

	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH \
0	0.045	45.0	170.0	1.0010	3.00
1	0.049	14.0	132.0	0.9940	3.30
2	0.050	30.0	97.0	0.9951	3.26
3	0.058	47.0	186.0	0.9956	3.19
4	0.058	47.0	186.0	0.9956	3.19

	sulphates	alcohol	quality_0	quality_1	quality_2	quality_3	quality_4
0	0.45	5	0	0	0	1	0
1	0.49	15	0	0	0	1	0
2	0.44	29	0	0	0	1	0
3	0.40	25	0	0	0	1	0
4	0.40	25	0	0	0	1	0

```
[40]: df_main.corr()
```

<ipython-input-40-b764c75a6398>:1: FutureWarning: The default value of

numeric_only in DataFrame.corr is deprecated. In a future version, it will default to False. Select only valid columns or specify the value of numeric_only to silence this warning.

```
df_main.corr()
```

```
[40]:
```

	fixed acidity	volatile acidity	citric acid	\
fixed acidity	1.000000	0.220172	0.323736	
volatile acidity	0.220172	1.000000	-0.378061	
citric acid	0.323736	-0.378061	1.000000	
residual sugar	-0.112319	-0.196702	0.142486	
chlorides	0.298421	0.377167	0.039315	
free sulfur dioxide	-0.283317	-0.353230	0.133437	
total sulfur dioxide	-0.329747	-0.414928	0.195218	
density	0.459204	0.271193	0.096320	
pH	-0.251814	0.260660	-0.328689	
sulphates	0.300380	0.225476	0.057613	
alcohol	-0.098798	-0.043591	-0.000616	
quality_0	0.033473	0.073398	-0.017692	
quality_1	0.010350	0.133305	-0.059262	
quality_2	0.060872	0.213069	-0.053040	
quality_3	-0.043064	-0.155956	0.036879	
quality_4	-0.030183	-0.137636	0.049299	

	residual sugar	chlorides	free sulfur dioxide	\
fixed acidity	-0.112319	0.298421	-0.283317	
volatile acidity	-0.196702	0.377167	-0.353230	
citric acid	0.142486	0.039315	0.133437	
residual sugar	1.000000	-0.128902	0.403439	
chlorides	-0.128902	1.000000	-0.195042	
free sulfur dioxide	0.403439	-0.195042	1.000000	
total sulfur dioxide	0.495820	-0.279580	0.720934	
density	0.552498	0.362594	0.025717	
pH	-0.267050	0.044806	-0.145191	
sulphates	-0.185745	0.395332	-0.188489	
alcohol	-0.344258	-0.260449	-0.172378	
quality_0	-0.004357	0.040817	0.033354	
quality_1	-0.050313	0.021742	-0.103325	
quality_2	0.052973	0.172449	-0.011363	
quality_3	0.018451	-0.073414	0.045233	
quality_4	-0.066606	-0.137205	-0.002594	

	total sulfur dioxide	density	pH	sulphates	\
fixed acidity	-0.329747	0.459204	-0.251814	0.300380	
volatile acidity	-0.414928	0.271193	0.260660	0.225476	
citric acid	0.195218	0.096320	-0.328689	0.057613	
residual sugar	0.495820	0.552498	-0.267050	-0.185745	
chlorides	-0.279580	0.362594	0.044806	0.395332	

free sulfur dioxide	0.720934	0.025717	-0.145191	-0.188489
total sulfur dioxide	1.000000	0.032395	-0.237687	-0.275381
density	0.032395	1.000000	0.011920	0.259454
pH	-0.237687	0.011920	1.000000	0.191248
sulphates	-0.275381	0.259454	0.191248	1.000000
alcohol	-0.257704	-0.683388	0.110382	-0.012222
quality_0	0.007579	0.023791	0.016652	-0.011392
quality_1	-0.040397	0.008411	0.015269	-0.032747
quality_2	0.063129	0.269154	-0.027682	-0.022592
quality_3	-0.003292	-0.085370	-0.001131	-0.000818
quality_4	-0.057214	-0.233773	0.026076	0.047432

	alcohol	quality_0	quality_1	quality_2	quality_3	\
fixed acidity	-0.098798	0.033473	0.010350	0.060872	-0.043064	
volatile acidity	-0.043591	0.073398	0.133305	0.213069	-0.155956	
citric acid	-0.000616	-0.017692	-0.059262	-0.053040	0.036879	
residual sugar	-0.344258	-0.004357	-0.050313	0.052973	0.018451	
chlorides	-0.260449	0.040817	0.021742	0.172449	-0.073414	
free sulfur dioxide	-0.172378	0.033354	-0.103325	-0.011363	0.045233	
total sulfur dioxide	-0.257704	0.007579	-0.040397	0.063129	-0.003292	
density	-0.683388	0.023791	0.008411	0.269154	-0.085370	
pH	0.110382	0.016652	0.015269	-0.027682	-0.001131	
sulphates	-0.012222	-0.011392	-0.032747	-0.022592	-0.000818	
alcohol	1.000000	-0.013644	-0.049315	-0.386320	0.131039	
quality_0	-0.013644	1.000000	-0.012631	-0.047700	-0.063752	
quality_1	-0.049315	-0.012631	1.000000	-0.129874	-0.173578	
quality_2	-0.386320	-0.047700	-0.129874	1.000000	-0.655529	
quality_3	0.131039	-0.063752	-0.173578	-0.655529	1.000000	
quality_4	0.338339	-0.030395	-0.082757	-0.312537	-0.417708	

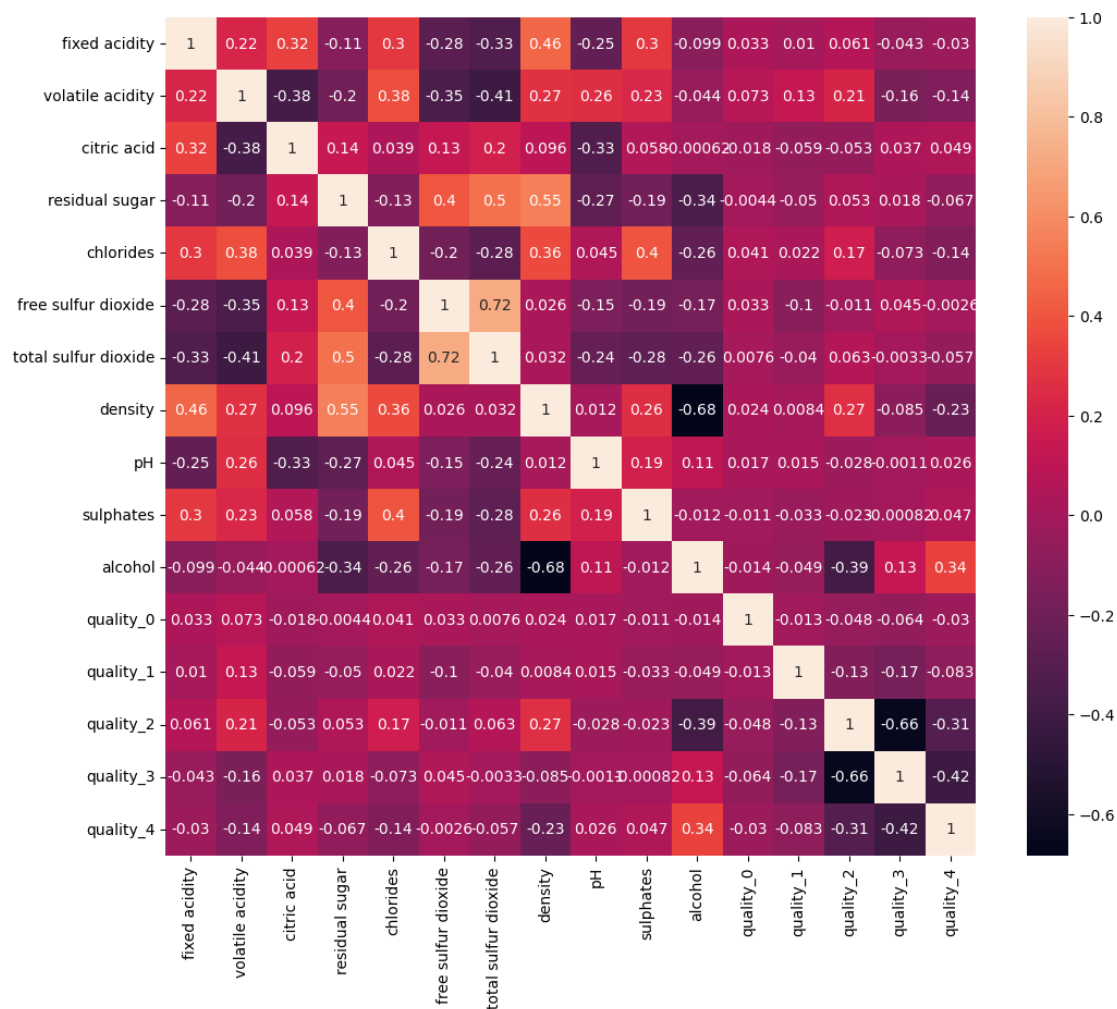
	quality_4
fixed acidity	-0.030183
volatile acidity	-0.137636
citric acid	0.049299
residual sugar	-0.066606
chlorides	-0.137205
free sulfur dioxide	-0.002594
total sulfur dioxide	-0.057214
density	-0.233773
pH	0.026076
sulphates	0.047432
alcohol	0.338339
quality_0	-0.030395
quality_1	-0.082757
quality_2	-0.312537
quality_3	-0.417708
quality_4	1.000000

```
[41]: plt.figure(figsize=(12,10))
sns.heatmap(df_main.corr(),annot =True)
```

<ipython-input-41-65b4d4cae710>:2: FutureWarning: The default value of numeric_only in DataFrame.corr is deprecated. In a future version, it will default to False. Select only valid columns or specify the value of numeric_only to silence this warning.

```
sns.heatmap(df_main.corr(),annot =True)
```

[41]: <Axes: >



```
[42]: df_main.corr().alcohol.sort_values(ascending=False)
```

<ipython-input-42-3bc757a85a29>:1: FutureWarning: The default value of numeric_only in DataFrame.corr is deprecated. In a future version, it will default to False. Select only valid columns or specify the value of numeric_only

to silence this warning.

```
df_main.corr().alcohol.sort_values(ascending=False)
```

```
[42]: alcohol          1.000000
      quality_4       0.338339
      quality_3       0.131039
      pH              0.110382
      citric acid     -0.000616
      sulphates       -0.012222
      quality_0       -0.013644
      volatile acidity -0.043591
      quality_1       -0.049315
      fixed acidity   -0.098798
      free sulfur dioxide -0.172378
      total sulfur dioxide -0.257704
      chlorides       -0.260449
      residual sugar  -0.344258
      quality_2       -0.386320
      density         -0.683388
      Name: alcohol, dtype: float64
```

```
[43]: df_main.head()
```

```
[43]:
```

	type	fixed acidity	volatile acidity	citric acid	residual sugar	\
0	white	7.0	0.27	0.36	20.7	
1	white	6.3	0.30	0.34	1.6	
2	white	8.1	0.28	0.40	6.9	
3	white	7.2	0.23	0.32	8.5	
4	white	7.2	0.23	0.32	8.5	

	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	\
0	0.045	45.0	170.0	1.0010	3.00	
1	0.049	14.0	132.0	0.9940	3.30	
2	0.050	30.0	97.0	0.9951	3.26	
3	0.058	47.0	186.0	0.9956	3.19	
4	0.058	47.0	186.0	0.9956	3.19	

	sulphates	alcohol	quality_0	quality_1	quality_2	quality_3	quality_4
0	0.45	5	0	0	0	1	0
1	0.49	15	0	0	0	1	0
2	0.44	29	0	0	0	1	0
3	0.40	25	0	0	0	1	0
4	0.40	25	0	0	0	1	0

0.2.9 x and y split

```
[44]: x =df.drop(columns =['quality'],axis =1)
```

```
[45]: x.head()
```

```
[45]:
```

	type	fixed acidity	volatile acidity	citric acid	residual sugar	\
0	white	7.0	0.27	0.36	20.7	
1	white	6.3	0.30	0.34	1.6	
2	white	8.1	0.28	0.40	6.9	
3	white	7.2	0.23	0.32	8.5	
4	white	7.2	0.23	0.32	8.5	

	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	\
0	0.045	45.0	170.0	1.0010	3.00	
1	0.049	14.0	132.0	0.9940	3.30	
2	0.050	30.0	97.0	0.9951	3.26	
3	0.058	47.0	186.0	0.9956	3.19	
4	0.058	47.0	186.0	0.9956	3.19	

	sulphates	alcohol
0	0.45	5
1	0.49	15
2	0.44	29
3	0.40	25
4	0.40	25

```
[46]: y =df.quality
      y.head()
```

```
[46]: 0    3
      1    3
      2    3
      3    3
      4    3
      Name: quality, dtype: int64
```

```
[47]: from sklearn.preprocessing import MinMaxScaler
      scale =MinMaxScaler()
```

0.2.10 Model Building

```
[71]: from sklearn.linear_model import LogisticRegression
      model = LogisticRegression()
```

0.2.11 Evaluation for classification model

```
[72]: from sklearn.metrics import accuracy_score, confusion_matrix, \
      ↪ classification_report, roc_auc_score, roc_curve
```

0.2.12 Random Forest Classifier

```
[73]: from sklearn.ensemble import RandomForestClassifier
      model = RandomForestClassifier(criterion='entropy')
```

```
[74]: df.head()
```

```
[74]:   type  fixed acidity  volatile acidity  citric acid  residual sugar \
0  white           7.0           0.27           0.36           20.7
1  white           6.3           0.30           0.34            1.6
2  white           8.1           0.28           0.40            6.9
3  white           7.2           0.23           0.32            8.5
4  white           7.2           0.23           0.32            8.5

      chlorides  free sulfur dioxide  total sulfur dioxide  density  pH \
0         0.045             45.0             170.0     1.0010  3.00
1         0.049             14.0             132.0     0.9940  3.30
2         0.050             30.0              97.0     0.9951  3.26
3         0.058             47.0             186.0     0.9956  3.19
4         0.058             47.0             186.0     0.9956  3.19

      sulphates  alcohol  quality
0         0.45         5         3
1         0.49        15         3
2         0.44        29         3
3         0.40        25         3
4         0.40        25         3
```

```
[77]: from sklearn.linear_model import LinearRegression
      lr = LinearRegression()
```