

```
In [1]: import numpy as np
import pandas as pd
import seaborn as sns
from sklearn.svm import SVC
import matplotlib.pyplot as plt
from sklearn.linear_model import SGDClassifier
from sklearn.preprocessing import LabelEncoder
from sklearn.model_selection import GridSearchCV
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import classification_report, confusion_matrix, accuracy_score

%matplotlib inline

In [2]: wine = pd.read_csv('winequality-red.csv')
wine.head()
```

Out[2]:

	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 [3]: wine.describe()

Out[3]:

	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 [4]: 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
```

```
In [5]: fig = plt.figure(figsize=(15,10))

plt.subplot(3,4,1)
sns.barplot(x='quality',y='fixed acidity',data=wine)

plt.subplot(3,4,2)
sns.barplot(x='quality',y='volatile acidity',data=wine)

plt.subplot(3,4,3)
sns.barplot(x='quality',y='citric acid',data=wine)

plt.subplot(3,4,4)
sns.barplot(x='quality',y='residual sugar',data=wine)

plt.subplot(3,4,5)
sns.barplot(x='quality',y='chlorides',data=wine)

plt.subplot(3,4,6)
sns.barplot(x='quality',y='free sulfur dioxide',data=wine)

plt.subplot(3,4,7)
sns.barplot(x='quality',y='total sulfur dioxide',data=wine)

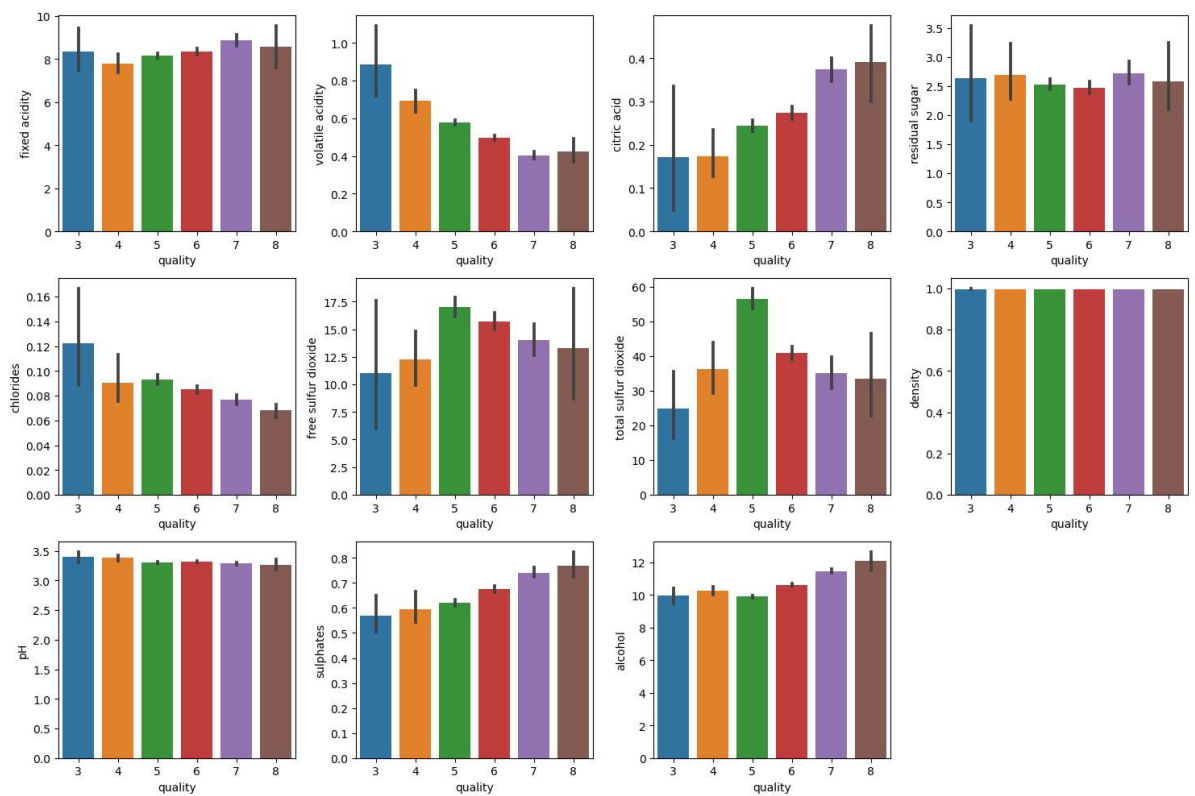
plt.subplot(3,4,8)
sns.barplot(x='quality',y='density',data=wine)

plt.subplot(3,4,9)
sns.barplot(x='quality',y='pH',data=wine)

plt.subplot(3,4,10)
sns.barplot(x='quality',y='sulphates',data=wine)

plt.subplot(3,4,11)
sns.barplot(x='quality',y='alcohol',data=wine)

plt.tight_layout()
```



```
In [6]: wine['quality'].value_counts()
```

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

```
In [7]: ranges = (2,6.5,8)
        groups = ['bad','good']
        wine['quality'] = pd.cut(wine['quality'],bins=ranges,labels=groups)
        le = LabelEncoder()
        wine['quality'] = le.fit_transform(wine['quality'])
        wine.head()
```

Out[7]:

	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 [8]: wine['quality'].value_counts()
```

```
Out[8]: 0    1382
        1     217
        Name: quality, dtype: int64
```

```
In [9]: good_quality = wine[wine['quality']==1]
bad_quality = wine[wine['quality']==0]

bad_quality = bad_quality.sample(frac=1)
bad_quality = bad_quality[:217]

new_df = pd.concat([good_quality,bad_quality])
new_df = new_df.sample(frac=1)
new_df
```

```
Out[9]:
```

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	alcol
1421	7.5	0.400	0.18	1.6	0.079	24.0	58.0	0.99650	3.34	0.58	
549	9.0	0.530	0.49	1.9	0.171	6.0	25.0	0.99750	3.27	0.61	
1193	6.4	0.885	0.00	2.3	0.166	6.0	12.0	0.99551	3.56	0.51	1
124	7.8	0.500	0.17	1.6	0.082	21.0	102.0	0.99600	3.39	0.48	
640	9.9	0.540	0.45	2.3	0.071	16.0	40.0	0.99910	3.39	0.62	
...
1204	7.2	0.360	0.46	2.1	0.074	24.0	44.0	0.99534	3.40	0.85	1
842	10.6	0.500	0.45	2.6	0.119	34.0	68.0	0.99708	3.23	0.72	1
290	8.7	0.520	0.09	2.5	0.091	20.0	49.0	0.99760	3.34	0.86	1
1440	7.2	0.370	0.32	2.0	0.062	15.0	28.0	0.99470	3.23	0.73	1
455	11.3	0.620	0.67	5.2	0.086	6.0	19.0	0.99880	3.22	0.69	1

434 rows × 12 columns

```
In [10]: new_df['quality'].value_counts()
```

```
Out[10]: 0    217
         1    217
         Name: quality, dtype: int64
```

```
In [11]: new_df.corr()['quality'].sort_values(ascending=False)
```

```
Out[11]: quality          1.000000
         alcohol        0.577559
         sulphates      0.355054
         citric acid     0.319310
         fixed acidity    0.174802
         residual sugar  -0.001258
         pH             -0.069247
         free sulfur dioxide -0.088221
         chlorides       -0.209602
         total sulfur dioxide -0.213407
         density        -0.245622
         volatile acidity -0.457369
         Name: quality, dtype: float64
```

```
In [18]: from sklearn.model_selection import train_test_split
```

```
X = new_df.drop('quality',axis=1)
y = new_df['quality']
```

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_st
```

```
In [19]: param = {'n_estimators':[100,200,300,400,500,600,700,800,900,1000]}

grid_rf = GridSearchCV(RandomForestClassifier(),param,scoring='accuracy',cv=10,)
grid_rf.fit(X_train, y_train)

print('Best parameters --> ', grid_rf.best_params_)

# Wine Quality Prediction
pred = grid_rf.predict(X_test)

print(confusion_matrix(y_test,pred))
print('\n')
print(classification_report(y_test,pred))
print('\n')
print(accuracy_score(y_test,pred))
```

Best parameters --> {'n_estimators': 900}
[[55 19]
 [3 54]]

	precision	recall	f1-score	support
0	0.95	0.74	0.83	74
1	0.74	0.95	0.83	57
accuracy			0.83	131
macro avg	0.84	0.85	0.83	131
weighted avg	0.86	0.83	0.83	131

0.8320610687022901

```
In [ ]:
```