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

In [24]: wine = pd.read_csv('E://Data Science//Sapient//Case_Onsite_Modeling_Wine//wine
_dataset.csv')
```

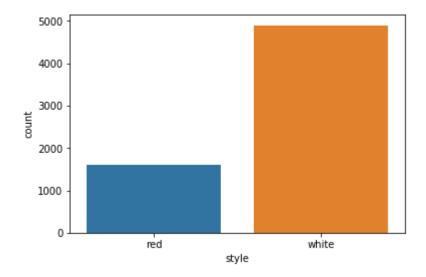
# Finding missing values in dataset

```
In [3]:
         wine.isna().sum()
Out[3]: fixed acidity
                                  0
         volatile_acidity
                                  0
         citric_acid
                                  0
                                  0
         residual sugar
         chlorides
                                  0
         free_sulfur_dioxide
                                  0
         total_sulfur_dioxide
                                  0
         density
                                  0
                                  0
         рΗ
                                  0
         sulphates
         alcohol
                                  0
         quality
                                  0
                                  0
         style
         dtype: int64
In [4]: wine.dtypes
Out[4]: fixed_acidity
                                  float64
         volatile_acidity
                                  float64
         citric_acid
                                  float64
         residual sugar
                                  float64
         chlorides
                                  float64
         free_sulfur_dioxide
                                  float64
         total_sulfur_dioxide
                                  float64
         density
                                  float64
                                  float64
         рΗ
         sulphates
                                  float64
         alcohol
                                  float64
         quality
                                    int64
                                   object
         style
         dtype: object
```

#### **EDA**

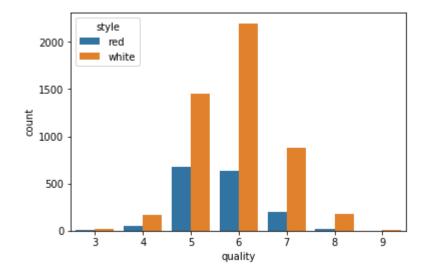
In [6]: sns.countplot(wine['style'])

Out[6]: <matplotlib.axes.\_subplots.AxesSubplot at 0x249389808d0>



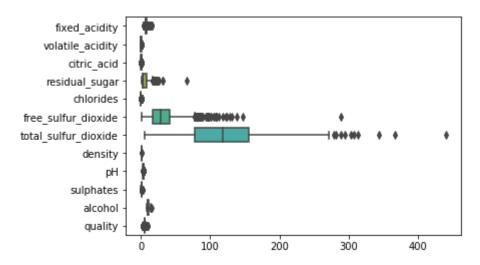
In [7]: sns.countplot(wine['quality'],hue = wine['style'])

Out[7]: <matplotlib.axes.\_subplots.AxesSubplot at 0x249389b1cc0>



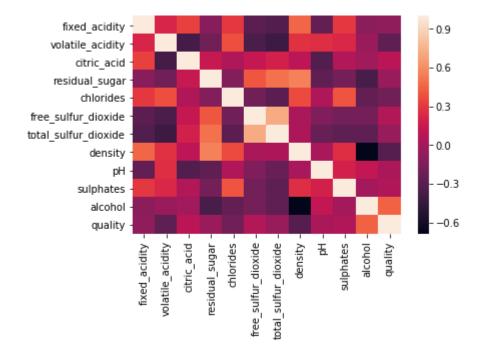
In [12]: temp = [x for x in wine.columns if x not in ['style']]
sns.boxplot(data = wine[temp], orient="h")

Out[12]: <matplotlib.axes.\_subplots.AxesSubplot at 0x2493a0cc668>



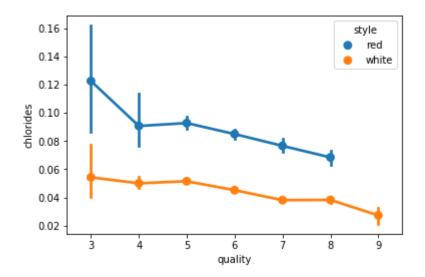
In [14]: corr = wine[temp].corr()
 sns.heatmap(corr)

Out[14]: <matplotlib.axes.\_subplots.AxesSubplot at 0x2493a285198>



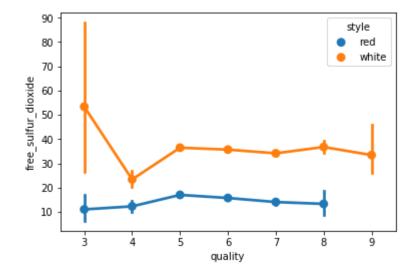
In [15]: sns.pointplot(x="quality",y="chlorides",hue="style",data = wine)

Out[15]: <matplotlib.axes.\_subplots.AxesSubplot at 0x2493a3399b0>



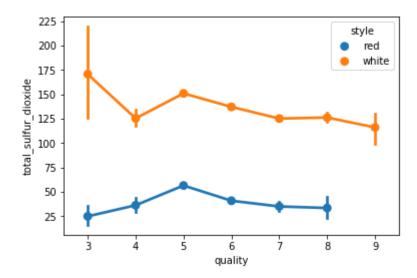
In [16]: sns.pointplot(x="quality",y="free\_sulfur\_dioxide",hue="style",data = wine)

Out[16]: <matplotlib.axes.\_subplots.AxesSubplot at 0x2493a370278>



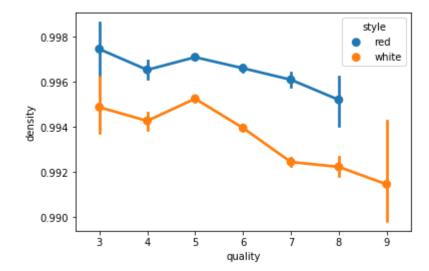
In [17]: sns.pointplot(x="quality",y="total\_sulfur\_dioxide",hue="style",data = wine)

Out[17]: <matplotlib.axes.\_subplots.AxesSubplot at 0x2493a464320>



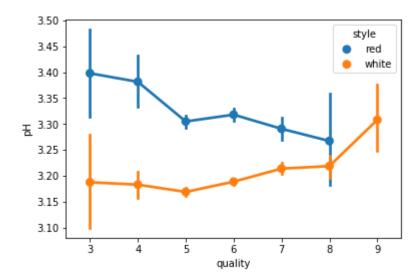
In [18]: sns.pointplot(x="quality",y="density",hue="style",data = wine)

Out[18]: <matplotlib.axes.\_subplots.AxesSubplot at 0x2493a500208>



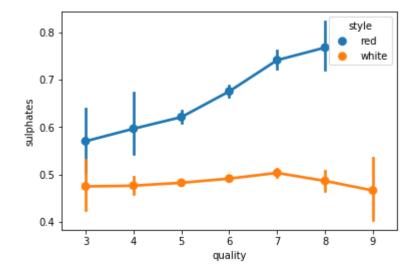
In [19]: sns.pointplot(x="quality",y="pH",hue="style",data = wine)

Out[19]: <matplotlib.axes.\_subplots.AxesSubplot at 0x2493a51cb00>



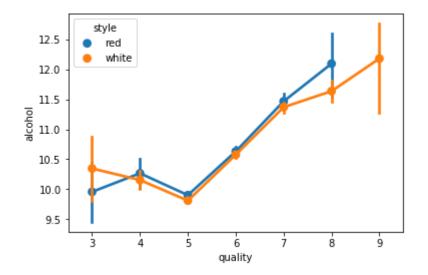
In [20]: sns.pointplot(x="quality",y="sulphates",hue="style",data = wine)

Out[20]: <matplotlib.axes.\_subplots.AxesSubplot at 0x2493a5ab400>



```
In [21]: sns.pointplot(x="quality",y="alcohol",hue="style",data = wine)
```

Out[21]: <matplotlib.axes. subplots.AxesSubplot at 0x2493a6ac470>



# converting style datatype to int

```
In [38]: from sklearn.preprocessing import LabelEncoder
    le = LabelEncoder()
    wine['style'] = le.fit_transform(wine['style'])
```

# checking the imbalance of dataset

```
In [25]: wine['style'].value_counts(normalize = True)
Out[25]: white    0.753886
    red    0.246114
    Name: style, dtype: float64
```

### **Modeling**

```
In [32]: from sklearn.model_selection import train_test_split
    from sklearn.model_selection import cross_val_score
    from sklearn.metrics import f1_score
    from sklearn.tree import DecisionTreeClassifier
    from sklearn.linear_model import LogisticRegression
    from sklearn.ensemble import RandomForestClassifier
    from xgboost import XGBClassifier
    from imblearn.over_sampling import SMOTE
```

```
In [27]: def data_sampling(X, y, method = 'smote', kind = 'regular', seed = 0):
    if method == 'smote':
        sm = SMOTE(kind = kind)
        X_res, y_res = sm.fit_sample(X, y)

    return {'X' : X_res, 'y' : y_res}

In [39]: feature_names = [x for x in wine.columns if x not in ['style']]
    target = wine['style']

In [40]: X = wine[feature_names].values
    y = wine['style'].values

In [41]: X_train, X_valid, y_train, y_valid= train_test_split(X,y,train_size = 0.7,str atify = y,random_state = 0)

    C:\Users\AA\Anaconda3\lib\site-packages\sklearn\model_selection\_split.py:202
    6: FutureWarning: From version 0.21, test_size will always complement train_s ize unless both are specified.
    FutureWarning)
```

### applying smote on data

```
In [42]: smote = data_sampling(X_train, y_train, method = 'smote')
    X_SMOTE_train = smote['X']
    y_SMOTE_train = smote['y']

In [43]: pd.DataFrame({'target' : y_SMOTE_train})['target'].value_counts(normalize=True)

Out[43]: 1    0.5
    0    0.5
    Name: target, dtype: float64
```

## **Logistic Regression**

#### kfold cross validation

```
In [47]: f1 = cross_val_score(lr,X_SMOTE_train, y_SMOTE_train, cv=3, scoring='f1')
In [48]: f1
Out[48]: array([0.97150373, 0.98601399, 0.98030635])
```

#### **Decission Tree**

## grid search with cross validation

```
In [51]: from sklearn.model_selection import GridSearchCV
    parameters = [{'max_depth' :[5,10,15,20]}]
    grid_search = GridSearchCV(clf,param_grid = parameters,scoring = 'f1',cv = 3,n
    _jobs = -1)

In [52]: grid_search = grid_search.fit(X_SMOTE_train, y_SMOTE_train)

In [53]: grid_search.best_score_
Out[53]: 0.9887692782409007

In [54]: grid_search.best_params_
Out[54]: {'max_depth': 10}
```

## **Random Forest**

## grid search with cross validation

```
In [56]: parameters = [{'max_depth' :[5,10,15,20] , 'max_features':[3,4,5,6]}]
    grid_search = GridSearchCV(rf,param_grid = parameters,scoring = 'f1',cv = 3,n_
    jobs = -1)

In [57]: grid_search = grid_search.fit(X_SMOTE_train, y_SMOTE_train)

In [69]: grid_search.best_score_
Out[69]: 0.9951893099026833

In [59]: grid_search.best_params_
Out[59]: {'max_depth': 10, 'max_features': 4}
```

## xgboost

```
In [97]: from sklearn.metrics import confusion matrix
         confusion matrix(y valid,xgb.predict(X valid))
         C:\Users\AA\Anaconda3\lib\site-packages\sklearn\preprocessing\label.py:151: D
         eprecationWarning: The truth value of an empty array is ambiguous. Returning
         False, but in future this will result in an error. Use `array.size > 0` to ch
         eck that an array is not empty.
           if diff:
Out[97]: array([[ 472,
                    8, 1462]], dtype=int64)
In [98]: f1 score(y valid, xgb.predict(X valid))
         C:\Users\AA\Anaconda3\lib\site-packages\sklearn\preprocessing\label.py:151: D
         eprecationWarning: The truth value of an empty array is ambiguous. Returning
         False, but in future this will result in an error. Use `array.size > 0` to ch
         eck that an array is not empty.
           if diff:
Out[98]: 0.9945578231292517
```

### grid search with cross validation

Out[105]:

	colname	importance
0	fixed_acidity	0.054598
1	volatile_acidity	0.117816
2	citric_acid	0.005747
3	residual_sugar	0.031609
4	chlorides	0.172414
5	free_sulfur_dioxide	0.005747
6	total_sulfur_dioxide	0.255747
7	density	0.080460
8	рН	0.106322
9	sulphates	0.120690
10	alcohol	0.048851
11	quality	0.000000

# applying parameters obtained form gridsearch to model

```
In [116]: xgb = XGBClassifier(random_state = 0, max_depth = 5,n_estimators=500)
xgb.fit(X_SMOTE_train, y_SMOTE_train)
```

```
In [117]: f1_score(y_valid, xgb.predict(X_valid))
```

C:\Users\AA\Anaconda3\lib\site-packages\sklearn\preprocessing\label.py:151: D
eprecationWarning: The truth value of an empty array is ambiguous. Returning
False, but in future this will result in an error. Use `array.size > 0` to ch
eck that an array is not empty.
 if diff:

Out[117]: 0.9959211420802174