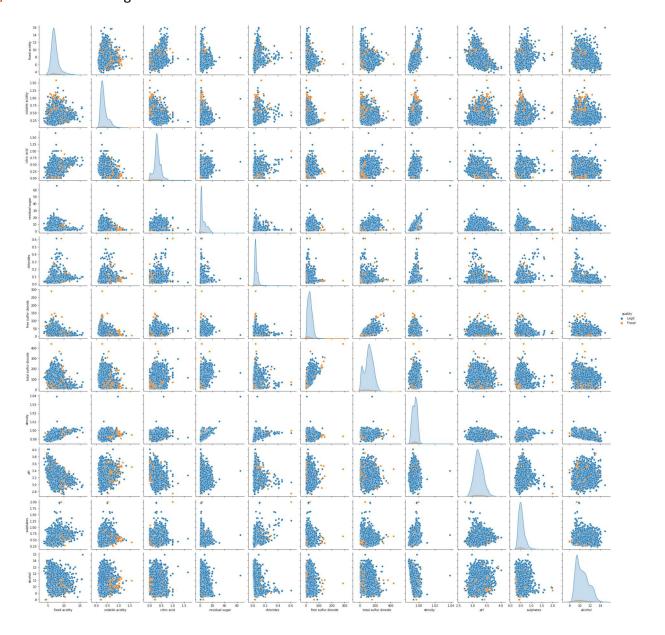
```
In [1]:
         import numpy as np
         import pandas as pd
         import matplotlib.pyplot as plt
         import seaborn as sns
In [2]: | df = pd.read_csv('wine_fraud.csv')
In [3]: | df.head()
Out[3]:
                                                                total
                                                         free
               fixed
                     volatile citric
                                   residual
                                            chlorides
                                                       sulfur
                                                               sulfur density
                                                                               pH sulphates alcohol
             acidity
                     acidity
                             acid
                                     sugar
                                                      dioxide
                                                              dioxide
          0
                7.4
                       0.70
                              0.00
                                        1.9
                                               0.076
                                                         11.0
                                                                 34.0
                                                                       0.9978 3.51
                                                                                        0.56
                                                                                                  9.4
                7.8
                             0.00
                                               0.098
                                                                       0.9968 3.20
          1
                       0.88
                                       2.6
                                                        25.0
                                                                 67.0
                                                                                        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
                                               0.076
                7.4
                       0.70
                             0.00
                                       1.9
                                                         11.0
                                                                 34.0
                                                                       0.9978 3.51
                                                                                         0.56
                                                                                                  9.4
In [4]: |df['type'].unique()
Out[4]: array(['red', 'white'], dtype=object)
In [5]: df['quality'].unique()
Out[5]: array(['Legit', 'Fraud'], dtype=object)
In [6]: df.isnull().sum()
Out[6]: fixed acidity
                                     0
                                     0
         volatile acidity
         citric acid
         residual sugar
                                     0
         chlorides
                                     0
         free sulfur dioxide
         total sulfur dioxide
                                     0
         density
                                     0
                                     0
         рΗ
         sulphates
                                     0
         alcohol
                                     0
         quality
                                     0
         type
         dtype: int64
In [7]: df.shape
Out[7]: (6497, 13)
```

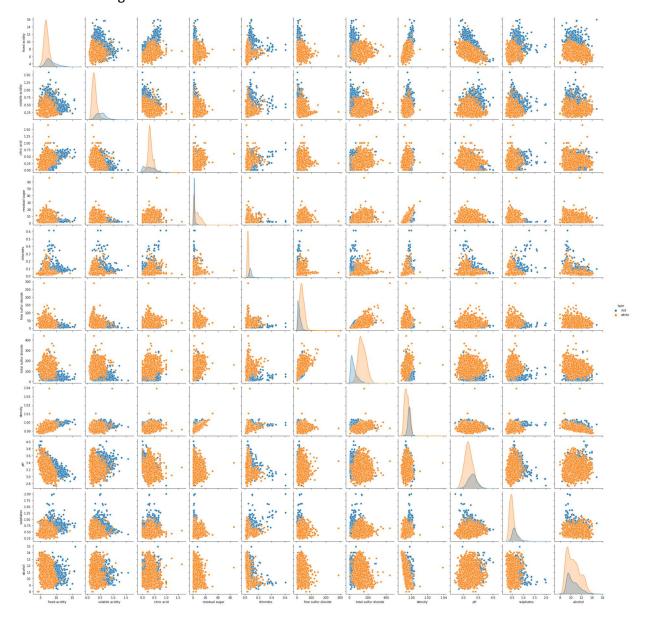
In [8]: | sns.pairplot(data=df,hue='quality')

Out[8]: <seaborn.axisgrid.PairGrid at 0x22de8b9e850>



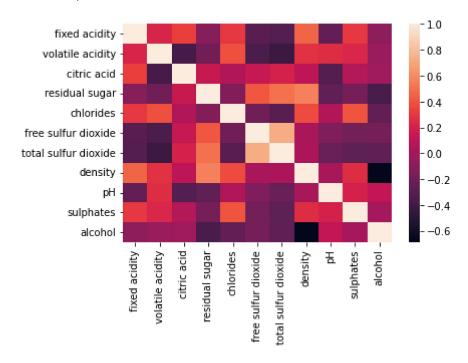
In [9]: sns.pairplot(data=df,hue='type')

Out[9]: <seaborn.axisgrid.PairGrid at 0x22df412ec70>



In [10]: sns.heatmap(data=df.corr())

Out[10]: <AxesSubplot:>

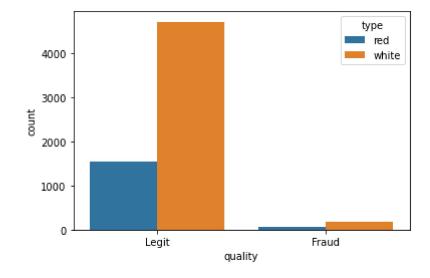


In [11]: sns.countplot('quality',data=df,hue='type')

C:\Users\aarav\anaconda3\lib\site-packages\seaborn_decorators.py:36: FutureWar ning: Pass the following variable as a keyword arg: x. From version 0.12, the o nly valid positional argument will be `data`, and passing other arguments witho ut an explicit keyword will result in an error or misinterpretation.

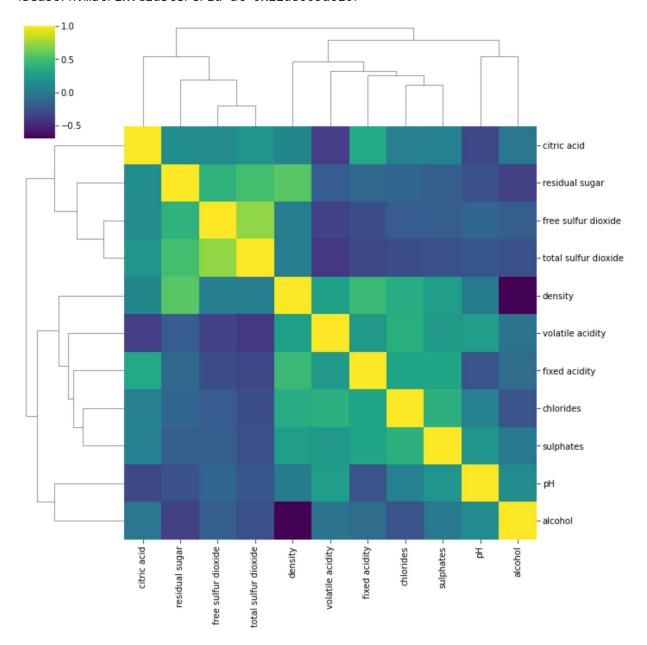
warnings.warn(

Out[11]: <AxesSubplot:xlabel='quality', ylabel='count'>



```
In [12]: sns.clustermap(df.corr(),cmap='viridis')
```

Out[12]: <seaborn.matrix.ClusterGrid at 0x22d8005a610>



```
In [13]: df['quality'] = df['quality'].map({'Legit':1,'Fraud':0})
In [14]: df['type'] = df['type'].map({'red':1,'white':0})
In [15]: X = df.drop('quality',axis=1)
In [16]: y = df['quality']
```

```
In [17]: from sklearn.model selection import train test split
In [18]:
          X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.1, random_
In [19]: from sklearn.preprocessing import StandardScaler
In [20]: from sklearn.pipeline import Pipeline
In [21]: from sklearn.svm import SVC
In [22]: from sklearn.model_selection import GridSearchCV
In [23]: from sklearn.metrics import classification_report,accuracy_score,plot_confusion_r
In [24]: | scaler = StandardScaler()
In [25]: svm = SVC(class weight='balanced') #c,kernel,gamma
In [26]: operations = [('scaler', scaler), ('svm', svm)]
In [27]: pipe = Pipeline(operations)
In [28]: C = [0.001, 0.01, 0.1, 0.5, 1.0]
         kernel = ['rbf','linear','poly','sigmoid']
         gamma = ['auto','scaled']
         param grid = {'svm C':C,'svm kernel':kernel,'svm gamma':gamma}
In [29]:
In [30]: full_grid_classifier = GridSearchCV(pipe,param_grid)
```

```
In [31]: full grid classifier.fit(X train,y train)
         C:\Users\aarav\anaconda3\lib\site-packages\sklearn\model_selection\_validation.
         py:372: FitFailedWarning:
         100 fits failed out of a total of 200.
         The score on these train-test partitions for these parameters will be set to na
         If these failures are not expected, you can try to debug them by setting error
         score='raise'.
         Below are more details about the failures:
         100 fits failed with the following error:
         Traceback (most recent call last):
           File "C:\Users\aarav\anaconda3\lib\site-packages\sklearn\model_selection\_val
         idation.py", line 680, in _fit_and_score
             estimator.fit(X_train, y_train, **fit_params)
           File "C:\Users\aarav\anaconda3\lib\site-packages\sklearn\pipeline.py", line 3
         94. in fit
             self._final_estimator.fit(Xt, y, **fit_params_last_step)
           File "C:\Users\aarav\anaconda3\lib\site-packages\sklearn\svm\_base.py", line
         243, in fit
             raise ValueError(
         ValueError: When 'gamma' is a string, it should be either 'scale' or 'auto'. Go
         t 'scaled' instead.
           warnings.warn(some fits failed message, FitFailedWarning)
         C:\Users\aarav\anaconda3\lib\site-packages\sklearn\model selection\ search.py:9
         69: UserWarning: One or more of the test scores are non-finite: [0.03745505 0.8
         0468908 0.95536078 0.15720924
                                                          nan
                            nan 0.72498812 0.7916893 0.92748115 0.78176921
                 nan
                                                   nan 0.81939491 0.78108559
                 nan
                                       nan
          0.89259298 0.61107909
                                       nan
                                                   nan
                                                              nan
                                                                         nan
          0.84197144 0.78142791 0.87121771 0.55498381
                                                              nan
                                                                         nan
                            nan 0.8546295 0.78074415 0.86728404 0.55601018
                 nan
                            nan
                                                   nan]
                 nan
                                       nan
           warnings.warn(
Out[31]: GridSearchCV(estimator=Pipeline(steps=[('scaler', StandardScaler()),
                                                 ('svm', SVC(class weight='balanced'))]),
                      param grid={'svm C': [0.001, 0.01, 0.1, 0.5, 1.0],
                                   'svm__gamma': ['auto', 'scaled'],
                                   'svm kernel': ['rbf', 'linear', 'poly', 'sigmoid']})
In [32]: |full_grid_classifier.best_params_
Out[32]: {'svm C': 0.001, 'svm gamma': 'auto', 'svm kernel': 'poly'}
In [33]: y pred = full grid classifier.predict(X test)
```

In [34]: print(classification_report(y_test,y_pred))

	precision	recall	f1-score	support
0 1	0.32 0.97	0.26 0.98	0.29 0.97	27 623
accuracy macro avg weighted avg	0.64 0.94	0.62 0.95	0.95 0.63 0.94	650 650 650

In [35]: plot_confusion_matrix(full_grid_classifier,X_test,y_test)

C:\Users\aarav\anaconda3\lib\site-packages\sklearn\utils\deprecation.py:87: Fut ureWarning: Function plot_confusion_matrix is deprecated; Function `plot_confusion_matrix` is deprecated in 1.0 and will be removed in 1.2. Use one of the class methods: ConfusionMatrixDisplay.from_predictions or ConfusionMatrixDisplay.from estimator.

warnings.warn(msg, category=FutureWarning)

