

```

import numpy as np
import pandas as pd
import graphviz
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score
from sklearn.metrics import confusion_matrix
from sklearn.metrics import confusion_matrix, ConfusionMatrixDisplay
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import roc_curve, roc_auc_score
from sklearn.svm import SVC
from sklearn.tree import DecisionTreeClassifier, export_graphviz
from sklearn.model_selection import GridSearchCV
from IPython.core.display import HTML
from sklearn.tree import export_text
import seaborn as sns
import matplotlib.pyplot as plt
%matplotlib inline
sns.set(style="ticks")

from sklearn import datasets
wine = datasets.load_wine()
wine_df = pd.DataFrame(wine.data, columns=wine.feature_names)
wine_df.head()

{"summary":{"\n  \"name\": \"wine_df\",\n  \"rows\": 178,\n  \"fields\": [\n    {\n      \"column\": \"alcohol\",\n      \"properties\": {\n        \"dtype\": \"number\",\n        \"std\": 0.8118265380058575,\n        \"min\": 11.03,\n        \"max\": 14.83,\n        \"num_unique_values\": 126,\n        \"samples\": [\n          11.62,\n          13.64,\n          13.69\n        ],\n        \"semantic_type\": \"\",\n        \"description\": \"\"\n      },\n      {\n        \"column\": \"malic_acid\",\n        \"properties\": {\n          \"dtype\": \"number\",\n          \"std\": 1.1171460976144627,\n          \"min\": 0.74,\n          \"max\": 5.8,\n          \"num_unique_values\": 133,\n          \"samples\": [\n            1.21,\n            2.83,\n            1.8\n          ],\n          \"semantic_type\": \"\",\n          \"description\": \"\"\n        },\n        {\n          \"column\": \"ash\",\n          \"properties\": {\n            \"dtype\": \"number\",\n            \"std\": 0.27434400906081485,\n            \"min\": 1.36,\n            \"max\": 3.23,\n            \"num_unique_values\": 79,\n            \"samples\": [\n              2.31,\n              2.43,\n              2.52\n            ],\n            \"semantic_type\": \"\",\n            \"description\": \"\"\n          },\n          {\n            \"column\": \"alcalinity_of_ash\",\n            \"properties\": {\n              \"dtype\": \"number\",\n              \"std\": 3.339563767173505,\n              \"min\": 10.6,\n              \"max\": 30.0,\n              \"num_unique_values\": 63,\n              \"samples\": [\n                25.5,\n                28.5,\n                15.6\n              ],\n              \"semantic_type\": \"\",\n              \"description\": \"\"\n            },\n            {\n              \"column\": \"magnesium\",\n              \"properties\": {\n                \"dtype\": \"number\",\n                \"std\": 14.282483515295665,\n                \"min\":

```

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70.0,\n          \"max\": 162.0,\n          \"num_unique_values\": 53,\n\"samples\": [\n          126.0,\n          85.0,\n          162.0\n],\n          \"semantic_type\": \"\",\n          \"description\": \"\"\n}\n    },\n    {\n        \"column\": \"total_phenols\",\n\"properties\": {\n        \"dtype\": \"number\",\n        \"std\": 0.6258510488339893,\n        \"min\": 0.98,\n        \"max\": 3.88,\n        \"num_unique_values\": 97,\n        \"samples\": [\n        1.68,\n        2.11,\n        1.35\n        ],\n        \"semantic_type\": \"\",\n        \"description\": \"\"\n    },\n    {\n        \"column\": \"flavanoids\",\n\"properties\": {\n        \"dtype\": \"number\",\n        \"std\": 0.9988586850169467,\n        \"min\": 0.34,\n        \"max\": 5.08,\n        \"num_unique_values\": 132,\n        \"samples\": [\n        3.18,\n        2.5,\n        3.17\n        ],\n        \"semantic_type\": \"\",\n        \"description\": \"\"\n    },\n    {\n        \"column\": \"nonflavanoid_phenols\",\n\"properties\": {\n        \"dtype\": \"number\",\n        \"std\": 0.12445334029667937,\n        \"min\": 0.13,\n        \"max\": 0.66,\n        \"num_unique_values\": 39,\n        \"samples\": [\n        0.58,\n        0.41,\n        0.39\n        ],\n        \"semantic_type\": \"\",\n        \"description\": \"\"\n    },\n    {\n        \"column\": \"proanthocyanins\",\n\"properties\": {\n        \"dtype\": \"number\",\n        \"std\": 0.5723588626747613,\n        \"min\": 0.41,\n        \"max\": 3.58,\n        \"num_unique_values\": 101,\n        \"samples\": [\n        0.75,\n        1.77,\n        1.42\n        ],\n        \"semantic_type\": \"\",\n        \"description\": \"\"\n    },\n    {\n        \"column\": \"color_intensity\",\n\"properties\": {\n        \"dtype\": \"number\",\n        \"std\": 2.318285871822413,\n        \"min\": 1.28,\n        \"max\": 13.0,\n        \"num_unique_values\": 132,\n        \"samples\": [\n        2.95,\n        3.3,\n        5.1\n        ],\n        \"semantic_type\": \"\",\n        \"description\": \"\"\n    },\n    {\n        \"column\": \"hue\",\n\"properties\": {\n        \"dtype\": \"number\",\n        \"std\": 0.22857156582982338,\n        \"min\": 0.48,\n        \"max\": 1.71,\n        \"num_unique_values\": 78,\n        \"samples\": [\n        1.22,\n        1.04,\n        1.45\n        ],\n        \"semantic_type\": \"\",\n        \"description\": \"\"\n    },\n    {\n        \"column\": \"od280/od315_of_diluted_wines\",\n\"properties\": {\n        \"dtype\": \"number\",\n        \"std\": 0.7099904287650504,\n        \"min\": 1.27,\n        \"max\": 4.0,\n        \"num_unique_values\": 122,\n        \"samples\": [\n        1.82,\n        1.59\n        ],\n        \"semantic_type\": \"\",\n        \"description\": \"\"\n    },\n    {\n        \"column\": \"proline\",\n\"properties\": {\n        \"dtype\": \"number\",\n        \"std\": 314.9074742768491,\n        \"min\": 278.0,\n        \"max\": 1680.0,\n        \"num_unique_values\": 121,\n        \"samples\": [\n        1375.0,\n        1270.0,\n        735.0\n        ],\n        \"semantic_type\": \"\",\n        \"description\": \"\"\n    }\n    }\n  ],\n  \"type\": \"dataframe\", \"variable_name\": \"wine_df\"}

```

```
wine_df.isna().sum()
```

```
alcohol          0
malic_acid       0
ash              0
alcalinity_of_ash 0
magnesium        0
total_phenols    0
flavanoids       0
nonflavanoid_phenols 0
proanthocyanins  0
color_intensity  0
hue              0
od280/od315_of_diluted_wines 0
proline          0
dtype: int64
```

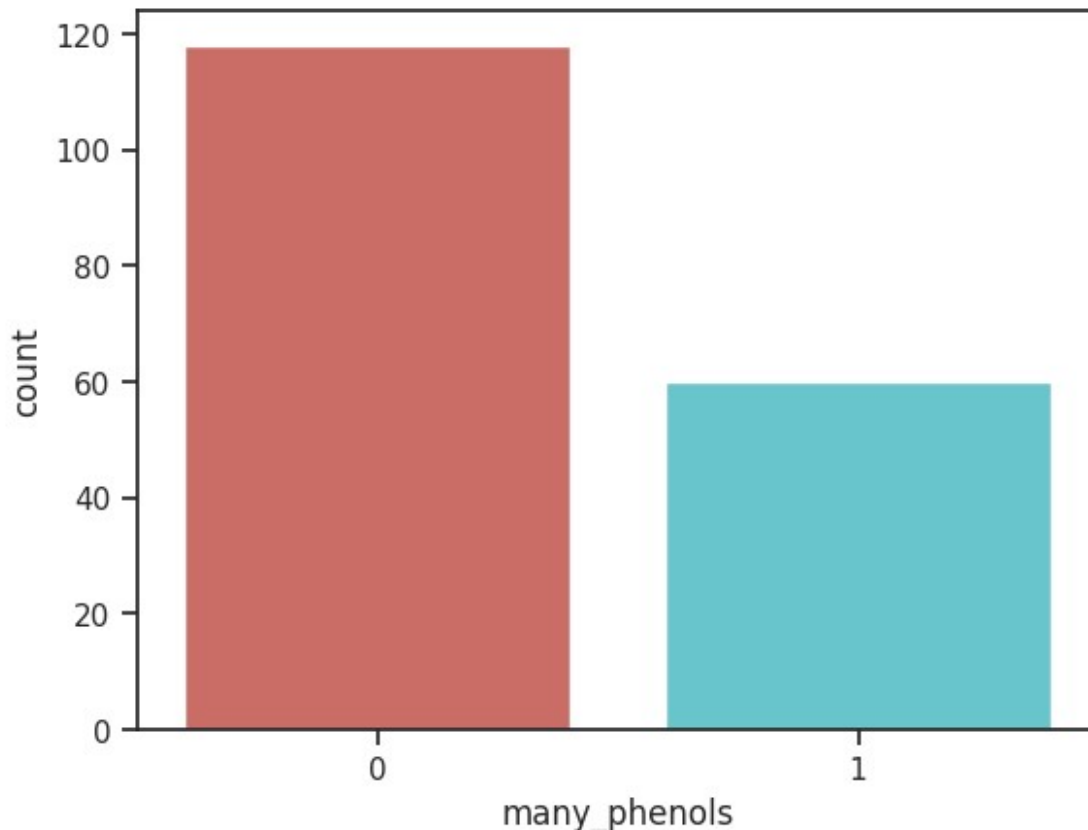
```
wine_df['many_phenols'] = np.where(wine_df['total_phenols'] >
wine_df['total_phenols'].quantile(0.65), 1, 0)
wine_df.drop(['total_phenols'], axis=1, inplace=True)
```

```
sns.countplot(x='many_phenols', data=wine_df, palette='hls')
plt.show()
```

```
<ipython-input-14-42956e5e8f46>:1: FutureWarning:
```

```
Passing `palette` without assigning `hue` is deprecated and will be
removed in v0.14.0. Assign the `x` variable to `hue` and set
`legend=False` for the same effect.
```

```
sns.countplot(x='many_phenols', data=wine_df, palette='hls')
```



```
last_column = wine_df.pop('many_phenols')
wine_df.insert(len(wine_df.columns), 'many_phenols', last_column)
X = wine_df.iloc[:, :-1].values
y = wine_df.iloc[:, -1].values

# Формирование обучающей и тестовой выборки
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size =
0.2, random_state = 1)

logreg = LogisticRegression()
logreg.fit(X_train, y_train)
y_pred_test_logreg = logreg.predict(X_test)
y_pred_train_logreg = logreg.predict(X_train)
ac1 = accuracy_score(y_train, y_pred_train_logreg),
accuracy_score(y_test, y_pred_test_logreg)
ac1

/usr/local/lib/python3.10/dist-packages/sklearn/linear_model/_logistic.py:458: ConvergenceWarning: lbfgs failed to converge
(status=1):
STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.

Increase the number of iterations (max_iter) or scale the data as
shown in:
```

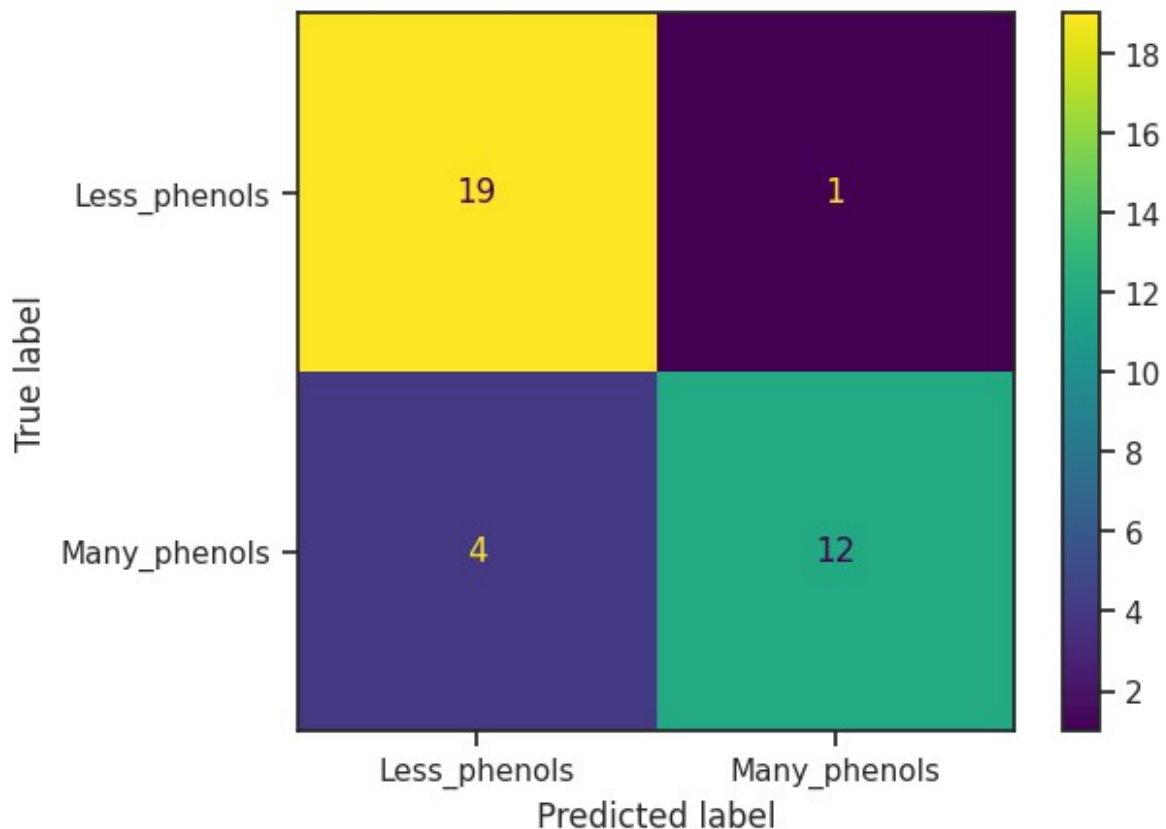
<https://scikit-learn.org/stable/modules/preprocessing.html>
Please also refer to the documentation for alternative solver options:

https://scikit-learn.org/stable/modules/linear_model.html#logistic-regression

```
n_iter_i = _check_optimize_result(  
(0.9084507042253521, 0.8611111111111112)
```

```
cm1 = confusion_matrix(y_test, y_pred_test_logreg, labels =  
logreg.classes_)  
disp = ConfusionMatrixDisplay(confusion_matrix=cm1,  
display_labels=['Less_phenols', 'Many_phenols'])  
disp.plot()
```

```
<sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at  
0x79bfef0a1ea0>
```



```
svc = SVC(kernel='poly')  
svc.fit(X_train, y_train)  
y_pred_test_svc = svc.predict(X_test)  
y_pred_train_svc = svc.predict(X_train)  
ac2 = accuracy_score(y_train, y_pred_train_svc),
```

```
accuracy_score(y_test, y_pred_test_svc)
ac2
```

```
(0.7887323943661971, 0.6666666666666666)
```

```
param_grid = {'degree': [1,2,3,4,5,6,7,8,9,10,11,12,13,14,15],
              'kernel':['poly']}
grid = GridSearchCV(SVC(), param_grid, verbose=2, scoring='accuracy')
grid.fit(X_train, y_train)
grid.best_params_
```

```
Fitting 5 folds for each of 15 candidates, totalling 75 fits
[CV] END .....degree=1, kernel=poly; total
time= 0.0s
[CV] END .....degree=1, kernel=poly; total
time= 0.0s
[CV] END .....degree=1, kernel=poly; total
time= 0.0s
[CV] END .....degree=1, kernel=poly; total
time= 0.0s
[CV] END .....degree=1, kernel=poly; total
time= 0.0s
[CV] END .....degree=2, kernel=poly; total
time= 0.0s
[CV] END .....degree=2, kernel=poly; total
time= 0.0s
[CV] END .....degree=2, kernel=poly; total
time= 0.0s
[CV] END .....degree=2, kernel=poly; total
time= 0.0s
[CV] END .....degree=2, kernel=poly; total
time= 0.0s
[CV] END .....degree=3, kernel=poly; total
time= 0.0s
[CV] END .....degree=3, kernel=poly; total
time= 0.0s
[CV] END .....degree=3, kernel=poly; total
time= 0.0s
[CV] END .....degree=3, kernel=poly; total
time= 0.0s
[CV] END .....degree=3, kernel=poly; total
time= 0.0s
[CV] END .....degree=4, kernel=poly; total
time= 0.0s
[CV] END .....degree=4, kernel=poly; total
time= 0.0s
[CV] END .....degree=4, kernel=poly; total
time= 0.0s
[CV] END .....degree=4, kernel=poly; total
time= 0.0s
```

```
[CV] END .....degree=4, kernel=poly; total
time= 0.0s
[CV] END .....degree=5, kernel=poly; total
time= 0.0s
[CV] END .....degree=5, kernel=poly; total
time= 0.0s
[CV] END .....degree=5, kernel=poly; total
time= 0.0s
[CV] END .....degree=5, kernel=poly; total
time= 0.0s
[CV] END .....degree=5, kernel=poly; total
time= 0.0s
[CV] END .....degree=6, kernel=poly; total
time= 0.0s
[CV] END .....degree=6, kernel=poly; total
time= 0.0s
[CV] END .....degree=6, kernel=poly; total
time= 0.0s
[CV] END .....degree=6, kernel=poly; total
time= 0.0s
[CV] END .....degree=6, kernel=poly; total
time= 0.0s
[CV] END .....degree=7, kernel=poly; total
time= 0.0s
[CV] END .....degree=7, kernel=poly; total
time= 0.0s
[CV] END .....degree=7, kernel=poly; total
time= 0.0s
[CV] END .....degree=7, kernel=poly; total
time= 0.0s
[CV] END .....degree=7, kernel=poly; total
time= 0.0s
[CV] END .....degree=8, kernel=poly; total
time= 0.0s
[CV] END .....degree=8, kernel=poly; total
time= 0.0s
[CV] END .....degree=8, kernel=poly; total
time= 0.0s
[CV] END .....degree=8, kernel=poly; total
time= 0.0s
[CV] END .....degree=8, kernel=poly; total
time= 0.0s
[CV] END .....degree=9, kernel=poly; total
time= 0.0s
[CV] END .....degree=9, kernel=poly; total
time= 0.0s
[CV] END .....degree=9, kernel=poly; total
time= 0.0s
[CV] END .....degree=9, kernel=poly; total
```

```
time= 0.1s
[CV] END .....degree=9, kernel=poly; total
time= 0.1s
[CV] END .....degree=10, kernel=poly; total
time= 0.0s
[CV] END .....degree=10, kernel=poly; total
time= 0.0s
[CV] END .....degree=10, kernel=poly; total
time= 0.1s
[CV] END .....degree=10, kernel=poly; total
time= 0.1s
[CV] END .....degree=10, kernel=poly; total
time= 0.1s
[CV] END .....degree=11, kernel=poly; total
time= 0.1s
[CV] END .....degree=11, kernel=poly; total
time= 0.1s
[CV] END .....degree=11, kernel=poly; total
time= 0.1s
[CV] END .....degree=11, kernel=poly; total
time= 0.1s
[CV] END .....degree=11, kernel=poly; total
time= 0.1s
[CV] END .....degree=12, kernel=poly; total
time= 0.1s
[CV] END .....degree=12, kernel=poly; total
time= 0.3s
[CV] END .....degree=12, kernel=poly; total
time= 0.1s
[CV] END .....degree=12, kernel=poly; total
time= 0.2s
[CV] END .....degree=12, kernel=poly; total
time= 0.2s
[CV] END .....degree=13, kernel=poly; total
time= 0.3s
[CV] END .....degree=13, kernel=poly; total
time= 0.2s
[CV] END .....degree=13, kernel=poly; total
time= 0.1s
[CV] END .....degree=13, kernel=poly; total
time= 0.1s
[CV] END .....degree=13, kernel=poly; total
time= 0.3s
[CV] END .....degree=14, kernel=poly; total
time= 0.4s
[CV] END .....degree=14, kernel=poly; total
time= 1.3s
[CV] END .....degree=14, kernel=poly; total
time= 0.5s
```



```
[CV] END .....degree=14, kernel=poly; total  
time= 0.3s  
[CV] END .....degree=14, kernel=poly; total  
time= 0.3s  
[CV] END .....degree=15, kernel=poly; total  
time= 1.5s  
[CV] END .....degree=15, kernel=poly; total  
time= 3.3s  
[CV] END .....degree=15, kernel=poly; total  
time= 1.8s  
[CV] END .....degree=15, kernel=poly; total  
time= 0.3s  
[CV] END .....degree=15, kernel=poly; total  
time= 1.2s
```

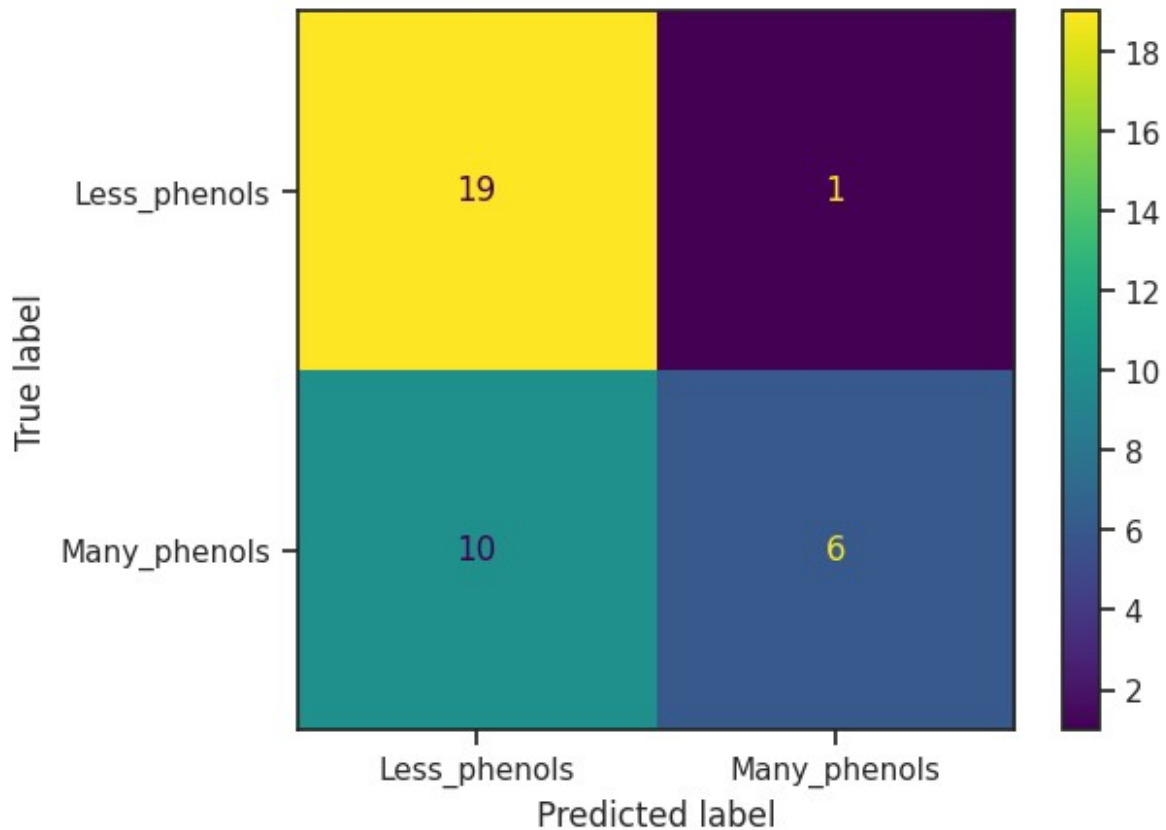
```
{'degree': 14, 'kernel': 'poly'}
```

```
svc = SVC(kernel='poly', degree=14)  
svc.fit(X_train, y_train)  
y_pred_test_svc = svc.predict(X_test)  
y_pred_train_svc = svc.predict(X_train)  
accuracy_score(y_train, y_pred_train_svc), accuracy_score(y_test,  
y_pred_test_svc)
```

```
(0.8380281690140845, 0.6944444444444444)
```

```
cm2 = confusion_matrix(y_test, y_pred_test_svc, labels = svc.classes_)  
disp = ConfusionMatrixDisplay(confusion_matrix=cm2,  
display_labels=['Less_phenols', 'Many_phenols'])  
disp.plot()
```

```
<sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at  
0x79bfeee4f940>
```

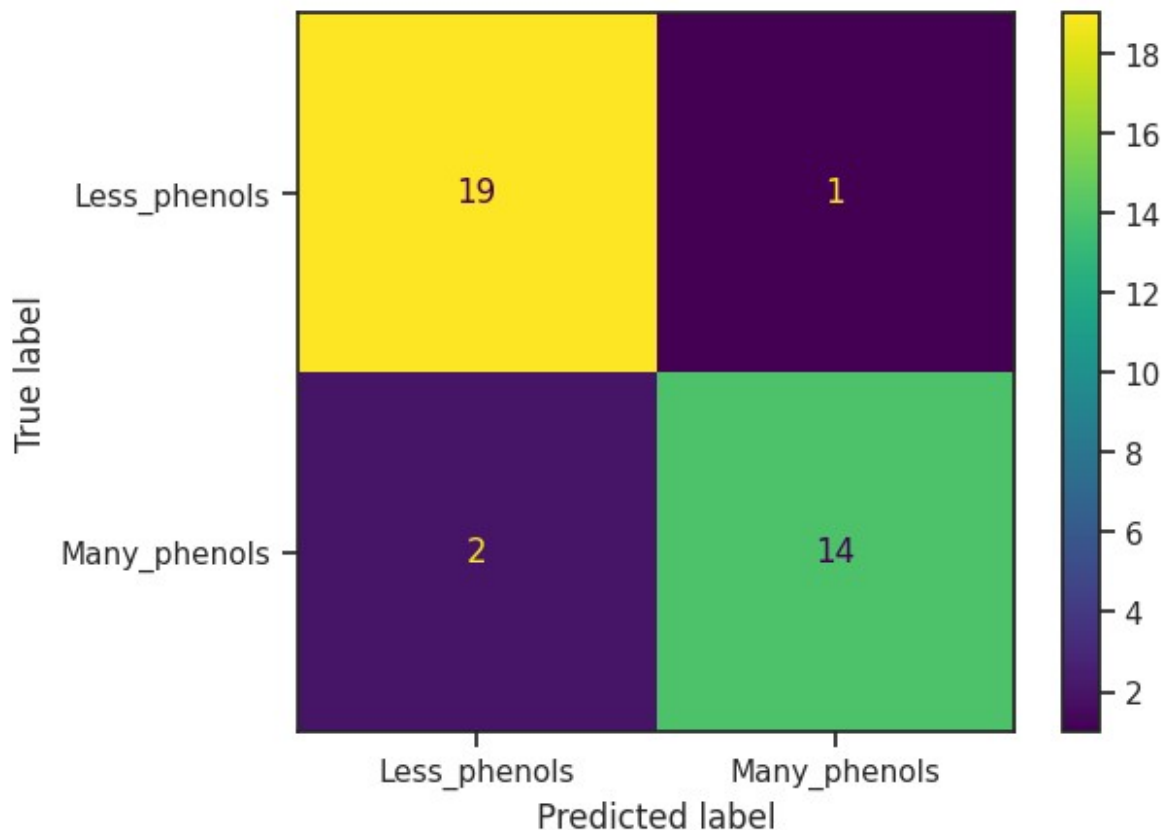


```
tree = DecisionTreeClassifier(random_state=1).fit(X_train, y_train)
y_pred_test_tree = tree.predict(X_test)
y_pred_train_tree = tree.predict(X_train)
ac3 = accuracy_score(y_train, y_pred_train_tree),
accuracy_score(y_test, y_pred_test_tree)
ac3
```

```
(1.0, 0.9166666666666666)
```

```
cm3 = confusion_matrix(y_test, y_pred_test_tree, labels =
tree.classes_)
disp = ConfusionMatrixDisplay(confusion_matrix=cm3,
display_labels=['Less_phenols', 'Many_phenols'])
disp.plot()
```

```
<sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at
0x79bfef0d75b0>
```



сравнение качества моделей по 2 метрикам

```
print('LogisticRegression: ', ac1)
```

```
print('SVM: ', ac2)
```

```
print('DecisionTreeClassifier: ', ac3)
```

```
LogisticRegression: (0.9084507042253521, 0.8611111111111112)
```

```
SVM: (0.7887323943661971, 0.6666666666666666)
```

```
DecisionTreeClassifier: (1.0, 0.9166666666666666)
```

```
fig, ax = plt.subplots(3,1)
```

```
ax[0].set_title("LogisticRegression")
```

```
ax[1].set_title("SVM")
```

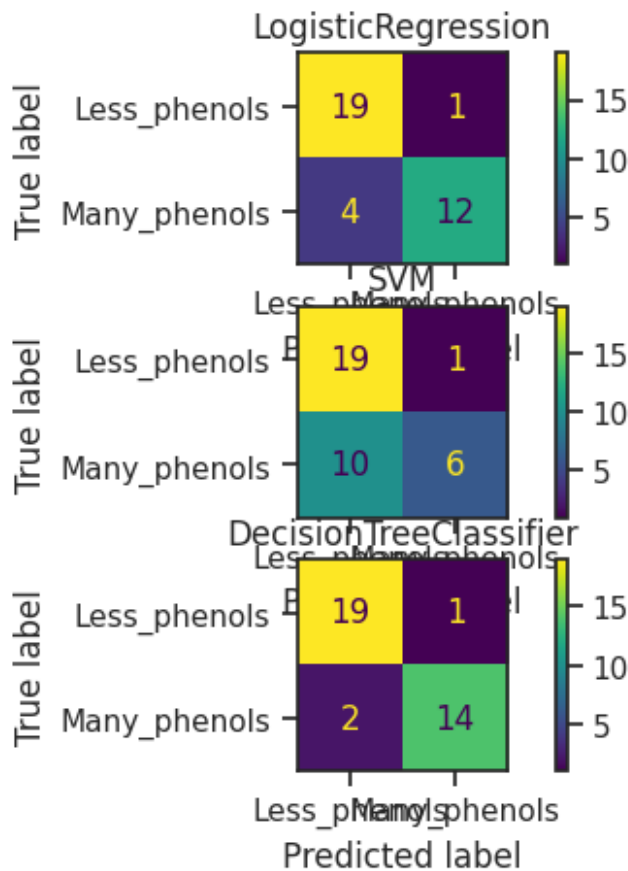
```
ax[2].set_title("DecisionTreeClassifier")
```

```
ConfusionMatrixDisplay(confusion_matrix=cm1,  
display_labels=['Less_phenols', 'Many_phenols']).plot(ax=ax[0])
```

```
ConfusionMatrixDisplay(confusion_matrix=cm2,  
display_labels=['Less_phenols', 'Many_phenols']).plot(ax=ax[1])
```

```
ConfusionMatrixDisplay(confusion_matrix=cm3,  
display_labels=['Less_phenols', 'Many_phenols']).plot(ax=ax[2])
```

```
<sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at 0x79bfeee10d30>
```



```
tree_rules = export_text(tree, feature_names=list(wine_df.iloc[:, :-1].columns))
HTML('<pre>' + tree_rules + '</pre>')
<IPython.core.display.HTML object>

dot_data = export_graphviz(tree, out_file=None,
                           feature_names=wine_df.iloc[:, :-1].columns,
                           class_names=['Less_phenols',
                                        'Many_phenols'],
                           filled=True, rounded=True,
                           special_characters=True)
graph = graphviz.Source(dot_data)
graph
```

```

from operator import itemgetter

def draw_feature_importances(tree_model, X_dataset, figsize=(18,5)):
    """
    Вывод важности признаков в виде графика
    """
    # Сортировка значений важности признаков по убыванию
    list_to_sort = list(zip(X_dataset.columns.values,
tree_model.feature_importances_))
    sorted_list = sorted(list_to_sort, key=itemgetter(1), reverse =
True)
    # Названия признаков
    labels = [x for x,_ in sorted_list]
    # Важности признаков
    data = [x for _,x in sorted_list]
    # Вывод графика
    fig, ax = plt.subplots(figsize=figsize)
    ind = np.arange(len(labels))
    plt.bar(ind, data)
    plt.xticks(ind, labels, rotation='vertical')
    # Вывод значений
    for a,b in zip(ind, data):
        plt.text(a-0.05, b+0.01, str(round(b,3)))
    plt.show()
    return labels, data

```

```
fl, fd = draw_feature_importances(tree, wine_df.iloc[:, :-1])
```

