Mushroom Dataset - 决策树

下面的示例数据需要我们通过一些蘑菇的若干属性判断这个品种是否有毒。数据链接: https://www.kaggle.com/uciml/mushroom-classification,

每个样本描述了蘑菇的 22 个属性,比如形状、气味等等响应为这个蘑菇是否可食用。

其中6513个样本做训练,1611个样本做测试。

导入工具包

```
import pandas as pd
import numpy as np
from matplotlib import pyplot as plt

from sklearn.model_selection import GridSearchCV

from sklearn.metrics import accuracy_score
from sklearn.metrics import roc_auc_score

#显示中文
plt.rcParams['font.sans-serif'] = ['Arial Unicode MS']
```

Reading the file

调用 head 函数看看每个特征的基本情况

```
dpath = "./data/"
df = pd. read_csv(dpath + "mushrooms.csv")
df. head()

In [3]:
```

In [2]:

#数据基本信息 df.info()

特征编码

特征全是类别型变量,很多模型需要数值型的输入(Logistic 回归、xgboost...),通常对类别行特征独热编码(OneHotEncoder) LableEncoder 虽然也可以将类别型变量变成数值,但结果是有序的,而颜色等特征是没有序关系,与实际不符。 决策树最好也将类型特征用独热编码转换,但这样会使得分支特别细。 LightGBM 里的决策树支持类别型特征,只需LableEncoder 编码即可 这里我们为了简单也用 LableEncoder(用 LabelEncoder 会使得特征分支少很多可能,有些情形下性能并不好,但这个例子很简单)

```
from sklearn.preprocessing import LabelEncoder
labelencoder=LabelEncoder()
for col in df. columns:
    df[col] = labelencoder.fit transform(df[col])
df. head()
import matplotlib.pyplot as plt
import seaborn as sns
%matplotlib inline
sns. countplot (x="gill-color", hue="class", data=df)
数据集是一个文件,我们自己分出一部分来做测试吧(不是校验集)
y = df['class']
X = df. drop('class', axis = 1)
from sklearn. model selection import train test split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, ra
ndom state=4)
columns = X_train.columns
体验一下分裂及 gini 计算
df1 = pd. DataFrame({'gill-color': X train['gill-color'], 'class':y trai
n})
dfl. sort values (by=['gill-color'], ascending=True)
gini=[]
for i in range (11):
    threshold = i + 0.5
    dl = df[df['gill-color'] <= threshold]</pre>
    dr = df[df['gill-color'] > threshold]
    d1 \ 0 = (d1['class'] == 0).sum()
    d1 1 = (d1['class'] == 1).sum()
    dr 0 = (dr['class'] == 0).sum()
    dr_1 = (dr['class'] == 1).sum()
    total 1 = d1 0 + d1 1
    total r = dr 0 + dr 1
    total = total_1 + total_r
    h_1 = 2.0 * dl_0/total_1 * dl_1/total_1
    h_r = 2.0 * dr_0/total_r * dr_1/total_r
    gini.append(h 1*total 1/total + h r*total r/total)
print(gini)
```

默认参数的决策树

```
from sklearn. tree import DecisionTreeClassifier
model tree = DecisionTreeClassifier()
model tree.fit(X train, y train)
y_prob = model_tree.predict_proba(X_test)[:,1] # This will give you posi
tive class prediction probabilities
y pred = np. where (y prob > 0.5, 1, 0) # This will threshold the probabil
ities to give class predictions.
model tree.score(X test, y pred)
print ('The AUC of default Desicion Tree is', roc_auc_score(y_test, y pre
d))
df = pd. DataFrame({"columns":list(columns), "importance":list(model_tree.
feature importances .T)})
df. sort values (by=['importance'], ascending=False)
plt.bar(range(len(model_tree.feature_importances_)), model_tree.feature_
importances )
plt. show()
```

决策树超参数调优

决策树的超参数有:

- 1. max_depth (树的深度) 或 max_leaf_nodes (叶子结点的数目)、
- 2. min_samples_leaf(叶子结点的最小样本数)、min_samples_split(中间结点的最小样本树)、min_weight_fraction_leaf(叶子节点的样本权重占总权重的比例)
- 3. min impurity split (最小不纯净度) 也可以调整
- 4. max features (最大特征数目)、

这个数据集的任务不难,深度设为3-10之间

```
from sklearn.tree import DecisionTreeClassifier

model_DD = DecisionTreeClassifier()

max_depth = range(3, 10, 1)
min_samples_leaf = range(5, 15, 1)
tuned_parameters = dict(max_depth=max_depth, min_samples_leaf=min_sample
s_leaf)
```

```
from sklearn.model_selection import GridSearchCV
gird = GridSearchCV (model DD, tuned parameters, scoring='roc auc', cv=10)
gird. fit (X train, y train)
print("Best: %f using %s" % (gird.best score , gird.best params ))
y_prob = gird.best_estimator_.predict_proba(X_test)[:,1] # This will giv
e you positive class prediction probabilities
y pred = np. where (y prob > 0.5, 1, 0) # This will threshold the probabil
ities to give class predictions.
gird. score (X test, y pred)
                                                              Out[50]:
print ('The AUC of GridSearchCV Desicion Tree is', roc_auc_score(y_test,
y_pred))
import graphviz
import pydotplus
from sklearn import tree
#dotfile = StringIO. StringIO()
dot_data = tree.export_graphviz(gird.best_estimator_, out_file='best_tre
e. dot', feature_names=columns, class_names='class',)
#$dot -Tpng best tree.dot -o best tree.png
test_means = gird.cv_results_[ 'mean_test_score' ]
#print(test_means. shape)
# plot results
test scores = np. array(test means).reshape(len(min samples leaf), len( m
ax depth))
for i, value in enumerate(min_samples_leaf):
    plt.plot(max_depth, test_scores[i], label= 'min_samples_leaf:' + s
tr(value))
plt.legend()
plt.xlabel('max_depth')
plt.ylabel('AUC')
plt. show()
```

```
#DD. grid scores
test_means = gird.cv_results_[ 'mean_test_score' ]
# plot results
test scores = np. array(test means).reshape(len(max depth), len(min sampl
es leaf ))
for i, value in enumerate (max depth):
    plt.plot(min samples leaf, test scores[i], label= 'max depth:' + s
tr(value))
plt. legend (loc='lower right')
plt.xlabel('min_samples_leaf')
plt.ylabel('AUC')
plt. show()
plt.bar(columns, gird.best_estimator_.feature_importances_)
plt. xticks (rotation=-80) # 设置 x 轴标签旋转角度
plt. show()
plt.barh(columns, gird.best_estimator_.feature_importances_)
#plt.xticks(rotation=-80)
                            # 设置 x 轴标签旋转角度
plt. xlabel(u'特征重要性', fontsize = 14)
plt. show()
df2 = pd. DataFrame({"columns":list(columns), "importance":list(gird.best
estimator .feature importances .T)})
df2. sort values (by=['importance'], ascending=False)
进一步根据特征重要性选择特征值
from numpy import sort
from sklearn.feature_selection import SelectFromModel
# Fit model using each importance as a threshold
thresholds = sort(gird.best estimator .feature importances)
for thresh in thresholds:
  # select features using threshold
  selection = SelectFromModel(gird.best_estimator_, threshold=thresh, pr
efit=True)
 select_X_train = selection. transform(X_train)
```

```
# train model
  selection_model = DecisionTreeClassifier()
  selection_model.fit(select_X_train, y_train)
# eval model
  select X test = selection.transform(X test)
  y pred = selection model.predict(select X test)
  predictions = [round(value) for value in y_pred]
  accuracy = accuracy score(y test, predictions)
  print ("Thresh=%. 3f, n=%d, Accuracy: %. 2f%%" % (thresh, select X train.
shape [1],
      accuracy*100.0))
最佳模型
from numpy import sort
from sklearn. feature selection import SelectFromModel
# Fit model using each importance as a threshold
thresholds = sort(gird.best estimator .feature importances)
idx = len(thresholds) - 7
thresh = thresholds[idx]
# select features using threshold
selection = SelectFromModel(gird.best estimator, threshold=thresh, pref
it=True)
select X train = selection.transform(X train)
# train model
selection model = DecisionTreeClassifier()
selection_model.fit(select_X_train, y_train)
# eval model
select X test = selection.transform(X test)
y pred = selection model.predict(select X test)
predictions = [round(value) for value in y_pred]
accuracy = accuracy score(y test, predictions)
print ("Thresh=%.3f, n=%d, Accuracy: %.2f%%" % (thresh, select_X_train.sh
ape[1], accuracy*100.0))
```

selection model. feature importances