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In [29]: import pandas as pd
from sklearn.datasets import load_breast_cancer
from sklearn.model_selection import train_test_split
from sklearn.naive_bayes import GaussianNB
from sklearn.metrics import confusion_matrix
from matplotlib import pyplot as plt
import seaborn as sns
from sklearn import metrics
from sklearn.metrics import classification_report

cancer=load_breast_cancer()
cancerdf=pd.DataFrame(cancer.data,columns=cancer.feature_names)
print(cancerdf.head()) # head() 默认显示前5行数据
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      mean radius    mean texture    mean perimeter    mean area    mean smoothness \
0      17.99      10.38      122.80      1001.0      0.11840
1      20.57      17.77      132.90      1326.0      0.08474
2      19.69      21.25      130.60      1203.0      0.10960
3      11.42      20.38       77.58       386.1      0.14250
4      20.29      14.34      135.10      1297.0      0.10030

      mean compactness    mean concavity    mean concave points    mean symmetry \
0      0.27760      0.3001      0.14710      0.2419
1      0.07864      0.0869      0.07017      0.1812
2      0.15990      0.1974      0.12790      0.2069
3      0.28390      0.2414      0.10520      0.2597
4      0.13280      0.1980      0.10430      0.1809

      mean fractal dimension    ...    worst radius    worst texture    worst perimeter \
0      0.07871    ...      25.38      17.33      184.60
1      0.05667    ...      24.99      23.41      158.80
2      0.05999    ...      23.57      25.53      152.50
3      0.09744    ...      14.91      26.50       98.87
4      0.05883    ...      22.54      16.67      152.20

      worst area    worst smoothness    worst compactness    worst concavity \
0      2019.0      0.1622      0.6656      0.7119
1      1956.0      0.1238      0.1866      0.2416
2      1709.0      0.1444      0.4245      0.4504
3       567.7      0.2098      0.8663      0.6869
4      1575.0      0.1374      0.2050      0.4000

      worst concave points    worst symmetry    worst fractal dimension
0      0.2654      0.4601      0.11890
1      0.1860      0.2750      0.08902
2      0.2430      0.3613      0.08758
3      0.2575      0.6638      0.17300
4      0.1625      0.2364      0.07678

[5 rows x 30 columns]
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In [30]: print("肿瘤的分类：",cancer['target_names'])
print("肿瘤的分类：",cancer['feature_names'])
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肿瘤的分类: ['malignant' 'benign']
肿瘤的分类: ['mean radius' 'mean texture' 'mean perimeter' 'mean area'
 'mean smoothness' 'mean compactness' 'mean concavity'
 'mean concave points' 'mean symmetry' 'mean fractal dimension'
 'radius error' 'texture error' 'perimeter error' 'area error'
 'smoothness error' 'compactness error' 'concavity error'
 'concave points error' 'symmetry error' 'fractal dimension error'
 'worst radius' 'worst texture' 'worst perimeter' 'worst area'
 'worst smoothness' 'worst compactness' 'worst concavity'
 'worst concave points' 'worst symmetry' 'worst fractal dimension']

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In [31]: x,y=cancer.data,cancer.target
x_train,x_test,y_train,y_test=train_test_split(x,y,test_size=0.2,random_state = 100)
print(x_train.shape)# 查看训练集数据形态
print(x_test.shape)# 查看测试集数据形态

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(455, 30)

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(114, 30)

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In [32]: clf=GaussianNB()
clf.fit(x_train,y_train)#对训练集进行拟合
Y_pred= clf.predict(x_test)
print("Accuracy:",metrics.accuracy_score(y_test, Y_pred))
print("Precision:",metrics.precision_score(y_test, Y_pred))
print("Recall:",metrics.recall_score(y_test, Y_pred))

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Accuracy: 0.9298246514035088
Precision: 0.9253731343283582
Recall: 0.9538461538461539

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In [33]: pred=clf.predict(x_test)
cm=confusion_matrix(pred,y_test)
sns.heatmap(cm,cmap=sns.color_palette("Blues"),annot=True,fmt='d')
plt.xlabel('real')
plt.ylabel('predict')
plt.show()

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