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1 正态性检验

1.1 单变量的多元正态性检验

单因变量正态是多因变量多元正态的必要非充分条件。

```
iris_d=iris[,1:4]

#检验每一个变量的qqplot

layout(matrix(1:8,nc=4))

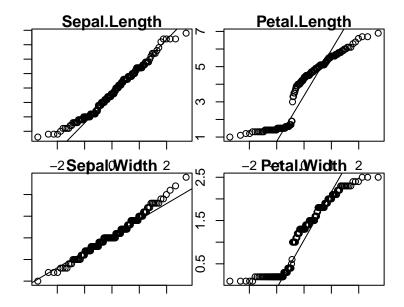
sapply(colnames(iris_d),function(x))

{qqnorm(iris_d[[x]],main=x)}

qqline(iris_d[[x]])

})
```

code 1: 单变量多元性检验



```
## $Sepal.Length
##
NULL
##
## $Sepal.Width
##
NULL
##
## $Petal.Length
##
## $Petal.Width
##
## $Petal.Width
```

Figure 1: 1.1

由上图可知, sepal.length 与 sepal.width 的概率图大致均匀分布在 y=x 线两侧, petal.length 与 petal.width 的图明显偏离 y=x 线, 左边朝直线下方弯曲, 右边朝直线上方弯曲, 具有长尾分布的特征。

1.2 二维散点图

判定指标:通过变量间的两两散点图,如果存在不服从线性关系的两个变量,也就是散点图不成直线,则说明数据不服从多元正态分布。

```
pairs(iris_d)
```

code 2: 变量之间相关性检验

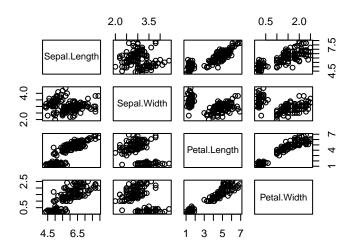


Figure 2: 1.2

1.3 马氏距离是否服从卡方分布

code 3: 马氏距离卡方分布检验

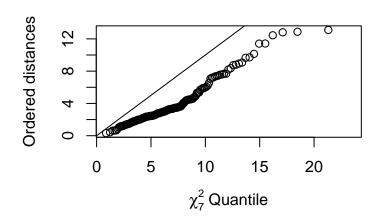


Figure 3: 1.3

从 qq 图可以看出,点距离标准线的偏离程度较大,说明可以拒绝数据服从正态分布的假设。

1.4 夏皮罗-威尔克检验 Shapiro-Wilk test

```
library(mvnormtest)
iris_d=-t(iris[,1:4])
mshapiro.test(iris_d)
```

code 4: Shapiro–Wilk test

```
##
## Shapiro-Wilk normality test
##
## data: Z
## W = 0.97935, p-value = 0.02342
```

Figure 4: 1.4

p-value<0.05, 说明可以在 0.05 的水平上拒绝数据服从多元正态分布的假设。

1.5 总结

正态性检验在样本量不大时是有必要的,当样本量足够大时,由中心极限定理,不必太过纠结正态性的问题。

2 分类

使用 python。

2.1 LDA

2.1.1 降维至三维

```
1 #导入各种程序包
2 import numpy as np
3 import pandas as pd
4 import seaborn as sns
5 import matplotlib.pyplot as plt
6 from sklearn.metrics import classification_report
7 from sklearn.discriminant_analysis import LinearDiscriminantAnalysis
8 clf = LinearDiscriminantAnalysis()
9 from sklearn.model_selection import train_test_split
10 ##划分训练集与测试集
from sklearn.datasets import load_iris
12 iris = load_iris()
13 X_train=iris.data
14 y_train=iris.target
15 X_train,X_test,y_train,y_test=train_test_split(X_train,y_train,
      test_size=0.4,random_state=0,stratify=y_train)
17 #三维 (数据可视化)
  def plot_LDA(converted_X,y):
      绘制经过 LDA 转换后的数据
20
21
      from mpl_toolkits.mplot3d import Axes3D
22
      fig=plt.figure()
23
      ax=Axes3D(fig)
24
      colors=['dodgerblue','lime','darkorange']
25
      target_names = iris.target_names
26
      markers='o*s'
27
      for target,color,marker,target_name in zip([0,1,2],colors,markers,target_names):
29
          pos=(y==target).ravel()
          X=converted_X[pos,:]
31
          ax.scatter(X[:,0], X[:,1], X[:,2],color=color,marker=marker,
33
             label=target_name)
34
      ax.legend(loc="best")
35
      fig.suptitle("After LDA")
36
  def plot_LDA1(converted_X,y):
37
38
      绘制未经 LDA 转换后的数据
39
40
```

6

```
from mpl_toolkits.mplot3d import Axes3D
42
      fig=plt.figure()
43
      ax=Axes3D(fig)
44
      colors=['dodgerblue','lime','darkorange']
45
      target_names = iris.target_names
46
      markers='o*s'
47
48
      for target,color,marker,target_name in zip([0,1,2],colors,markers,target_names):
49
          pos=(y==target).ravel()
          X=converted_X[pos,:]
          ax.scatter(X[:,0], X[:,1], X[:,2],color=color,marker=marker,
53
             label=target_name)
54
      ax.legend(loc="best")
55
      fig.suptitle("Before LDA")
56
      clf.fit(X_train,y_train)
57
58 # converted_X(150,3) 对数据降维了 权值lda.coef_(3, 4) lda.intercept_(3,)
59 converted_X=np.dot(X_train,np.transpose(clf.coef_))+clf.intercept_ # X*权值+偏置b 就是输出值
60 plot_LDA(converted_X,y_train)
61 #最初未转化的三维的情况
62 plot_LDA1(X_train,y_train)
63 y_pred = clf.predict(X_test)
64 print(y_pred)
65 print(y_test)
66 #评估
67 # 分类报告: precision/recall/fi-score/均值/分类个数
68 from sklearn.metrics import classification_report
69 target_names = ['class 0', 'class 1', 'class 2']
70 print(classification_report(y_test, y_pred, target_names=target_names))
71 #准确率
72 from sklearn.metrics import accuracy_score
73 accuracy_score(y_test, y_pred)
74 # 混淆矩阵
75 from sklearn.metrics import confusion_matrix
76 confusion_matrix(y_test, y_pred)
```

code 5: 3—dimension-LDA

运行结果:

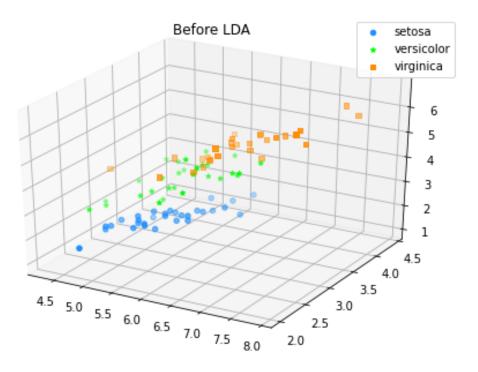


Figure 5: Before 3-dimension-LDA

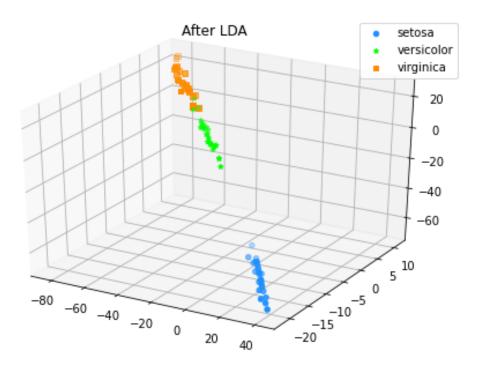


Figure 6: After 2-dimension-LDA

2.1.2 降维至二维

大致过程与上述相同,下面只给出了关键部分的代码。

1 #二维

code 6: 2—dimension-LDA

2.1.3 运行结果

Tabel 1: 评估分类报告 LDA

	precision	recall	f1-score	support
class 0	1.00	1.00	1.00	20
class 1	1.00	1.00	1.00	20
class 2	1.00	1.00	1.00	20
accuracy	1.00	60		
macro avg	1.00	1.00	1.00	60
weighted avg	1.00	1.00	1.00	60

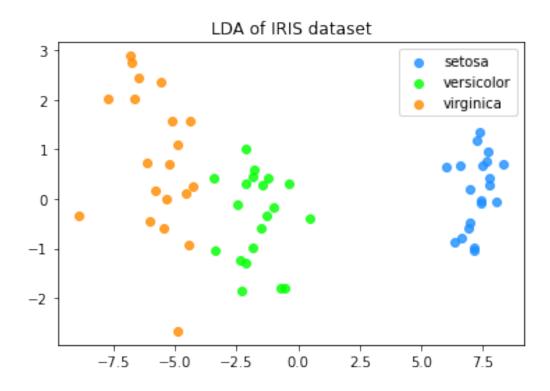


Figure 7: After 2-dimension-LDA

2.2 SVM

```
#导入各种程序包
1 import numpy as np
3 import pandas as pd
4 import seaborn as sns
5 import matplotlib.pyplot as plt
6 from sklearn.metrics import classification_report
7 from sklearn.model_selection import train_test_split
8 from sklearn.model_selection import GridSearchCV
9 from sklearn.metrics import classification_report
10 from sklearn.svm import SVC
11 #网格搜索调参
tuned_parameters = [{'kernel': ['rbf'], 'gamma': [1e-3, 1e-4],
                      'C': [1, 10, 100, 1000]},
13
                      {'kernel': ['linear'], 'C': [1, 10, 100, 1000]}]
14
15 #评分方法定义
16 scores = ['precision', 'recall']
  for score in scores:
      print("# Tuning hyper-parameters for %s" % score)
18
19
20
       # 调用 GridSearchCV, 将 SVC(), tuned_parameters, cv=5, 还有 scoring 传递进去,
21
      clf = GridSearchCV(SVC(), tuned_parameters, cv=5, scoring='%s_macro' % score)
      # 用训练集训练这个学习器 clf
```

```
clf.fit(X_train, y_train)
24
25
      print("Best parameters set found on development set:")
26
      print()
27
28
      # 再调用 clf.best_params_ 就能直接得到最好的参数搭配结果
29
      print(clf.best_params_)
30
31
      print()
32
      print("Grid scores on development set:")
      print()
      means = clf.cv_results_['mean_test_score']
      stds = clf.cv_results_['std_test_score']
36
37
      # 看一下具体的参数间不同数值的组合后得到的分数是多少
38
      for mean, std, params in zip(means, stds, clf.cv_results_['params']):
39
          print("%0.3f (+/-%0.03f) for %r"
40
                % (mean, std * 2, params))
41
      y_{-}pred=clf.predict(X_{-}test)
42
43
```

code 7: SVM

2.2.1 运行结果

Tabel 2: 评估分类报告 SVM

	precision	recall	f1-score	support
class 0	1.00	1.00	1.00	20
class 1	1.00	0.90	0.95	20
class 2	0.91	1.00	0.95	20
accuracy	0.97	60		
macro avg	0.97	0.97	0.97	60
weighted avg	0.97	0.97	0.97	60

准确率: 0.966666666666667

混淆矩阵:

 $20 \ 0 \ 0$

 $0 \ 18 \ 2$

0 0 20

2.3 总结

从上述评估结果对比可知, LDA 在 iris 数据集上的分类效果优于 SVM 的分类效果。