Exercise 6

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R Markdown

#一、分别采用至少5种分类算法对"鸢尾花"数据集进行分析,并计算其分类结果的混淆矩阵、准确率 (Accuracy)、查准率/精确率(Precision)、查全率/召回率(Recall)和F1值(F1-Score)解:

##[策略]

- (1) 读取iris数据,进行数据概览,并作预处理 (将分类变量Species因子化)
- (2) 用sample函数划分好测试集和训练集,比例为0.7:0.3
- (3) 完成模型构建,用到的包与函数如下:
- ①party包的ctree函数建立C4.5决策树模型
- ②tree包的tree函数建立Cart决策树模型
- ③C50包的C5.0函数建立C5.0决策树模型
- ④nnet包的nnet函数建立BP神经网络模型
- ⑤e1071包的naiveBayes函数建立朴素贝叶斯分类模型
- ⑥kknn包的kknn函数建立knn模型
- ⑦e1071包的svm函数建立SVM模型
- ⑧MASS包的ida函数建立IDA模型
- ⑨randomForest包的randomForest函数建立随机森林模型
- (4) 用predict函数,结合分类模型分别对训练数据及和测试数据及进行分类预测
- (5) 用cbind函数,将原始数据与预测数据结合,得到分类矩阵
- (6) 用table函数,传入真实列与预测列,分别作为actual与predictedclass的值,构建混淆矩阵 (7) 利用自定义的prf函数,计算多分类模型的Accuracy、Precision、Recall和F1-Score,对模型进行评价

##[过程|结果] 0、编写自定义函数,用于多分类问题的P、R、F值计算

```
prf <- function(test confusion)</pre>
  Accuracy <- sum(diag(test_confusion))/sum(test_confusion)
  for (i in 1:nrow(test_confusion))
    confusion <- test confusion
    classname <- colnames(confusion)[i]</pre>
    TP <- confusion[i, i]</pre>
    FN <- sum(confusion[i,])-TP
    FP <- sum(confusion[, i])-TP
    TN <- sum(diag(confusion))-confusion[i,i]</pre>
    Precision <- TP/(TP+FP)
    Recall <- TP/sum(confusion[i,])</pre>
    F1 <- 2*Precision*Recall/(Precision+Recall)
    print(paste(as.character(classname), "'s Precision:", as.character(Precision), sep="", coll
    print(paste(as.character(classname), "'s Recall:", as.character(Recall), sep="", collapse=NULL
))
    print(paste(as.character(classname), "'s F1-Score:", as.character(F1), sep="", collapse=NULL))
  print(paste('Accuracy: ',Accuracy))
```

1、决策树

```
#iris
# 设置随机种子
set. seed (1234)
#分类变量因子化
Data <- iris
Data$Species = as.factor(Data$Species)
# 随机抽取70%定义为训练数据集,30%为测试数据集
ind <- sample(2, nrow(Data), replace=TRUE, prob=c(0.7, 0.3))
traindata <- Data[ind==1, ]
testdata <- Data[ind==2, ]
#一、建立决策树模型进行预测
library(grid)
library (mvtnorm)
library (stats4)
library(modeltools)
library (sandwich)
library (strucchange)
```

```
## Loading required package: zoo
```

```
##
## Attaching package: 'zoo'
```

```
## The following objects are masked from 'package:base':
##
       as. Date, as. Date. numeric
library(zoo)
#1、C4.5决策树
#构建C4.5决策树
library (party)
ctree.model <- ctree(Species~., data=traindata)</pre>
#输出C4.5决策树图
#plot(ctree.model, type="simple")
#plot(ctree.model, type="extended")
print("***C4.5***")
## [1] "***C4.5***"
#预测C4.5决策树结果
train predict <- predict(ctree.model) #训练数据集
test_predict <- predict(ctree.model, newdata=testdata) #测试数据集
print("***traindata***")
## [1] "***traindata***"
#输出训练集的混淆矩阵,Accuracy、P、R、F1值
train_predictdata <- cbind(traindata, predictedclass = train_predict)</pre>
(train_confusion <- table(actual = traindata$Species, predictedclass = train_predict))</pre>
               predictedclass
##
                setosa versicolor virginica
## actual
##
    setosa
                    40
                               0
                    0
                              37
                                         1
##
   versicolor
    virginica
                                         31
prf(train_confusion)
## [1] "setosa's Precision:1"
## [1] "setosa's Recall:1"
## [1] "setosa's F1-Score:1"
## [1] "versicolor's Precision:0.925"
## [1] "versicolor's Recall:0.973684210526316"
## [1] "versicolor's F1-Score:0.948717948717949"
## [1] "virginica's Precision:0.96875"
## [1] "virginica's Recall:0.911764705882353"
## [1] "virginica's F1-Score:0.939393939393939"
## [1] "Accuracy: 0.964285714285714"
```

```
print("***testdata***")
## [1] "***testdata***"
#输出测试集的混淆矩阵, Accuracy、P、R、F1值
test predictdata <- cbind(testdata, predictedclass=test predict)
(test confusion <- table(actual = testdata$Species, predictedclass = test predict))</pre>
##
              predictedclass
## actual
               setosa versicolor virginica
##
                   10
                              0
   setosa
                    0
                             12
                                        ()
##
   versicolor
##
   virginica
                    0
                              2
                                       14
prf(test_confusion)
## [1] "setosa's Precision:1"
## [1] "setosa's Recall:1"
## [1] "setosa's F1-Score:1"
## [1] "versicolor's Precision: 0.857142857142857"
## [1] "versicolor's Recall:1"
## [1] "versicolor's F1-Score:0.923076923076923"
## [1] "virginica's Precision:1"
## [1] "virginica's Recall:0.875"
## [1] "Accuracy: 0.947368421052632"
#2、Cart决策树
library(tree)
#建立CART决策树模型
tree.model <- tree(Species~., data=traindata)
#决策树图输出
#plot(tree.model, type="uniform")
#text(tree.mode1)
print("***cart***")
## [1] "***cart***"
#预测结果
train_predict <- predict(tree.model, newdata=traindata, type="class")</pre>
test_predict <- predict(tree.model, newdata=testdata, type="class")</pre>
print("***traindata***")
## [1] "***traindata***"
#输出训练数据的分类结果与混淆矩阵
train_predictdata <- cbind(traindata, predictedclass=train_predict)</pre>
(train confusion <- table(actual=traindata$Species, predictedclass=train predict))
```

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```
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##
              predictedclass
## actual
               setosa versicolor virginica
##
    setosa
                   40
                               0
                    0
                              37
                                        1
##
    versicolor
##
    virginica
                    0
                               3
                                       31
#输出训练结果的Accuracy、P、R、F1值
prf(train_confusion)
## [1] "setosa's Precision:1"
## [1] "setosa's Recall:1"
## [1] "setosa's F1-Score:1"
## [1] "versicolor's Precision:0.925"
## [1] "versicolor's Recall:0.973684210526316"
## [1] "versicolor's F1-Score:0.948717948717949"
## [1] "virginica's Precision:0.96875"
## [1] "virginica's Recall:0.911764705882353"
## [1] "virginica's F1-Score:0.939393939393939"
## [1] "Accuracy: 0.964285714285714"
print("***testdata***")
## [1] "***testdata***"
#输出测试数据的分类结果与混淆矩阵
test_predictdata <- cbind(testdata, predictedclass = test_predict)</pre>
(test confusion <- table(actual=testdata$Species, predictedclass=test predict))</pre>
              predictedclass
##
## actual
               setosa versicolor virginica
##
    setosa
                   10
                              0
                                        0
                                        0
                    0
                              12
##
    versicolor
                    ()
                               2
                                       14
##
    virginica
#输出测试结果的Accuracy、P、R、F1值
prf(test confusion)
## [1] "setosa's Precision:1"
## [1] "setosa's Recall:1"
## [1] "setosa's F1-Score:1"
## [1] "versicolor's Precision:0.857142857142857"
## [1] "versicolor's Recall:1"
## [1] "versicolor's F1-Score:0.923076923076923"
## [1] "virginica's Precision:1"
## [1] "virginica's Recall:0.875"
```

[1] "Accuracy: 0.947368421052632"

```
#3、C5.0决策树
library (C50)
#构建决策树模型
c50.model <- C5.0(Species~., data=traindata)
#plot (c50. model)
print("***C5.0***")
## [1] "***C5.0***"
#预测结果
train predict <- predict(c50.model, newdata=traindata, type="class")
test_predict <- predict(c50.model, newdata=testdata, type="class")
print("***traindata***")
## [1] "***traindata***"
#输出训练数据的分类结果与混淆矩阵
train_predictdata <- cbind(traindata, predictedclass=train_predict)</pre>
(train\_confusion \leftarrow table(actual=traindata\$Species, predictedclass=train\_predict))
##
               predictedclass
## actual
                setosa versicolor virginica
                    40
                               0
##
    setosa
##
    versicolor
                    0
                               37
                                          1
                     0
                               3
                                         31
##
    virginica
#输出训练结果的Accuracy、P、R、F1值
prf(train confusion)
## [1] "setosa's Precision:1"
## [1] "setosa's Recall:1"
## [1] "setosa's F1-Score:1"
## [1] "versicolor's Precision:0.925"
## [1] "versicolor's Recall:0.973684210526316"
## [1] "versicolor's F1-Score:0.948717948717949"
## [1] "virginica's Precision:0.96875"
## [1] "virginica's Recall:0.911764705882353"
## [1] "virginica's F1-Score:0.939393939393939"
## [1] "Accuracy: 0.964285714285714"
print("***testdata***")
## [1] "***testdata***"
#输出测试数据的分类结果与混淆矩阵
test_predictdata <- cbind(testdata, predictedclass = test_predict)</pre>
(test_confusion <- table(actual=testdata$Species, predictedclass=test_predict))</pre>
```

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```
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##
              predictedclass
## actual
               setosa versicolor virginica
##
   setosa
                   10
                               0
                                        0
                    0
                              12
##
    versicolor
##
   virginica
                    0
                               2
                                        14
#输出测试结果的Accuracy、P、R、F1值
prf(test_confusion)
## [1] "setosa's Precision:1"
## [1] "setosa's Recall:1"
## [1] "setosa's F1-Score:1"
## [1] "versicolor's Precision:0.857142857142857"
## [1] "versicolor's Recall:1"
## [1] "versicolor's F1-Score:0.923076923076923"
## [1] "virginica's Precision:1"
## [1] "virginica's Recall:0.875"
## [1] "Accuracy: 0.947368421052632"
#4、ID3决策树
library(rpart)
print("***ID3***")
## [1] "***ID3***"
ID3. model <-rpart (Species~., data = traindata, method = "class")
train predict <- predict(ID3. model, data=traindata, type="class")
test_predict <- predict(ID3.model, newdata=testdata, type="class")</pre>
print("***traindata***")
## [1] "***traindata***"
train_predictdata <- cbind(traindata, predictedclass=train_predict)</pre>
(train confusion <- table(actual=traindata$Species, pretictedclass=train predict))
##
              pretictedclass
## actual
               setosa versicolor virginica
##
    setosa
                   40
                               0
                                         0
##
                    0
                              37
                                        1
    versicolor
##
   virginica
                    0
                               3
                                        31
```

```
prf(test_confusion)
```

```
## [1] "setosa's Precision:1"

## [1] "setosa's Recall:1"

## [1] "versicolor's Precision:0.857142857142857"

## [1] "versicolor's Recall:1"

## [1] "versicolor's F1-Score:0.923076923"

## [1] "virginica's Precision:1"

## [1] "virginica's Recall:0.875"

## [1] "virginica's F1-Score:0.9333333333333"

## [1] "Accuracy: 0.947368421052632"
```

```
print("***testdata***")
```

```
## [1] "***testdata***"
```

```
test_predictdata <- cbind(testdata, predictedclass=test_predict)
(test_confusion <- table(actual=testdata$Species, pretictedclass=test_predict))</pre>
```

```
##
              pretictedclass
## actual
                setosa versicolor virginica
##
                   10
                               0
   setosa
                    0
                              12
                                          0
##
   versicolor
                                2
##
    virginica
                    0
                                         14
```

```
prf(test_confusion)
```

2、BP神经网络

```
#BP神经网络
library(nnet)

#设置参数
size <- 10 #设置节点数
decay <- 0.05 #设置权值衰减系数
nnet.model <- nnet(Species~., traindata, size=size, decay=decay) #建立BP模型
```

```
## # weights: 83
## initial value 200.717468
## iter 10 value 56.810758
        20 value 21.636468
## iter
## iter 30 value 16.717897
## iter 40 value 15.223857
## iter 50 value 14.534830
## iter 60 value 14.352886
## iter 70 value 14.272249
## iter 80 value 14.209957
## iter 90 value 14.195980
## iter 100 value 14.188723
## final value 14.188723
## stopped after 100 iterations
#summary(nnet.model) #输出模型概要
```

```
#summary(nnet.model) #输出模型概要

#预测结果
train_predict <- predict(nnet.model, newdata=traindata, type="class")
test_predict <- predict(nnet.model, newdata=testdata, type="class")

#输出训练数据的分类结果与混淆矩阵
train_predictdata <- cbind(traindata, predictedclass=train_predict)
(train_confusion <- table(actual=traindata$Species, predictedclass=train_predict))
```

```
##
               predictedclass
## actual
                 setosa versicolor virginica
                     40
##
                                 ()
                                            0
   setosa
                      0
                                36
                                            2
##
    versicolor
##
     virginica
                      0
                                  1
                                           33
```

```
#输出训练结果的Accuracy、P、R、FI值
prf(train_confusion)
```

```
## [1] "setosa's Precision:1"
## [1] "setosa's Recall:1"
## [1] "setosa's F1-Score:1"
## [1] "versicolor's Precision:0.972972972973"
## [1] "versicolor's Recall:0.947368421052632"
## [1] "versicolor's F1-Score:0.96"
## [1] "virginica's Precision:0.942857142857143"
## [1] "virginica's Recall:0.970588235294118"
## [1] "virginica's F1-Score:0.956521739130435"
## [1] "Accuracy: 0.973214285714286"
```

```
#输出测试数据的分类结果与混淆矩阵
test_predictdata <- cbind(testdata,predictedclass = test_predict)
(test_confusion <- table(actual=testdata$Species, predictedclass=test_predict))
```

```
##
                predictedclass
## actual
                 setosa versicolor virginica
##
    setosa
                     10
                                  ()
                                            0
                                 12
##
     versicolor
                      0
                                           16
##
    virginica
                      0
                                  0
```

```
#输出测试结果的Accuracy、P、R、F1值
prf(test_confusion)
```

```
## [1] "setosa's Precision:1"
## [1] "setosa's Recall:1"
## [1] "versicolor's Precision:1"
## [1] "versicolor's Recall:1"
## [1] "versicolor's F1-Score:1"
## [1] "virginica's Precision:1"
## [1] "virginica's Recall:1"
## [1] "virginica's F1-Score:1"
## [1] "Accuracy: 1"
```

3、贝叶斯分类

```
#利用e1071包的naiveBayes函数建立朴素贝叶斯分类模型
library(e1071)
naiveBayes.model <- naiveBayes(Species~., data=traindata)

# 预测结果
train_predict <- predict(naiveBayes.model, newdata = traindata) # 训练数据集
test_predict <- predict(naiveBayes.model, newdata = testdata) # 测试数据集
# 输出训练数据的分类结果、混淆矩阵
train_predictdata <- cbind(traindata, predictedclass = train_predict)
(train_confusion <- table(actual = traindata$Species, predictedclass = train_predict))
```

```
##
                predictedclass
## actual
                 setosa versicolor virginica
##
    setosa
                     40
                                  0
                      0
                                 36
                                            2
##
    versicolor
                      0
                                  2
                                           32
##
     virginica
```

```
##输出训练结果的Accuracy、P、R、F1值
prf(train_confusion)
```

```
## [1] "setosa's Precision:1"

## [1] "setosa's Recall:1"

## [1] "setosa's F1-Score:1"

## [1] "versicolor's Precision:0.947368421052632"

## [1] "versicolor's Recall:0.947368421052632"

## [1] "versicolor's F1-Score:0.947368421052632"

## [1] "virginica's Precision:0.941176470588235"

## [1] "virginica's Recall:0.941176470588235"

## [1] "virginica's F1-Score:0.941176470588235"

## [1] "Accuracy: 0.964285714285714"
```

```
# 输出测试数据的分类结果、混淆矩阵
test_predictdata <- cbind(testdata, predictedclass = test_predict)
(test_confusion <- table(actual = testdata$Species, predictedclass = test_predict))
```

```
##
                predictedclass
                 setosa versicolor virginica
## actual
##
                     10
                                 0
    setosa
                                 12
                                            0
##
    versicolor
                      0
                      0
                                  2
                                           14
##
     virginica
```

```
# 输出测试数据的Precision、Recall、Accuracy以及F1-SCOREs
prf(test_confusion)
```

4、KNN

```
#载入数据
Data <- iris
Data$Species <- as. factor(Data$Species)
set. seed (123)
#随机抽取70%定义为训练数据集,30%为测试数据集
ind <- sample(2, nrow(Data), replace=TRUE, prob=c(0.7, 0.3))
traindata <- Data[ind==1, ]
testdata <- Data[ind==2, ]</pre>
#使用kknn函数构建knn模型
library (kknn)
kknn.model <- kknn(Species~., train=traindata, test=traindata, k=5) #训练数据
kknn.model2 <- kknn(Species~., train=traindata, test=testdata, k=5) #测试数据
#预测结果
train_predict <- predict(kknn.model) #训练
test predict <- predict(kknn.model2) #测试
#输出训练集的混淆矩阵, Accuracy、P、R、F1值
(train_confusion <- table(actual = traindata$Species, predictedclass=train_predict))</pre>
```

```
##
                predictedclass
## actual
                 setosa versicolor virginica
##
     setosa
                     35
                                 0
                                            0
                                            0
                      0
                                 36
##
    versicolor
                      0
                                 0
                                           35
##
     virginica
```

prf(train_confusion)

```
## [1] "setosa's Precision:1"
## [1] "setosa's Recall:1"
## [1] "versicolor's Precision:1"
## [1] "versicolor's Recall:1"
## [1] "versicolor's F1-Score:1"
## [1] "virginica's Precision:1"
## [1] "virginica's Recall:1"
## [1] "virginica's Recall:1"
## [1] "virginica's F1-Score:1"
## [1] "Accuracy: 1"
```

#输出测试集的混淆矩阵,Accuracy、P、R、F1值 (test_confusion <- table(actual=testdata\$Species,predictedclass=test_predict))

```
##
               predictedclass
                setosa versicolor virginica
## actual
                     15
                                 0
                                            0
##
    setosa
##
    versicolor
                      0
                                10
                                            4
##
     virginica
                      0
                                 2
                                           13
```

prf(test confusion)

5、利用SVM模型进行分类与预测

```
# 利用SVM建模
library(e1071)
svm.model <- svm(Species~., data = traindata)

#预测结果
train_predict <- predict(svm.model, newdata = traindata, type="class") # 训练数据集
test_predict <- predict(svm.model, newdata = testdata, type="class") #测试数据集

#输出训练集的混淆矩阵, Accuracy、P、R、F1值
(train_confusion <- table(actual = traindata$Species, predictedclass=train_predict))
```

```
##
                predictedclass
## actual
                 setosa versicolor virginica
##
     setosa
                     35
                                  0
                                             1
                      0
                                 35
##
     versicolor
##
                                            35
     virginica
                      0
                                  0
```

```
prf(train_confusion)
```

```
## [1] "setosa's Precision:1"

## [1] "setosa's Recall:1"

## [1] "setosa's F1-Score:1"

## [1] "versicolor's Precision:1"

## [1] "versicolor's Recall:0.97222222222222"

## [1] "versicolor's F1-Score:0.985915492957746"

## [1] "virginica's Precision:0.97222222222222"

## [1] "virginica's Recall:1"

## [1] "virginica's F1-Score:0.985915492957746"

## [1] "Accuracy: 0.990566037735849"
```

```
#输出测试集的混淆矩阵,Accuracy、P、R、F1值
(test_confusion <- table(actual=testdata$Species,predictedclass=test_predict))
```

```
##
                predictedclass
                 setosa versicolor virginica
## actual
##
     setosa
                     15
                                  0
                                             0
##
     versicolor
                      0
                                 11
                                             3
                      0
##
     virginica
                                  1
                                            14
```

prf(test confusion)

6、利用LDA模型分类预测

```
# 建立1da分类模型
library (MASS)
lda.model <- lda(Species ~., data = traindata)
#预测结果
train_predict <- predict(lda.model, newdata = traindata, type = "class") # 训练数据集
test predict <- predict(lda.model, newdata = testdata, type = "class") #测试数据集
#输出训练集的混淆矩阵, Accuracy、P、R、F1值
(train confusion <- table(actual = traindata$Species, predictedclass=train predict$class))
##
             predictedclass
## actual
              setosa versicolor virginica
                            0
##
                 35
    setosa
##
                  0
                           36
                                     ()
   versicolor
                  0
                            0
                                    35
##
    virginica
prf(train_confusion)
## [1] "setosa's Precision:1"
## [1] "setosa's Recall:1"
## [1] "setosa's F1-Score:1"
## [1] "versicolor's Precision:1"
## [1] "versicolor's Recall:1"
## [1] "versicolor's F1-Score:1"
## [1] "virginica's Precision:1"
## [1] "virginica's Recall:1"
## [1] "virginica's F1-Score:1"
## [1] "Accuracy: 1"
#输出测试集的混淆矩阵, Accuracy、P、R、F1值
(test confusion <- table(actual=testdata$Species, predictedclass=test predict$class))</pre>
##
             predictedclass
              setosa versicolor virginica
## actual
##
    setosa
                 15
                            0
                  0
                           13
                                     1
##
    versicolor
                  0
                                    14
##
    virginica
                            1
prf(test_confusion)
## [1] "setosa's Precision:1"
## [1] "setosa's Recall:1"
## [1] "setosa's F1-Score:1"
## [1] "versicolor's Precision:0.928571428571429"
## [1] "versicolor's Recall:0.928571428571429"
## [1] "versicolor's F1-Score:0.928571428571429"
## [1] "Accuracy: 0.954545454545455"
```

7、利用随机森林模型分类预测

```
#建立randomForest模型
library(randomForest)
## randomForest 4.6-14
## Type rfNews() to see new features/changes/bug fixes.
randomForest.model <- randomForest(Species ~ ., data = traindata)
# 预测结果
test predict <- predict(randomForest.model, newdata = testdata) #测试数据集
# 输出训练数据的混淆矩阵
(train_confusion <- randomForest.model$confusion)</pre>
##
             setosa versicolor virginica class.error
                             0
                                       0 0.00000000
## setosa
                 35
                  0
                            35
                                       1 0.02777778
## versicolor
                  0
                             2
                                      33 0.05714286
## virginica
prf(train confusion)
## [1] "setosa's Precision:1"
## [1] "setosa's Recall:1"
## [1] "setosa's F1-Score:1"
## [1] "versicolor's Precision:0.945945945945946"
## [1] "versicolor's Recall:0.971472629144179"
## [1] "versicolor's F1-Score:0.958539368581209"
## [1] "virginica's Precision:0.970588235294118"
## [1] "virginica's Recall:0.941320293398533"
## [1] "virginica's F1-Score:0.955730244104262"
## [1] "Accuracy: 0.970920272019272"
# 输出测试数据的混淆矩阵
(test confusion <- table(actual = testdata$Species, predictedclass = test predict))</pre>
##
              predictedclass
## actual
               setosa versicolor virginica
##
                   15
                               0
   setosa
##
   versicolor
                    0
                              11
                                         3
    virginica
                    0
                               1
                                        14
```

```
prf(test confusion)
```

```
## [1] "setosa's Precision:1"
## [1] "setosa's Recall:1"
## [1] "setosa's F1-Score:1"
## [1] "versicolor's Precision:0.916666666666667"
## [1] "versicolor's Recall:0.785714285714286"
## [1] "versicolor's F1-Score:0.846153846153846"
## [1] "virginica's Precision:0.823529411764706"
## [1] "virginica's Recall:0.93333333333333"
## [1] "virginica's F1-Score:0.875"
## [1] "Accuracy: 0.909090909090909"
```

#二、观察分类结果中的误差,尝试分析误差产生的原因与改进措施。

- 1、决策树(ID3)(1)模型概述以信息增益度量属性选择,选择分裂后信息增益最大的属性进行分裂(2)误差分析 实验结果表明,ID3算法对于iris数据集的分类精度较高(accuracy为0.947),而且对于traindata和 testdata的评估指数几乎一致,模型训练效果较好,存在的误差主要在versicolor与virginica两类中,误差产生的原因可能是ID3算法用信息增益过分偏向取值多的属性。(3)改进措施 利用C4.5算法,在生成过程中用信息增益比来选择特征;同时在树的构造过程中进行剪枝。
- 2、BP神经网络 (1) 模型概述 利用输出后的误差来估计输出层的直接前导层的误差,再用这个误差估计更前一层的误差,如此一层一层的反传下去,就获得了所有其他各层的误差估计。 (2) 误差分析 在iris数据集中,BP神经网络的整体精度(Accuracy)高于决策树,其中train数据集出现部分少量的误差,test则完全命中,训练集中的误差可能与设置的节点数与权重衰减有关。 (3) 改进措施 不断调整节点数和权重衰减系数,"炼丹"
- 3、朴素贝叶斯 (1) 模型概述 确定属性特征,获取训练样本,对每个类别计算P (yi) ,对每个特征属性计算所有划分的条件概率,对每个类别根据先验概率得出后验,取最大值确定分类 (2) 误差分析 朴素贝叶斯模型在iris数据集中,训练集的精度高于测试数据集,误差可能来源于其对于类条件独立的假设,未考虑到不同属性之间的相关性。 (3) 改进措施 使用半朴素贝叶斯分类器,放松一定的独立性假定
- 4、SVM (1)模型概述 在线性可分以及线性不可分的情况下,获得最优分类面 (2)误差分析 在iris数据集中,SVM模型之于train数据集的精度高于test数据集,误差主要在于"versicolor"和"virginica"两类中,原因可能在于①不能直接适用于多类线性不可分②容易被偏性样本误导 (3)改进措施 调整模型,选用不同的核函数拟合
- 5、KNN (1) 模型概述 根据k近邻对未知元组xq的距离贡献,计算预测值 (2) 误差分析 相对而言误差较大,尤其是测试集数据,原因可能在于KNN算法容易产生维数灾难,相邻距离可能由不想关的属性主导,从而降低模型精度 (3) 改进措施 通过轴拉伸,删除不相关的属性
- 6、LDA算法 (1) 模型概述 给定训练样例集,设法将样例投影到一条直线上,使得同类样例的投影点尽可能接近,异样样例的投影点尽可能远离;在对新样本进行分类时,将其投影到同样的直线上,再根据投影点的位置来确定新样本的类别。 (2) 误差分析 LDA算法比较朴素,但是对于iris数据集而言分类精度较高。
- 7、随机森林 (1) 模型概述 它基于决策树集合的多数投票来进行预测。单个决策树往往是不稳定的,不能提供稳定的预测。而通过许多树的预测,随机森林方法能更加稳定,且与其他分类器相比具有更好的性能。 (2) 误差分析 在针对iris数据集的分类中,随机森林模型精度适中,train的精度大于test的精度,误差原因比较难解释,因为随机森林模型过程是难以解释的黑箱,不过可能与样本的分布有关。
- (3) 改进措施 调整模型参数: 随机森林中树的数量, 以及用于分裂的随机采样的基因数量