

Naive Bayes Classifier in R

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R 中的包 `e1071` 提供了运用朴素贝叶斯分类器的教程: <http://cran.r-project.org/web/packages/e1071/e1071.pdf>

(一) 输入数据都是分类型数据

首先安装并载入这个包:

```
# install.packages('e1071', dependencies = TRUE)
library(e1071)
```

所使用的数据集是包 `mlbench` 中的 `HouseVotes84` 数据集。

```
data(HouseVotes84, package = "mlbench")
```

这个数据集是 1984 年美国国会的投票记录。共有 435 个观测值和 17 个分类型变量, 其中的 `Class` 变量记录的是投票者是共和党 (republican) 还是民主党 (democrat), 剩下的 16 个分类变量记录的是这 435 个投票者对 16 个问题的投票结果, 同意是 “y”, 反对是 “n”, 由于种种原因没有投票记录的话是缺失值 “NA”。可以用 `structure` 函数看一下这个数据集的结构:

```
str(HouseVotes84)

## 'data.frame':   435 obs. of  17 variables:
##  $ Class: Factor w/ 2 levels "democrat","republican": 2 2 1 1 1 1 1 1 2 2 1 ...
##  $ V1    : Factor w/ 2 levels "n","y": 1 1 NA 1 2 1 1 1 1 2 ...
##  $ V2    : Factor w/ 2 levels "n","y": 2 2 2 2 2 2 2 2 2 2 ...
##  $ V3    : Factor w/ 2 levels "n","y": 1 1 2 2 2 2 1 1 1 2 ...
##  $ V4    : Factor w/ 2 levels "n","y": 2 2 NA 1 1 1 2 2 2 1 ...
##  $ V5    : Factor w/ 2 levels "n","y": 2 2 2 NA 2 2 2 2 2 1 ...
##  $ V6    : Factor w/ 2 levels "n","y": 2 2 2 2 2 2 2 2 2 1 ...
##  $ V7    : Factor w/ 2 levels "n","y": 1 1 1 1 1 1 1 1 1 2 ...
##  $ V8    : Factor w/ 2 levels "n","y": 1 1 1 1 1 1 1 1 1 2 ...
##  $ V9    : Factor w/ 2 levels "n","y": 1 1 1 1 1 1 1 1 1 2 ...
##  $ V10   : Factor w/ 2 levels "n","y": 2 1 1 1 1 1 1 1 1 1 ...
##  $ V11   : Factor w/ 2 levels "n","y": NA 1 2 2 2 1 1 1 1 1 ...
##  $ V12   : Factor w/ 2 levels "n","y": 2 2 1 1 NA 1 1 1 2 1 ...
```

```
## $ V13 : Factor w/ 2 levels "n","y": 2 2 2 2 2 2 NA 2 2 1 ...
## $ V14 : Factor w/ 2 levels "n","y": 2 2 2 1 2 2 2 2 2 1 ...
## $ V15 : Factor w/ 2 levels "n","y": 1 1 1 1 2 2 2 NA 1 NA ...
## $ V16 : Factor w/ 2 levels "n","y": 2 NA 1 2 2 2 2 2 2 NA ...
```

我们的目标是根据对这 16 个项目的投票结果通过朴素贝叶斯分类器预测投票者是共和党还是民主党。首先抽取 60% 的数据作为训练集，剩下的作为测试集：

```
set.seed(2016)
n = nrow(HouseVotes84) # sample size
p = 0.6 # proportion of training set
idx = sample(n, n*p, replace = FALSE) # index set for training set
# training set
train = HouseVotes84[idx, ]
# test set
test = HouseVotes84[-idx, ]
```

然后用朴素贝叶斯分类器进行训练，需要用到的函数是 `naiveBayes`

```
# train a naive bayes model
model = naiveBayes(Class~., data = train)
```

最后用我们的模型进行预测：

```
# make predictions
result = predict(model, test[, -1], type = "class")
```

在 `predict` 函数中，`type` 变量可以设为 `class` 或 `raw`，前者得到的结果是预测的类别，后者得到的是各个类别的后验概率。默认值为 `class`。

看一下预测的准确率：

```
# confusion matrix
table(result, test$Class)
```

```
##
## result      democrat republican
## democrat      89             5
## republican    13            67
```

```
prop.table(table(result, test$Class))
```

```
##  
## result      democrat republican  
## democrat  0.51149425 0.02873563  
## republican 0.07471264 0.38505747
```

```
mean(result == test$Class) # accuracy
```

```
## [1] 0.8965517
```

准确率达到 $(89 + 67)/(89 + 67 + 5 + 13) = 90\%$ 。

(二) 输入数据都是连续型数据

我们使用的数据是 `kernlab` 包中的 `spam` 数据集。该数据集搜集了 4601 份电子邮件，并人工分类为垃圾邮件 (spam) 和非垃圾邮件 (non-spam)，记录在数据表最后一列 `type` 内。并记录了这些邮件中 57 个单词或符号的频率，作为 57 个特征。可以载入数据，并看一下数据的形式：

```
# install.packages("kernlab")
data(spam, package = "kernlab")
head(spam, 4)
```

```
##  make address  all num3d  our over remove internet order mail receive
## 1 0.00    0.64 0.64    0 0.32 0.00    0.00    0.00 0.00 0.00    0.00
## 2 0.21    0.28 0.50    0 0.14 0.28    0.21    0.07 0.00 0.94    0.21
## 3 0.06    0.00 0.71    0 1.23 0.19    0.19    0.12 0.64 0.25    0.38
## 4 0.00    0.00 0.00    0 0.63 0.00    0.31    0.63 0.31 0.63    0.31
##  will people report addresses free business email  you credit your font
## 1 0.64    0.00 0.00    0.00 0.32    0.00 1.29 1.93 0.00 0.96    0
## 2 0.79    0.65 0.21    0.14 0.14    0.07 0.28 3.47 0.00 1.59    0
## 3 0.45    0.12 0.00    1.75 0.06    0.06 1.03 1.36 0.32 0.51    0
## 4 0.31    0.31 0.00    0.00 0.31    0.00 0.00 3.18 0.00 0.31    0
##  num000 money hp hpl george num650 lab labs telnet num857 data num415
## 1 0.00 0.00 0 0 0 0 0 0 0 0 0 0
## 2 0.43 0.43 0 0 0 0 0 0 0 0 0 0
## 3 1.16 0.06 0 0 0 0 0 0 0 0 0 0
## 4 0.00 0.00 0 0 0 0 0 0 0 0 0 0
##  num85 technology num1999 parts pm direct cs meeting original project
## 1 0 0 0.00 0 0 0.00 0 0 0.00 0
## 2 0 0 0.07 0 0 0.00 0 0 0.00 0
## 3 0 0 0.00 0 0 0.06 0 0 0.12 0
## 4 0 0 0.00 0 0 0.00 0 0 0.00 0
##  re  edu table conference charSemicolon charRoundbracket
## 1 0.00 0.00 0 0 0.00 0.000
## 2 0.00 0.00 0 0 0.00 0.132
## 3 0.06 0.06 0 0 0.01 0.143
## 4 0.00 0.00 0 0 0.00 0.137
##  charSquarebracket charExclamation charDollar charHash capitalAve
## 1 0 0.778 0.000 0.000 3.756
## 2 0 0.372 0.180 0.048 5.114
## 3 0 0.276 0.184 0.010 9.821
## 4 0 0.137 0.000 0.000 3.537
```

```
##   capitalLong capitalTotal type
## 1          61          278 spam
## 2         101         1028 spam
## 3         485         2259 spam
## 4          40          191 spam
```

仍旧同之前一样，将数据分为训练集和测试集，然后用朴素贝叶斯模型进行训练：

```
set.seed(2016)
n = nrow(spam) # sample size
p = 0.6 # proportion of training set
idx = sample(n, n*p, replace = FALSE) # index set for training set
# training set
train = spam[idx, ]
# test set
test = spam[-idx, ]
# train a naive bayes model
model = naiveBayes(type~., data = train)
# make predictions
result = predict(model, test[, -ncol(spam)], type = "class")
```

最后看一下准确率

```
# confusion matrix
table(result, test$type)
```

```
##
## result      nonspam spam
## nonspam      639   49
## spam         471  682
```

```
prop.table(table(result, test$type))
```

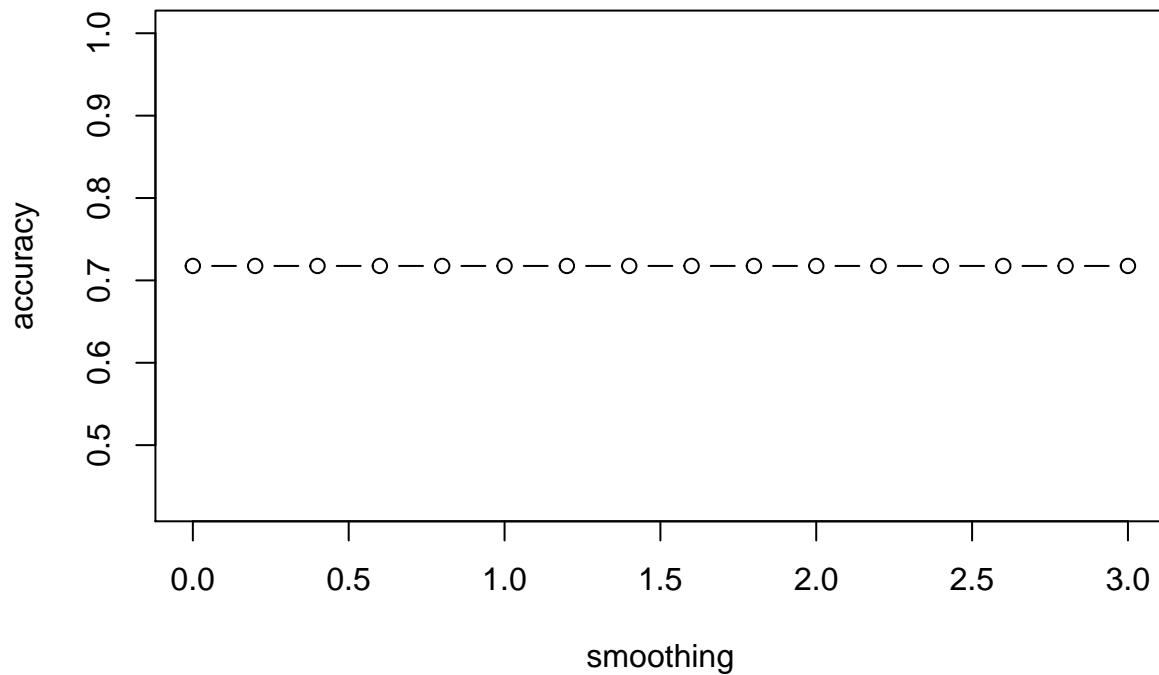
```
##
## result      nonspam      spam
## nonspam 0.34709397 0.02661597
## spam    0.25583922 0.37045084
```

```
mean(result == test$type) # accuracy
```

```
## [1] 0.7175448
```

准确率并不高，只有 72%。我们还可以看一下不同的 *smoothing* 参数对预测效果的影响

```
s = seq(from = 0, to = 3, by = 0.2)
accuracy = numeric(length(s))
for(i in seq_along(s)){
  model = naiveBayes(type~., data = train, laplace = s[i])
  result = predict(model, test[, -ncol(spam)], type = "class")
  accuracy[i] = mean(result == test$type)
}
plot(s, accuracy, type = "b", xlab = "smoothing")
```



可以看出对这个数据集而言，*smoothing* 与否并没有太大的影响。

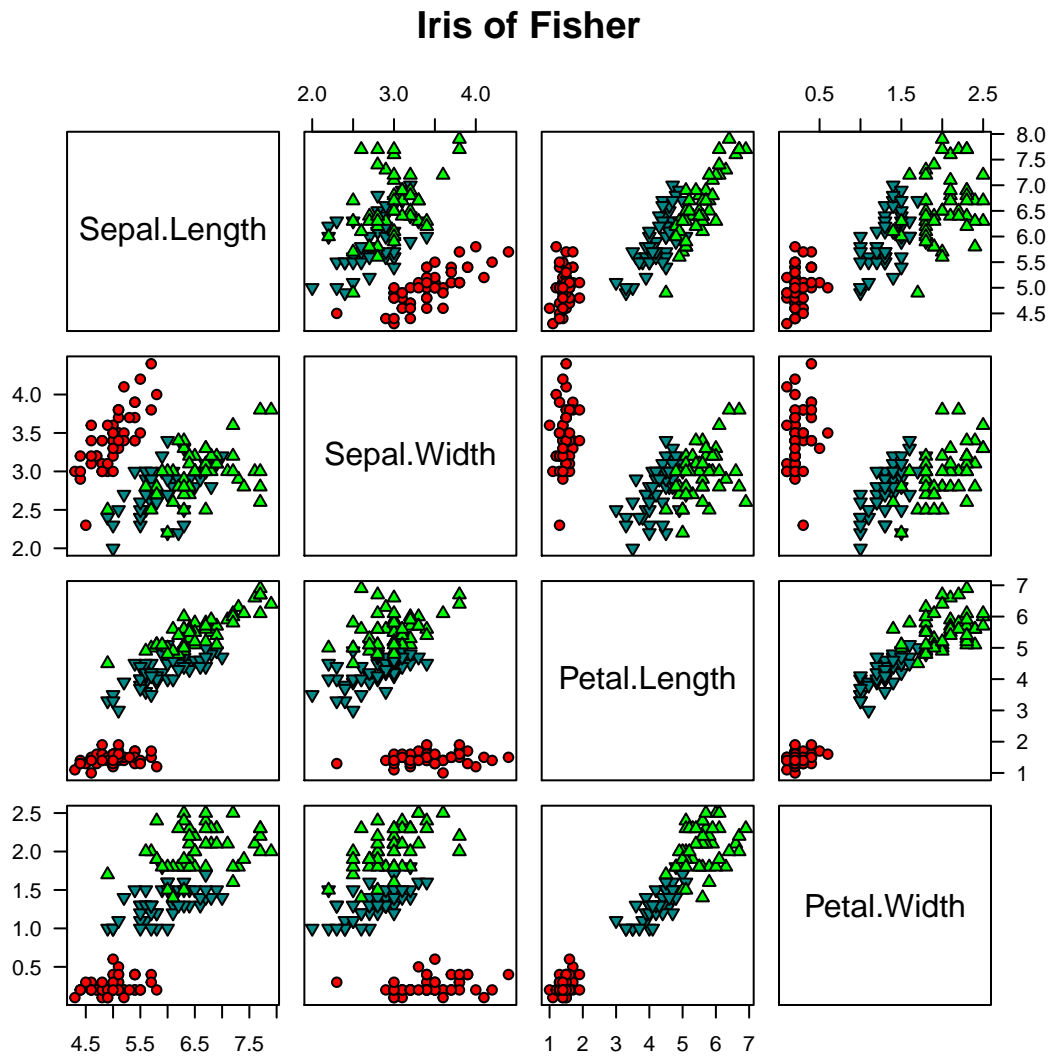
用 caret 包进行朴素贝叶斯分类

我们使用的是 R 自带的 `iris` 数据集。`iris` 给出了三种类型的鸢尾花 `Iris setosa`, `versicolor` 和 `virginica` 在四种特征变量上的差异。可以看一下这个数据集，并画出散点图：

```
summary(iris)
```

```
##      Sepal.Length      Sepal.Width      Petal.Length      Petal.Width
## Min.       :4.300    Min.       :2.000    Min.       :1.000    Min.       :0.100
## 1st Qu.:5.100    1st Qu.:2.800    1st Qu.:1.600    1st Qu.:0.300
## Median :5.800    Median :3.000    Median :4.350    Median :1.300
## Mean   :5.843    Mean   :3.057    Mean   :3.758    Mean   :1.199
## 3rd Qu.:6.400    3rd Qu.:3.300    3rd Qu.:5.100    3rd Qu.:1.800
## Max.   :7.900    Max.   :4.400    Max.   :6.900    Max.   :2.500
##           Species
## setosa      :50
## versicolor:50
## virginica  :50
##
##
##
```

```
pairs(iris[1:4], main = "Iris of Fisher",
      pch = c(21, 25, 24)[iris$Species],
      bg = c("red", "cyan4", "green")[iris$Species], las = 1)
```



载入 `caret` 包。`caret` 其实使用的是 `klaR` 包来进行朴素贝叶斯分类的。将数据分为训练集和测试集：

```
library(caret)
set.seed(2016)
# 60%/40% train/test split of the dataset
idx = createDataPartition(y = iris$Species,
                           p = .6,
                           list = FALSE)

# training set
training = iris[idx, ]
# test set
test = iris[-idx, ]
```


然后用朴素贝叶斯模型进行训练，只需要将参数设为"nb" 即可。选用了 10 重交叉验证：

```
# train a naive bayes model
model = train(training[, -5], training[, 5], "nb",
              trControl = trainControl(method = "cv", number = 10))
```

用最终得到的模型在测试集上进行预测：

```
# make predictions
result = predict(model$finalModel, test[, -5])$class
confusionMatrix(result, test[, 5])
```

```
## Confusion Matrix and Statistics
##
##              Reference
## Prediction  setosa versicolor virginica
##   setosa      20           0           0
##   versicolor   0          19           1
##   virginica    0           1          19
##
## Overall Statistics
##
##              Accuracy : 0.9667
##              95% CI : (0.8847, 0.9959)
##   No Information Rate : 0.3333
##   P-Value [Acc > NIR] : < 2.2e-16
##
##              Kappa : 0.95
##   McNemar's Test P-Value : NA
##
## Statistics by Class:
##
##              Class: setosa Class: versicolor Class: virginica
## Sensitivity          1.0000          0.9500          0.9500
## Specificity          1.0000          0.9750          0.9750
## Pos Pred Value       1.0000          0.9500          0.9500
## Neg Pred Value       1.0000          0.9750          0.9750
## Prevalence           0.3333          0.3333          0.3333
## Detection Rate       0.3333          0.3167          0.3167
## Detection Prevalence 0.3333          0.3333          0.3333
## Balanced Accuracy    1.0000          0.9625          0.9625
```

```
mean(result == test[, 5]) # accuracy
```

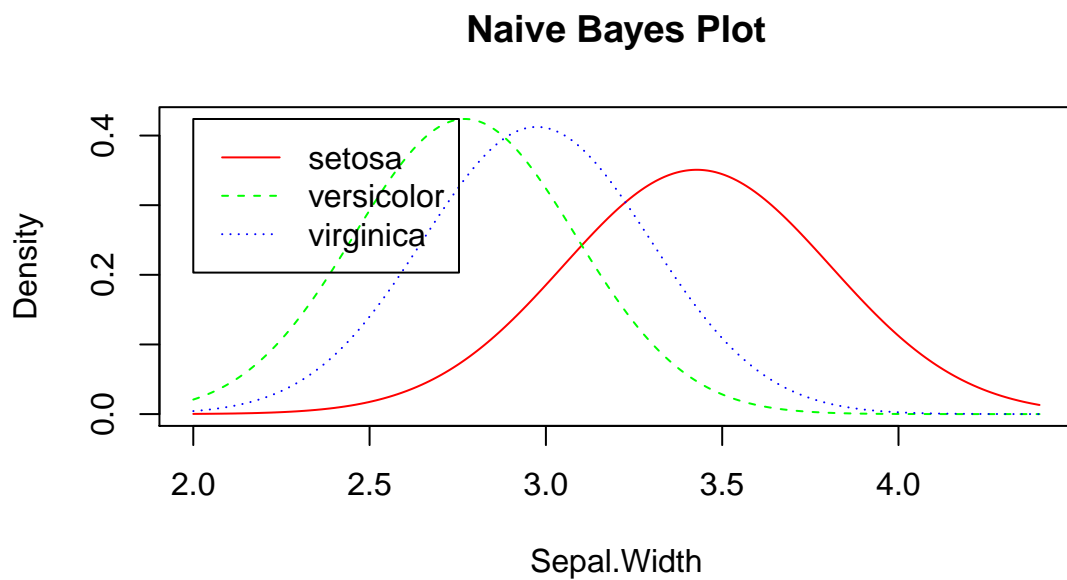
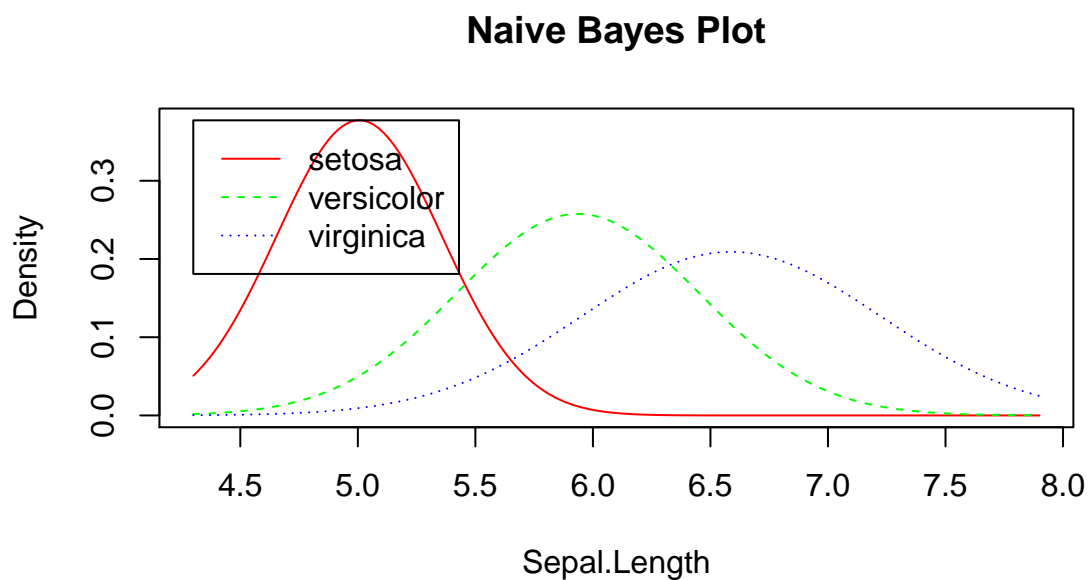
```
## [1] 0.9666667
```

准确率达到 97%。

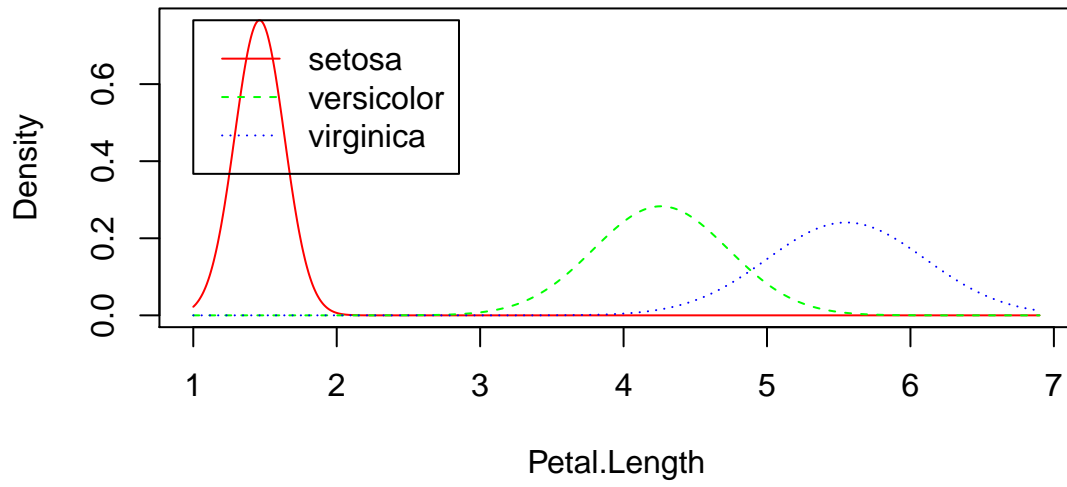
用 `klaR` 包进行朴素贝叶斯分类

步骤都是类似的，另外可以画出高斯朴素贝叶斯分类后，不同类别的鸢尾花对应的高斯分布的密度曲线。

```
library(klaR)
plot(NaiveBayes(iris$Species ~ ., data = iris))
```



Naive Bayes Plot



Naive Bayes Plot

