

hw3
501 hw2.1
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Contents

Executive Summary	1
Full Report	1
Data Cleansing	1
Descriptive Statistics	2
Build model	3
Use the training set to see model efficiency	6
Test the model with test sets	7
Summarize	9

Executive Summary

To understand whether there are significant differences in gender and age factors with different medication efficiency, age and gender were divided into four categories, and the naive Bayes model was established, and the training set and test set were used to test. It is found that the accuracy of the training set and the test set are very close, indicating that the model has a good effect, but the accuracy is not very high. In particular, men were more accurate and women were less accurate, older men (over 40) had the highest accuracy. It indicated that these drugs had obvious differences in gender and age factors, and the effect was more significant in men.

Full Report

Data Cleansing

The data in the first 18 columns of the original data were selected and classified according to gender and age. Taking 40 years old as the cut-off point, the data were roughly divided into four categories, namely, young men, old men, young women and old women, which were saved as group factor variables. Finally, the TRAINING set and test set are divided by a ratio of 4:1 and named training and TESTING. There are many missing values in the original data. Since the naiveBayes function allows missing values when doing naiveBayes analysis, it will not calculate the true terms, so the missing terms are not processed.

```
data=read.csv("R_cleaned_data.csv")

data$group=1*((data$Age=="13-19"|data$Age=="20-40")&data$Gender=="Female")+2*(!(data$Age=="13-19"|data$Age=="20-40")&data$Gender=="Female")

data=data[,c(3:18,26)]
data$group=as.factor(data$group)
```

Descriptive Statistics

By setting random seeds and making descriptive statistics on the training set, information such as minimum value, first quantile, median value, mean value, third quantile, maximum value and number of missing values of each variable can be seen. The training set has a total of 13416 rows, of which the first group has 2911 rows, the second group has 7126 rows, the third group has 1005 rows, and the fourth group has 2374 rows. The second group accounted for more in the discovery group.

```
set.seed(88)
sampling=sample(1:nrow(data),nrow(data)*0.8,replace = FALSE)
TRAINING=data[sampling,]
TESTING=data[-sampling,]
nrow(TRAINING)
```

```
## [1] 13416
```

```
summary(TRAINING)
```

##	AMITRIPTYLINE	BUPROPION	CITALOPRAM	DESVENLAFAXINE	
##	Min. :0.000	Min. :0.000	Min. :0.047	Min. :0.046	
##	1st Qu.:0.008	1st Qu.:0.024	1st Qu.:0.361	1st Qu.:0.203	
##	Median :0.018	Median :0.040	Median :0.475	Median :0.511	
##	Mean :0.029	Mean :0.068	Mean :0.478	Mean :0.442	
##	3rd Qu.:0.037	3rd Qu.:0.058	3rd Qu.:0.617	3rd Qu.:0.635	
##	Max. :0.319	Max. :0.698	Max. :0.912	Max. :0.823	
##	NA's :12778	NA's :11310	NA's :11455	NA's :13236	
##	DOXEPIN	DULOXETINE	ESCITALOPRAM	FLUOXETINE	
##	Min. :0.025	Min. :0.038	Min. :0.019	Min. :0.029	
##	1st Qu.:0.119	1st Qu.:0.275	1st Qu.:0.116	1st Qu.:0.311	
##	Median :0.164	Median :0.378	Median :0.162	Median :0.450	
##	Mean :0.212	Mean :0.389	Mean :0.209	Mean :0.446	
##	3rd Qu.:0.290	3rd Qu.:0.505	3rd Qu.:0.273	3rd Qu.:0.592	
##	Max. :0.726	Max. :0.779	Max. :0.948	Max. :0.909	
##	NA's :13299	NA's :12370	NA's :10974	NA's :11417	
##	MIRTAZAPINE	NORTRIPTYLINE	PAROXETINE	ROPINIROLE	
##	Min. :0.029	Min. :0.010	Min. :0.065	Min. :0.065	
##	1st Qu.:0.109	1st Qu.:0.045	1st Qu.:0.324	1st Qu.:0.235	
##	Median :0.139	Median :0.065	Median :0.415	Median :0.382	
##	Mean :0.182	Mean :0.099	Mean :0.443	Mean :0.352	
##	3rd Qu.:0.221	3rd Qu.:0.127	3rd Qu.:0.560	3rd Qu.:0.449	
##	Max. :0.704	Max. :0.745	Max. :0.942	Max. :0.716	
##	NA's :13001	NA's :13196	NA's :12363	NA's :13260	
##	SERTRALINE	TRAZODONE	VENLAFAXINE	OTHER	group
##	Min. :0.045	Min. :0.000	Min. :0.051	Min. :0.019	1:2911

```
## 1st Qu.:0.357 1st Qu.:0.008 1st Qu.:0.309 1st Qu.:0.182 2:7126
## Median :0.484 Median :0.015 Median :0.515 Median :0.333 3:1005
## Mean :0.482 Mean :0.035 Mean :0.475 Mean :0.346 4:2374
## 3rd Qu.:0.625 3rd Qu.:0.026 3rd Qu.:0.636 3rd Qu.:0.487
## Max. :0.960 Max. :0.586 Max. :0.908 Max. :0.885
## NA's :10671 NA's :12225 NA's :12084 NA's :7156
```

Build model

Using the training set data to do Naive Bayes classification training, the parameter data of each variable are obtained as follows:

```
#install.packages("klaR")
#install.packages("caret")
library(klaR)
library(MASS)
library(tidyverse)
library(e1071)
library(caret)
model <- naiveBayes(group~., data = TRAINING, laplace = 0)
summary(model)
```

```
##           Length Class  Mode
## apriori      4      table numeric
## tables      16      -none- list
## levels       4      -none- character
## isnumeric   16      -none- logical
## call         4      -none- call
```

```
model[1:2]
```

```
## $apriori
## Y
##      1      2      3      4
## 2911 7126 1005 2374
##
## $tables
## $tables$AMITRIPTYLINE
##      AMITRIPTYLINE
## Y           [,1]           [,2]
## 1 0.01593162 0.02297333
## 2 0.03021216 0.03727609
## 3 0.02461720 0.04393033
## 4 0.04232392 0.04906467
##
## $tables$BUPROPION
##      BUPROPION
## Y           [,1]           [,2]
## 1 0.06345101 0.10452770
## 2 0.06648531 0.10737051
## 3 0.06247740 0.08988754
## 4 0.08043100 0.12637988
```

```

##
## $tables$CITALOPRAM
##      CITALOPRAM
## Y      [,1]      [,2]
##  1 0.3969153 0.1595273
##  2 0.5259181 0.1711053
##  3 0.3614962 0.1510808
##  4 0.5152602 0.1693922
##
## $tables$DESVENLAFAXINE
##      DESVENLAFAXINE
## Y      [,1]      [,2]
##  1 0.4005846 0.2049971
##  2 0.5041807 0.2307743
##  3 0.3084157 0.1576679
##  4 0.4014441 0.2260122
##
## $tables$DOXEPIIN
##      DOXEPIIN
## Y      [,1]      [,2]
##  1 0.1471834 0.06732850
##  2 0.2206078 0.14341429
##  3 0.1468870 0.06475589
##  4 0.2483045 0.13022295
##
## $tables$DULOXETINE
##      DULOXETINE
## Y      [,1]      [,2]
##  1 0.3234018 0.1445106
##  2 0.4190225 0.1614224
##  3 0.3072140 0.1328874
##  4 0.3764414 0.1458542
##
## $tables$ESCITALOPRAM
##      ESCITALOPRAM
## Y      [,1]      [,2]
##  1 0.1742376 0.1102565
##  2 0.2316238 0.1549544
##  3 0.1740031 0.1142118
##  4 0.2302593 0.1409038
##
## $tables$FLUOXETINE
##      FLUOXETINE
## Y      [,1]      [,2]
##  1 0.3662099 0.1569459
##  2 0.5127779 0.1783902
##  3 0.3197576 0.1528554
##  4 0.5143816 0.1661522
##
## $tables$MIRTAZAPINE
##      MIRTAZAPINE
## Y      [,1]      [,2]
##  1 0.1462789 0.09932235
##  2 0.1809437 0.12423292

```

```

## 3 0.1283701 0.07253908
## 4 0.2113635 0.12926088
##
## $tables$NORTRIPTYLINE
## NORTRIPTYLINE
## Y      [,1]      [,2]
## 1 0.06591028 0.05544560
## 2 0.10449437 0.10138691
## 3 0.08325420 0.02843395
## 4 0.13444744 0.12716724
##
## $tables$PAROXETINE
## PAROXETINE
## Y      [,1]      [,2]
## 1 0.3573463 0.1314918
## 2 0.4807117 0.1673142
## 3 0.3519902 0.1402756
## 4 0.4902271 0.1765165
##
## $tables$ROPINIROLE
## ROPINIROLE
## Y      [,1]      [,2]
## 1 0.2405408 0.09187148
## 2 0.3419170 0.13846804
## 3 0.3310911 0.08443993
## 4 0.3916550 0.12692624
##
## $tables$SERTRALINE
## SERTRALINE
## Y      [,1]      [,2]
## 1 0.4097992 0.1672594
## 2 0.5445272 0.1824576
## 3 0.3594761 0.1698660
## 4 0.5393300 0.1837558
##
## $tables$TRAZODONE
## TRAZODONE
## Y      [,1]      [,2]
## 1 0.03924059 0.08121022
## 2 0.03430328 0.07965588
## 3 0.03787187 0.07892185
## 4 0.03244389 0.07165869
##
## $tables$VENLAFAXINE
## VENLAFAXINE
## Y      [,1]      [,2]
## 1 0.4185478 0.1835313
## 2 0.5092101 0.1909295
## 3 0.3967992 0.1821960
## 4 0.5011050 0.1922947
##
## $tables$OTHER
## OTHER
## Y      [,1]      [,2]

```

```
## 1 0.2474623 0.1543639
## 2 0.3839827 0.1824118
## 3 0.2207556 0.1505837
## 4 0.3719530 0.1805411
```

Use the training set to see model efficiency

The confusion matrix was established to check the fitting efficiency of the training set, and it was found that the accuracy rate was 53.14%, and the accuracy rate of the second group (elderly men) was as high as 88.9%, and the accuracy rate of women was very low, indicating that these drugs had a significant effect on elderly men (over 40 years old).

```
pred <- predict(model, TRAINING)

cm=table(TRAINING$group, pred)
cm
```

```
##      pred
##      1    2    3    4
## 1  737 2104   45   25
## 2  716 6338   28   44
## 3  327  645   25    8
## 4  235 2099   11   29
```

```
confusionMatrix(cm)
```

```
## Confusion Matrix and Statistics
##
##      pred
##      1    2    3    4
## 1  737 2104   45   25
## 2  716 6338   28   44
## 3  327  645   25    8
## 4  235 2099   11   29
##
## Overall Statistics
##
##              Accuracy : 0.5314
##              95% CI : (0.5229, 0.5399)
##      No Information Rate : 0.8338
##      P-Value [Acc > NIR] : 1
##
##              Kappa : 0.1032
##
##      McNemar's Test P-Value : <2e-16
##
## Statistics by Class:
##
##              Class: 1 Class: 2 Class: 3 Class: 4
## Sensitivity          0.36576   0.5666 0.229358 0.273585
## Specificity          0.80931   0.6466 0.926355 0.823817
## Pos Pred Value       0.25318   0.8894 0.024876 0.012216
```

```
## Neg Pred Value      0.87834    0.2293 0.993232 0.993027
## Prevalence          0.15019    0.8338 0.008125 0.007901
## Detection Rate      0.05493    0.4724 0.001863 0.002162
## Detection Prevalence 0.21698    0.5312 0.074911 0.176953
## Balanced Accuracy    0.58754    0.6066 0.577856 0.548701
```

```
#install.packages("gmodels")
library(gmodels)

CrossTable(pred, TRAINING$group,
           prop.chisq = FALSE, prop.t = FALSE, prop.r = FALSE,
           dnn = c('predicted', 'actual'))
```

```
##
##
##      Cell Contents
## |-----|
## |                      N |
## |          N / Col Total |
## |-----|
##
##
## Total Observations in Table:  13416
##
##
##      | actual
## predicted |      1 |      2 |      3 |      4 | Row Total |
## -----|-----|-----|-----|-----|-----|
##      1 |      737 |      716 |      327 |      235 |      2015 |
##      |      0.253 |      0.100 |      0.325 |      0.099 |
## -----|-----|-----|-----|-----|
##      2 |      2104 |      6338 |      645 |      2099 |      11186 |
##      |      0.723 |      0.889 |      0.642 |      0.884 |
## -----|-----|-----|-----|-----|
##      3 |        45 |        28 |        25 |        11 |        109 |
##      |      0.015 |      0.004 |      0.025 |      0.005 |
## -----|-----|-----|-----|-----|
##      4 |        25 |        44 |         8 |        29 |        106 |
##      |      0.009 |      0.006 |      0.008 |      0.012 |
## -----|-----|-----|-----|-----|
## Column Total |      2911 |      7126 |      1005 |      2374 |      13416 |
##      |      0.217 |      0.531 |      0.075 |      0.177 |
## -----|-----|-----|-----|-----|
##
##
```

Test the model with test sets

Using the test set to test, it is found that the accuracy is 52.06%, which is very close to the accuracy of the training set. Other results are consistent with the data of the training set, indicating that the model is accurate and the fitting result is good.

```
pred <- predict(model,TESTING)
```

```
cm=table(TESTING$group,pred)
cm
```

```
##      pred
##      1    2    3    4
## 1 184 512 13 10
## 2 172 1542 11 13
## 3 74 160 11 1
## 4 93 544 5 9
```

```
confusionMatrix(cm)
```

```
## Confusion Matrix and Statistics
```

```
##
##      pred
##      1    2    3    4
## 1 184 512 13 10
## 2 172 1542 11 13
## 3 74 160 11 1
## 4 93 544 5 9
```

```
##
```

```
## Overall Statistics
```

```
##
##              Accuracy : 0.5206
##              95% CI : (0.5035, 0.5376)
##      No Information Rate : 0.8223
##      P-Value [Acc > NIR] : 1
```

```
##
```

```
##              Kappa : 0.1083
```

```
##
```

```
## McNemar's Test P-Value : <2e-16
```

```
##
```

```
## Statistics by Class:
```

```
##
```

```
##              Class: 1 Class: 2 Class: 3 Class: 4
## Sensitivity      0.35182 0.5591 0.27500 0.272727
## Specificity      0.81102 0.6711 0.92909 0.806685
## Pos Pred Value    0.25591 0.8872 0.04472 0.013825
## Neg Pred Value    0.87135 0.2475 0.99067 0.991121
## Prevalence        0.15593 0.8223 0.01193 0.009839
## Detection Rate    0.05486 0.4597 0.00328 0.002683
## Detection Prevalence 0.21437 0.5182 0.07335 0.194097
## Balanced Accuracy 0.58142 0.6151 0.60204 0.539706
```

```
table(TESTING$group)
```

```
##
##      1    2    3    4
## 719 1738 246 651
```



```
library(gmodels)

CrossTable(pred, TESTING$group,
  prop.chisq = FALSE, prop.t = FALSE, prop.r = FALSE,
  dnn = c('predicted', 'actual'))
```

```
##
##
##      Cell Contents
## |-----|
## |                      N |
## |          N / Col Total |
## |-----|
##
##
## Total Observations in Table:  3354
##
##
##      | actual
## predicted |          1 |          2 |          3 |          4 | Row Total |
## -----|-----|-----|-----|-----|-----|
##          1 |          184 |          172 |          74 |          93 |          523 |
##          |          0.256 |          0.099 |          0.301 |          0.143 |
## -----|-----|-----|-----|-----|
##          2 |          512 |         1542 |          160 |          544 |         2758 |
##          |          0.712 |          0.887 |          0.650 |          0.836 |
## -----|-----|-----|-----|-----|
##          3 |           13 |           11 |           11 |           5 |           40 |
##          |          0.018 |          0.006 |          0.045 |          0.008 |
## -----|-----|-----|-----|-----|
##          4 |           10 |           13 |           1 |           9 |           33 |
##          |          0.014 |          0.007 |          0.004 |          0.014 |
## -----|-----|-----|-----|-----|
## Column Total |          719 |         1738 |          246 |          651 |         3354 |
##          |          0.214 |          0.518 |          0.073 |          0.194 |
## -----|-----|-----|-----|-----|
##
##
```

Summarize

According to the above results, it can be found that both the training set and the test set have the highest accuracy of about 88% for the second group, namely, the older male group. It was followed by young men, meaning younger than 40, with an accuracy rate of about 25 percent. The accuracy rate for women is very low, all of which are less than 5%, indicating that these drugs have no significant change in women, so it can be said that the effect on women is not great.