

Practice Problems 4

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```
library(tidyverse)
```

```
## -- Attaching packages -----  
  
## v ggplot2 3.3.3      v purrr  0.3.4  
## v tibble  3.0.3      v dplyr  1.0.2  
## v tidyr   1.1.2      v stringr 1.4.0  
## v readr   1.3.1      v forcats 0.5.0  
  
## -- Conflicts -----  
## x dplyr::filter() masks stats::filter()  
## x dplyr::lag()    masks stats::lag()
```

Problem 1 (50 Points)

Build an R Notebook of the SMS message filtering example in the textbook on pages 103 to 123 ([data set](#)). Show each step and add appropriate documentation. Note that the attached data set differs slightly from the one used on the book; the number of cases differ.

Step 1: Collecting Data

Here we are looking at different types of text (ham vs spam), and we want to develop a Naive Bayes classifier to determine what type of text fits what class.

Step 2: Exploring and Preparing the Data

```
sms_raw <- read_csv('da5030.spammsgdataset.csv')
```

```
## Parsed with column specification:  
## cols(  
##   type = col_character(),  
##   text = col_character()  
## )
```

```
sms_raw
```

```
## # A tibble: 5,574 x 2
##   type text
##   <chr> <chr>
## 1 ham   Go until jurong point, crazy.. Available only in bugis n great world l~
## 2 ham   Ok lar... Joking wif u oni...
## 3 spam  Free entry in 2 a wkly comp to win FA Cup final tkts 21st May 2005. Te~
## 4 ham   U dun say so early hor... U c already then say...
## 5 ham   Nah I don't think he goes to usf, he lives around here though
## 6 spam  FreeMsg Hey there darling it's been 3 week's now and no word back! I'd~
## 7 ham   Even my brother is not like to speak with me. They treat me like aids ~
## 8 ham   As per your request 'Melle Melle (Oru Minnaminunginte Nurungu Vettam)'~
## 9 spam  WINNER!! As a valued network customer you have been selected to receiv~
## 10 spam Had your mobile 11 months or more? U R entitled to Update to the lates~
## # ... with 5,564 more rows
```

```
# Convert to factor
sms_raw$type <- factor(sms_raw$type)

str(sms_raw$type)
```

```
## Factor w/ 2 levels "ham","spam": 1 1 2 1 1 2 1 1 2 2 ...
```

```
table(sms_raw$type)
```

```
##
##   ham spam
## 4827  747
```

Cleaning and Standardizing the Data: Here we start to remove numbers, punctuation, and uninteresting words so we can get to the bottom of what ham vs spam text is like.

```
library(tm)
```

```
## Loading required package: NLP
```

```
##
## Attaching package: 'NLP'
```

```
## The following object is masked from 'package:ggplot2':
##
##   annotate
```

```
# Create a corpus, which is a collection of text documents
sms_corpus <- VCorpus(VectorSource(sms_raw$text))

# Inspect the corpus
inspect(sms_corpus[1:2])
```

```
## <<VCorpus>>
## Metadata: corpus specific: 0, document level (indexed): 0
```

```
## Content:  documents: 2
##
## [[1]]
## <<PlainTextDocument>>
## Metadata:  7
## Content:  chars: 111
##
## [[2]]
## <<PlainTextDocument>>
## Metadata:  7
## Content:  chars: 29
```

```
# View the text
as.character(sms_corpus[[1]])
```

```
## [1] "Go until jurong point, crazy.. Available only in bugis n great world la e buffet... Cine there g
```

```
# View multiple documents
lapply(sms_corpus[1:2], as.character)
```

```
## $'1'
## [1] "Go until jurong point, crazy.. Available only in bugis n great world la e buffet... Cine there g
##
## $'2'
## [1] "Ok lar... Joking wif u oni..."
```

```
# Standardize to all lowercase
sms_corpus_clean <- tm_map(sms_corpus, content_transformer(tolower))

# Make sure tm_map() function worked
as.character(sms_corpus[[1]])
```

```
## [1] "Go until jurong point, crazy.. Available only in bugis n great world la e buffet... Cine there g
```

```
as.character(sms_corpus_clean[[1]])
```

```
## [1] "go until jurong point, crazy.. available only in bugis n great world la e buffet... cine there g
```

```
# Remove all the numbers from the corpus
sms_corpus_clean <- tm_map(sms_corpus_clean, removeNumbers)

# Remove the filler words like and, if, but, etc using the stopwords() function
sms_corpus_clean <- tm_map(sms_corpus_clean, removeWords, stopwords())

# Remove punctuation to complete the standardization
sms_corpus_clean <- tm_map(sms_corpus_clean, removePunctuation)

# Reduce words to their root form, a.k.a. stemming
library(SnowballC)
sms_corpus_clean <- tm_map(sms_corpus_clean, stemDocument)
```

```
# Strip the additional whitespace to finalize the data cleaning
sms_corpus_clean <- tm_map(sms_corpus_clean, stripWhitespace)
```

```
# Before cleaning
as.character(sms_corpus[[1]])
```

```
## [1] "Go until jurong point, crazy.. Available only in bugis n great world la e buffet... Cine there g
```

```
# After cleaning
as.character(sms_corpus_clean[[1]])
```

```
## [1] "go jurong point crazi avail bugi n great world la e buffet cine got amor wat"
```

Splitting Text Documents into Words: We want to create a document term matrix (DTM) using the `DocumentTermMatrix()` function. a DTM is a data structure where the rows indicate documents (SMS messages) and columns indicate terms (words).

```
# original dtm
sms_dtm <- DocumentTermMatrix(sms_corpus_clean)
sms_dtm
```

```
## <<DocumentTermMatrix (documents: 5574, terms: 6592)>>
## Non-/sparse entries: 42608/36701200
## Sparsity           : 100%
## Maximal term length: 40
## Weighting           : term frequency (tf)
```

```
# modify preprocessing steps
sms_dtm2 <- DocumentTermMatrix(sms_corpus, control = list(
  tolower = TRUE,
  removeNumbers = TRUE,
  stopwords = TRUE,
  removePunctuation = TRUE,
  stemming = TRUE
))
sms_dtm2
```

```
## <<DocumentTermMatrix (documents: 5574, terms: 6995)>>
## Non-/sparse entries: 43713/38946417
## Sparsity           : 100%
## Maximal term length: 40
## Weighting           : term frequency (tf)
```

Create training/test datasets: We always do this, just like we have done before.

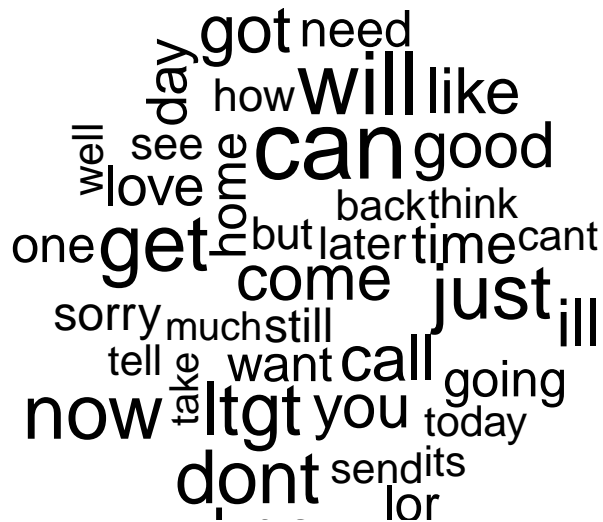
```
sms_dtm_train <- sms_dtm[1:4169, ]
sms_dtm_test  <- sms_dtm[4170:5559, ]

sms_train_labels <- sms_raw[1:4169, ]$type
sms_test_labels  <- sms_raw[4170:5559, ]$type

prop.table(table(sms_train_labels))
```




```
# Ham word cloud  
wordcloud(ham$text, max.words = 40, scale = c(3, 0.5))
```



Creating indicator features for frequent words: We want to find the frequent words we can count as a reference for the classifier.

```
sms_freq_words <- findFreqTerms(sms_dtm_train, 5)
str(sms_freq_words)
```

```
## chr [1:1157] "&wk" "abiola" "abl" "abt" "accept" "access" "account" ...
```

```
sms_dtm_freq_train<- sms_dtm_train[ , sms_freq_words]
sms_dtm_freq_test <- sms_dtm_test[ , sms_freq_words]
```

```
sms_dtm_freq_train
```

```
## <<DocumentTermMatrix (documents: 4169, terms: 1157)>>
## Non-/sparse entries: 25173/4798360
## Sparsity          : 99%
## Maximal term length: 13
## Weighting          : term frequency (tf)
```

```
sms_dtm_freq_test
```

```
## <<DocumentTermMatrix (documents: 1390, terms: 1157)>>
## Non-/sparse entries: 8165/1600065
```

```
## Sparsity          : 99%
## Maximal term length: 13
## Weighting         : term frequency (tf)

# Convert counts to yes/no strings
convert_counts <- function(x) {
  x <- ifelse(x > 0, "Yes", "No")
}

# Apply the function to the new training and test datasets
sms_train <- apply(sms_dtm_freq_train, MARGIN = 2,
                  convert_counts)
sms_test <- apply(sms_dtm_freq_test, MARGIN = 2,
                 convert_counts)
```

Step 3, Train a model on the data

```
library(e1071)
sms_classifier <- naiveBayes(sms_train, sms_train_labels)
```

Step 4, Evaluate model performance

We use the same stuff that we used for kNN, where we use CrossTable to look at the accuracy of the model.

```
sms_test_pred <- predict(sms_classifier, sms_test)

# Use CrossTable from gmodels
library(gmodels)
```

```
## Warning: package 'gmodels' was built under R version 4.0.3
```

```
CrossTable(sms_test_pred, sms_test_labels, prop.chisq = FALSE, prop.t = FALSE, dnn = c('predicted
```

```
##
##
##      Cell Contents
## |-----|
## |                      N |
## |          N / Row Total |
## |          N / Col Total |
## |-----|
##
##
## Total Observations in Table: 1390
##
##
##          | actual
## predicted |      ham |      spam | Row Total |
## -----|-----|-----|-----|
```



```
##      ham |      1200 |      20 |      1220 |
##      |      0.984 |      0.016 |      0.878 |
##      |      0.993 |      0.110 |      |
## -----|-----|-----|-----|
##      spam |        9 |      161 |      170 |
##      |      0.053 |      0.947 |      0.122 |
##      |      0.007 |      0.890 |      |
## -----|-----|-----|-----|
## Column Total |      1209 |      181 |      1390 |
##      |      0.870 |      0.130 |      |
## -----|-----|-----|-----|
##
##
```

Step 5, Improve model performance

This time we changed the Laplace estimator to 1, which reduced our false positives.

```
sms_classifier2 <- naiveBayes(sms_train, sms_train_labels, laplace = 1)
sms_test_pred2 <- predict(sms_classifier2, sms_test)
CrossTable(sms_test_pred2, sms_test_labels,
            prop.chisq = FALSE, prop.t = FALSE, prop.r = FALSE,
            dnn = c('predicted', 'actual'))
```

```
##
##
##      Cell Contents
## |-----|
## |              N |
## |      N / Col Total |
## |-----|
##
##
## Total Observations in Table:  1390
##
##
##      | actual
## predicted |      ham |      spam | Row Total |
## -----|-----|-----|-----|
##      ham |      1182 |        10 |      1192 |
##      |      0.978 |      0.055 |      |
## -----|-----|-----|-----|
##      spam |        27 |       171 |      198 |
##      |      0.022 |      0.945 |      |
## -----|-----|-----|-----|
## Column Total |      1209 |      181 |      1390 |
##      |      0.870 |      0.130 |      |
## -----|-----|-----|-----|
##
##
```

Problem 2 (50 Points)

Install the requisite packages to execute the following code that classifies the built-in **iris** data using Naive Bayes. Build an R Notebook and explain in detail what each step does. Be sure to look up each function to understand how it is used.

```
library(klaR)
data(iris)

nrow(iris)
summary(iris)
head(iris)

testidx <- which(1:length(iris[, 1]) %% 5 == 0)

# separate into training and testing datasets
iristrain <- iris[-testidx,]
iristest <- iris[testidx,]

# apply Naive Bayes
nbmodel <- NaiveBayes(Species~., data=iristrain)

# check the accuracy
prediction <- predict(nbmodel, iristest[, -5])
table(prediction$class, iristest[, 5])
```

Data Preparation

There isn't too much we have to do for data preparation since the data is nicely set up for us in R.

```
# Install packages and show data
library(klaR)

## Loading required package: MASS

##
## Attaching package: 'MASS'

## The following object is masked from 'package:dplyr':
##
##      select

data(iris)

# Get rows, summary, and head of data
nrow(iris)

## [1] 150
```

```
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
##
##
##
```

```
head(iris)
```

```
##      Sepal.Length Sepal.Width Petal.Length Petal.Width Species
## 1           5.1         3.5         1.4         0.2  setosa
## 2           4.9         3.0         1.4         0.2  setosa
## 3           4.7         3.2         1.3         0.2  setosa
## 4           4.6         3.1         1.5         0.2  setosa
## 5           5.0         3.6         1.4         0.2  setosa
## 6           5.4         3.9         1.7         0.4  setosa
```

```
# get all multiples of 150 that are divisible by 5
testidx <- which(1:length(iris[, 1]) %% 5 == 0)
testidx
```

```
## [1]  5 10 15 20 25 30 35 40 45 50 55 60 65 70 75 80 85 90 95
## [20] 100 105 110 115 120 125 130 135 140 145 150
```

```
# separate into training and testing datasets
iristrain <- iris[-testidx,]
iristest <- iris[testidx,]
```

Naive Bayes Model

Basically for this model, we use the species as the given part of Bayes Theorem where we look for a probability given a condition. The condition here is the species. For instance, we will get the sepal length, petal length, etc given each species (setosa, versicolor, virginica).

```
# apply Naive Bayes
library(naivebayes)
```

```
## naivebayes 0.9.7 loaded
```

```
nbmodel <- naive_bayes(Species~., data=iristrain)
nbmodel
```

```
##
## ===== Naive Bayes =====
##
## Call:
## naive_bayes.formula(formula = Species ~ ., data = iristrain)
##
## -----
##
## Laplace smoothing: 0
##
## -----
##
## A priori probabilities:
##
##      setosa versicolor  virginica
## 0.3333333  0.3333333  0.3333333
##
## -----
##
## Tables:
##
## -----
## ::: Sepal.Length (Gaussian)
## -----
##
## Sepal.Length      setosa versicolor virginica
##      mean 4.9975000  5.9900000 6.6100000
##      sd   0.3675892  0.5295378 0.6647922
##
## -----
## ::: Sepal.Width (Gaussian)
## -----
##
## Sepal.Width      setosa versicolor virginica
##      mean 3.4175000  2.7775000 2.9700000
##      sd   0.3960623  0.3415556 0.3081791
##
## -----
## ::: Petal.Length (Gaussian)
## -----
##
## Petal.Length      setosa versicolor virginica
##      mean 1.4425000  4.3100000 5.5575000
##      sd   0.1583367  0.4850588 0.5930743
##
## -----
## ::: Petal.Width (Gaussian)
## -----
##
## Petal.Width      setosa versicolor virginica
```

```
##          mean 0.2525000  1.3325000  2.0300000
##          sd   0.1109111  0.2080280  0.2355572
##
## -----
```

Accuracy of Model

This just tracks the predictions against the actual results which are in the `iristest` data. We use the `table` function to count how many accurate predictions we have.

```
# check the accuracy
prediction <- predict(nbmodel, iristest[,5])
prediction
```

```
## [1] setosa      setosa      setosa      setosa      setosa      setosa
## [7] setosa      setosa      setosa      setosa      versicolor versicolor
## [13] versicolor versicolor versicolor versicolor versicolor versicolor
## [19] versicolor versicolor virginica  virginica  virginica  versicolor
## [25] virginica  virginica  versicolor virginica  virginica  virginica
## Levels: setosa versicolor virginica
```

```
table(prediction, iristest[,5])
```

```
##
## prediction  setosa versicolor virginica
## setosa      10         0         0
## versicolor   0         10        2
## virginica    0         0         8
```

It looks like setosas have a 100% accuracy, versicolors had 10/12 correct guesses, and virginicas had 8/10 correct guesses. Not bad.