

Practical -8

Write a program for measuring similarity among documents and detecting passages which have been reused.

Installation of required packages before executing program:-

```
install.packages("tm")  
require("tm")  
install.packages("ggplot2")  
install.packages("textreuse")  
install.packages("devtools")
```

Source Code 1:-

```
my.corpus <- Corpus(DirSource("c:/msc/r-corpus"))  
  
my.corpus <- tm_map(my.corpus, removeWords, stopwords("english"))  
  
my.tdm <- TermDocumentMatrix(my.corpus)  
  
#inspect(my.tdm)  
  
my.dtm <- DocumentTermMatrix(my.corpus, control = list(weighting =  
weightTfIdf, stopwords = TRUE))  
  
#inspect(my.dtm)  
  
my.df <- as.data.frame(inspect(my.tdm))  
  
my.df.scale <- scale(my.df)  
  
d <- dist(my.df.scale,method="euclidean")  
  
fit <- hclust(d, method="ward")  
  
plot(fit)
```

OutPut:-

<<TermDocumentMatrix (terms: 69, documents: 6)>>

Non-/sparse entries: 97/317

Sparsity : 77%

Maximal term length: 12

Weighting : term frequency (tf)

Terms	Docs					
	File1.txt	File2.txt	File3.txt	File4.txt	File5.txt	File6.txt
also	0	1	1	1	0	0
bed	0	0	0	1	0	0
better	0	0	0	1	0	0
call	0	1	0	0	0	0
can	0	0	1	1	0	0
cat	0	0	0	1	0	0
cats	0	0	0	1	0	0
couch.	0	0	0	1	0	0
danger	0	0	1	0	0	0
dangerous	0	0	0	0	1	1
different	0	0	0	1	0	0
document	0	1	0	0	0	0
document,	0	1	0	0	0	0
dog,	0	0	0	1	0	0
dogs	0	0	3	1	0	0
dogs,	0	0	0	1	0	0
dogs.	0	0	0	1	0	0
eat	0	0	1	0	0	0
even	0	0	0	0	1	1
fairness,	0	0	0	1	0	0
fall	0	0	1	0	0	0
felines	0	0	0	1	0	0
fun.	0	0	1	0	0	0
gives	0	0	0	0	1	1
help	0	0	1	0	0	0
hide.	0	0	0	1	0	0
home,	0	0	0	1	0	0
however,	0	0	0	0	1	1
instead	0	1	0	0	0	0
instructions	0	0	0	0	1	1
intruder	0	0	0	1	0	0
jump	0	0	1	1	0	0
jump.	0	0	0	0	1	1
just	0	0	0	1	0	0
like	0	0	1	0	0	0
line.	1	0	0	0	0	1
lot.	0	0	0	2	0	0
lots	0	0	0	1	0	0
minerals,	0	0	0	0	1	1
mining	0	0	0	0	1	1
mining,	0	1	0	0	0	0
mining?	0	1	0	0	0	0
much,	0	0	0	1	0	0
one	1	0	0	0	0	1
operation,	0	0	0	0	1	1
pick	0	0	1	0	0	0

playing	0	0	0	1	0	0
poop	0	0	1	0	0	0
precise	0	0	0	0	1	1
probably	0	0	0	1	0	0
protect	0	0	1	0	0	0
rather	0	0	0	1	0	0
romance.	1	0	0	0	0	1
run	0	0	2	2	1	1
short,	1	0	0	0	0	1
since	0	0	0	1	0	0
sit	0	0	0	1	0	0
sleep	0	0	0	2	0	0
stinky,	0	0	1	0	0	0
text	0	1	0	0	1	1
text.	2	1	0	0	0	2
things.	0	0	1	0	0	0
time.	0	0	1	0	0	0
unlike	0	0	0	1	0	0
usually	0	0	0	1	0	0
valuable.	0	0	0	0	1	1
well.	0	0	1	0	0	0
whales	1	0	0	0	0	1
will	0	0	0	1	0	0

```

> barplot(as.matrix(my.tdm))

```

```

> my.df.scale <- scale(my.df)

```

```

> d <- dist(my.df.scale,method="euclidean")

```

```

> fit <- hclust(d, method="ward")

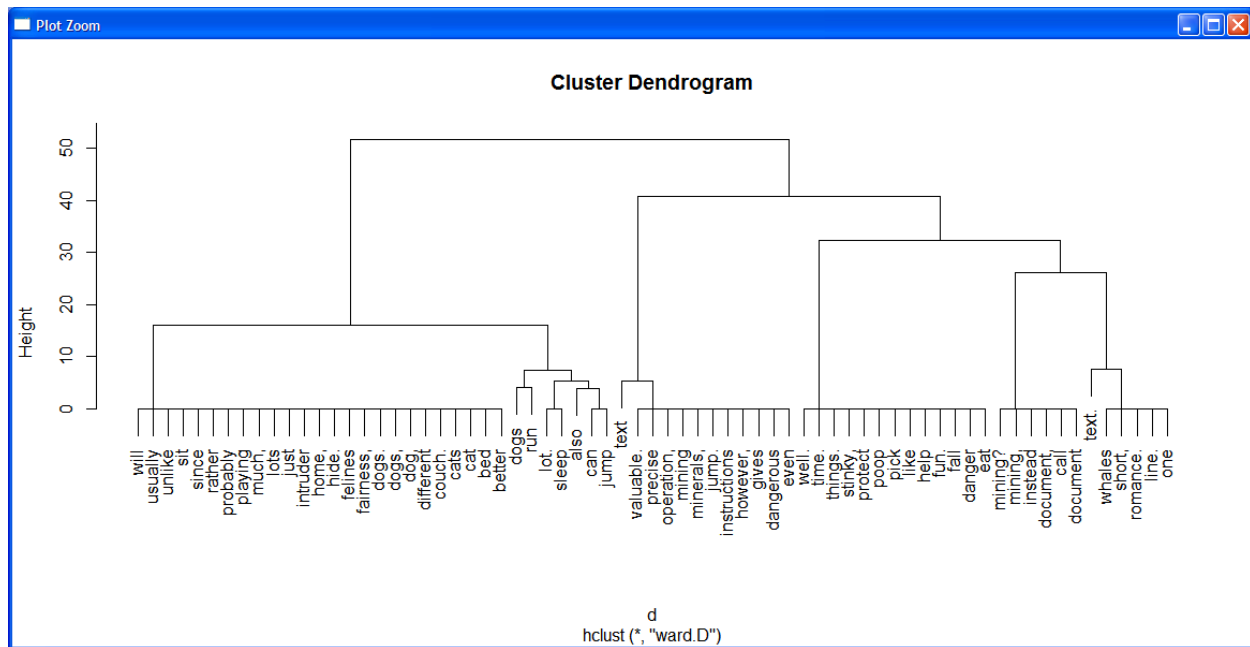
```

The "ward" method has been renamed to "ward.D"; note new "ward.D2"

```

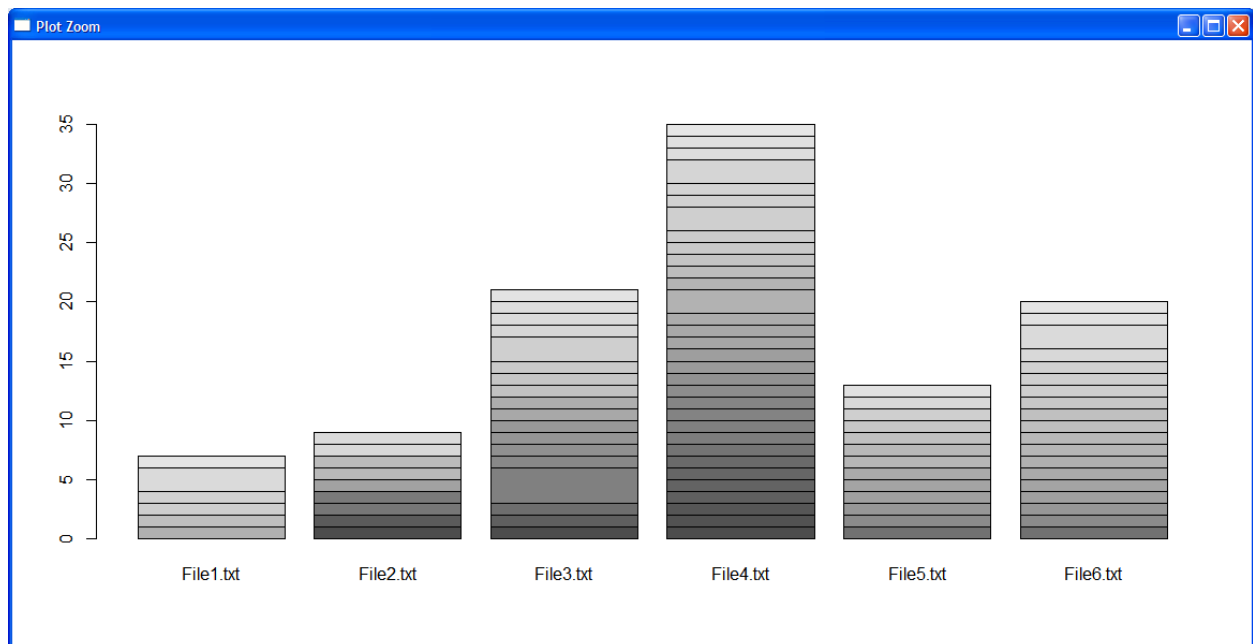
> plot(fit)

```

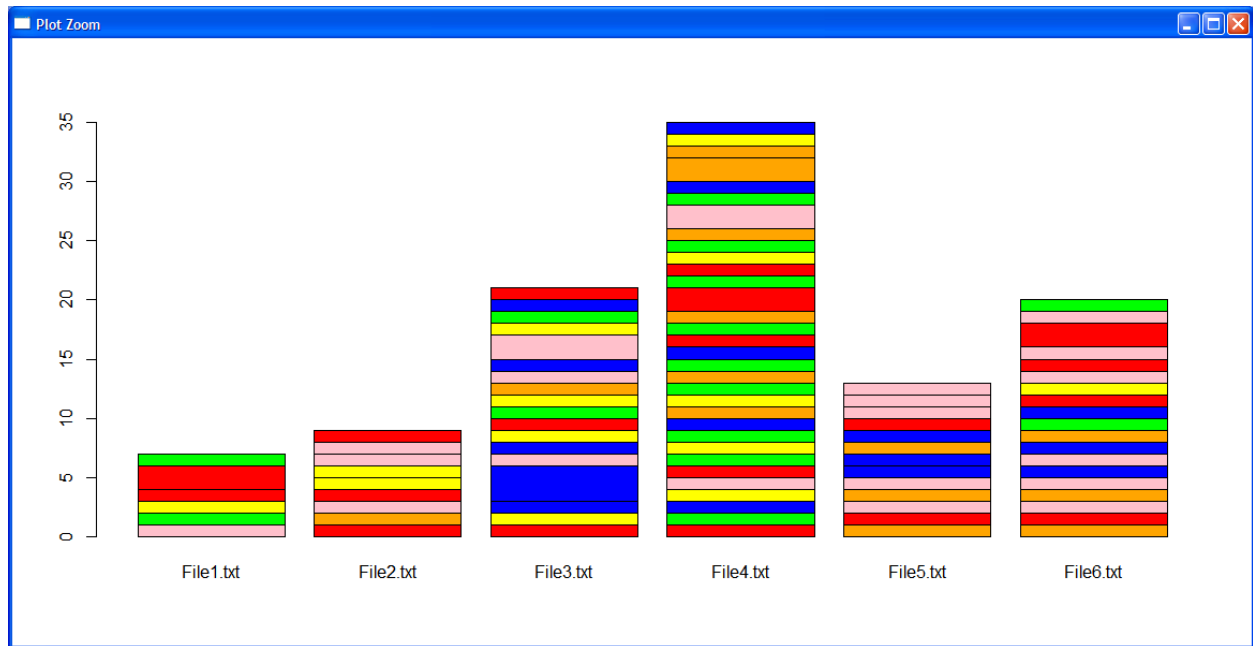


Sourec code 2 (using bar plot with and without color):-

```
my.corpus <- Corpus(DirSource("c:/msc/r-corpus"))  
my.corpus <- tm_map(my.corpus, removeWords, stopwords("english"))  
my.tdm <- TermDocumentMatrix(my.corpus)  
inspect(my.tdm)  
my.df <- as.data.frame(inspect(my.tdm))  
barplot(as.matrix(my.tdm))  
#barplot(as.matrix(my.tdm),col = color)
```

OutPut:-

```
barplot(as.matrix(my.tdm),col = color)
```



Source code 3 (using minhash and jaccard similarity):-

```
library(textreuse)
```

Source Code:-

```
minhash <- minhash_generator(200, seed = 235)
ats <- TextReuseCorpus(dir = "c:/msc/r-corpus", tokenizer = tokenize_ngrams, n =
5, minhash_func = minhash)
buckets <- lsh(ats, bands = 50, progress = interactive())
candidates <- lsh_candidates(buckets)
scores <- lsh_compare(candidates, ats, jaccard_similarity, progress = FALSE)
scores
color <- c("red","green","blue","orange","yellow","pink")
barplot(as.matrix(scores),col = color)
```

Output:

```
      a      b      score
<chr> <chr>  <dbl>
1 File 1 File 6 0.4651163
2 File 5 File 6 0.4418605
```

