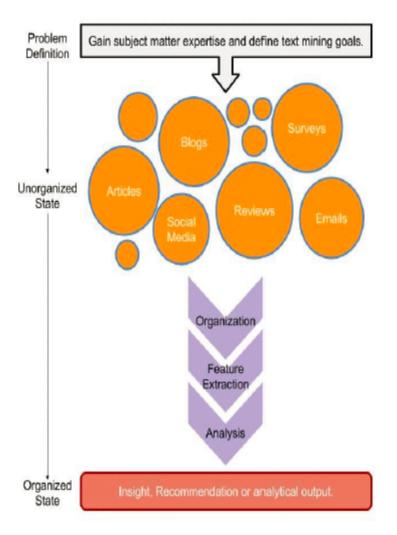
### Text Data Analysis

The Process of distilling actionable insights from text

## **Text mining workflow**

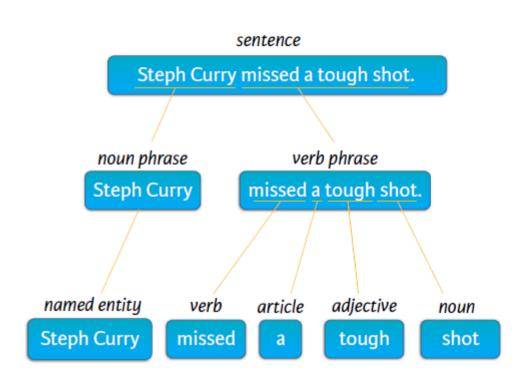


1 - Problem definition & specific goals

2 - Identify text to be collected

- 3 Text organization
- 4 Feature extraction
- 5 Analysis
- 6 Reach an insight, recommendation or output

### Semantic parsing vs. bag of words

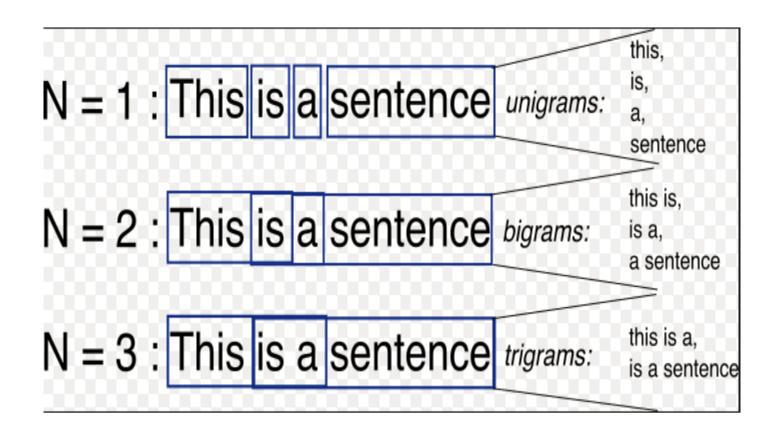




### Bag of words

- bag of words text mining represents a way to count terms, or *n-grams*, across a collection of documents.
   Consider the following sentences:
- text<- "data analytics is the new sensation of the modern times, often misunderstood as only graphs and dashboards. Its time for businesses to think beyond charts and start leveraging the science behind the data"
- Manually counting words in the sentences above is a pain!

### N-grams



### Steps to follow

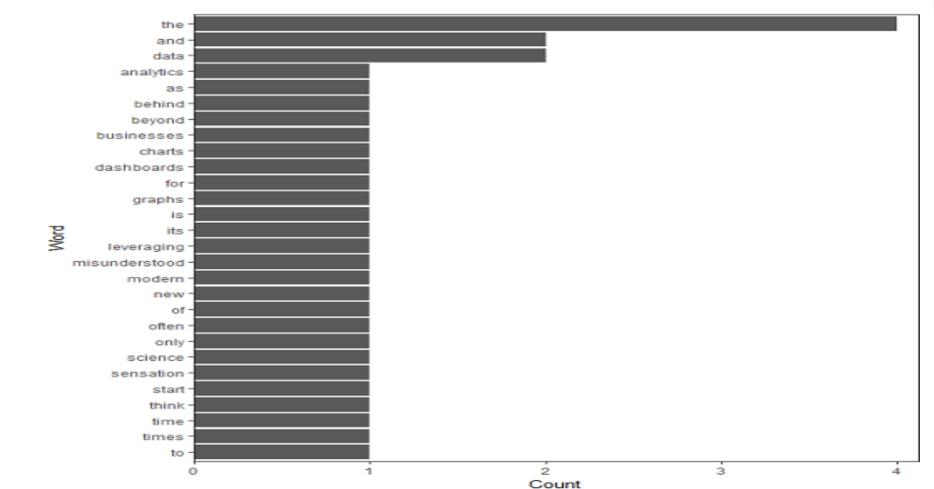
# Load qdap

# Print text to the console

# Find the 4 most frequent terms: term\_count

# Plot term\_count

```
> text <- "data analytics is the new sensation of the modern times, o
ften misunderstood as only graphs and dashboards. Its time for busi
nesses to think beyond charts and start leveraging the science behin
d the data"
> library(qdap)
> freq_terms<-freq_terms(text,4)
> plot(freq_terms)
> |
```



#### Load some text

- # Import text data
- tweets<-read.csv(" XiaomiIndiaTweets.csv",stringsAsFactors</li>= FALSE)
- # View the structure of tweets
- str(tweets)
- # Print out the number of rows in tweets
- nrow(tweets)
- # Isolate text from tweets: xiaomi\_tweets
- xiaomi\_tweets <- tweets\$text</li>
- xiaomi\_tweets

### Make the vector a VCorpus object

- We have loaded your text data as a vector called xiaomi\_tweets in the last exercise.
- Your next step is to convert this vector containing the text data to a corpus.
- Corpus is a collection of documents, but it's also important to know that in the tm domain, R recognizes it as a data type.
- There are two kinds of the corpus data type, the *permanent* corpus, PCorpus, and the *volatile corpus*, VCorpus. In essence, the difference between the two has to do with how the collection of documents is stored in your computer. In this course, we will use the volatile corpus, which is held in your computer's RAM rather than saved to disk, just to be more memory efficient.
- To make a volatile corpus, R needs to interpret each element in our vector of text, xiaomi\_tweets, as a document.
- And the tm package provides what are called Source functions to do just that!
- In this exercise, we'll use a Source function called VectorSource() because our text data is contained in a vector.
- The output of this function is called a Source object.

```
Console \sim/ML/bangaloresession1/ \stackrel{>}{lpha}
```

- > library(tm)
- > xiaomi\_source<-VectorSource(xiaomi\_tweets)</pre>

### **VCorpus**

- As we've converted our vector to a Source object, we pass it to another tm function, VCorpus(), to create our volatile corpus.
- The VCorpus object is a nested list, or list of lists. At each index of the VCorpus object, there is a PlainTextDocument object, which is essentially a list that contains the actual text data (content), as well as some corresponding metadata (meta).
- It can help to visualize a VCorpus object to conceptualize the whole thing.
- For example, to examine the contents of the second tweet in xiaomi\_corpus, you'd subset twice.
- Once to specify the second PlainTextDocument corresponding to the second tweet and again to extract the first (or content) element of that PlainTextDocument:
- xiaomi\_corpus[[15]][1]

### Steps

- ## xiaomi\_source is already in your workspace
- # Make a volatile corpus: xiaomi\_corpus
- xiaomi\_corpus <- VCorpus(xiaomi\_source)</li>
- # Print out xiaomi\_corpus
- xiaomi\_corpus
- # Print data on the 15th tweet in xiaomi\_corpus
- xiaomi\_corpus[[15]]
- # Print the content of the 15th tweet in xiaomi\_corpus
- xiaomi\_corpus[[15]][1]

DMTHyFh"

> library(tm) > xiaomi\_source<-VectorSource(xiaomi\_tweets)</pre> > xiaomi\_corpus <- VCorpus(xiaomi\_source)</pre> > xiaomi\_corpus <<VCorpus>> Metadata: corpus specific: 0, document level (indexed): 0 Content: documents: 1561 > xiaomi\_corpus[[15]] <<PlainTextDocument>> Metadata: 7 Content: chars: 140 > xiaomi\_corpus[[15]][1] \$content [1] "Mi fans! We bring you super saving deals on your favourite Mi pr oducts. The Republic Day sale is live. Check it now... https://t.co/PGA

### Make a VCorpus from a data frame

- Because another common text source is a data frame, there is a Source function called DataframeSource().
   The DataframeSource() function must have a specific structure:
- Column one must be called doc\_id and contain a unique string for each row.
- Column **two** must be called text with "UTF-8" encoding (pretty standard).
- Any other columns, **3+** are considered meta-data and will be retained as such.

## **Common preprocessing functions**

TM Function	Description	Before	After
tolower()	Makes all text lowercase	Starbucks is from Seattle.	starbucks is from seattle.
removePunctuation()	Removes punctuation like periods and exclamation points	Watch out! That coffee is going to spill!	Watch out That coffee is going to spill
removeNumbers()	Removes numbers	I drank 4 cups of coffee 2 days	I drank cups of coffee days ago.
stripWhiteSpace()	Removes tabs and extra spaces	I like coffee.	I like coffee.
removeWords()	Removes specific words (e.g. "the", "of") defined by the data scientist	The coffee house and barista he visited were nice, she said hello.	The coffee house barista visited nice, said hello.

## Preprocessing in practice



### Common cleaning functions from tm

- Common preprocessing functions include:
- tolower(): Make all characters lowercase
- removePunctuation(): Remove all punctuation marks
- removeNumbers(): Remove numbers
- stripWhitespace(): Remove excess whitespace

- > text <- "data analytics is the new sensation of the modern times, o ften misunderstood as !!!! : only graphs and dashboards. Its time f or businesses to think beyond charts and start leveraging the scienc e behind the data in 2018 "
- > removeNumbers(text)
- [1] "data analytics is the new sensation of the modern times, often misunderstood as !!!! : only graphs and dashboards. Its time for bus inesses to think beyond charts and start leveraging the science behind the data in
- > removePunctuation(text)
- [1] "data analytics is the new sensation of the modern times often misunderstood as only graphs and dashboards Its time for businesses to think beyond charts and start leveraging the science behind the data in 2018 "
- > stripWhitespace(text)
- [1] "data analytics is the new sensation of the modern times, often m isunderstood as !!!! : only graphs and dashboards. Its time for busin esses to think beyond charts and start leveraging the science behind the data in 2018 "

### Cleaning with qdap

- The qdap package offers other text cleaning functions.
   Each is useful in its own way and is particularly powerful when combined with the others.
- bracketX(): Remove all text within brackets (e.g. "It's (so) cool" becomes "It's cool")
- replace\_number(): Replace numbers with their word equivalents (e.g. "2" becomes "two")
- replace\_abbreviation(): Replace abbreviations with their full text equivalents (e.g. "Sr" becomes "Senior")
- replace\_contraction(): Convert contractions back to their base words (e.g. "shouldn't" becomes "should not")
- replace\_symbol() Replace common symbols with their word equivalents (e.g. "\$" becomes "dollar")

### **Stop Words**

 Often there are words that are frequent but provide little information. So you may want to remove these so-called *stop words*. Some common English stop words include "I", "she'll", "the", etc. In the tm package, there are 174 stop words on this common list.

### **Stop Words**

- In fact, when you are doing an analysis you will likely need to add to this list. In our xiaomi tweet example, all tweets contain "xiaomi", so it's important to pull out that word in addition to the common stop words. Leaving it in doesn't add any insight and will cause it to be overemphasized in a frequency analysis.
- Using the c() function allows you to add new words (separated by commas) to the stop words list. For example, the following would add "word1" and "word2" to the default list of English stop words:
- all\_stops <- c("word1", "word2", stopwords("en"))</li>
- Once you have a list of stop words that makes sense, you will use the removeWords() function on your text.
- RemoveWords()takes two arguments: the text object to which it's being applied and the list of words to remove

### Tweets example

- stopwords("en")
- removeWords(xiaomi\_tweets, stopwords("en"))
- new\_stops <- c("xiaomi", "redmi", stopwords("en"))
- data<-removeWords(xiaomi\_tweets, new\_stops)

# Intro to word stemming and stem completion

- One useful preprocessing step involves word stemming and stem completion.
- The tm package provides the stemDocument() function to get to a word's root. This function either takes in a character vector and returns a character vector, or takes in a PlainTextDocument and returns a PlainTextDocument.

#### For example,

 stemDocument(c("computational", "computers", "computation"))

```
> complicate <- c("complicated", "complication", "complicatedly")</pre>
> # Perform word stemming: stem_doc
```

> stem\_doc <- stemDocument(complicate)</pre>

> # Create the completion dictionary: comp\_dict > comp\_dict <- "complicate"</pre>

> # Perform stem completion: complete\_text

> complete\_text <- stemCompletion(stem\_doc, comp\_dict)</pre>

> # Print complete\_text

> complete\_text complic complic complic

"complicate" "complicate" "complicate"

# Word stemming and stem completion on a sentence

**text\_data**<- "In a complicated haste, Ram rushed to fix a new complication, too complicatedly."

This sentence contains the same three forms of the word "complicate" that we saw in the previous exercise. The difference here is that even if you called stemDocument() on this sentence, it would return the sentence without stemming any words. Take a moment and try it out in the console. Be sure to include the punctuation marks.

### Stemming on sentence

- This happens because stemDocument() treats the whole sentence as one word.
- In other words, our document is a character vector of length 1, instead of length n, where n is the number of words in the document.
- To solve this problem, we first remove the punctation marks with the removePunctuation()function, which you learned a few exercises back.
- We then strsplit()this character vector of length 1 to length n,
- unlist(), then proceed to stem and re-complete.

```
Console ~/ML/bangaloresession1/ @
> text_data<- "In a complicated haste, Tom rushed to fix a new compl
ication, too complicatedly."
> # Remove punctuation: rm_punc
> rm_punc <- removePunctuation(text_data)</pre>
> # Create character vector: n char vec
> n_char_vec <- unlist(strsplit(rm_punc, split = ' '))</pre>
> #
> # Perform word stemming: stem_doc
> stem_doc <- stemDocument(n_char_vec)</pre>
> # Print stem_doc
> stem_doc
 [1] "In"
                           "complic" "hast"
                                                "Tom"
                                                            "rush"
 [7] "to"
            "fix"
                                      "new"
                                                 "complic" "too"
[13] "complic"
> # Re-complete stemmed document: complete_doc
> complete_doc <- stemCompletion(stem_doc,comp_dict)</pre>
> # Print complete_doc
> complete_doc
                                  complic
                                                   hast
                                                                   Tom
           In
                         "" "complicate"
           11 11
                                                                    11 11
                                      fix
        rush
                         to
                                                                   new
                                                       a
                         11 11
                                                      11 11
                                                                    11 11
                                  complic
     complic
                        too
```

"complicate" "" "complicate"

### Apply preprocessing steps to a corpus

```
clean_corpus <- function(corpus){</pre>
  corpus <- tm_map(corpus, stripWhitespace)</pre>
  corpus <- tm_map(corpus, removePunctuation)
  corpus <- tm_map(corpus, content_transformer(tolower))</pre>
  corpus <- tm_map(corpus, content_transformer(replace_abbreviation))</pre>
  corpus <- tm_map(corpus, removeNumbers)
  corpus <- tm_map(corpus, removeWords, c(stopwords("en"), "xiaomi", "redmi"))
  return(corpus)
```

```
Console ~/ML/bangaloresession1/ 🖒
> clean_corpus <- function(corpus){</pre>
      corpus <- tm_map(corpus, stripWhitespace)</pre>
      corpus <- tm_map(corpus, removePunctuation)</pre>
      corpus <- tm_map(corpus, content_transformer(tolower))</pre>
      corpus <- tm_map(corpus, content_transformer(replace_abbreviat</pre>
ion))
      corpus <- tm_map(corpus, removeNumbers)</pre>
     corpus <- tm_map(corpus, removeWords, c(stopwords("en"), "xiao
mi", "redmi"))
    return(corpus)
+ }
> # Apply your customized function to the tweet_corp: clean_corp
> clean_corp<-clean_corpus(xiaomi_corpus)</pre>
> # Print out a cleaned up tweet
> clean_corp[[227]][1]
$content
[1] "check shotonmia entries mi fans one think
                                                              best
                                                                       V0
ure u... httpstcojizojhm"
> # Print out the same tweet in original form
> tweets$text[227]
[1] "Check out some of the #ShotOnMiA1 entries by our Mi fans! Whic
h one do you think is the best in these?\nIf you're u... https://t.co/
ji087ZoJHM"
```

## TDM vs. DTM

	Tweet 1	Tweet 2	Tweet 3		Tweet N
Term 1	0	0	0	0	0
Term 2	1	1	0	0	0
Term 3	1	0	0	0	0
	0	0	3	1	1
Term M	0	0	0	1	0

	Term 1	Term 2	Term 3		Term M
Tweet 1	0	1	1	0	0
Tweet 2	0	1	0	0	0
Tweet 3	0	0	0	3	0
	0	0	0	1	1
Tweet N	0	0	0	1	0

Term Document Matrix (TDM)

Document Term Matrix (DTM)

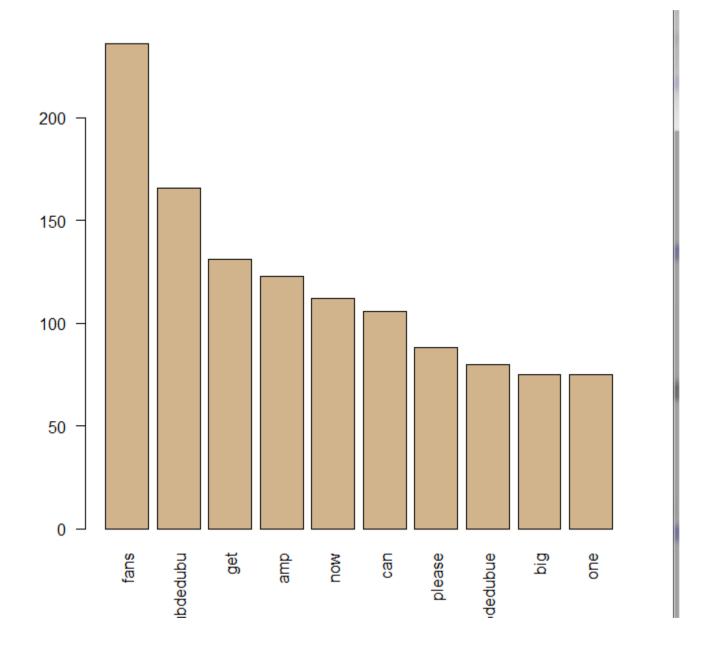
```
Console ~/ML/bangaloresession1/ 🖒
> xiaomi_tdm<-TermDocumentMatrix(clean_corp)</pre>
> xiaomi_dtm<-DocumentTermMatrix(clean_corp)</pre>
> xiaomi_m<- as.matrix(xiaomi_dtm)</pre>
> dim(xiaomi_m)
[1] 1561 4401
> xiaomi_m[100:103,2390:2393]
     Terms
      hugo hundreds hungama hungamacom
  100
  101
  102
  103
> xiaomi_m[148:150,2587:2590]
     Terms
Docs justoverspoken jyotigujarati kabiradvani kai
  148
  149
  150
```

Console ~/ML/bangaloresession1/ 🖒

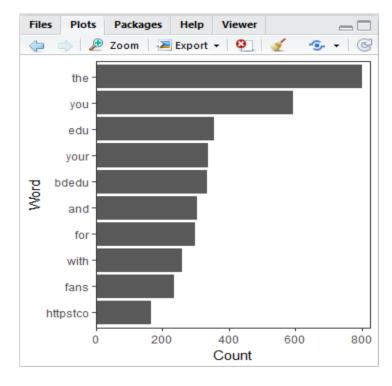
```
> xiaomi_t<-as.matrix(xiaomi_tdm)</pre>
> xiaomi_t[2587:2590,148:150]
                Docs
                  148 149 150
Terms
  justoverspoken
  iyotigujarati
  kabiradvani
  kai
> xiaomi_t[2087:2090,95:100]
                    Docs
                     95 96 97 98 99 100
Terms
  httpstcornbwkfhou
  httpstcornplffwfk 0 0 0 0
                                  0
  httpstcornsligl
  httpstcoroydv
```

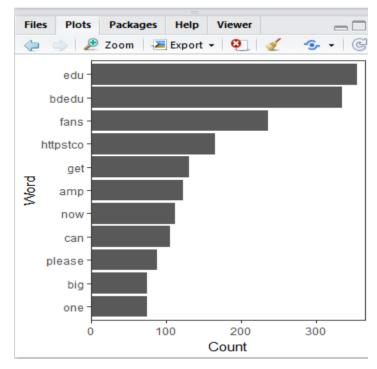
### Frequent terms with tm

```
00
Console ~/ML/bangaloresession1/ 🖒
> ## xiaomi_tdm is still loaded in your workspace
> # Create a matrix: xiaomi m
> xiaomi_m <- as.matrix(xiaomi_tdm)</pre>
> # Calculate the rowSums: term_frequency
> term_frequency <- rowSums(xiaomi_m)</pre>
> # Sort term_frequency in descending order
> term_frequency <- sort(term_frequency, decreasing = TRUE)</pre>
> # View the top 10 most common words
> term_frequency[1:10]
         fans eduaubdedubu
                                         get
                                                         amp
           236
                          166
                                         131
                                                         123
                                      please eduaubdedubue
                          can
          now
           112
                          106
                                          88
                                                          80
           big
                          one
                           75
> # Plot a barchart of the 10 most common words
> barplot(term_frequency[1:10], col = "tan", las = 2)
```



### Frequent terms with qdap





### Distance matrix and dendrogram

- A simple way to do word cluster analysis is with a dendrogram on your term-document matrix. Once you have a TDM, you can call dist() to compute the differences between each row of the matrix.
- Next, you call hclust() to perform cluster analysis on the dissimilarities of the distance matrix. Lastly, you can visualize the word frequency distances using a dendrogram and plot().

# Make a distance matrix and dendrogram from a TDM

- We can apply them to text. But first, you have to limit the number of words in your TDM using removeSparseTerms() from tm.
- Why would you want to adjust the sparsity of the TDM/DTM?
- TDMs and DTMs are sparse, meaning they contain mostly zeros.
- Remember that 1000 tweets can become a TDM with over 3000 terms! You won't be able to easily interpret a dendrogram that is so cluttered, especially if you are working on more text.
- A good TDM has between 25 and 70 terms.
- The lower the sparse value, the more terms are kept.
- The closer it is to 1, the fewer are kept.
- This value is a percentage cutoff of zeros for each term in the TDM.

```
Console ~/ML/bangaloresession1/ 🖒
> dim(xiaomi_tdm)
[1] 4401 1561
> tdm1 <- removeSparseTerms(xiaomi_tdm, sparse = 0.95)</pre>
> tdm1
<<TermDocumentMatrix (terms: 8, documents: 1561)>>
Non-/sparse entries: 1006/11482
Sparsity
                    : 92%
Maximal term length: 13
Weighting
          : term frequency (tf)
> tdm2 <- removeSparseTerms(xiaomi_tdm, sparse = 0.975)</pre>
> tdm2
<<TermDocumentMatrix (terms: 38, documents: 1561)>>
Non-/sparse entries: 2629/56689
Sparsity
                    : 96%
Maximal term length: 14
Weighting : term frequency (tf)
```

# Put it all together: a text based dendrogram

- Let's work to make your first text-based dendrogram. Remember, dendrograms reduce information to help you make sense of the data.
- This is much like how an average tells you something, but not everything, about a population. Both can be misleading.
- With text, there are often a lot of nonsensical clusters, but some valuable clusters may also appear.
- A peculiarity of TDM and DTM objects is that you have to convert them first to matrices (with as.matrix()), then to data frames (with as.data.frame()), before using them with the dist()function.

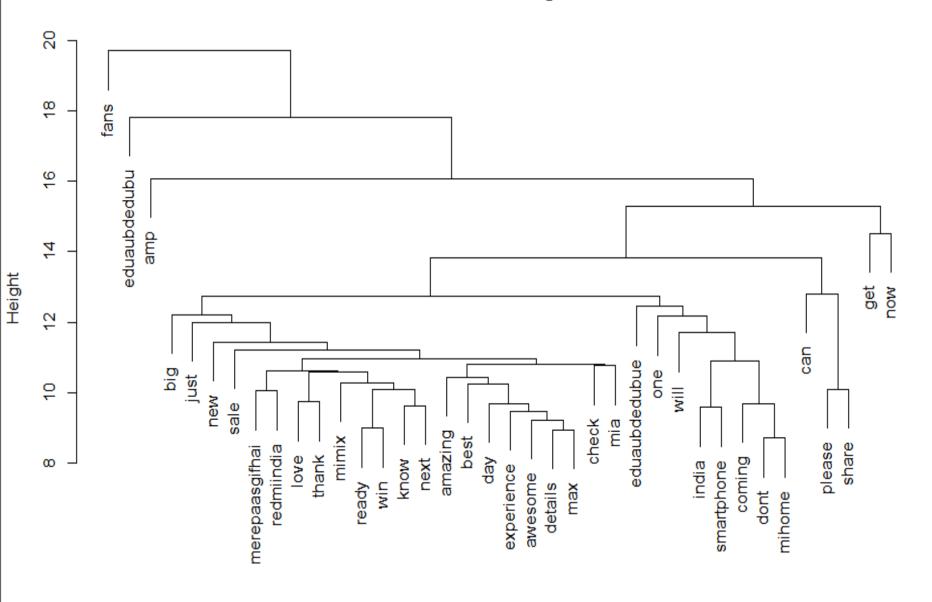
```
Console ~/ML/bangaloresession1/ 🖒
```



```
> tdm2 <- removeSparseTerms(xiaomi_tdm, sparse = 0.975)</pre>
```

- > tdm\_m<-as.matrix(tdm2)</pre>
- > tdm\_df<-as.data.frame(tdm\_m)</pre>
- > tweets\_dist<-dist(tdm\_df)</pre>
- > hc<-hclust(tweets\_dist)</pre>
- > plot(hc)

#### **Cluster Dendrogram**

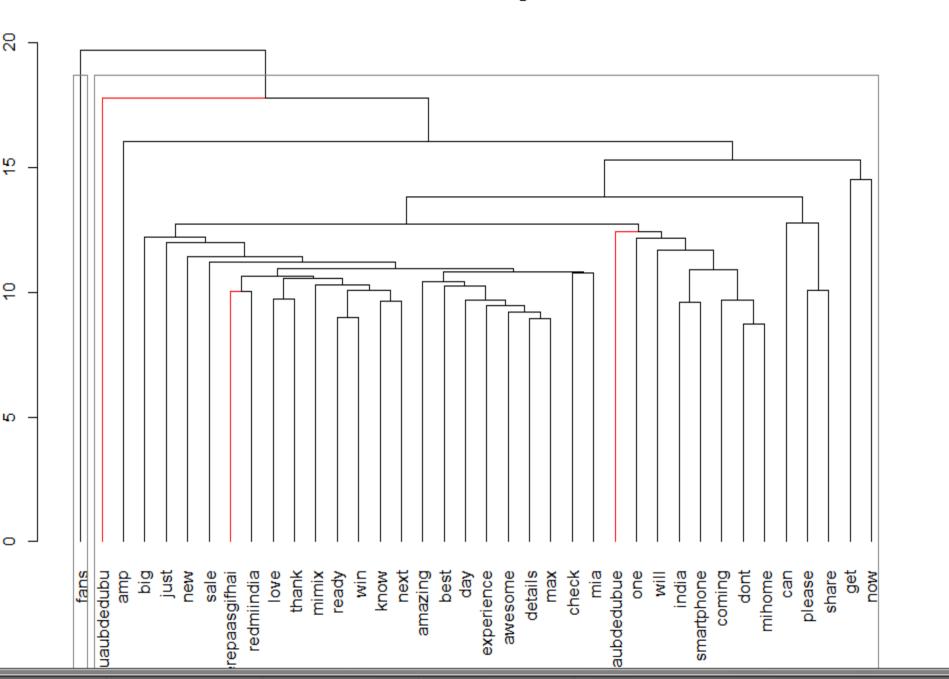


tweets\_dist hclust (\*, "complete")

### **Dendrogram aesthetics**

```
Source
Console ~/ML/bangaloresession1/ A
> library(dendextend)
> hcd<-as.dendrogram(hc)</pre>
> hcd <- branches_attr_by_labels(hcd, c("eduaubdedubu", "eduaubdedubu")</pre>
e", "merepaasgifhai"), "red")
> plot(hcd, main = "Better Dendrogram")
> rect.dendrogram(hcd, k = 2, border = "grey50")
```

#### **Better Dendrogram**



### Using word association

- Another way to think about word relationships is with the findAssocs() function in the tm package.
- For any given word, findAssocs() calculates its correlation with every other word in a TDM or DTM.
- Scores range from 0 to 1. A score of 1 means that two words always appear together, while a score of 0 means that they never appear together.
- To use findAssocs() pass in a TDM or DTM, the search term, and a minimum correlation.
- The function will return a list of all other terms that meet or exceed the minimum threshold.
- findAssocs(tdm, "word", 0.25) Minimum correlation values are often relatively low because of word diversity.
- Don't be surprised if 0.10 demonstrates a strong pairwise term association.

- # Create associations
- associations <- findAssocs(xiaomi\_tdm, "smart", 0.2)</li>
- # View the associations
- associations
- # Create associations\_df
- associations\_df <- list\_vect2df(associations)[, 2:3]</li>
- # Plot the associations\_df values (don't change this)ggplot(associations\_df, aes(y = associations\_df[, 1])) + geom\_point(aes(x = associations\_df[, 2]), data = associations\_df, size = 3)