Week 2 - Milestone Report

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Introduction and Preprocessing

For this assignment, I first loaded the necessary packages and the data. I also provided some basic information about the english dataset, like size in megabytes, number of lines, longest line in each file and number of words per line.

```
# Read the data in R
con <- file("./final/en US/en US.twitter.txt", "r")</pre>
twitter<-readLines(con,skipNul=TRUE,encoding="UTF-8")</pre>
close(con)
con<-file("./final/en_US/en_US.blogs.txt","r")</pre>
blogs<-readLines(con,skipNul=TRUE,encoding="UTF-8")</pre>
close(con)
con<-file("./final/en_US/en_US.news.txt","r")</pre>
news<-readLines(con,skipNul=TRUE,encoding="UTF-8")</pre>
close(con)
# some info about the dataset
# rounded size of files in megabytes
twitter_size <- round(file.info("./final/en_US/en_US.twitter.txt")$size / (1024^2))</pre>
blogs_size <- round(file.info("./final/en_US/en_US.blogs.txt")$size / (1024^2))
news_size <- round(file.info("./final/en_US/en_US.news.txt")$size / (1024^2))</pre>
# lines of files
#length(twitter)
#length(blogs)
#length(news)
# longest line in three files
#max(nchar(twitter))
#max(nchar(blogs))
#max(nchar(news))
# create a dataframe consisting of size, number of lines, max characters
df<-data.frame(file = c("twitter", "blogs", "news"),</pre>
                size_MB = c(twitter_size, blogs_size, news_size),
               num_lines = c(length(twitter), length(blogs), length(news)),
               longest_line_chars = c(max(nchar(twitter)), max(nchar(blogs)), max(nchar(news))),
               number_words = c(sum(stri_count_words(twitter)),sum(stri_count_words(blogs)),sum(stri_co
df
```

3 news 196 1010242 11384 34762395

Exploratory Analysis

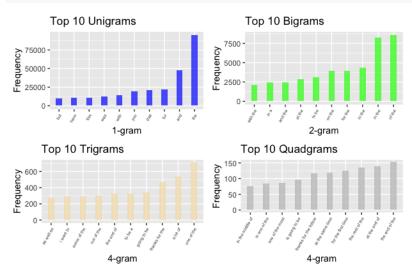
To continue, I proceeded with some exploratory analysis. I sampled 2% of the total data, cleaned it and plot the most frequent n-grams in histograms.

```
# Exploratory analysis to find the most frequent n-grams in a sample of data
# I sampled 2% of the data
set.seed(12345)
twitter_sample <- sample(twitter, 0.02 * length(twitter))</pre>
blogs_sample <- sample(blogs, 0.02 * length(blogs))</pre>
news_sample <- sample(news, 0.02 * length(news))</pre>
###$$$$$$$$$$$############
twitter_sample <- sample(twitter, size=10000, replace=TRUE)</pre>
blogs_sample <- sample(blogs, size=10000, replace=TRUE)</pre>
news_sample <- sample(news, size=10000, replace=TRUE)</pre>
# The data should be cleaned. Remove alpha-numeric characters and create a dataframe containing the sam
df_sample <- c(twitter_sample, blogs_sample, news_sample)</pre>
df_sample <- gsub("[^[:alnum:]']", " ",df_sample)</pre>
corpus sample <- Corpus(VectorSource(df sample))</pre>
corpus_sample <- tm_map(corpus_sample, tolower)</pre>
corpus sample <- tm map(corpus sample, removePunctuation)</pre>
corpus_sample <- tm_map(corpus_sample, removeNumbers)</pre>
corpus_sample <- tm_map(corpus_sample, stripWhitespace)</pre>
corpus_sample <- tm_map(corpus_sample, PlainTextDocument)</pre>
\# n-qrams
uniGramTokenizer <- function(x) NGramTokenizer(x, Weka_control(min = 1, max = 1))
biGramTokenizer <- function(x) NGramTokenizer(x, Weka_control(min = 2, max = 2))
triGramTokenizer <- function(x) NGramTokenizer(x, Weka_control(min = 3, max = 3))</pre>
fourGramTokenizer <- function(x) NGramTokenizer(x, Weka_control(min = 4, max = 4))</pre>
uniGrams <- TermDocumentMatrix(corpus_sample, control = list(tokenize = uniGramTokenizer))
biGrams <- TermDocumentMatrix(corpus_sample, control = list(tokenize = biGramTokenizer))</pre>
triGrams <- TermDocumentMatrix(corpus_sample, control = list(tokenize = triGramTokenizer))</pre>
fourGrams <- TermDocumentMatrix(corpus_sample, control = list(tokenize = fourGramTokenizer))</pre>
#####
bigram <- NGramTokenizer(corpus_sample, Weka_control(min = 2, max = 2,delimiters = " \\r\\n\\t.,;:\"()?
bigram <- data.frame(table(bigram))</pre>
bigram <- bigram[order(bigram$Freq,decreasing = TRUE),]</pre>
names(bigram) <- c("words", "freq")</pre>
head(bigram)
bigram$words <- as.character(bigram$words)</pre>
str2 <- strsplit(bigram$words,split=" ")</pre>
bigram <- transform(bigram,</pre>
                     one = sapply(str2,"[[",1),
                     two = sapply(str2,"[[",2))
bigram <- data.frame(word1 = bigram$one,word2 = bigram$two,freq = bigram$freq,stringsAsFactors=FALSE)
## saving files
```

```
#write.csv(bigram[bigram$freq > 1,],"bigram.csv",row.names=F)
#bigram <- read.csv("bigram.csv", stringsAsFactors = F)</pre>
saveRDS(bigram, "bigram.RData")
trigram <- NGramTokenizer(corpus_sample, Weka_control(min = 3, max = 3,delimiters = " \\r\\n\\t.,;:\"()</pre>
trigram <- data.frame(table(trigram))</pre>
trigram <- trigram[order(trigram$Freq,decreasing = TRUE),]</pre>
names(trigram) <- c("words", "freq")</pre>
head(trigram)
######################
trigram$words <- as.character(trigram$words)</pre>
str3 <- strsplit(trigram$words,split=" ")</pre>
trigram <- transform(trigram,</pre>
                      one = sapply(str3,"[[",1),
                      two = sapply(str3,"[[",2),
                      three = sapply(str3,"[[",3))
# trigram$words <- NULL
trigram <- data.frame(word1 = trigram$one,word2 = trigram$two,</pre>
                       word3 = trigram$three, freq = trigram$freq,stringsAsFactors=FALSE)
# saving files
#write.csv(trigram[trigram$freq > 1,],"trigram.csv",row.names=F)
#trigram <- read.csv("trigram.csv",stringsAsFactors = F)</pre>
saveRDS(trigram, "trigram.RData")
#####
quadgram <- NGramTokenizer(corpus_sample, Weka_control(min = 4, max = 4,delimiters = " \\r\\n\\t.,;:\"(
quadgram <- data.frame(table(quadgram))</pre>
quadgram <- quadgram[order(quadgram$Freq,decreasing = TRUE),]</pre>
names(quadgram) <- c("words", "freq")</pre>
quadgram$words <- as.character(quadgram$words)</pre>
str4 <- strsplit(quadgram$words,split=" ")</pre>
quadgram <- transform(quadgram,</pre>
                       one = sapply(str4,"[[",1),
                       two = sapply(str4,"[[",2),
                       three = sapply(str4,"[[",3),
                       four = sapply(str4,"[[",4))
# quadqram$words <- NULL
quadgram <- data.frame(word1 = quadgram$one,
                        word2 = quadgram$two,
                        word3 = quadgram$three,
                        word4 = quadgram$four,
                        freq = quadgram$freq, stringsAsFactors=FALSE)
# saving files
#write.csv(quadgram[quadgram$freq > 1,], "quadgram.csv",row.names=F)
#quadgram <- read.csv("quadgram.csv",stringsAsFactors = F)</pre>
saveRDS(quadgram, "quadgram.RData")
saveRDS(uniGrams,file="uniGrams.RData")
saveRDS(biGrams,file="biGrams.RData")
saveRDS(triGrams,file="triGrams.RData")
saveRDS(fourGrams,file="fourGrams.RData")
```

```
# plots together
# 1-grams
frequent terms <- findFreqTerms(uniGrams, lowfreq = 50)</pre>
frequency_terms <- rowSums(as.matrix(uniGrams[frequent_terms,]))</pre>
frequency_terms <- data.frame(unigram=names(frequency_terms), frequency=frequency_terms)</pre>
frequency_terms <- frequency_terms[order(-frequency_terms$frequency),][1:10,]</pre>
g1 <- ggplot(frequency_terms, aes(x=reorder(unigram, frequency), y=frequency)) +
  geom_bar(width=0.4,stat = "identity", fill = "blue", alpha=0.7) + xlab("1-gram") + ylab("Frequency")
  theme(axis.text.x = element_text(angle = 60, size = 5, hjust = 1)) +
  labs(title = "Top 10 Unigrams")
# 2-grams
frequent_terms <- findFreqTerms(biGrams, lowfreq = 50)</pre>
frequency_terms <- rowSums(as.matrix(biGrams[frequent_terms,]))</pre>
frequency_terms <- data.frame(bigram=names(frequency_terms), frequency=frequency_terms)</pre>
frequency_terms <- frequency_terms[order(-frequency_terms$frequency),][1:10,]</pre>
g2 <- ggplot(frequency_terms, aes(x=reorder(bigram, frequency), y=frequency)) +</pre>
  geom_bar(width=0.4,stat = "identity", fill = "green", alpha=0.7) + xlab("2-gram") + ylab("Frequency")
  theme(axis.text.x = element_text(angle = 60, size = 5, hjust = 1)) +
  labs(title = "Top 10 Bigrams")
# 3-grams
frequent_terms <- findFreqTerms(triGrams, lowfreq = 50)</pre>
frequency_terms <- rowSums(as.matrix(triGrams[frequent_terms,]))</pre>
frequency_terms <- data.frame(trigram=names(frequency_terms), frequency=frequency_terms)</pre>
frequency_terms <- frequency_terms[order(-frequency_terms$frequency),][1:10,]</pre>
g3 <- ggplot(frequency_terms, aes(x=reorder(trigram, frequency), y=frequency)) +
  geom_bar(width=0.4,stat = "identity", fill = "wheat", alpha=0.7)+
  theme(axis.text.x = element_text(angle = 60, size = 5, hjust = 1)) + xlab("4-gram") + ylab("Frequency
  labs(title = "Top 10 Trigrams")
# 4-grams
frequent_terms <- findFreqTerms(fourGrams, lowfreq = 50)</pre>
frequency_terms <- rowSums(as.matrix(fourGrams[frequent_terms,]))</pre>
frequency_terms <- data.frame(fourgram=names(frequency_terms), frequency=frequency_terms)</pre>
frequency_terms <- frequency_terms[order(-frequency_terms$frequency),][1:10,]</pre>
g4 <- ggplot(frequency_terms, aes(x=reorder(fourgram, frequency), y=frequency)) +
  geom_bar(width=0.4,stat = "identity", fill = "grey", alpha=0.7) +
  theme(axis.text.x = element_text(angle = 60, size = 5, hjust = 1)) +
  xlab("4-gram") + ylab("Frequency") +
  labs(title = "Top 10 Quadgrams")
grid.arrange(g1, g2, g3, g4, nrow = 2)
```

The calculations has already been made and saved to a png file. Therefore I loaded just the png fil
img <- load.image("/Users/iopetrid/Desktop/Coursera/Data Science/10_Capstone/Rplot.png")
plot(img,axes=FALSE)</pre>



Plans for the future

The following steps would be to create a shiny app that will predict the next word based on the word that has given as input.