# Data Science Capstone: Exploratory Analysis Milestone Report

#### **Load Data**

The first steps are to load the required libraries and set up the work environment.

```
Sys.setenv(JAVA_HOME="C:/Program Files (x86)/Java/jre1.8.0_91/")
library(RWeka)
library(dplyr)
library(stringi)
library(tm)
library(RWeka)
library(ggplot2)
```

It is assumed that the data has been dowloaded from here and unzipped: https://d396gusza40orc.cloudfront.net/dsscapstone/dataset/Coursera-SwiftKey.zip

Once it is, I can go ahead and read in the text files.

```
blogs <- readLines("P:/Documents/DataScience/Capstone/Data/Coursera-SwiftKey/final.blo
gs.txt", encoding = "UTF-8", skipNul = TRUE)

news <- readLines("P:/Documents/DataScience/Capstone/Data/Coursera-SwiftKey/final.news
.txt", encoding = "UTF-8", skipNul = TRUE)

twitter <- readLines("P:/Documents/DataScience/Capstone/Data/Coursera-SwiftKey/final.t
witter.txt", encoding = "UTF-8", skipNul = TRUE)</pre>
```

#### **Summary Statistics**

To get a sense of what the data looks like, I determine the number of lines, number of characters, and number of words for each of the 3 datasets (twitter, blogs, and news). I also calculate some basic statistics on the number of words per line (min, mean, and max).

```
WPL=sapply(list(blogs,news,twitter),function(x) summary(stri_count_words(x))[c('Min.',
'Mean','Max.')])
rownames(WPL)=c('WPL_Min','WPL_Mean','WPL_Max')
stats=data.frame(
    Dataset=c("blogs","news","twitter"),
```

```
t(rbind(
sapply(list(blogs,news,twitter),stri_stats_general)[c('Lines','Chars'),],
Words=sapply(list(blogs,news,twitter),stri_stats_latex)['Words',],
WPL)
))
head(stats)
```

```
## Dataset Lines Chars Words WPL_Min WPL_Mean WPL_Max

## 1 blogs 899288 206824382 37570839 0 41.75 6726

## 2 news 77259 15639408 2651432 1 34.62 1123

## 3 twitter 2360148 162096241 30451170 1 12.75 47
```

As we can see above, blogs tend to have the most words per line and tweets tend to have the least. This is what we would expect to see, given the character limit to tweets.

### Clean and sample data

I first go ahead and remove all non-English characters and then go ahead and compile a sample dataset that is composed of 1% of each of the 3 original datasets.

#### **Build corpus**

Next I will use the functions within the tm package to build and clean my corpus that will be analyzed. After building the corpus, I convert everything to lower case, remove punctuation and numbers, strip white space, and then convert it to plain text.

```
corpus <- VCorpus(VectorSource(sample_data))
corpus <- tm_map(corpus, tolower)
corpus <- tm_map(corpus, removePunctuation)</pre>
```

```
corpus <- tm_map(corpus, removeNumbers)

corpus <- tm_map(corpus, stripWhitespace)

corpus <- tm_map(corpus, PlainTextDocument)</pre>
```

#### **Tokenize and Calculate Frequencies of N-Grams**

I use the RWeka package to construct functions that tokenize the sample and construct matrices of uniqrams, bigrams, and trigrams.

```
uni_tokenizer <- function(x) NGramTokenizer(x, Weka_control(min = 1, max = 1))
bi_tokenizer <- function(x) NGramTokenizer(x, Weka_control(min = 2, max = 2))
tri_tokenizer <- function(x) NGramTokenizer(x, Weka_control(min = 3, max = 3))

uni_matrix <- TermDocumentMatrix(corpus, control = list(tokenize = uni_tokenizer))
bi_matrix <- TermDocumentMatrix(corpus, control = list(tokenize = bi_tokenizer))
tri_matrix <- TermDocumentMatrix(corpus, control = list(tokenize = tri_tokenizer))</pre>
```

Then I find the frequency of terms in each of these 3 matrices and construct dataframes of these frequencies. ###Calculate frequency of n-grams

```
uni_corpus <- findFreqTerms(uni_matrix,lowfreq = 50)
bi_corpus <- findFreqTerms(bi_matrix,lowfreq=50)
tri_corpus <- findFreqTerms(tri_matrix,lowfreq=50)

uni_corpus_freq <- rowSums(as.matrix(uni_matrix[uni_corpus,]))
uni_corpus_freq <- data.frame(word=names(uni_corpus_freq), frequency=uni_corpus_freq)
bi_corpus_freq <- rowSums(as.matrix(bi_matrix[bi_corpus,]))
bi_corpus_freq <- data.frame(word=names(bi_corpus_freq), frequency=bi_corpus_freq)
tri_corpus_freq <- rowSums(as.matrix(tri_matrix[tri_corpus,]))
tri_corpus_freq <- data.frame(word=names(tri_corpus_freq), frequency=tri_corpus_freq)
head(tri_corpus_freq)
## word frequency</pre>
```

```
## word frequency
## a couple of a couple of 89
## a little bit a little bit 50
```

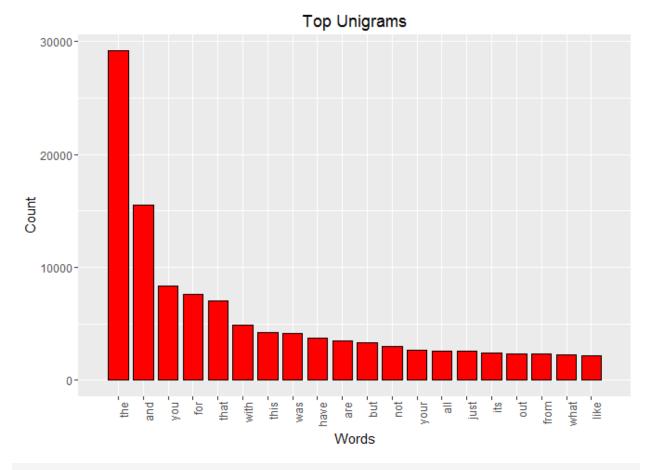
```
## a lot of a lot of 183
## all of the all of the 62
## as well as as well as 85
## back to the back to the 55
```

## Make plots

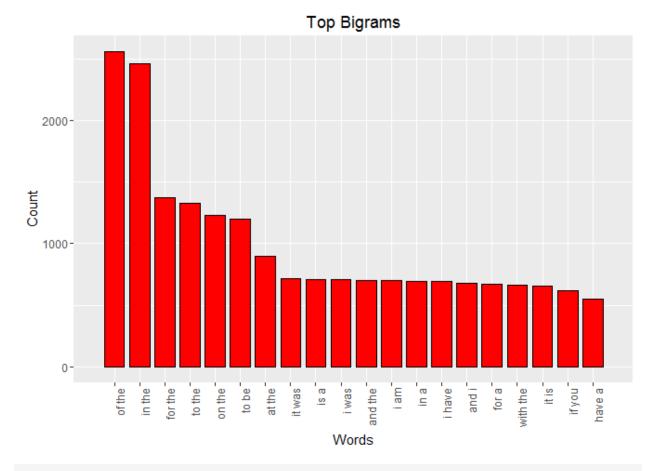
Lastly, I write a function to plot the n-gram frequency and go ahead and plot the 20 most frequent Unigrams, Bigrams, and Trigrams.

```
plot_n_grams <- function(data, title, num) {
    df2 <- data[order(-data$frequency),][1:num,]
    ggplot(df2, aes(x = seq(1:num), y = frequency)) +
        geom_bar(stat = "identity", fill = "red", colour = "black", width = 0.80) +
        coord_cartesian(xlim = c(0, num+1)) +
        labs(title = title) +
        xlab("Words") +
        ylab("Count") +
        scale_x_discrete(breaks = seq(1, num, by = 1), labels = df2$word[1:num]) +
        theme(axis.text.x = element_text(angle = 90, hjust = 1))
}

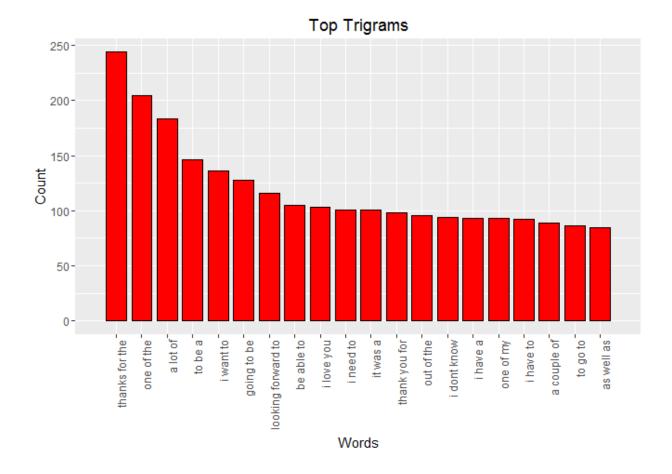
plot_n_grams(uni_corpus_freq, "Top Unigrams", 20)</pre>
```



plot\_n\_grams(bi\_corpus\_freq,"Top Bigrams",20)



plot\_n\_grams(tri\_corpus\_freq,"Top Trigrams",20)



## **Next steps**

This concludes the initial exploratory analysis. The next steps will be to build a predictive algorithm that uses an n-gram model with a frequency lookup similar to the analysis above. This algorithm will then be deployed in a Shiny app and will suggest the most likely next word after a phrase is typed.