Lab 6

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Data Setup

```
knitr::opts_chunk$set(eval = FALSE, include = TRUE)
 library(readr)
 library(dplyr)
Attaching package: 'dplyr'
The following objects are masked from 'package:stats':
    filter, lag
The following objects are masked from 'package:base':
    intersect, setdiff, setequal, union
mt_samples <- read_csv("https://raw.githubusercontent.com/USCbiostats/data-science-data/</pre>
New names:
• `` -> `...1`
Rows: 4999 Columns: 6
— Column specification
Delimiter: ","
chr (5): description, medical_specialty, sample_name, transcription, keywords
dbl (1): ...1
i Use `spec()` to retrieve the full column specification for this data.
i Specify the column types or set `show_col_types = FALSE` to quiet this message.
mt_samples <- mt_samples %>%
   select(description, medical_specialty, transcription)
 head(mt_samples)
# A tibble: 6 \times 3
  description
                                                  medical_specialty transcription
  <chr>
                                                   <chr>
                                                                     <chr>
1 A 23-year-old white female presents with comp... Allergy / Immuno... "SUBJECTIVE:...
2 Consult for laparoscopic gastric bypass.
                                                                     "PAST MEDICA...
                                                   Bariatrics
3 Consult for laparoscopic gastric bypass.
                                                                     "HISTORY OF ...
                                                   Bariatrics
4 2-D M-Mode. Doppler.
                                                  Cardiovascular /... "2-D M-MODE:...
5 2-D Echocardiogram
                                                  Cardiovascular /... "1. The lef...
6 Morbid obesity. Laparoscopic antecolic anteg... Bariatrics
                                                                     "PREOPERATIV...
```

Question 1

```
library(dplyr)
mt_samples %>%
  count(medical_specialty, sort = TRUE)
# surgery is the most repeated category among the medical specialties
```

Question 2

```
library(tm)
library(ggplot2)
corpus <- Corpus(VectorSource(mt_samples$transcription))</pre>
corpus <- tm_map(corpus, content_transformer(tolower))</pre>
corpus <- tm_map(corpus, removePunctuation)</pre>
corpus <- tm_map(corpus, removeNumbers)</pre>
corpus <- tm_map(corpus, removeWords, stopwords("english"))</pre>
corpus <- tm_map(corpus, stripWhitespace)</pre>
# from this we get shortened versions of what happnened, smiliar to short hand form when
library(Matrix)
dtm <- DocumentTermMatrix(corpus)</pre>
word_freq <- rowSums(as.matrix(dtm))</pre>
word_freq_df <- data.frame(word = names(word_freq), freq = word_freq)</pre>
word_freq_df <- word_freq_df[order(-word_freq_df$freq), ]</pre>
top_words <- head(word_freq_df, 20)</pre>
ggplot(top\_words, aes(x = reorder(word, -freq), y = freq)) +
 geom_bar(stat = "identity", fill = "blue") +
 coord_flip() +
 labs(x = "Words", y = "Frequency", title = "Top 20 Most Frequent Words") +
 theme_minimal() +
```

Question 3

theme(axis.text.y = element_text(size = 12))

```
corpus <- tm_map(corpus, removeWords, stopwords("english"))</pre>
corpus <- tm_map(corpus, removeNumbers)</pre>
dtm <- DocumentTermMatrix(corpus)</pre>
word_freq <- rowSums(as.matrix(dtm))</pre>
word_freq_df <- data.frame(word = names(word_freq), freq = word_freq)</pre>
word_freq_df <- word_freq_df[order(-word_freq_df$freq), ]</pre>
top_words <- head(word_freq_df, 20)</pre>
ggplot(top\_words, aes(x = reorder(word, -freq), y = freq)) +
 geom_bar(stat = "identity", fill = "blue") +
 coord_flip() +
 labs(x = "Words", y = "Frequency", title = "Top 20 Most Frequent Words (Stopwords and
 theme_minimal() +
 theme(axis.text.y = element_text(size = 12))
```

Question 4

```
bigram_tokenizer <- function(x) {</pre>
  unlist(lapply(ngrams(words(x), 2), paste, collapse = " "))
corpus_bigrams <- Corpus(VectorSource(corpus))</pre>
corpus_bigrams <- tm_map(corpus_bigrams, content_transformer(tolower))</pre>
corpus_bigrams <- tm_map(corpus_bigrams, removePunctuation)</pre>
corpus_bigrams <- tm_map(corpus_bigrams, removeNumbers)</pre>
corpus_bigrams <- tm_map(corpus_bigrams, removeWords, stopwords("english"))</pre>
corpus_bigrams <- tm_map(corpus_bigrams, stripWhitespace)</pre>
corpus_bigrams <- tm_map(corpus_bigrams, content_transformer(bigram_tokenizer))</pre>
trigram_tokenizer <- function(x) {</pre>
  unlist(lapply(ngrams(words(x), 3), paste, collapse = " "))
corpus_trigrams <- Corpus(VectorSource(corpus))</pre>
corpus_trigrams <- tm_map(corpus_trigrams, content_transformer(tolower))</pre>
corpus_trigrams <- tm_map(corpus_trigrams, removePunctuation)</pre>
corpus_trigrams <- tm_map(corpus_trigrams, removeNumbers)</pre>
corpus_trigrams <- tm_map(corpus_trigrams, removeWords, stopwords("english"))</pre>
corpus_trigrams <- tm_map(corpus_trigrams, stripWhitespace)</pre>
corpus_trigrams <- tm_map(corpus_trigrams, content_transformer(trigram_tokenizer))</pre>
# the bi gram gives very little information outside the race and sex of the person, the
```

Question 5

```
count_words_around_target <- function(corpus, female) {</pre>
   before counts <- numeric(♥)
   after_counts <- numeric(0)</pre>
   for (doc in corpus) {
     tokens <- unlist(strsplit(as.character(doc$content), " "))</pre>
     target_positions <- which(tokens == target_word)</pre>
     for (position in target_positions) {
       if (position > 1) {
          before_word <- tokens[position - 1]</pre>
         before_counts <- append(before_counts, before_word)</pre>
       if (position < length(tokens)) {</pre>
         after_word <- tokens[position + 1]</pre>
         after_counts <- append(after_counts, after_word)</pre>
   }}
Question 6
```

library(dplyr)

```
library(tm)
 corpus <- Corpus(VectorSource(mt samples))</pre>
 corpus <- tm_map(corpus, content_transformer(tolower))</pre>
 corpus <- tm_map(corpus, removePunctuation)</pre>
 corpus <- tm_map(corpus, removeNumbers)</pre>
 corpus <- tm_map(corpus, removeWords, stopwords("english"))</pre>
 corpus <- tm_map(corpus, stripWhitespace)</pre>
 corpus <- tm_map(corpus, content_transformer(tokenize))</pre>
 top_words_5 <- head(corpus, 5)</pre>
Question 7
```

```
library(dplyr)
library(tm)
corpus <- Corpus(VectorSource(corpus$content))</pre>
corpus <- tm_map(corpus, content_transformer(tolower))</pre>
corpus <- tm_map(corpus, removePunctuation)</pre>
corpus <- tm_map(corpus, removeNumbers)</pre>
corpus <- tm_map(corpus, removeWords, stopwords("english"))</pre>
corpus <- tm_map(corpus, stripWhitespace)</pre>
top_words_2 <- head(corpus, 2)</pre>
```