Lab 6

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Lab 06 - Text Mining

```
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
library(ggplot2)
library(tidytext)
library(readr)
mt_samples <- read_csv("https://raw.githubusercontent.com/USCbiostats/data-science-data/maste</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 x 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.
                                                 Bariatrics
                                                                    "PAST MEDICA~
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: What specialties do we have?

There are 40 different medical specialties. Specialties such as Cosmetic / Plastic Surgery and Dentistry both have 27 counts. Diets and Nutrition and Rheumatology specialties both have counts of 10. Autopsy and Lab Medicine - Pathology specialties both have counts of 8. There does not appear to be an even distribution between the medical specialties as we can see that Surgery has 1103 counts compared to

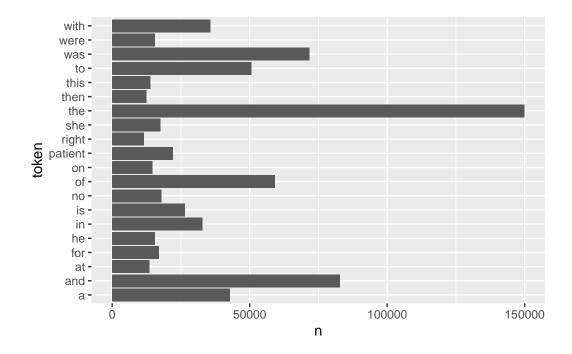
Hospice - Palliative Care with only 6 counts.

```
med_specialty_counts <- mt_samples |>
  count(medical_specialty, name = "n", sort = TRUE)
print(med_specialty_counts)
```

```
# A tibble: 40 x 2
  medical_specialty
                                      n
   <chr>
                                  <int>
                                   1103
1 Surgery
2 Consult - History and Phy.
                                    516
3 Cardiovascular / Pulmonary
                                    372
4 Orthopedic
                                    355
5 Radiology
                                    273
6 General Medicine
                                    259
7 Gastroenterology
                                    230
8 Neurology
                                    223
9 SOAP / Chart / Progress Notes
                                    166
```

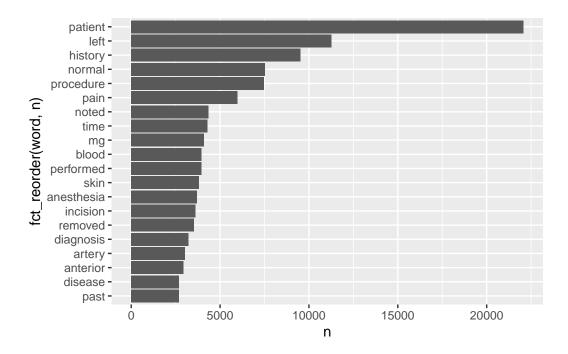
The list shows that the word "the" appears the most (149888 times) in the text. This makes sense because stop words usually appear the most in English text. Looking at the top tenth word that appears the most, patient, which appears 22065 times, it does give us an insight that the text is focused on medical transcripts mainly revolving around patient interactions.

```
mt_samples |>
  unnest_tokens(token, transcription) |>
  count(token) |>
  top_n(20, n) |>
  ggplot(aes(n, token)) +
  geom_col()
```



Now that we have removed stop words, we can see that the word 'patient' appears the most which is more fitting given that this is a medical transcript. Looking at the rest of the top 20 words, it is clear that this text is about patient procedures or charting.

```
library(forcats)
library(tidytext)
mt_samples |>
  unnest_tokens(word, transcription) |>
  anti_join(stop_words, by = "word") |>
  filter(!grepl("[0-9]", word)) |>
  count(word, sort = TRUE) |>
  top_n(20, n) |>
  ggplot(aes(n, fct_reorder(word, n))) +
  geom_col()
```



We have a lot more insight into what the text is about when tokenizing into tri-grams rather than bi-grams. Bi-grams is mostly stop words but looking at tri-grams, we can see more insight into procedures and even patient symptoms.

```
mt_samples |>
unnest_tokens(bigram, transcription, token = "ngrams", n = 2) |>
count(bigram, sort = TRUE)
```

```
# A tibble: 301,415 x 2
  bigram
                    n
   <chr>
               <int>
1 the patient 20307
2 of the
               19062
3 in the
               12790
4 to the
               12374
5 was then
                6956
6 and the
                6350
7 patient was
                6293
8 the right
                5509
9 on the
                5241
```

```
10 the left
                4860
# i 301,405 more rows
mt samples |>
  unnest_tokens(trigram, transcription, token = "ngrams", n = 3) |>
  count(trigram, sort = TRUE)
# A tibble: 655,441 x 2
   trigram
                          n
   <chr>
                       <int>
                       6104
 1 the patient was
 2 the patient is
                       3075
                       2243
 3 as well as
 4 there is no
                       1678
 5 the operating room 1532
 6 patient is a
                       1491
 7 prepped and draped 1490
 8 was used to
                       1480
 9 and draped in
                       1372
10 at this time
                       1333
# i 655,431 more rows
```

```
library(stringr)
library(tidyr)
word_to_analyze <- "patient"
bi_grams <- mt_samples|>
unnest_tokens(bigram, transcription, token = "ngrams", n = 2)
print(head(bi_grams))
```

```
before_after <- bi_grams|>
filter(str_detect(bigram, word_to_analyze))
before_after <- before_after |>
separate(bigram, into = c("word1", "word2"), sep = " ")
before_count <- before_after |>
filter(word2 == word_to_analyze) |>
count(word1, sort = TRUE) |>
rename(before = word1)
after_count <- before_after |>
filter(word1 == word_to_analyze) |>
count(word2, sort = TRUE) |>
rename(after = word2)
print("Words Before 'patient':")
[1] "Words Before 'patient':"
print(before_count)
# A tibble: 269 x 2
  before
                n
  <chr> <int>
           20307
 1 the
 2 this
             470
 3 history 101
 4 a
              67
               47
 5 and
 6 procedure
             32
 7 female
              26
 8 with
               25
 9 use
               24
10 old
# i 259 more rows
print("Words After 'patient':")
```

[1] "Words After 'patient':"

print(after_count)

```
# A tibble: 588 x 2
  after
                 n
   <chr>
             <int>
              6293
1 was
2 is
              3332
3 has
              1417
4 tolerated
               994
5 had
               888
6 will
               616
7 denies
               552
8 and
               377
9 states
               363
10 does
               334
# i 578 more rows
```

Question 6

The most used word in allergy/immunology is 'history.' Autopsy is 'left,' Bariatrics is 'patient,' etc. The top 5 most used words include 'patient' 'left' 'history' '2', and '1'.

```
most_used_words <- mt_samples |>
  unnest_tokens(word, transcription) |>
  anti_join(stop_words, by = "word") |>
  group_by(medical_specialty, word) |>
  count(n = n(), sort = TRUE) |>
  arrange(medical_specialty, desc(n))
```

Storing counts in `nn`, as `n` already present in input i Use `name = "new_name"` to pick a new name.

```
2 Allergy / Immunology noted
                                    1263045
                                               23
3 Allergy / Immunology patient
                                    1263045
                                               22
4 Allergy / Immunology allergies
                                    1263045
                                               21
5 Allergy / Immunology nasal
                                    1263045
                                                13
6 Allergy / Immunology past
                                                13
                                    1263045
7 Allergy / Immunology bilaterally 1263045
                                                12
8 Allergy / Immunology masses
                                    1263045
                                                12
9 Allergy / Immunology asthma
                                    1263045
                                                11
10 Allergy / Immunology medical
                                    1263045
                                                11
# i 149,963 more rows
```

```
most_used_words <- mt_samples|>
  unnest_tokens(word, transcription)|>
  anti_join(stop_words, by = "word") |>
  count(word, sort = TRUE) |>
  top_n(5, n)
print(most_used_words)
```