Topic Modeling

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Packages

```
library(tidyverse)
-- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
v dplyr 1.1.4 v readr 2.1.5
1.0.2
v purrr
-- Conflicts ----- tidyverse_conflicts() --
x dplyr::filter() masks stats::filter()
x dplyr::lag()
               masks stats::lag()
i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicted.r-lib.org/</a>) to force all conflicts to become
  library(tm)
Loading required package: NLP
Attaching package: 'NLP'
The following object is masked from 'package:ggplot2':
   annotate
  library(topicmodels)
  library(ldatuning)
```

```
library(tidytext)
library(Rtsne)
library(ggplot2)
library(wordcloud)

Loading required package: RColorBrewer
library(RColorBrewer)
```

Data Cleaning

```
movie_data = read_csv("movie_plots.csv")
view(movie_data)

plots_by_word <- movie_data %>% unnest_tokens(word, Plot)
plot_word_counts <- plots_by_word %>%
    anti_join(stop_words,by = join_by(word)) %>%
    count("Movie Name", word, sort = TRUE)

corpus <- VCorpus(VectorSource(movie_data$Plot))

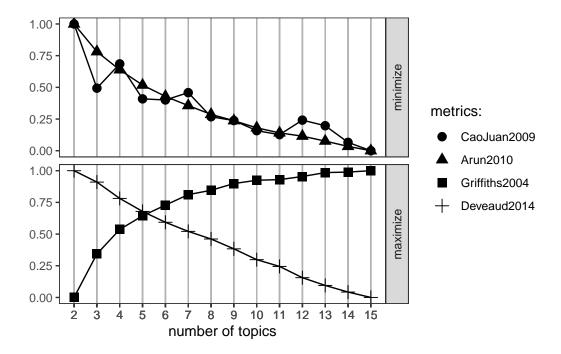
corpus <- corpus %>%
    tm_map(content_transformer(tolower)) %>%
    tm_map(removePunctuation) %>%
    tm_map(removeWords, stopwords("english")) %>%
    tm_map(stripWhitespace)

dtm <- DocumentTermMatrix(corpus)</pre>
```

- 1. Tokenization: split the "Plot" column into individual words.
- 2. Removing Stop Words: remove common stop words (like "the," "and," "is") which do not contribute to meaningful topics.
- 3. Counting Words: count the occurrences of each word, sorted by frequency.

Visualizations

```
result <- FindTopicsNumber(</pre>
       topics = seq(2, 15, by = 1),
       metrics = c("CaoJuan2009", "Arun2010", "Griffiths2004", "Deveaud2014"),
       method = "Gibbs",
       control = list(seed = 123456),
       mc.cores = 1L,
       verbose = TRUE
  )
fit models... done.
calculate metrics:
  CaoJuan2009... done.
  Arun2010... done.
  Griffiths2004... done.
  Deveaud2014... done.
  FindTopicsNumber_plot(result)
Warning: The `<scale>` argument of `guides()` cannot be `FALSE`. Use "none" instead as
of ggplot2 3.3.4.
i The deprecated feature was likely used in the ldatuning package.
  Please report the issue at <a href="https://github.com/nikita-moor/ldatuning/issues">https://github.com/nikita-moor/ldatuning/issues</a>.
```



For each metric, identify the point at which the metric either stabilizes or reaches an optimal value. The intersection of these indicators provides a recommendation for the best number of topics. Around k=6 to k=8, most metrics start to stabilize or reach optimal values, we could choose one of these as the ideal number of topics. Let's try k=8.

```
#set k to 8
k <- 8
lda_model <- LDA(dtm, k = k, control = list(seed = 123456))

gamma_matrix <- posterior(lda_model)$topics

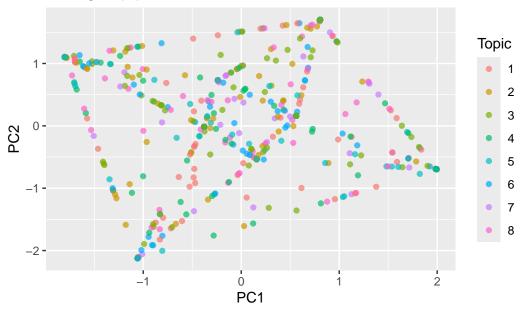
pca_model <- prcomp(gamma_matrix, center = TRUE, scale. = TRUE)
pca_data <- as.data.frame(pca_model$x)

document_topics <- tidy(lda_model, matrix = "gamma")
doc_topic <- document_topics %>%
    group_by(document) %>%
    slice_max(gamma, n = 1) %>%
    ungroup()

pca_data$Topic <- factor(doc_topic$topic)</pre>
```

```
ggplot(pca_data, aes(x = PC1, y = PC2, color = Topic)) +
geom_point(alpha = 0.7) +
labs(title = "PCA group plot", x = "PC1", y = "PC2")
```

PCA group plot



The code fits an LDA model with k=8 topics on a document-term matrix derived from movie plots, extracts the document-topic probabilities (gamma matrix), and uses PCA to reduce the dimensionality for visualization. Each document is assigned to its dominant topic based on the highest probability, and a scatter plot is generated to display the documents in a 2D space with colors representing different topics. The resulting PCA plot reveals how documents cluster by topic, illustrating the distinctiveness and relationships between topics by showing how closely documents with similar themes group together.

```
document_topics <- tidy(lda_model, matrix = "gamma")

# Gamma plot
ggplot(document_topics, aes(x = gamma)) +
    geom_histogram(bins = 30, fill = "steelblue", color = "black", alpha = 0.7) +
    facet_wrap(~ topic, scales = "free_y") +
    labs(title = "Gamma Plot", x = "Gamma value", y = "file number")</pre>
```

2 1 3 750 750 750 500 -500 500 250 -250 -250 0 0 0 6 4 5 file number 750 -750 **-**750 -500 -500 -500 -250 -250 -250 -0 0 0 0.00 0.25 0.50 0.75 1.00

750 **-**

500 -

250 **-**

1.00

8

0.00 0.25 0.50 0.75 1.00

Gamma value

Gamma Plot

7

0.00 0.25 0.50 0.75

750 500

250

The Gamma Plot visualizes the distribution of topic probabilities (gamma values) for each document across the eight topics in the LDA model. Each subplot shows the gamma values for a specific topic, revealing that most documents have low gamma values for most topics but a high gamma value for one topic, indicating strong association with a single dominant topic. This pattern confirms that the model effectively assigns documents to distinct topics, as expected in well-separated topic modeling, where each document is primarily linked to one topic rather than spread across multiple topics.

```
topic_terms <- tidy(lda_model, matrix = "beta")

top_terms <- topic_terms %>%
    group_by(topic) %>%
    slice_max(beta, n = 10) %>%
    ungroup() %>%
    arrange(topic, -beta)

ggplot(top_terms, aes(x = reorder(term, beta), y = beta, fill = factor(topic))) +
    geom_bar(stat = "identity", show.legend = FALSE) +
    facet_wrap(~ topic, scales = "free") +
    coord_flip() +
    labs(title = "Beta Plot", x = "words", y = "Beta value")
```



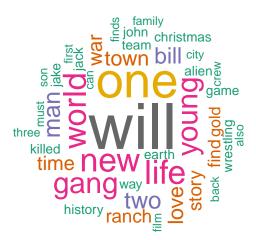
The Beta Plot displays the top 10 terms for each of the eight topics in the LDA model, with each term's importance measured by its beta value. Beta values represent the probability of a word being associated with a particular topic, so higher beta values indicate words that are more representative of the topic. Each subplot corresponds to one topic, showing the most significant words in descending order of their beta values. This plot provides insight into the defining terms of each topic, making it easier to interpret and label the topics based on the prominent words associated with them. This visualization helps understand the thematic structure of each topic in the model.

```
library(RColorBrewer)

all_terms <- top_terms %>%
    group_by(term) %>%
    summarize(total_beta = sum(beta)) %>%
    arrange(desc(total_beta))

palette <- brewer.pal(8, "Dark2")
wordcloud(words = all_terms$term,
    freq = all_terms$total_beta,
    min.freq = 0.001,
    colors = palette,
    random.order = FALSE,
    rot.per = 0.35,</pre>
```

```
scale = c(4, 0.5),
main = "wordcloud")
```



It generates a word cloud that visualizes the most important terms across all topics in the LDA model based on their aggregated beta values, which represent each term's overall significance in the topic structure. Terms are sized according to their total beta score, with larger words like "will," "one," "world," and "life" indicating higher importance across multiple topics. A color palette from library RColorBrewer is used to add visual distinction to the words, enhancing readability. This word cloud provides a quick, intuitive overview of the central themes and recurring terms in the movie plot dataset.