

Package ‘conversim’

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Title Conversation Similarity Analysis

Version 0.1.0

Description Analyze and compare conversations using various similarity measures including topic, lexical, semantic, structural, stylistic, sentiment, participant, and timing similarities. Supports both pairwise conversation comparisons and analysis of multiple dyads. Methods are based on established research: Topic modeling: Blei et al. (2003) <[doi:10.1162/jmlr.2003.3.4-5.993](https://doi.org/10.1162/jmlr.2003.3.4-5.993)>; Landauer et al. (1998) <[doi:10.1080/01638539809545028](https://doi.org/10.1080/01638539809545028)>; Lexical similarity: Jaccard (1912) <[doi:10.1111/j.1469-8137.1912.tb05611.x](https://doi.org/10.1111/j.1469-8137.1912.tb05611.x)>; Semantic similarity: Salton & Buckley (1988) <[doi:10.1016/0306-4573\(88\)90021-0](https://doi.org/10.1016/0306-4573(88)90021-0)>; Mikolov et al. (2013) <[doi:10.48550/arXiv.1301.3781](https://doi.org/10.48550/arXiv.1301.3781)>; Pennington et al. (2014) <[doi:10.3115/v1/D14-1162](https://doi.org/10.3115/v1/D14-1162)>; Structural and stylistic analysis: Graesser et al. (2004) <[doi:10.1075/target.21131.ryu](https://doi.org/10.1075/target.21131.ryu)>; Sentiment analysis: Rinker (2019) <<https://github.com/trinker/sentimentr>>.

License GPL (>= 3)

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agg_seq	<i>Aggregate Similarity Sequence</i>
---------	--------------------------------------

Description

Aggregate similarity sequence for a single dyad

Usage

```
agg_seq(sequence, num_segments)
```

Arguments

sequence A numeric vector of similarity scores for a single dyad
num_segments The number of segments to aggregate into

Details

This function aggregates a similarity sequence into a specified number of segments for a single dyad.

Value

A numeric vector of aggregated similarity scores

Examples

```
seq <- c(0.5, 0.6, 0.7, 0.6, 0.8, 0.7, 0.9, 0.8, 0.7, 0.8)
# Aggregate the sequence into 3 segments
agg_3 <- agg_seq(seq, 3)
print(agg_3)

# Aggregate the sequence into 5 segments
agg_5 <- agg_seq(seq, 5)
print(agg_5)
```

calc_sim_cor	<i>Calculate correlation between similarity measures</i>
--------------	--

Description

This function calculates the correlation between different similarity measures.

Usage

```
calc_sim_cor(comparison_df)
```

Arguments

comparison_df A data frame output from compare_sim_meas()

Value

A correlation matrix

Examples

```
topic_similarities <- list("1" = c(0.5, 0.6, 0.7), "2" = c(0.4, 0.5, 0.6))
lexical_similarities <- list("1" = c(0.6, 0.7, 0.8), "2" = c(0.5, 0.6, 0.7))
comparison_df <- compare_sim_meas(
  list(topic_similarities, lexical_similarities),
  c("Topic", "Lexical")
)
calc_sim_cor(comparison_df)
print(plot)
```

calc_sim_seq	<i>Calculate similarity sequence</i>
--------------	--------------------------------------

Description

This function calculates a sequence of similarities between consecutive windows in a conversation.

Usage

```
calc_sim_seq(conversation, window_size, similarity_func)
```

Arguments

conversation A dataframe containing the conversation, with a column named 'processed_text'.
 window_size An integer specifying the size of each window.
 similarity_func A function that calculates similarity between two text strings.

Value

A list containing two elements:

sequence A numeric vector of similarity scores between consecutive windows
 average The mean of the similarity scores

Examples

```
conversation <- data.frame(processed_text = c("hello", "world", "how", "are", "you"))
result <- calc_sim_seq(conversation, 2, function(x, y) sum(x == y) / max(length(x), length(y)))
```

calc_sum_stats	<i>Calculate summary statistics for similarities</i>
----------------	--

Description

This function calculates summary statistics for the similarities of multiple dyads.

Usage

```
calc_sum_stats(similarities)
```

Arguments

`similarities` A list of similarity sequences for each dyad

Value

A matrix with summary statistics for each dyad

Examples

```
similarities <- list(  
  "1" = c(0.5, 0.6, 0.7),  
  "2" = c(0.4, 0.5, 0.6)  
)  
calc_sum_stats(similarities)  
print(plot)
```

combine_sims	<i>Utility and visualization functions for speech similarity analysis</i>
--------------	---

Description

This file contains utility functions and visualization tools to complement the main similarity calculation functions for comparing two speeches. Combine multiple similarity measures

Usage

```
combine_sims(similarities, weights = NULL)
```

Arguments

`similarities` A named list of similarity scores
`weights` A named list of weights for each similarity measure (optional)

Details

This function combines multiple similarity measures into a single score.

Value

A single combined similarity score

Examples

```
sims <- list(topic = 0.8, lexical = 0.6, semantic = 0.7, structural = 0.9)
combine_sims(sims)
combine_sims(sims, weights = list(topic = 2, lexical = 1, semantic = 1.5, structural = 1))
print(plot)
```

combine_sim_seq

Combine Similarity Measures

Description

Combine similarity measures for a single dyad

Usage

```
combine_sim_seq(similarities, weights = NULL)
```

Arguments

`similarities` A list of similarity measures for a single dyad
`weights` A numeric vector of weights for each similarity measure (default is equal weights)

Details

This function combines multiple similarity measures into a single overall similarity score for a single dyad.

Value

A list containing the combined sequence and average similarity

Examples

```
sim1 <- list(sequence = c(0.8, 0.7, 0.9), average = 0.8)
sim2 <- list(sequence = c(0.6, 0.8, 0.7), average = 0.7)
combine_sim_seq(list(sim1, sim2))
print(plot)
```

compare_sim_meas	<i>Compare multiple similarity measures</i>
------------------	---

Description

This function compares multiple similarity measures for the same set of dyads.

Usage

```
compare_sim_meas(similarity_list, measure_names)
```

Arguments

`similarity_list` A list of lists, where each inner list contains similarities for each dyad

`measure_names` A vector of names for each similarity measure

Value

A data frame with all similarity measures for each dyad

Examples

```
topic_similarities <- list("1" = c(0.5, 0.6, 0.7), "2" = c(0.4, 0.5, 0.6))
lexical_similarities <- list("1" = c(0.6, 0.7, 0.8), "2" = c(0.5, 0.6, 0.7))
compare_sim_meas(
  list(topic_similarities, lexical_similarities),
  c("Topic", "Lexical")
)
print(plot)
```

compare_style	<i>Compare stylistic features</i>
---------------	-----------------------------------

Description

This function visualizes the comparison of stylistic features between two speeches.

Usage

```
compare_style(stylistic_result)
```

Arguments

`stylistic_result` The result from `stylistic_similarity` function

Value

A ggplot object

Examples

```
text1 <- "The quick brown fox jumps over the lazy dog. It's a sunny day."  
text2 <- "A lazy cat sleeps on the warm windowsill. Birds chirp outside."  
result <- stylistic_similarity(text1, text2)  
compare_style(result)  
print(plot)
```

cor_sim_seq	<i>Calculate Correlation Between Similarity Measures for a Single Dyad</i>
-------------	--

Description

Calculate Correlation Between Similarity Measures for a Single Dyad

Usage

```
cor_sim_seq(similarities, method = "pearson")
```

Arguments

`similarities` A list of similarity measures for a single dyad
`method` The correlation method to use (default is "pearson")

Details

This function calculates the correlation between different similarity measures for a single dyad.

Value

A correlation matrix

Examples

```
sim1 <- list(sequence = c(0.8, 0.7, 0.9), average = 0.8)  
sim2 <- list(sequence = c(0.6, 0.8, 0.7), average = 0.7)  
cor_sim_seq(list(sim1, sim2))  
print(plot)
```

create_windows	<i>Create windows from a conversation</i>
----------------	---

Description

This function creates a list of windows from a conversation dataframe.

Usage

```
create_windows(conversation, window_size)
```

Arguments

conversation A dataframe containing the conversation, with a column named 'processed_text'.
window_size An integer specifying the size of each window.

Value

A list of character vectors, where each vector represents a window of text.

Examples

```
conversation <- data.frame(processed_text = c("hello", "world", "how", "are", "you"))  
windows <- create_windows(conversation, 3)
```

gen_sim_report	<i>Generate similarity report</i>
----------------	-----------------------------------

Description

This function generates a comprehensive report of all similarity measures.

Usage

```
gen_sim_report(  
  speech1,  
  speech2,  
  topic_method = "lda",  
  semantic_method = "tfidf",  
  glove_path = NULL  
)
```

Arguments

speech1	A character string representing the first speech
speech2	A character string representing the second speech
topic_method	Method for topic similarity calculation ("lda" or "lsa")
semantic_method	Method for semantic similarity calculation ("tfidf", "word2vec", or "glove")
glove_path	Path to pre-trained GloVe file (if using "glove" method)

Value

A list containing all similarity measures and visualizations

Examples

```
speech1 <- "This is the first speech. It talks about important topics."
speech2 <- "This is the second speech. It covers similar subjects."
report <- gen_sim_report(speech1, speech2)
```

heatmap_sim

Create Similarity Heatmap

Description

Create a heatmap of similarity measures for a single dyad

Usage

```
heatmap_sim(similarities, titles)
```

Arguments

similarities	A list of similarity measures for a single dyad
titles	A character vector of titles for each similarity measure

Details

This function creates a heatmap of multiple similarity measures for a single dyad.

Value

A ggplot object

Examples

```
sim1 <- list(sequence = c(0.5, 0.6, 0.7, 0.6, 0.8), average = 0.64)
sim2 <- list(sequence = c(0.4, 0.5, 0.6, 0.7, 0.7), average = 0.58)
similarities <- list(sim1, sim2)
titles <- c("Measure 1", "Measure 2")

# Plot multiple similarity measures
plot <- plot_sim_multi(similarities, titles)
print(plot)
```

lexical_similarity	<i>Calculate lexical similarity between two conversations</i>
--------------------	---

Description

This function calculates the lexical similarity between two conversations based on the overlap of unique words.

Usage

```
lexical_similarity(conv1, conv2)
```

Arguments

conv1	A character string representing the first conversation
conv2	A character string representing the second conversation

Value

A numeric value representing the lexical similarity

Examples

```
conv1 <- "The quick brown fox jumps over the lazy dog"
conv2 <- "The lazy dog sleeps under the quick brown fox"
lexical_similarity(conv1, conv2)
```

lexical_sim_dyads *Calculate lexical similarity for multiple dyads*

Description

This function calculates lexical similarity over a sequence of conversation exchanges for multiple dyads.

Usage

```
lexical_sim_dyads(conversations, window_size = 3)
```

Arguments

conversations A data frame with columns 'dyad_id', 'speaker', and 'processed_text'
window_size An integer specifying the size of the sliding window

Value

A list containing the sequence of similarities for each dyad and the overall average similarity

Examples

```
library(lme4)
convs <- data.frame(
  dyad_id = c(1, 1, 1, 1, 2, 2, 2, 2),
  speaker = c("A", "B", "A", "B", "C", "D", "C", "D"),
  processed_text = c("i love pizza", "me too favorite food",
                    "whats your favorite topping", "enjoy pepperoni mushrooms",
                    "i prefer pasta", "pasta delicious like spaghetti carbonara",
                    "ever tried making home", "yes quite easy make")
)
lexical_sim_dyads(convs, window_size = 2)
```

lex_sim_seq *Calculate lexical similarity sequence for a single dyad*

Description

This function calculates lexical similarity over a sequence of conversation exchanges within a single dyad.

Usage

```
lex_sim_seq(conversation, window_size = 3)
```

Arguments

conversation A data frame representing the conversation
window_size An integer specifying the size of the sliding window

Value

A list containing the sequence of similarities and the average similarity

Examples

```
conversation <- data.frame(  
  processed_text = c("Hello world", "World of programming",  
                    "Programming is fun", "Fun world of coding")  
)  
result <- lex_sim_seq(conversation, window_size = 2)  
print(result)
```

norm_sim	<i>Normalize Similarity Scores</i>
----------	------------------------------------

Description

Normalize similarity scores

Usage

```
norm_sim(similarities)
```

Arguments

similarities A numeric vector of similarity scores

Details

This function normalizes similarity scores to a 0-1 range.

Value

A numeric vector of normalized similarity scores

Examples

```
similarities <- c(0.2, 0.5, 0.8, 1.0, 0.3)  
norm_sim(similarities)  
print(plot)
```

participant_sim_dyads *Calculate participant similarity for multiple dyads*

Description

This function calculates an extended measure of participant similarity for multiple dyads.

Usage

```
participant_sim_dyads(conversations)
```

Arguments

conversations A data frame with columns 'dyad_id', 'speaker', and 'processed_text'

Value

A list containing participant similarity for each dyad and the overall average similarity

Examples

```
convs <- data.frame(
  dyad_id = c(1, 1, 1, 1, 2, 2, 2, 2),
  speaker = c("A", "B", "A", "B", "C", "D", "C", "D"),
  processed_text = c("i love pizza", "me too favorite food",
                    "whats your favorite topping", "enjoy pepperoni mushrooms",
                    "i prefer pasta", "pasta delicious like spaghetti carbonara",
                    "ever tried making home", "yes quite easy make")
)
participant_sim_dyads(convs)
```

plot_cor_heatmap *Plot Correlation Heatmap for a Single Dyad*

Description

Plot Correlation Heatmap for a Single Dyad

Usage

```
plot_cor_heatmap(cor_matrix, titles)
```

Arguments

cor_matrix A correlation matrix for a single dyad
titles A character vector of titles for each similarity measure

Details

This function creates a heatmap of correlations between similarity measures for a single dyad.

Value

A ggplot object

Examples

```
sim1 <- list(sequence = c(0.8, 0.7, 0.9), average = 0.8)
sim2 <- list(sequence = c(0.6, 0.8, 0.7), average = 0.7)
cor_matrix <- cor_sim_seq(list(sim1, sim2))
plot_cor_heatmap(cor_matrix, c("Topic", "Lexical"))
print(plot)
```

plot_sims

Visualize similarity scores

Description

This function creates a bar plot of similarity scores.

Usage

```
plot_sims(similarities)
```

Arguments

`similarities` A named list of similarity scores

Value

A ggplot object

Examples

```
sims <- list(topic = 0.8, lexical = 0.6, semantic = 0.7, structural = 0.9)
plot_sims(sims)
```

plot_sim_comp *Plot comparison of multiple similarity measures*

Description

This function creates a ggplot object comparing multiple similarity measures for the same set of dyads.

Usage

```
plot_sim_comp(comparison_df, title)
```

Arguments

comparison_df A data frame output from compare_sim_meas()
title A string specifying the plot title

Value

A ggplot object

Examples

```
topic_similarities <- list("1" = c(0.5, 0.6, 0.7), "2" = c(0.4, 0.5, 0.6))  
lexical_similarities <- list("1" = c(0.6, 0.7, 0.8), "2" = c(0.5, 0.6, 0.7))  
comparison_df <- compare_sim_meas(  
  list(topic_similarities, lexical_similarities),  
  c("Topic", "Lexical")  
)  
plot_sim_comp(comparison_df, "Comparison of Similarity Measures")  
print(plot)
```

plot_sim_cor_heatmap *Plot heatmap of similarity measure correlations*

Description

This function creates a ggplot object showing a heatmap of correlations between similarity measures.

Usage

```
plot_sim_cor_heatmap(cor_matrix, title)
```

Arguments

cor_matrix A correlation matrix output from calc_sim_cor()
title A string specifying the plot title

Value

A ggplot object

Examples

```
topic_similarities <- list("1" = c(0.5, 0.6, 0.7), "2" = c(0.4, 0.5, 0.6))  
lexical_similarities <- list("1" = c(0.6, 0.7, 0.8), "2" = c(0.5, 0.6, 0.7))  
comparison_df <- compare_sim_meas(  
  list(topic_similarities, lexical_similarities),  
  c("Topic", "Lexical")  
)  
cor_matrix <- calc_sim_cor(comparison_df)  
plot_sim_cor_heatmap(cor_matrix, "Correlation of Similarity Measures")  
print(plot)
```

plot_sim_multi *Plot Multiple Similarity Measures*

Description

Plot multiple similarity measures for a single dyad

Usage

```
plot_sim_multi(similarities, titles)
```

Arguments

similarities A list of similarity measures for a single dyad
titles A character vector of titles for each similarity measure

Details

This function creates a faceted plot of multiple similarity measures for a single dyad.

Value

A ggplot object

Examples

```
sim1 <- list(sequence = c(0.5, 0.6, 0.7, 0.6, 0.8), average = 0.64)
sim2 <- list(sequence = c(0.4, 0.5, 0.6, 0.7, 0.7), average = 0.58)
similarities <- list(sim1, sim2)
titles <- c("Measure 1", "Measure 2")

# Plot multiple similarity measures
plot <- plot_sim_multi(similarities, titles)
print(plot)
```

plot_sim_seq

Plot Similarity Sequence

Description

Plot similarity sequence for a single dyad

Usage

```
plot_sim_seq(similarity, title)
```

Arguments

similarity	A list containing the sequence of similarities and the average similarity
title	A character string for the plot title

Details

This function creates a line plot of the similarity sequence for a single dyad.

Value

A ggplot object

Examples

```
sim_list <- list(
  sequence = c(0.5, 0.6, 0.7, 0.6, 0.8),
  average = 0.64
)

# Plot the similarity sequence
plot <- plot_sim_seq(sim_list, "Dyad Similarity Sequence")
print(plot)
```

plot_sim_time	<i>Plot similarity over time for multiple dyads</i>
---------------	---

Description

This function creates a ggplot object showing the similarity over time for multiple dyads.

Usage

```
plot_sim_time(similarities, title, y_label)
```

Arguments

similarities	A list of similarity sequences for each dyad
title	A string specifying the plot title
y_label	A string specifying the y-axis label

Value

A ggplot object

Examples

```
similarities <- list(  
  "1" = c(0.5, 0.6, 0.7),  
  "2" = c(0.4, 0.5, 0.6)  
)  
plot_sim_time(similarities, "Topic Similarity", "Similarity Score")  
print(plot)
```

plot_sum_stats	<i>Plot summary statistics for similarities</i>
----------------	---

Description

This function creates a ggplot object showing summary statistics for similarities of multiple dyads.

Usage

```
plot_sum_stats(summary_stats, title)
```

Arguments

summary_stats	A data frame with summary statistics for each dyad
title	A string specifying the plot title

Value

A ggplot object

Examples

```
similarities <- list(
  "1" = c(0.5, 0.6, 0.7),
  "2" = c(0.4, 0.5, 0.6)
)
stats <- calc_sum_stats(similarities)
plot_sum_stats(stats, "Summary Statistics of Similarities")
print(plot)
```

preprocess_dyads

Preprocess multiple dyad conversations

Description

This function preprocesses conversations from multiple dyads by applying text cleaning to each utterance.

Usage

```
preprocess_dyads(conversations)
```

Arguments

`conversations` A data frame with columns 'dyad_id', 'speaker', and 'text'

Value

A data frame with an additional 'processed_text' column, removing any rows with empty processed text

Examples

```
convs <- data.frame(
  dyad_id = c(1, 1, 2, 2),
  speaker = c("A", "B", "C", "D"),
  text = c("Hello!", "Hi there!", "How are you?", "I'm fine, thanks!")
)
preprocess_dyads(convs)
```

preprocess_text	<i>This file contains core similarity calculation functions such as topic similarity, lexical similarity, semantic similarity, structural similarity, stylistic similarity, sentiment similarity, participant similarity, and timing similarity.</i>
-----------------	--

Description

Preprocess text for analysis

Usage

```
preprocess_text(text)
```

Arguments

text	A character string to be preprocessed
------	---------------------------------------

Details

This function preprocesses the input text by converting to lowercase, removing punctuation and digits, and trimming whitespace.

Value

A preprocessed character string

Examples

```
text <- "Hello, World! This is an example text (with 123 numbers)."  
preprocess_text(text)
```

print_sim_report	<i>Print similarity report</i>
------------------	--------------------------------

Description

This function prints a formatted summary of the similarity report.

Usage

```
print_sim_report(report)
```

Arguments

report	A similarity report generated by gen_sim_report function
--------	--

Value

NULL (invisibly). This function is called for its side effect of printing to the console.

Examples

```
speech1 <- "This is the first speech. It talks about important topics."  
speech2 <- "This is the second speech. It covers similar subjects."  
report <- gen_sim_report(speech1, speech2)  
print_sim_report(report)
```

radar_sim

Create Radar Chart of Average Similarities

Description

Create a radar chart of average similarities for a single dyad

Usage

```
radar_sim(similarities, titles)
```

Arguments

`similarities` A list of similarity measures for a single dyad
`titles` A character vector of titles for each similarity measure

Details

This function creates a radar chart of average similarities for multiple measures of a single dyad.

Value

A ggplot object

Examples

```
sim1 <- list(sequence = c(0.5, 0.6, 0.7, 0.6, 0.8), average = 0.64)  
sim2 <- list(sequence = c(0.4, 0.5, 0.6, 0.7, 0.7), average = 0.58)  
sim3 <- list(sequence = c(0.6, 0.7, 0.8, 0.7, 0.9), average = 0.74)  
sim4 <- list(sequence = c(0.3, 0.4, 0.5, 0.6, 0.6), average = 0.48)  
similarities <- list(sim1, sim2, sim3, sim4)  
titles <- c("Measure 1", "Measure 2", "Measure 3", "Measure 4")  
  
# Create radar chart  
radar <- radar_sim(similarities, titles)  
print(radar)
```

run_example	<i>Run package examples</i>
-------------	-----------------------------

Description

Run package examples

Usage

```
run_example(example_name)
```

Arguments

example_name Name of the example file to run

Value

No return value, called for side effects.

Examples

```
run_example("sequence_multidyads_examples.R")
run_example("main_functions_examples.R")
```

semantic_similarity	<i>Calculate semantic similarity between two conversations</i>
---------------------	--

Description

This function calculates the semantic similarity between two conversations using either TF-IDF, Word2Vec, or GloVe embeddings approach.

Usage

```
semantic_similarity(
  conversation1,
  conversation2,
  method = "tfidf",
  model_path = NULL,
  dim = 100,
  window = 5,
  iter = 5
)
```

Arguments

conversation1	A character string representing the first conversation
conversation2	A character string representing the second conversation
method	A character string specifying the method to use: "tfidf", "word2vec", or "glove"
model_path	A character string specifying the path to pre-trained GloVe file (required for "glove" method)
dim	An integer specifying the dimensionality for Word2Vec embeddings (default: 100)
window	An integer specifying the window size for Word2Vec (default: 5)
iter	An integer specifying the number of iterations for Word2Vec (default: 5)

Value

A numeric value representing the semantic similarity (between 0 and 1)

Examples

```
conv1 <- "The quick brown fox jumps over the lazy dog"
conv2 <- "A fast auburn canine leaps above an idle hound"
semantic_similarity(conv1, conv2, method = "tfidf")
```

semantic_sim_dyads *Calculate semantic similarity for multiple dyads*

Description

This function calculates semantic similarity over a sequence of conversation exchanges for multiple dyads.

Usage

```
semantic_sim_dyads(conversations, method = "tfidf", window_size = 3, ...)
```

Arguments

conversations	A data frame with columns 'dyad_id', 'speaker', and 'processed_text'
method	A character string specifying the method to use: "tfidf", "word2vec", or "glove"
window_size	An integer specifying the size of the sliding window
...	Additional arguments passed to semantic_similarity

Value

A list containing the sequence of similarities for each dyad and the overall average similarity

Examples

```
library(lme4)
convs <- data.frame(
  dyad_id = c(1, 1, 1, 1, 2, 2, 2, 2),
  speaker = c("A", "B", "A", "B", "C", "D", "C", "D"),
  processed_text = c("i love pizza", "me too favorite food",
                    "whats your favorite topping", "enjoy pepperoni mushrooms",
                    "i prefer pasta", "pasta delicious like spaghetti carbonara",
                    "ever tried making home", "yes quite easy make")
)
semantic_sim_dyads(convs, method = "tfidf", window_size = 2)
```

`sem_sim_seq`*Calculate semantic similarity sequence for a single dyad*

Description

This function calculates semantic similarity over a sequence of conversation exchanges within a single dyad.

Usage

```
sem_sim_seq(conversation, method = "tfidf", window_size = 3, ...)
```

Arguments

<code>conversation</code>	A data frame representing the conversation
<code>method</code>	A character string specifying the method to use: "tfidf", "word2vec", or "glove"
<code>window_size</code>	An integer specifying the size of the sliding window
<code>...</code>	Additional arguments passed to <code>semantic_similarity</code>

Value

A list containing the sequence of similarities and the average similarity

Examples

```
conversation <- data.frame(
  processed_text = c("The weather is nice", "It's a beautiful day",
                    "The sun is shining", "Perfect day for a picnic")
)
result <- sem_sim_seq(conversation, method = "tfidf", window_size = 2)
print(result)
```

`sentiment_similarity` *Calculate sentiment similarity between two conversations*

Description

This function calculates the sentiment similarity between two conversations using the `sentimentr` package.

Usage

```
sentiment_similarity(conv1, conv2)
```

Arguments

`conv1` A character string representing the first conversation
`conv2` A character string representing the second conversation

Value

A numeric value representing the sentiment similarity

Examples

```
conv1 <- "I love this product! It's amazing and works great."  
conv2 <- "This item is okay. It does the job but could be better."  
sentiment_similarity(conv1, conv2)
```

`sentiment_sim_dyads` *Calculate sentiment similarity for multiple dyads*

Description

This function calculates sentiment similarity over a sequence of conversation exchanges for multiple dyads.

Usage

```
sentiment_sim_dyads(conversations, window_size = 3)
```

Arguments

`conversations` A data frame with columns 'dyad_id', 'speaker', and 'processed_text'
`window_size` An integer specifying the size of the sliding window

Value

A list containing the sequence of similarities for each dyad and the overall average similarity

Examples

```
library(lme4)
convs <- data.frame(
  dyad_id = c(1, 1, 1, 1, 2, 2, 2, 2),
  speaker = c("A", "B", "A", "B", "C", "D", "C", "D"),
  processed_text = c("i love pizza", "me too favorite food",
                    "whats your favorite topping", "enjoy pepperoni mushrooms",
                    "i prefer pasta", "pasta delicious like spaghetti carbonara",
                    "ever tried making home", "yes quite easy make")
)
sentiment_sim_dyads(convs, window_size = 2)
```

sent_sim_seq

Calculate sentiment similarity sequence for a single dyad

Description

This function calculates sentiment similarity over a sequence of conversation exchanges within a single dyad.

Usage

```
sent_sim_seq(conversation, window_size = 3)
```

Arguments

conversation A data frame representing the conversation
window_size An integer specifying the size of the sliding window

Value

A list containing the sequence of similarities and the average similarity

Examples

```
conversation <- data.frame(
  processed_text = c("I love this movie!", "It's really amazing.",
                    "The acting is superb.", "I couldn't agree more.")
)
result <- sent_sim_seq(conversation, window_size = 2)
print(result)
```

structural_similarity *Calculate structural similarity between two conversations*

Description

This function calculates the structural similarity between two conversations based on their length and average turn length.

Usage

```
structural_similarity(conv1, conv2)
```

Arguments

conv1	A character vector representing the first conversation
conv2	A character vector representing the second conversation

Value

A numeric value representing the structural similarity

Examples

```
conv1 <- c("Hello", "Hi there", "How are you?", "I'm fine, thanks")
conv2 <- c("Good morning", "Hello", "Nice day, isn't it?", "Yes, indeed")
structural_similarity(conv1, conv2)
```

structural_sim_dyads *Calculate structural similarity for multiple dyads*

Description

This function calculates an extended measure of structural similarity for multiple dyads.

Usage

```
structural_sim_dyads(conversations)
```

Arguments

conversations	A data frame with columns 'dyad_id', 'speaker', and 'processed_text'
---------------	--

Value

A list containing structural similarity for each dyad and the overall average similarity

Examples

```
convs <- data.frame(
  dyad_id = c(1, 1, 1, 1, 2, 2, 2, 2),
  speaker = c("A", "B", "A", "B", "C", "D", "C", "D"),
  processed_text = c("i love pizza", "me too favorite food",
                    "whats your favorite topping", "enjoy pepperoni mushrooms",
                    "i prefer pasta", "pasta delicious like spaghetti carbonara",
                    "ever tried making home", "yes quite easy make")
)
structural_sim_dyads(convs)
```

`style_sim_seq`*Calculate stylistic similarity sequence for a single dyad*

Description

This function calculates stylistic similarity over a sequence of conversation exchanges within a single dyad.

Usage

```
style_sim_seq(conversation, window_size = 3)
```

Arguments

`conversation` A data frame representing the conversation
`window_size` An integer specifying the size of the sliding window

Value

A list containing the sequence of similarities and the average similarity

Examples

```
conversation <- data.frame(
  processed_text = c("How are you doing?", "I'm doing great, thanks!",
                    "That's wonderful to hear.", "I'm glad you're doing well.")
)
result <- style_sim_seq(conversation, window_size = 2)
print(result)
```

`stylistic_similarity` *Calculate stylistic similarity between two conversations*

Description

This function calculates various stylistic features and their similarity between two conversations.

Usage

```
stylistic_similarity(text1, text2)
```

Arguments

`text1` A character string representing the first conversation
`text2` A character string representing the second conversation

Value

A list containing stylistic features and similarity measures

Examples

```
text1 <- "The quick brown fox jumps over the lazy dog. It's a sunny day."  
text2 <- "A lazy cat sleeps on the warm windowsill. Birds chirp outside."  
stylistic_similarity(text1, text2)
```

`stylistic_sim_dyads` *Calculate stylistic similarity for multiple dyads*

Description

This function calculates stylistic similarity over a sequence of conversation exchanges for multiple dyads.

Usage

```
stylistic_sim_dyads(conversations, window_size = 3)
```

Arguments

`conversations` A data frame with columns 'dyad_id', 'speaker', and 'processed_text'
`window_size` An integer specifying the size of the sliding window

Value

A list containing the sequence of similarities for each dyad and the overall average similarity

Examples

```
convs <- data.frame(
  dyad_id = c(1, 1, 1, 1, 2, 2, 2, 2),
  speaker = c("A", "B", "A", "B", "C", "D", "C", "D"),
  processed_text = c("i love pizza", "me too favorite food",
                    "whats your favorite topping", "enjoy pepperoni mushrooms",
                    "i prefer pasta", "pasta delicious like spaghetti carbonara",
                    "ever tried making home", "yes quite easy make")
)
stylistic_sim_dyads(convs, window_size = 2)
```

timing_sim_dyads

Calculate timing similarity for multiple dyads

Description

This function calculates an extended measure of timing similarity for multiple dyads.

Usage

```
timing_sim_dyads(conversations)
```

Arguments

`conversations` A data frame with columns 'dyad_id', 'speaker', and 'processed_text'

Value

A list containing timing similarity for each dyad and the overall average similarity

Examples

```
convs <- data.frame(
  dyad_id = c(1, 1, 1, 1, 2, 2, 2, 2),
  speaker = c("A", "B", "A", "B", "C", "D", "C", "D"),
  processed_text = c("i love pizza", "me too favorite food",
                    "whats your favorite topping", "enjoy pepperoni mushrooms",
                    "i prefer pasta", "pasta delicious like spaghetti carbonara",
                    "ever tried making home", "yes quite easy make")
)
timing_sim_dyads(convs)
```

topic_similarity	<i>Calculate topic similarity between two conversations</i>
------------------	---

Description

This function calculates the topic similarity between two conversations using either Latent Dirichlet Allocation (LDA) or Latent Semantic Analysis (LSA).

Usage

```
topic_similarity(conv1, conv2, method = "lda", num_topics = 2)
```

Arguments

conv1	A character vector representing the first conversation
conv2	A character vector representing the second conversation
method	A character string specifying the method to use: "lda" or "lsa"
num_topics	An integer specifying the number of topics to use in the model

Value

A numeric value representing the topic similarity

Examples

```
conv1 <- c("I love pizza", "Pizza is my favorite food")
conv2 <- c("I prefer pasta", "Pasta is delicious")
topic_similarity(conv1, conv2, method = "lda", num_topics = 2)
topic_similarity(conv1, conv2, method = "lsa", num_topics = 2)
```

topic_sim_dyads	<i>Calculate topic similarity for multiple dyads</i>
-----------------	--

Description

This function calculates topic similarity over a sequence of conversation exchanges for multiple dyads. It uses the Latent Dirichlet Allocation (LDA) method for topic modeling and the "slam" package for efficient handling of sparse matrices.

Usage

```
topic_sim_dyads(conversations, method = "lda", num_topics = 2, window_size = 3)
```

Arguments

conversations	A data frame with columns 'dyad_id', 'speaker', and 'processed_text'
method	A character string specifying the method to use: currently only "lda" is supported
num_topics	An integer specifying the number of topics to use in the LDA model
window_size	An integer specifying the size of the sliding window

Value

A list containing the sequence of similarities for each dyad and the overall average similarity

Examples

```

convs <- data.frame(
  dyad_id = c(1, 1, 1, 2, 2, 2, 2),
  speaker = c("A", "B", "A", "B", "C", "D", "C", "D"),
  processed_text = c("i love pizza", "me too favorite food",
                    "whats your favorite topping", "enjoy pepperoni mushrooms",
                    "i prefer pasta", "pasta delicious like spaghetti carbonara",
                    "ever tried making home", "yes quite easy make")
)
topic_sim_dyads(convs, method = "lda", num_topics = 2, window_size = 2)

```

topic_sim_seq	<i>Calculate topic similarity sequence for a single dyad</i>
---------------	--

Description

This function calculates topic similarity over a sequence of conversation exchanges within a single dyad.

Usage

```
topic_sim_seq(conversation, method = "lda", num_topics = 2, window_size = 3)
```

Arguments

conversation	A data frame representing the conversation
method	A character string specifying the method to use: "lda" or "lsa"
num_topics	An integer specifying the number of topics to use in the model
window_size	An integer specifying the size of the sliding window

Value

A list containing the sequence of similarities and the average similarity

Examples

```
conversation <- data.frame(  
  processed_text = c("The cat sat on the mat", "The dog chased the cat",  
                    "The mat was comfortable", "The cat liked the mat")  
)  
result <- topic_sim_seq(conversation, method = "lda", num_topics = 2, window_size = 2)  
print(result)
```

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