

# Package ‘BTM’

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**Type** Package

**Title** Biterm Topic Models for Short Text

**Version** 0.3.8

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**Description** Biterm Topic Models find topics in collections of short texts.

It is a word co-occurrence based topic model that learns topics by modeling word-word co-occurrences patterns which are called biterms.

This in contrast to traditional topic models like Latent Dirichlet Allocation and Probabilistic Latent Semantic Analysis

which are word-document co-occurrence topic models.

A biterm consists of two words co-occurring in the same short text window.

This context window can for example be a twitter message, a short answer on a survey, a sentence of a text or a document identifier.

The techniques are explained in detail in the paper 'A Biterm Topic Model For Short Text' by Xiaohui Yan, Jiafeng Guo, Yanyan Lan, Xueqi Cheng (2013) <<https://github.com/xiaohuiyan/xiaohuiyan.github.io/blob/master/paper/BTM-WWW13.pdf>>.

**License** Apache License 2.0

**URL** <https://github.com/bnosac/BTM>

**Encoding** UTF-8

**Imports** Rcpp, utils

**Suggests** udpipe, data.table

**LinkingTo** Rcpp

**RoxygenNote** 7.3.2

**NeedsCompilation** yes

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## Contents

BTM . . . . .	2
logLik.BTM . . . . .	6
predict.BTM . . . . .	7
terms.BTM . . . . .	8
terms.data.frame . . . . .	9
<b>Index</b>	<b>11</b>

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 BTM

*Construct a Biterm Topic Model on Short Text*


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### Description

The Biterm Topic Model (BTM) is a word co-occurrence based topic model that learns topics by modeling word-word co-occurrences patterns (e.g., biterns)

- A biterm consists of two words co-occurring in the same context, for example, in the same short text window.
- BTM models the biterm occurrences in a corpus (unlike LDA models which model the word occurrences in a document).
- It's a generative model. In the generation procedure, a biterm is generated by drawing two words independently from a same topic  $z$ . In other words, the distribution of a biterm  $b = (w_i, w_j)$  is defined as:  $P(b) = \sum_k P(w_i|z) * P(w_j|z) * P(z)$  where  $k$  is the number of topics you want to extract.
- Estimation of the topic model is done with the Gibbs sampling algorithm. Where estimates are provided for  $P(w|k) = \phi$  and  $P(z) = \theta$ .

### Usage

```
BTM(
  data,
  k = 5,
  alpha = 50/k,
  beta = 0.01,
  iter = 1000,
  window = 15,
  background = FALSE,
  trace = FALSE,
  biterns,
  detailed = FALSE
)
```

**Arguments**

<code>data</code>	a tokenised data frame containing one row per token with 2 columns <ul style="list-style-type: none"> <li>• the first column is a context identifier (e.g. a tweet id, a document id, a sentence id, an identifier of a survey answer, an identifier of a part of a text)</li> <li>• the second column is a column called of type character containing the sequence of words occurring within the context identifier</li> </ul>
<code>k</code>	integer with the number of topics to identify
<code>alpha</code>	numeric, indicating the symmetric dirichlet prior probability of a topic $P(z)$ . Defaults to $50/k$ .
<code>beta</code>	numeric, indicating the symmetric dirichlet prior probability of a word given the topic $P(w z)$ . Defaults to 0.01.
<code>iter</code>	integer with the number of iterations of Gibbs sampling
<code>window</code>	integer with the window size for biterm extraction. Defaults to 15.
<code>background</code>	logical if set to TRUE, the first topic is set to a background topic that equals to the empirical word distribution. This can be used to filter out common words. Defaults to FALSE.
<code>trace</code>	logical indicating to print out evolution of the Gibbs sampling iterations. Defaults to FALSE.
<code>biterms</code>	optionally, your own set of biterms to use for modelling. This argument should be a data.frame with column names <code>doc_id</code> , <code>term1</code> , <code>term2</code> and <code>cooc</code> , indicating how many times each biterm (as indicated by terms <code>term1</code> and <code>term2</code> ) is occurring within a certain <code>doc_id</code> . The field <code>cooc</code> indicates how many times this biterm happens with the <code>doc_id</code> . Note that <code>doc_id</code> 's which are not in <code>data</code> are not allowed, as well as terms (in <code>term1</code> and <code>term2</code> ) which are not also in <code>data</code> . See the examples. If provided, the <code>window</code> argument is ignored and the <code>data</code> argument will only be used to calculate the background word frequency distribution.
<code>detailed</code>	logical indicating to return detailed output containing as well the vocabulary and the biterms used to construct the model. Defaults to FALSE.

**Value**

an object of class BTM which is a list containing

- `model`: a pointer to the C++ BTM model
- `K`: the number of topics
- `W`: the number of tokens in the data
- `alpha`: the symmetric dirichlet prior probability of a topic  $P(z)$
- `beta`: the symmetric dirichlet prior probability of a word given the topic  $P(w|z)$
- `iter`: the number of iterations of Gibbs sampling
- `background`: indicator if the first topic is set to the background topic that equals the empirical word distribution.
- `theta`: a vector with the topic probability  $p(z)$  which is determined by the overall proportions of biterms in it

- `phi`: a matrix of dimension  $W \times K$  with one row for each token in the data. This matrix contains the probability of the token given the topic  $P(w|z)$ . the rownames of the matrix indicate the token  $w$
- `vocab`: a `data.frame` with columns `token` and `freq` indicating the frequency of occurrence of the tokens in data. Only provided in case argument `detailed` is set to `TRUE`
- `biterms`: the result of a call to `terms` with type set to `biterms`, containing all the biterms used in the model. Only provided in case argument `detailed` is set to `TRUE`

### Note

A biterm is defined as a pair of words co-occurring in the same text window. If you have as an example a document with sequence of words 'A B C B', and assuming the window size is set to 3, that implies there are two text windows which can generate biterms namely text window 'A B C' with biterms 'A B', 'B C', 'A C' and text window 'B C B' with biterms 'B C', 'C B', 'B B' A biterm is an unordered word pair where 'B C' = 'C B'. Thus, the document 'A B C B' will have the following biterm frequencies:

- 'A B': 1
- 'B C': 3
- 'A C': 1
- 'B B': 1

These biterms are used to create the model.

### References

Xiaohui Yan, Jiafeng Guo, Yanyan Lan, Xueqi Cheng. A Biterm Topic Model For Short Text. WWW2013, <https://github.com/xiaohuiyan/BTM>, <https://github.com/xiaohuiyan/xiaohuiyan.github.io/blob/master/paper/BTM-WWW13.pdf>

### See Also

[predict.BTM](#), [terms.BTM](#), [logLik.BTM](#)

### Examples

```
library(udpipe)
data("brussels_reviews_anno", package = "udpipe")
x <- subset(brussels_reviews_anno, language == "nl")
x <- subset(x, xpos %in% c("NN", "NNP", "NNS"))
x <- x[, c("doc_id", "lemma")]
model <- BTM(x, k = 5, alpha = 1, beta = 0.01, iter = 10, trace = TRUE)
model
terms(model)
scores <- predict(model, newdata = x)

## Another small run with first topic the background word distribution
set.seed(123456)
model <- BTM(x, k = 5, beta = 0.01, iter = 10, background = TRUE)
```

```

model
terms(model)

##
## You can also provide your own set of biterns to cluster upon
## Example: cluster nouns and adjectives in the neighbourhood of one another
##
library(data.table)
library(udpipe)
x <- subset(brussels_reviews_anno, language == "nl")
x <- head(x, 5500) # take a sample to speed things up on CRAN
biterns <- as.data.table(x)
biterns <- biterns[, cooccurrence(x = lemma,
                                relevant = xpos %in% c("NN", "NNP", "NNS", "JJ"),
                                skipgram = 2),
                  by = list(doc_id)]

head(biterns)
set.seed(123456)
x <- subset(x, xpos %in% c("NN", "NNP", "NNS", "JJ"))
x <- x[, c("doc_id", "lemma")]
model <- BTM(x, k = 5, beta = 0.01, iter = 10, background = TRUE,
            biterns = biterns, trace = 10, detailed = TRUE)

model
terms(model)
biternset <- terms(model, "biterns")
head(biternset$biterns, 100)

biternset$n
sum(biterns$cooc)

## Not run:
##
## Visualisation either using the textplot or the LDAvis package
##
library(textplot)
library(ggraph)
library(concaveman)
plot(model, top_n = 4)

library(LDAvis)
docsize <- table(x$doc_id)
scores <- predict(model, x)
scores <- scores[names(docsize), ]
json <- createJSON(
  phi = t(model$phi),
  theta = scores,
  doc.length = as.integer(docsize),
  vocab = model$vocabulary$token,
  term.frequency = model$vocabulary$freq)
serVis(json)

## End(Not run)

```

---

`logLik.BTM`*Get the likelihood of biterms in a BTM model*

---

**Description**

Get the likelihood how good biterms are fit by the BTM model

**Usage**

```
## S3 method for class 'BTM'  
logLik(object, data = terms.BTM(object, type = "biterms"), ...)
```

**Arguments**

<code>object</code>	an object of class BTM as returned by <a href="#">BTM</a>
<code>data</code>	a data.frame with 2 columns <code>term1</code> and <code>term2</code> containing biterms. Defaults to the biterms used to construct the model.
<code>...</code>	other arguments not used

**Value**

a list with elements

- `likelihood`: a vector with the same number of rows as `data` containing the likelihood of the biterms alongside the BTM model. Calculated as  $\sum(\phi[\text{term1}, ] * \phi[\text{term2}, ] * \theta)$ .
- `ll` the sum of the log of the biterm likelihoods

**See Also**

[BTM](#), [predict.BTM](#), [terms.BTM](#)

**Examples**

```
library(udpipe)  
data("brussels_reviews_anno", package = "udpipe")  
x <- subset(brussels_reviews_anno, language == "nl")  
x <- subset(x, xpos %in% c("NN", "NNP", "NNS"))  
x <- x[, c("doc_id", "lemma")]  
  
model <- BTM(x, k = 5, iter = 5, trace = TRUE, detailed = TRUE)  
fit <- logLik(model)  
fit$ll
```

---

predict.BTM

*Predict function for a Biterm Topic Model*

---

### Description

Classify new text alongside the biterm topic model.

To infer the topics in a document, it is assumed that the topic proportions of a document is driven by the expectation of the topic proportions of biterms generated from the document.

### Usage

```
## S3 method for class 'BTM'  
predict(object, newdata, type = c("sum_b", "sub_w", "mix"), ...)
```

### Arguments

object	an object of class BTM as returned by <a href="#">BTM</a>
newdata	a tokenised data frame containing one row per token with 2 columns <ul style="list-style-type: none"><li>• the first column is a context identifier (e.g. a tweet id, a document id, a sentence id, an identifier of a survey answer, an identifier of a part of a text)</li><li>• the second column is a column called of type character containing the sequence of words occurring within the context identifier</li></ul>
type	character string with the type of prediction. Either one of 'sum_b', 'sub_w' or 'mix'. Default is set to 'sum_b' as indicated in the paper, indicating to sum over the the expectation of the topic proportions of biterms generated from the document. For the other approaches, please inspect the paper.
...	not used

### Value

a matrix containing containing  $P(z|d)$  - the probability of the topic given the biterms. The matrix has one row for each unique doc\_id (context identifier) which contains words part of the dictionary of the BTM model and has K columns, one for each topic.

### References

Xiaohui Yan, Jiafeng Guo, Yanyan Lan, Xueqi Cheng. A Biterm Topic Model For Short Text. WWW2013, <https://github.com/xiaohuiyan/BTM>, <https://github.com/xiaohuiyan/xiaohuiyan.github.io/blob/master/paper/BTM-WWW13.pdf>

### See Also

[BTM](#), [terms.BTM](#), [logLik.BTM](#)

## Examples

```
library(udpipe)
data("brussels_reviews_anno", package = "udpipe")
x <- subset(brussels_reviews_anno, language == "nl")
x <- subset(x, xpos %in% c("NN", "NNP", "NNS"))
x <- x[, c("doc_id", "lemma")]
model <- BTM(x, k = 5, iter = 5, trace = TRUE)
scores <- predict(model, newdata = x, type = "sum_b")
scores <- predict(model, newdata = x, type = "sub_w")
scores <- predict(model, newdata = x, type = "mix")
head(scores)
```

---

terms.BTM	<i>Get highest token probabilities for each topic or get biterns used in the model</i>
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---

## Description

Get highest token probabilities for each topic or get biterns used in the model

## Usage

```
## S3 method for class 'BTM'
terms(x, type = c("tokens", "biterns"), threshold = 0, top_n = 5, ...)
```

## Arguments

x	an object of class BTM as returned by <a href="#">BTM</a>
type	a character string, either 'tokens' or 'biterns'. Defaults to 'tokens'.
threshold	threshold in 0-1 range. Only the terms which are more likely than the threshold are returned for each topic. Only used in case type = 'tokens'.
top_n	integer indicating to return the top n tokens for each topic only. Only used in case type = 'tokens'.
...	not used

## Value

Depending if type is set to 'tokens' or 'biterns' the following is returned:

- If type='tokens': Get the probability of the token given the topic  $P(w|z)$ . It returns a list of data.frames (one for each topic) where each data.frame contains columns token and probability ordered from high to low. The list is the same length as the number of topics.
- If type='biterns': a list containing 2 elements:
  - n which indicates the number of biterns used to train the model
  - biterns which is a data.frame with columns term1, term2 and topic, indicating for all biterns found in the data the topic to which the bitern is assigned to

Note that a bitern is unordered, in the output of type='biterns' term1 is always smaller than or equal to term2.

**See Also**

[BTM](#), [predict.BTM](#), [logLik.BTM](#)

**Examples**

```
library(udpipe)
data("brussels_reviews_anno", package = "udpipe")
x <- subset(brussels_reviews_anno, language == "nl")
x <- subset(x, xpos %in% c("NN", "NNP", "NNS"))
x <- x[, c("doc_id", "lemma")]
model <- BTM(x, k = 5, iter = 5, trace = TRUE)
terms(model)
terms(model, top_n = 10)
terms(model, threshold = 0.01, top_n = +Inf)
bi <- terms(model, type = "bitersms")
str(bi)
```

---

terms.data.frame

*Get the set of Bitersms from a tokenised data frame*

---

**Description**

This extracts words occurring in the neighbourhood of one another, within a certain window range. The default setting provides the bitersms used when fitting [BTM](#) with the default window parameter.

**Usage**

```
## S3 method for class 'data.frame'
terms(x, type = c("tokens", "bitersms"), window = 15, ...)
```

**Arguments**

x	a tokenised data frame containing one row per token with 2 columns <ul style="list-style-type: none"> <li>the first column is a context identifier (e.g. a tweet id, a document id, a sentence id, an identifier of a survey answer, an identifier of a part of a text)</li> <li>the second column is a column called of type character containing the sequence of words occurring within the context identifier</li> </ul>
type	a character string, either 'tokens' or 'bitersms'. Defaults to 'tokens'.
window	integer with the window size for bitersms extraction. Defaults to 15.
...	not used

**Value**

Depending if type is set to 'tokens' or 'biterms' the following is returned:

- If type='tokens': a list containing 2 elements:
  - n which indicates the number of tokens
  - tokens which is a data.frame with columns id, token and freq, indicating for all tokens found in the data the frequency of occurrence
- If type='biterms': a list containing 2 elements:
  - n which indicates the number of biterms used to train the model
  - biterms which is a data.frame with columns term1 and term2, indicating all biterms found in the data. The same biterm combination can occur several times.

Note that a biterm is unordered, in the output of type='biterms' term1 is always smaller than or equal to term2.

**Note**

If x is a data.frame which has an attribute called 'terms', it just returns that 'terms' attribute

**See Also**

[BTM](#), [predict.BTM](#), [logLik.BTM](#)

**Examples**

```
library(udpipe)
data("brussels_reviews_anno", package = "udpipe")
x <- subset(brussels_reviews_anno, language == "nl")
x <- subset(x, xpos %in% c("NN", "NNP", "NNS"))
x <- x[, c("doc_id", "lemma")]
biterms <- terms(x, window = 15, type = "biterms")
str(biterms)
tokens <- terms(x, type = "tokens")
str(tokens)
```

# Index

BTM, [2](#), [6–10](#)

logLik.BTM, [4](#), [6](#), [7](#), [9](#), [10](#)

predict.BTM, [4](#), [6](#), [7](#), [9](#), [10](#)

terms.BTM, [4](#), [6](#), [7](#), [8](#)

terms.data.frame, [9](#)