

wam package: computing Word association measure

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October 28, 2015

Abstract

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1 Introduction: Indicators of word association

```
> library(wam);
```

1.1 Introduction

This package contains various functions for computing word association measure as well as a high level function for conveniently applying these functions on all the lexical types of a corpora.

Word association can serve two goals: analyzing the association strength between a word and a subcorpora or analyzing the association strength between two words. The two are modeled with the same quantitative framework: association between two words is simply the association between a word and the subcorpus made of all its cooccurrences with the other word.

1.2 Value

All association measure functions are prefixed with "wam." and return a numeric vector indicating the association strength between the word(s) under scrutiny and subcorpus/ora.

These association measures, unless otherwise stated in the help pages of the functions, are positive when the word is over-represented ("attracted"), and negative when the word is under-represented.

In absolute value, the more the word is over-represented or under-represented, the more the association measure given is high.

1.3 Arguments

All association measure functions have the following signature: (N, n, K, k) , where:

1. N is the total size of the corpora
2. n is the size of the subcorpora
3. K is the total frequency of the form under scrutiny in the corpora
4. k is the sub-frequency of the form under scrutiny in the subcorpora

This can be easily turn into the "contingency table" representation used in some presentation (according to Stefan Evert UCS documentation) :

	word	$\neg word$	T
subcorpus	11	12	R1
$\neg subcorpus$	21	22	R2
Totals	C1	C2	N

where :

- N = total words in corpus (or subcorpus or restriction, but they are not implemented yet)
- $C1$ = frequency of the collocate in the whole corpus
- $C2$ = frequency of words that aren't the collocate in the corpus
- $R1$ = total words in window
- $R2$ = total words outside of window
- 11 = how many of collocate there are in the window
- 12 = how many words other than the collocate there are in the window (calculated from row total)
- 21 = how many of collocate there are outside the window
- 22 = how many words other than the collocate there are outside the window

Conversion from N, n, K, k notation :

	word	$\neg word$	T
subcorpus	k	$n - k$	n
$\neg subcorpus$	$K - k$	$N - K - (n - k)$	$N - n$
Totals	K	$N - K$	N

Conversion to N, n, K, k notation :

- $N = N$
- $n = O11 + O12$
- $K = O11 + O21$
- $k = O11$

See the functions `make.contingency` and `make.list` for an easy way of doing these conversions.

1.4 Recycling arguments

For all functions arguments are recycled.

2 The indicators

2.1 Introduction

All word association functions will be illustrated with data from the `robespierre` dataset. We will consider the subfrequency of the lexical types

peuple in a subcorpora containing the fourth discourse by Robespierre. Is this type over- or under-represented in this discourse, according to its frequency in the corpus of all the discourses?

```
> data(robespierre, package="wam")
> head(robespierre)
```

	types	parts	k	N	K	n
1	de	D1	464	61449	3173	8395
2	la	D1	365	61449	2788	8395
3	les	D1	281	61449	2123	8395
4	et	D1	227	61449	1708	8395
5	le	D1	200	61449	1351	8395
6	l	D1	188	61449	1287	8395

```
> peuple_D4 <- robspierre[robaspierre$types=="peuple" & robspierre$parts == "D4",]
> peuple_D4
```

	types	parts	k	N	K	n
495	peuple	D4	14	61449	296	6903

```
> N <- peuple_D4$N
> n <- peuple_D4$n
> K <- peuple_D4$K
> k <- peuple_D4$k
```

A graph of the function is provided for all the possible values of k . The interval of the possible value of K is $[0, \min(k, n)]$. (it is not possible to have more occurrences of the lexical type than the size of the subcorpus or more than the total frequency of the type in the whole corpus).

```
> maxk <- min(K,n)
> maxk
```

```
[1] 296
```

```
> allk <- 0:maxk
```

The expected subfrequency of "peuple" in the subcorpus D4 (the fourth discourse by Robespierre) is: $K \times n/N$.

```
> expected = round(K * n / N)
> expected
```

```
[1] 33
```

A form is over-used (attracted) if the subfrequency k in the subcorpus is greater than the expected frequency and under-used otherwise. Here, the form *peuple* is under-represented.

2.2 Log-likelihood

See Dunning 1993.

```
> wam.loglikelihood(N, n, K, k);
```

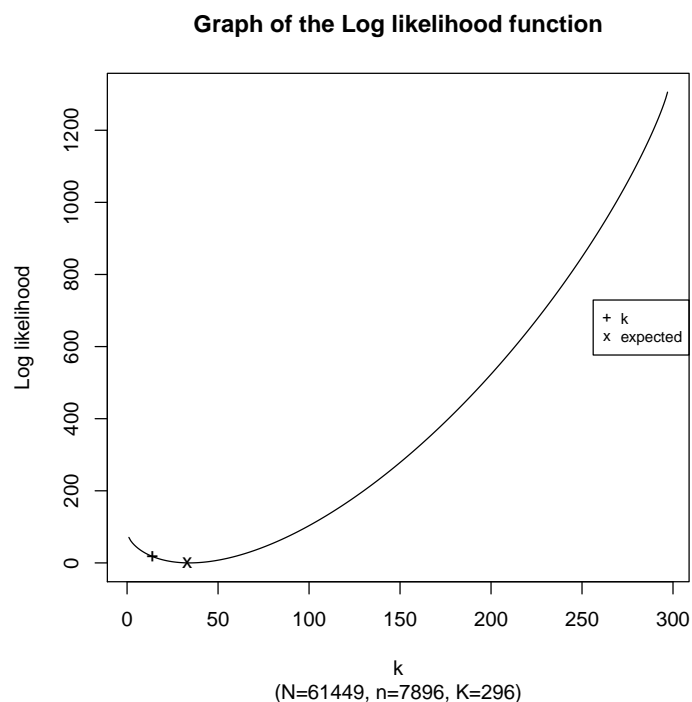
```
[1] 15.7202
```

```
> wam.loglikelihood(N, n, K, expected);
```

```
[1] 0.002162702
```

Graph of the function :

```
> plot(wam.loglikelihood(N, n, K, allk),  
+      type="l", xlab="k", ylab="Log likelihood",  
+      main="Graph of the Log likelihood function",  
+      sub="(N=61449, n=7896, K=296)")  
> points(k, wam.loglikelihood(N, n, K, k), pch="+")  
> points(expected, wam.loglikelihood(N, n, K, expected), pch="x")  
> legend("right", legend=c("k", "expected"), pch=c("+", "x"), cex=0.75)
```



2.3 Specificities

See Lafon 1980.

```
> wam.specificities(N, n, K, k, method="base");
```

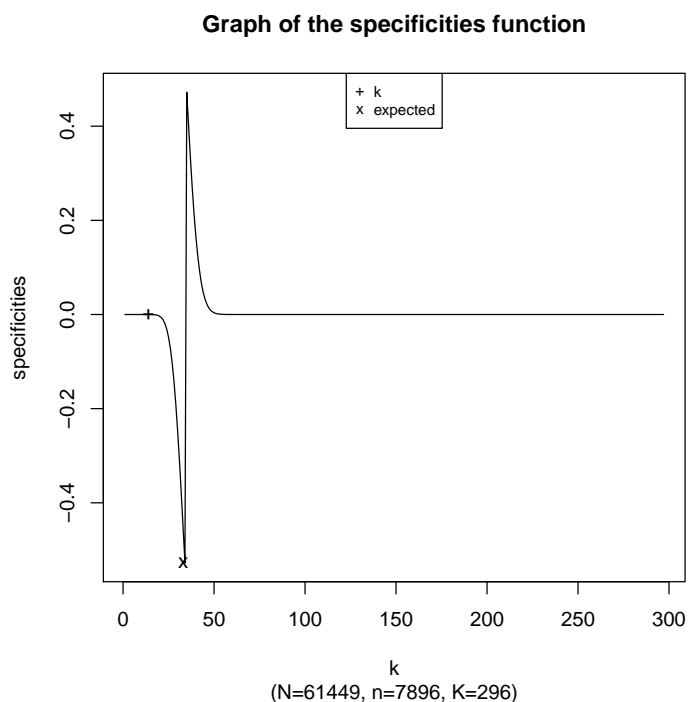
```
[1] -6.693709e-05
```

```
> wam.specificities(N, n, K, expected, method="base");
```

```
[1] -0.5276713
```

Graph of the function:

```
> plot(wam.specificities(N, n, K, allk, method="base"),  
+      type="l", xlab="k", ylab="specificities",  
+      main="Graph of the specificities function",  
+      sub="(N=61449, n=7896, K=296)")  
> points(k, wam.specificities(N, n, K, k, method="base"), pch="+")  
> points(expected, wam.specificities(N, n, K, expected, method="base"), pch="x")  
> legend("top", legend=c("k", "expected"), pch=c("+", "x"), cex=0.75)
```



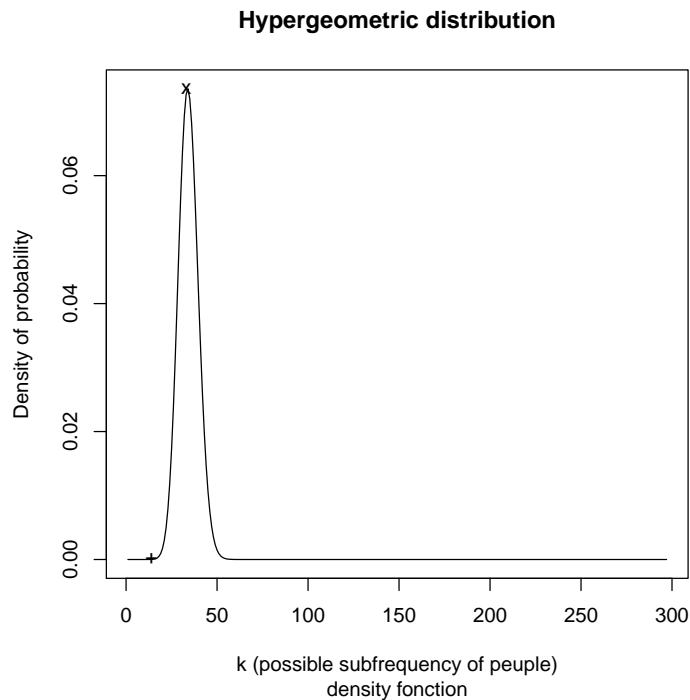
2.3.1 Analysis of the specificities indicator : Standard indicator (method="base")

The presentation below follows (Lafon, 1980).

The specificities indicator is based on the hypergeometric distribution. This distribution give the probability associated with a drawing without replacement.

For all the possible subfrequencys of *peuple* in the fourth discourses we can compute the density of probability in the hypergeometric distribution. The graph contains also the observed frequency as well as the mode. The mode is the closest positive integers to the expected frequency.

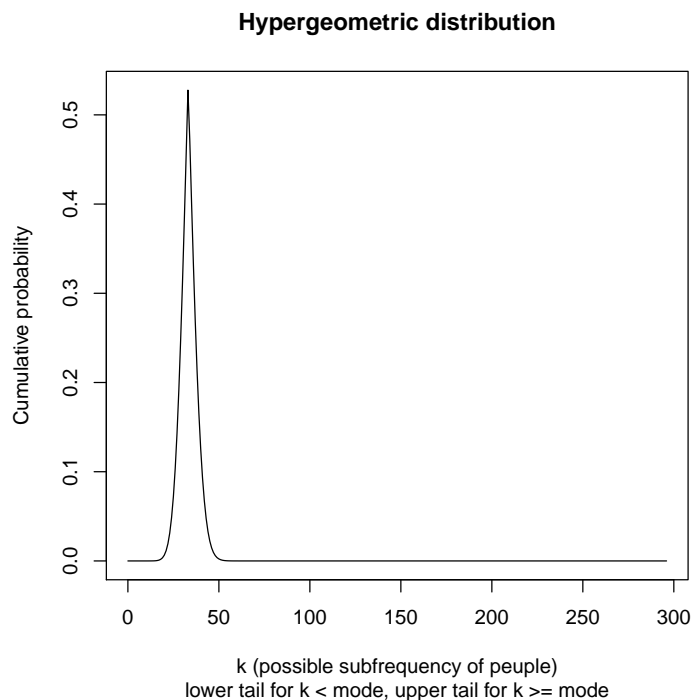
```
> mode <- floor((n+1)*(K+1)/(N+2));  
> mode  
  
[1] 33  
  
> plot(dhyper(allk, K, N-K, n),  
+ type="l", xlab="k (possible subfrequency of peuple)", ylab="Density of probability",  
+ main="Hypergeometric distribution", sub="density fonction")  
> points(k, dhyper(k, K, N-K, n), pch="+")  
> points(mode, dhyper(mode, K, N-K, n), pch="x")
```



If the observed frequency is less than the expected frequency, we compute the sum of the probability for a frequency lesser or equal to the observed frequency ($Prob(X \leq k)$) – that is, the cumulative probability.

If the observed frequency is greater than the expected frequency, we compute the sum of the probability for a frequency greater to the observed frequency ($Prob(X > k)$) (Lafon 1980 : 141) – that is, the cumulative probability for the upper tail of the distribution.

```
> y <- ifelse(allk <= mode, phyper(allk, K, N-K, n),
+               phyper(allk-1, K, N-K, n, lower.tail=FALSE))
> plot(allk, y,
+       type="l", xlab="k (possible subfrequency of peuple)",
+       ylab="Cumulative probability",
+       main="Hypergeometric distribution",
+       sub="lower tail for k < mode, upper tail for k >= mode")
```



We add a sign: negative if the frequency is lower than expected, positive if it is greater.

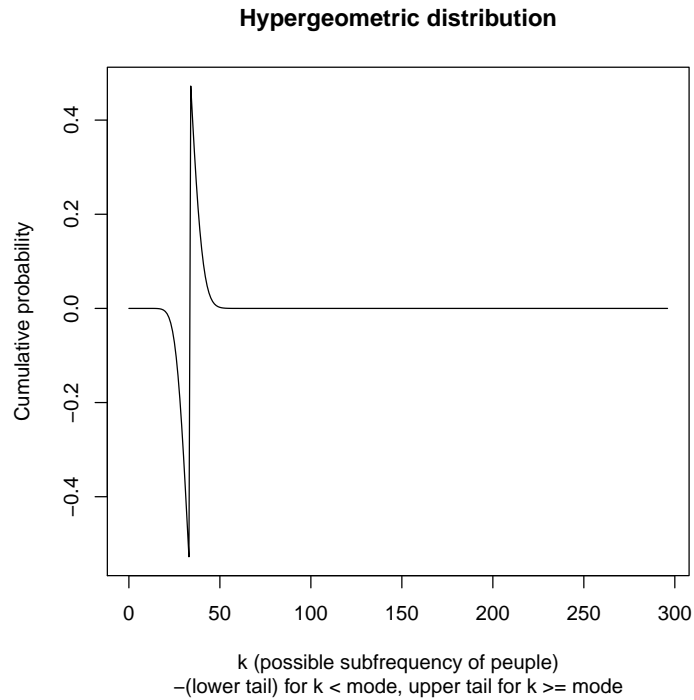
```
> y <- ifelse(allk <= mode, phyper(allk, K, N-K, n),
+               phyper(allk-1, K, N-K, n, lower.tail=FALSE))
> y <- ifelse(allk <= mode, -y, y);
```



```

> plot(allk, y,
+      type="l", xlab="k (possible subfrequency of peuple)",
+      ylab="Cumulative probability",
+      main="Hypergeometric distribution",
+      sub="-(lower tail) for k < mode, upper tail for k >= mode")

```

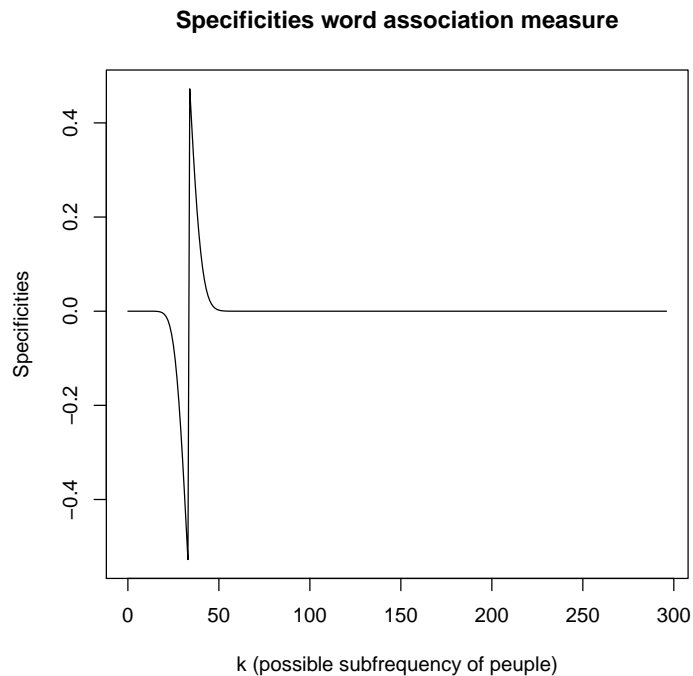


It is the standard Specificities function (Lafon 1980), as implemented in the function `wam.specificities` with `method="base"` :

```

> plot(allk, wam.specificities(N, n, K, allk, method="base"),
+      type="l", xlab="k (possible subfrequency of peuple)",
+      ylab="Specificities",
+      main="Specificities word association measure")

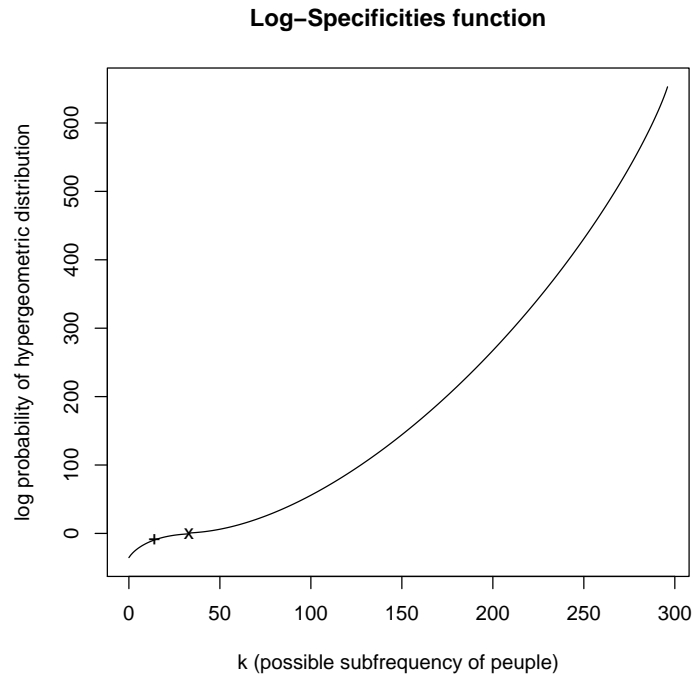
```



2.3.2 Analysis of the specificities indicator : log (method="log")

In order to ease the reading, log are used:

```
> y <- ifelse(allk <= mode, phyper(allk, K, N-K, n, log.p=TRUE),
+           phyper(allk-1, K, N-K, n, lower.tail=FALSE, log.p=TRUE))
> y <- ifelse(allk <= mode, -abs(y), abs(y));
> plot(allk, y,
+       type="l", xlab="k (possible subfrequency of peuple)",
+       ylab="log probability of hypergeometric distribution",
+       main="Log-Specificities function");
> points(k, phyper(k, K, N-K, n, log.p=TRUE), , pch="+")
> points(mode, phyper(mode, K, N-K, n, log.p=TRUE), pch="x")
```



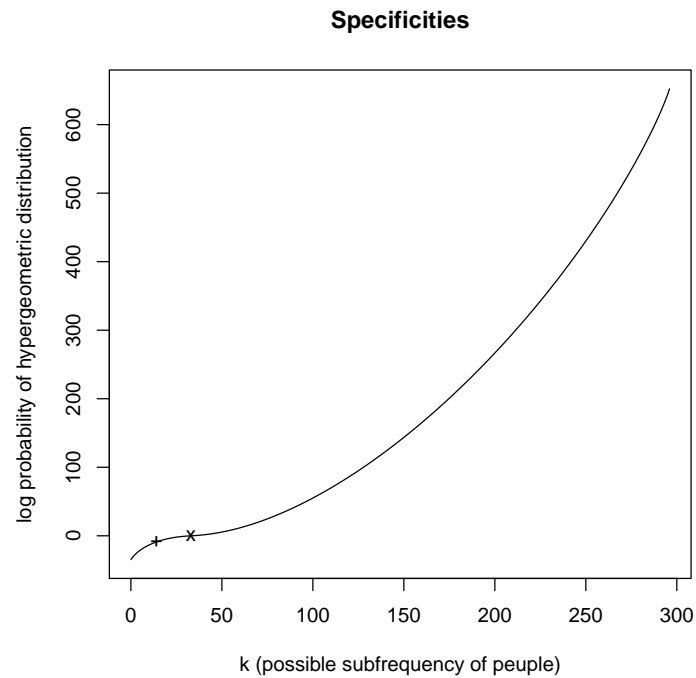
Another issue is that the mode is different from 0:

```
> phyper(mode, K, N-K, n, log.p=TRUE)
```

```
[1] -0.6392818
```

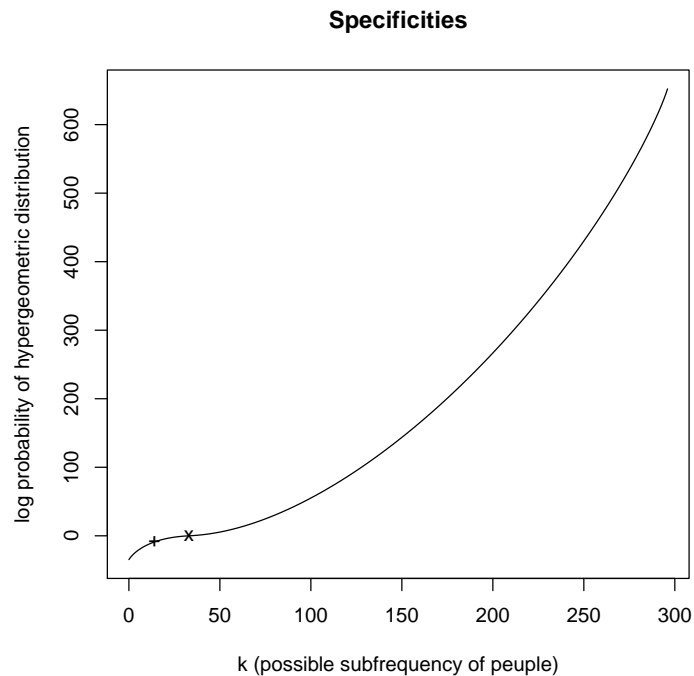
in order to have $mode = 0$, the value of the mode is substracted from all values:

```
> y <- ifelse(allk <= mode, phyper(allk, K, N-K, n, log.p=TRUE),
+             phyper(allk-1, K, N-K, n, lower.tail=FALSE, log.p=TRUE))
> cdm0 <- phyper(mode, K, N-K, n, log.p=TRUE);
> y <- ifelse(allk <= mode, -abs(cdm0-y), abs(cdm0-y));
> plot(allk, y,
+       type="l", xlab="k (possible subfrequency of peuple)",
+       ylab="log probability of hypergeometric distribution",
+       main="Specificities")
> points(k, wam.specificities(N, n, K, k, method="log"), pch="+")
> points(mode, wam.specificities(N, n, K, mode, method="log"), pch="x")
```



That is the `wam.specificities` function with `method="log"`:

```
> plot(allk, wam.specificities(N, n, K, allk, method="log"),
+       type="l", xlab="k (possible subfrequency of people)",
+       ylab="log probability of hypergeometric distribution",
+       main="Specificities")
> points(k, wam.specificities(N, n, K, k, method="log"), pch="+")
> points(mode, wam.specificities(N, n, K, mode, method="log"), pch="x")
```



where the mode is 0 :

```
> wam.specificities(N, n, K, mode, method="log");
[1] 0
```

2.4 z

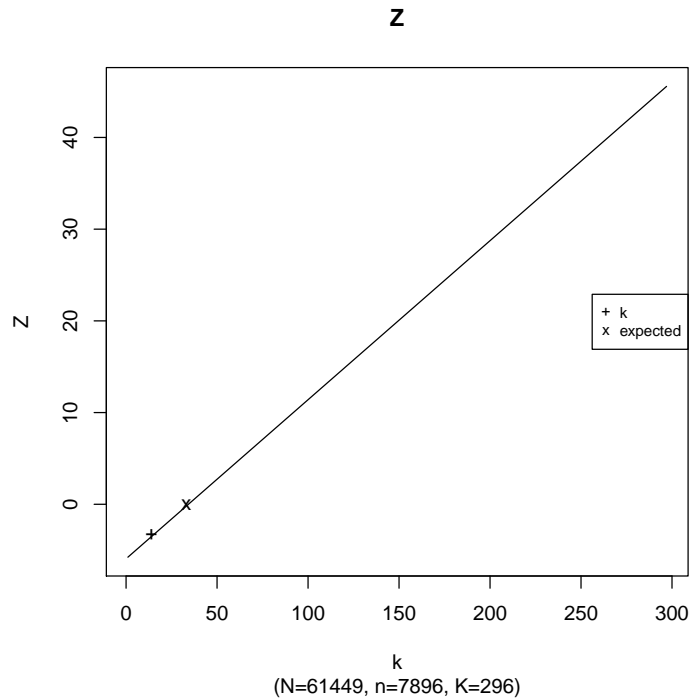
```
> wam.z(N, n, K, k);
[1] -3.338591

> wam.z(N, n, K, mode);
[1] -0.04366125
```

Graph of the function :

```
> plot(wam.z(N, n, K, allk),
+       type="l", xlab="k", ylab="Z",
+       main="Z",
+       sub="(N=61449, n=7896, K=296)")
> points(k, wam.z(N, n, K, k), pch="+")
```

```
> points(mode, wam.z(N, n, K, mode), pch="x")
> legend("right", legend=c("k", "expected"), pch=c("+", "x"), cex=0.75)
```



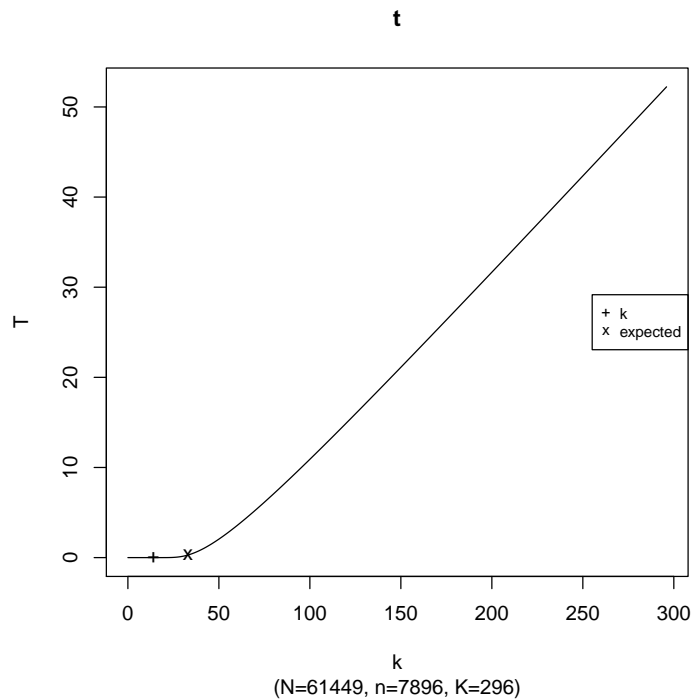
2.5 t

```
> wam.t(N, n, K, k);
[1] 5.801351e-08
> wam.t(N, n, K, mode);
```

```
[1] 0.2861072
```

Graph of the function :

```
> plot(allk, wam.t(N, n, K, allk),
+       type="l", xlab="k", ylab="T",
+       main="t",
+       sub="(N=61449, n=7896, K=296)")
> points(k, wam.t(N, n, K, k), pch="+")
> points(mode, wam.t(N, n, K, mode), pch="x")
> legend("right", legend=c("k", "expected"), pch=c("+", "x"), cex=0.75)
```



2.6 chisq

```
> wam.chisq(N, n, K, k);
```

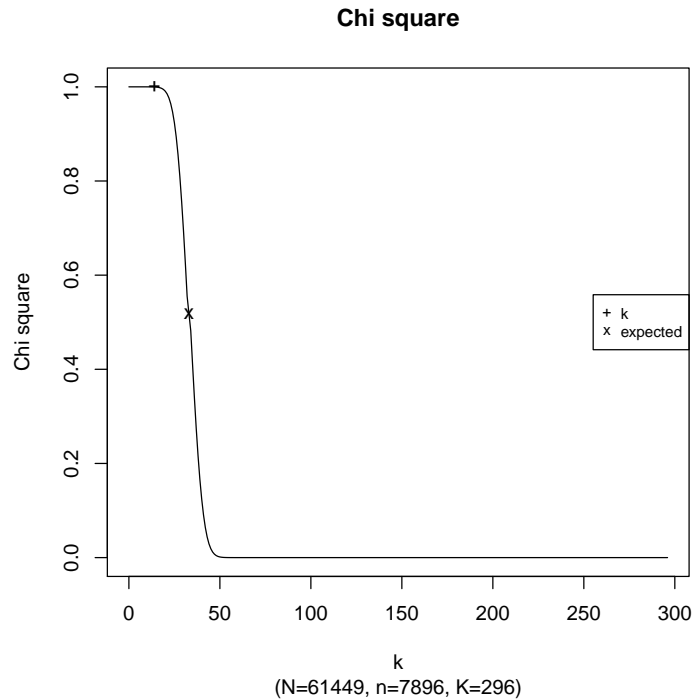
```
[1] 0.9997298
```

```
> wam.chisq(N, n, K, mode);
```

```
[1] 0.5182654
```

Graph of the function :

```
> plot(allk, wam.chisq(N, n, K, allk),
+       type="l", xlab="k", ylab="Chi square",
+       main="Chi square",
+       sub="(N=61449, n=7896, K=296)")
> points(k, wam.chisq(N, n, K, k), pch="+")
> points(mode, wam.chisq(N, n, K, mode), pch="x")
> legend("right", legend=c("k", "expected"), pch=c("+", "x"), cex=0.75)
```



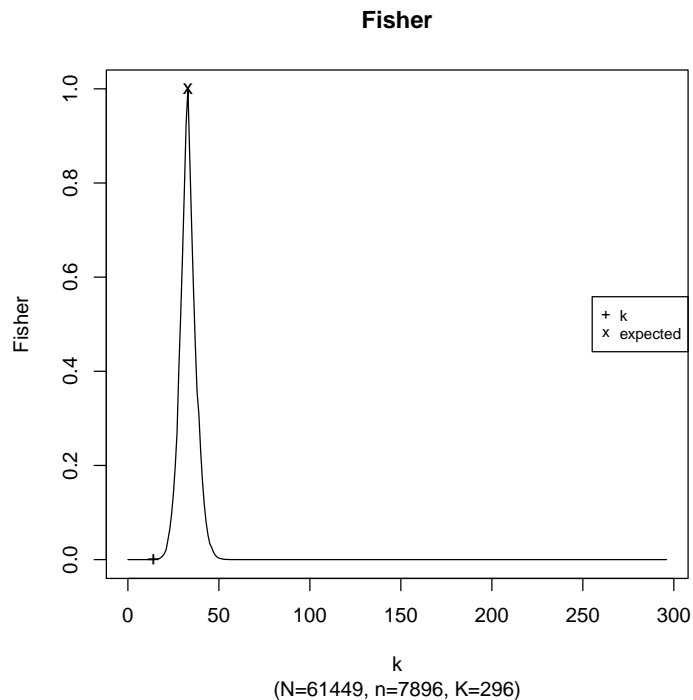
2.7 fisher

```
> wam.fisher(N, n, K, k);
[1] 0.0001353407

> wam.fisher(N, n, K, expected);
[1] 1
```

Graph of the function :

```
> plot(allk, wam.fisher(N, n, K, allk),
+       type="l", xlab="k", ylab="Fisher",
+       main="Fisher",
+       sub="(N=61449, n=7896, K=296)")
> points(k, wam.fisher(N, n, K, k), pch="+")
> points(expected, wam.fisher(N, n, K, expected), pch="x")
> legend("right", legend=c("k", "expected"), pch=c("+", "x"), cex=0.75)
```

3 High level interface

The function *wam* provide a more high level interface to the actual functions.

This function allows for computing several indicators at the same time.

It takes also as argument the subcorpus name and lexical types.

```
> rm(list=ls())
> data(robespierre, package="wam")
> attach(robespierre)
> wam.res <- wam(N, n, K, k, measure=c("loglikelihood", "specificities"),
+ types=types, parts=parts)
```

function allows for retrieving the basic information (see the manual page for WordAssociation):

```
> head(k(wam.res));

[1] 464 365 281 227 200 188

> head(association(wam.res));
```

```

      loglikelihood specificities
[1,]      2.5758482      2.1848933
[2,]      0.8133694     -0.9840740
[3,]      0.3408187     -0.5627793
[4,]      0.2069236     -0.3991217
[5,]      1.4940429      1.4588965
[6,]      0.9787803      1.0712966

```

```
> indicator.name(wam.res)
```

```
[1] "loglikelihood" "specificities"
```

The print function allows for an easy to use reading of the results. See :

```

> wam.res <- wam(N, n, K, k, measure=c("loglikelihood", "specificities"),
+ types=types, parts=parts)
> print(wam.res, from=1, to=10, parts="D4");

```

Printing association measure for 1 part(s); from: 1 to: 10

Corpus size: 61449

Sorted by: loglikelihood

```

-----
word                | sub freq | tot freq |   loglikelihood |   specificities
-----
.....
Part name: D4
Part size: 6903 tokens.
Positive specificities printed: 10
Negative specificities printed: 0
bourdon             |      20 |      20 |      87.50 |      43.25
nous                 |       3 |     430 |      79.45 |     -40.51
vous                 |       6 |     424 |      63.13 |     -32.58
salut                |      33 |      91 |      39.00 |      21.12
comité              |      35 |     103 |      37.31 |      20.20
fabre                |      13 |      19 |      34.59 |      18.43
convention           |      37 |     126 |      30.52 |      16.88
public               |      32 |     108 |      26.84 |      14.98
ils                  |      21 |     419 |      20.14 |     -11.35
il                   |     105 |     605 |      20.12 |      11.70

```

4 Bibliographie

Chaudhari, D. L., Damani, O. P. & Laxman, S. 2011. "Lexical co-occurrence, statistical significance, and word association". In: Conference on Empirical Methods in Natural Language Processing (Edinburgh, Scotland, UK, July 27-31). pp. 1058-68

<http://www.aclweb.org/anthology-new/D/D11/D11-1098.pdf>

Dunning, T. 1993. "Accurate methods for the statistics of surprise and coincidence." In: Computational Linguistics. 19(1). Pp 61–74.

<http://acl.ldc.upenn.edu/J/J93/J93-1003.pdf>

Hofland, K. and Johanssen, S. 1989. Frequency analysis of English vocabulary and grammar, based on the LOB corpus. Oxford: Clarendon.

Kilgariff, A. 1996. "Which words are particularly characteristic of a text? A survey of statistical approaches." In: Proceedings, ALLC-ACH '96. Bergen, Norway.

http://www.cse.iitb.ac.in/~shwetaghonge/prec_recall.pdf

Lafon P. 1980. "Sur la variabilité de la fréquence des formes dans un corpus". *Mots*, 1, 1980, 127–165.

http://www.persee.fr/web/revues/home/prescript/article/mots_0243-6450_1980_num_1_1_1008.