

# Package ‘languageR’

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**Type** Package

**Title** Data sets and functions with “Analyzing Linguistic Data: A practical introduction to statistics”.

**Version** 1.4

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**Description** Data sets exemplifying statistical methods, and some  
facilitatory utility functions used in “Analyzing Linguistic  
Data: A practical introduction to statistics using R”, Cambridge University Press, 2008.

**LazyData** yes

**License** GPL (>= 2)

**Depends** R(>= 2.10), methods

**Suggests** cluster, e1071, rms, Hmisc, MASS, rpart, lattice, zipfR, lme4, multcomp

**Imports** methods, lattice, lme4

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languageR-package	<i>Data sets and functions for 'Analyzing Linguistic Data'</i>
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## Description

Data sets and functions accompanying 'Analyzing Linguistic Data: A practical introduction to statistics', Cambridge University Press, 2007.

## Details

Package:	languageR
Type:	Package
Version:	1.0
Date:	2007-01-15
License:	GNU public license

The main function of this package is to make available the data sets discussed and analyzed in 'Analyzing Linguistic Data: A practical introduction to statistics using R', to appear with Cambridge University Press. The following packages should be installed, as ancillary functions in this package depend on them.

zipfR for word frequency distributions  
 lme4 for mixed-effects models  
 coda for Markov-Chain Monte Carlo estimation  
 lattice for trellis graphics  
 Matrix for mixed-effects modeling

The following packages need to be installed for working through specific examples.

rms for regression modeling  
 rpart for CART trees  
 e1071 for support vector machines  
 MASS for many useful functions  
 ape for phylogenetic clustering

The main convenience functions in this library are, by category:

**correspondence analysis** (extending code by Murtagh, 2005)

corres.fnc correspondence analysis  
 corsup.fnc supplementary data

**vocabulary richness** (supplementing current zipfR functionality)

compare.richness.fnc for two texts, compare richness  
 growth.fnc empirical vocabulary growth data for text  
 growth2vgc conversion to vgc object of zipfR  
 spectrum.fnc creates frequency spectrum  
 text2spc.fnc conversion to spc object of zipfR

**lmer functions** (p-values for mixed-effects models with lme4)

pvals.fnc p-values for table of coefficients including MCMC  
 aovlmer.fnc p-values for anova tables and/or MCMC p-value for specified factor

**simulation functions** (for comparing mixed models with traditional techniques including F1, F2, and F1+F2)

simulateRegression.fnc simulate simple regression design  
 simulateQuasif.fnc simulate data for Quasi-F ratios  
 simulateLatinsquare.fnc simulating simple Latin-square design

**miscellaneous** (convenience functions)

pairscore.fnc scatterplot matrix with correlation tests  
 collin.fnc collinearity diagnostics  
 pvals.fnc p-values and MCMC confidence intervals for mixed models  
 plot.logistic.fit.fnc diagnostic visualization for logistic models  
 xylowess.fnc trellis scatterplots with smoother  
 mvrnormplot.fnc scatterplot for bivariate standard normal random numbers with regression line  
 lmerPlotInt.fnc offers choice of four ways to visualize an interaction between two numeric predictors in an lmer model

**Author(s)**

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 Maintainer: harald.baayen@gmail.com

**References**

R. H. Baayen (2007) *Analyzing Linguistic Data: A practical introduction to statistics using R*, Cambridge: Cambridge University Press.

**Examples**

```
## Not run:
library(languageR)
data(package="languageR")

## End(Not run)
```

---

`acf.fnc`*Autocorrelation trellis graph*

---

**Description**

This function creates a trellis plot with autocorrelation functions for by-subject sequential dependencies in response latencies.

**Usage**

```
acf.fnc(dat, group="Subject", time="Trial", x = "RT", plot=TRUE, ...)
```

**Arguments**

<code>dat</code>	A data frame with (minimally) a grouping factor, an index for successive trails/events, and a behavioral measure
<code>group</code>	A grouping factor such as Subject
<code>time</code>	A sequential time measure such as Trial number in the experimental list
<code>x</code>	The dependent variable, usually a chronometric measure such as RT
<code>plot</code>	If true, a trellis graph is produced, otherwise a data frame with the data on which the trellis graph is based is returned
<code>...</code>	other optional arguments, such as layout

**Value**

If `plot=TRUE`, a trellis graph, otherwise a data frame with as column names

Lag	Autocorrelation lag
Acf	Autocorrelation
Subject	The grouping factor, typically Subject
ci	The (approximate) 95% confidence interval.

**Author(s)**

R. H. Baayen

**References**

R. H. Baayen (2001) *Word Frequency Distributions*, Dordrecht: Kluwer.

**See Also**

[lags.fnc](#)

**Examples**

```
## Not run:
data(beginningReaders)
acf.fnc(beginningReaders, x="LogRT") # autocorrelations even though nonword responses not included

## End(Not run)
```

---

addMCMCci.fnc	<i>Add MCMC confidence intervals</i>
---------------	--------------------------------------

---

**Description**

Adds MCMC confidence intervals to the plot data frames for `plotLMER.fnc()`.

**Usage**

```
addMCMCci.fnc(mcmcM, model, m, fun, pred, predname = NA, factor = FALSE)
```

**Arguments**

mcmcM	An MCMC matrix as obtained with <code>mcmcsamp()</code> or <code>pvals.fnc()\$mcmc</code>
model	A (generalized) linear mixed effects model obtained with <code>lmer()</code>
m	A design matrix constructed by <code>plotLMER.fnc()</code>
fun	A function for transforming the dependent variable
pred	The predictor to be plotted
predname	The name of the predictor
factor	Boolean for whether predictor is factor

**Details**

Not intended for independent use.

**Value**

A data frame with components

x	values for horizontal axis
lower	lower bound 95% prediction interval
upper	upper bound of 95% prediction interval

**Note**

Under development

**Author(s)**

R. H. Baayen

**See Also**

See Also as [plotLMER.fnc](#)

**Examples**

```
## Not run:  
  Not intended for independent use.  
  
## End(Not run)
```

---

affixProductivity	<i>Affix productivity</i>
-------------------	---------------------------

---

**Description**

Affix productivity, gauged by the P\* productivity measure, for 27 English affixes in 44 texts.

**Usage**

```
data(affixProductivity)
```

**Format**

A data frame with 44 observations on the following 30 variables.

semi a numeric vector of P\*-values  
anti a numeric vector of P\*-values  
ee a numeric vector of P\*-values  
ism a numeric vector of P\*-values  
ian a numeric vector of P\*-values  
ful a numeric vector of P\*-values  
y a numeric vector of P\*-values  
ness a numeric vector of P\*-values  
able a numeric vector of P\*-values  
ly a numeric vector of P\*-values  
unV a numeric vector of P\*-values  
unA a numeric vector of P\*-values  
ize a numeric vector of P\*-values  
less a numeric vector of P\*-values

erA a numeric vector of P\*-values  
 erC a numeric vector of P\*-values  
 ity a numeric vector of P\*-values  
 super a numeric vector of P\*-values  
 est a numeric vector of P\*-values  
 ment a numeric vector of P\*-values  
 ify a numeric vector of P\*-values  
 re a numeric vector of P\*-values  
 ation a numeric vector of P\*-values  
 in. a numeric vector of P\*-values  
 ex a numeric vector of P\*-values  
 en a numeric vector of P\*-values  
 be a numeric vector of P\*-values  
 AuthorCodes a factor with levels  
     BLu (King James Version: Luke-Acts)  
     BMo (Book of Mormon)  
     CAs (Aesop's fables, translation by Townsend)  
     CBo (Baum, The Marvelous Land of Oz)  
     CBp (Barrie, Peter Pan and Wendy)  
     CBw (Baum, The Wonderful Wizard of Oz)  
     CCa (Carroll, Alice's Adventures in Wonderland)  
     CCt (Carroll, Through the Looking Glass and what Alice Found There)  
     CGr (Grimm Fairy Tales, translations)  
     CKj (Kipling, The Jungle Book)  
     LAp (Austen, Pride and Prejudice)  
     LBp (Burroughs, A Princess of Mars)  
     LBw (Bronte, Wuthering Heights)  
     LC1 (Conrad, Lord Jim)  
     LCn (Conrad, Nigger of the Narcissus)  
     LDb (Doyle, The Casebook of Sherlock Holmes)  
     LDc (Dickens, The Chimes: a Goblin Story)  
     LDC (Dickens, A Christmas Carol)  
     LDh (Doyle, The Hound of the Baskervilles)  
     LDv (Doyle, The Valley of Fear)  
     LJc (James, Confidence)  
     LJe (James, The Europeans)  
     LLc (London, The Call of the Wild)  
     LLs (London, The Sea Wolf)  
     LMa (Montgomery, Anne of Avonlea)  
     LMm (Melville, Moby Dick)

LMn (Morris, News from Nowhere)  
 LMp (Milton, Paradise Lost)  
 LOs (Orczy, The Scarlet Pimpernel)  
 LSd (Stoker, Dracula)  
 LSS (Chu, More than a Chance Meeting (Startrek))  
 LTA (Trollope, Ayala's Angel)  
 LTe (Trollope, The Eustace Diamonds)  
 LTF (Trollope, Can you Forgive her?)  
 LTy (Twain, A Connecticut Yankee in King Arthur's Court)  
 LWi (Wells, The Invisible Man)  
 LWT (Wells, The Time Machine)  
 LWw (Wells, The War of the Worlds)  
 OAf (The Federalist Papers)  
 OCh (Texts sampled from Congress Hearings)  
 OCl (Texts sampled from Clinton's Election Speeches)  
 ODo (Darwin, On the Origin of the Species)  
 OGa (Selected Texts from the Government Accounting Office)  
 OJe (James, Essays in Radical Empiricism)

Registers a factor with levels B (Biblical texts) C (Children's books) L (Literary texts) O (other)

Birth a numeric vector for the author's year of birth (where available)

## Source

Most texts were obtained from the Gutenberg Project ([http://www.gutenberg.org/wiki/Main\\_Page](http://www.gutenberg.org/wiki/Main_Page)) and the Oxford Text Archive (<http://ota.ahds.ac.uk/>).

## References

Baayen, R. H. (1994) Derivational Productivity and Text Typology, *Journal of Quantitative Linguistics*, 1, 16-34.

## Examples

```
## Not run:
data(affixProductivity)
affixes.pr = prcomp(affixProductivity[,1:(ncol(affixProductivity)-3)],
  center = TRUE, scale. = TRUE)
library(lattice)
trellis.device()
super.sym = trellis.par.get("superpose.symbol")
splom(data.frame(affixes.pr$x[,1:3]),
  groups = affixProductivity$Registers,
  panel = panel.superpose,
  key = list(title = "texts in productivity space",
  text = list(c("Religious", "Children", "Literary", "Other")),
  points = list(pch = super.sym$pch[1:4], col = super.sym$col[1:4])))

## End(Not run)
```

---

alice	<i>Alice's Adventures in Wonderland</i>
-------	---

---

**Description**

The text of Lewis Carroll's 'Alice's Adventures in Wonderland', with punctuation marks removed.

**Usage**

```
data(alice)
```

**Format**

A character vector with 27269 words.

**Source**

The project Gutenberg at [http://www.gutenberg.org/wiki/Main\\_Page](http://www.gutenberg.org/wiki/Main_Page)

**Examples**

```
data(alice)
alice[1:5]
```

---

aovlmer.fnc	<i>Compute p-values for factors in mixed models</i>
-------------	---

---

**Description**

This function computes p-values for factors in a mixed-effects model on the basis of an MCMC sample.

**Usage**

```
aovlmer.fnc(object, mcmc, which, noMCMC, ...)
```

**Arguments**

object	An lmer or glmer model for a response variable fitted with lmer.
mcmc	A Markov chain Monte Carlo sample obtained with <code>pvals.fnc(..., withMCMC=TRUE)\$mcmc</code> for the lmer model.
which	A vector of integers or strings denoting the rows in the table of coefficients that specify the coefficients for the factor.
noMCMC	In earlier versions of lmer, a single-argument anova was supported. In more recent versions, this is no longer possible. Hence the p-values based on F-tests are no longer supported.
...	Other optional arguments.

**Value**

A list with components

`p` the empirical p-value for the hypothesis that the columns of an MCMC sample specified by `which` have mean zero versus a general multivariate distribution with elliptical contours

`which` the rownames specified by `which`

.

**Author(s)**

R. H. Baayen, D. Bates

**See Also**

See also `lmer`, `mcmc`, `mcmc`.

**Examples**

```
## Not run:
library(lme4, keep.source=FALSE)
data(latinsquare)
l.lmer = lmer(RT~SOA+(1|Word)+(1|Subject), data=latinsquare)
mcmc = pvals.fnc(l.lmer, nsim=10000, withMCMC=TRUE)
aovlmer.fnc(l.lmer, mcmc$mcmc, c("SOAmedium", "SOAshort"))

## End(Not run)
```

---

auxiliaries

*Auxiliaries for regular and irregular verbs in Dutch*

---

**Description**

For 285 regular and irregular Dutch verbs, the auxiliary for the present and past perfect is listed together with the count of verbal synsets in WordNet. Regular and irregular verbs are matched in the mean for lemma frequency.

**Usage**

```
data(auxiliaries)
```

**Format**

A data frame with 285 observations on the following 4 variables.

Verb a factor with 285 monomorphemic Dutch verbs.

Aux a factor with as levels the auxiliaries hebben, zijn and zijnheb (for verbs allowing both auxiliaries).

VerbalSynsets a numeric vector with the number of verbal synonym sets in WordNet in which the verb is listed.

Regularity a factor with levels irregular and regular.

**References**

Baayen, R. H. and Moscoso del Prado Martin, F. (2005) Semantic density and past-tense formation in three Germanic languages, *Language*, 81, 666-698.

**Examples**

```
data(auxiliaries)
kruskal.test(auxiliaries$VerbalSynsets, auxiliaries$Aux)
```

---

beginningReaders      *Visual lexical decision with beginning readers*

---

**Description**

Visual lexical decision latencies for beginning readers (8 year-old Dutch children).

**Usage**

```
data(beginningReaders)
```

**Format**

A data frame with 7923 observations on the following 13 variables.

Word a factor for the words.

Subject a factor for the subjects.

LogRT a numeric vector with the log-transformed reaction time (in ms).

Trial a numeric vector coding the rank of the trial in the experimental list.

OrthLength a numeric vector coding the word's length in letters.

LogFrequency a numeric vector with log-transformed frequency in Vermeer's frequency dictionary of Dutch children's texts.

LogFamilySize a numeric vector with the log-transformed morphological family size count (with family members judged to be unknown to young children removed).

ReadingScore a numeric vector with a score for reading proficiency.

- ProportionOfErrors a numeric vector for the proportion of error responses for the word.
- PC1 a numeric vector for the first principal component of a PCA orthogonalization of the preceding 4 reaction times
- PC2 a numeric vector for the second principal component of a PCA orthogonalization of the preceding 4 reaction times
- PC3 a numeric vector for the third principal component of a PCA orthogonalization of the preceding 4 reaction times
- PC4 a numeric vector for the fourth principal component of a PCA orthogonalization of the preceding 4 reaction times

## References

Perdijk, K., Schreuder, R., Verhoeven, L. and Baayen, R. H. (2006) *Tracing individual differences in reading skills of young children with linear mixed-effects models*. Manuscript, Radboud University Nijmegen.

## Examples

```
## Not run:
data(beginningReaders)
library(lme4, keep.source = FALSE)
beginningReaders.lmer = lmer(LogRT ~ PC1 + PC2 + PC3 + ReadingScore +
  OrthLength + I(OrthLength^2) + LogFrequency + LogFamilySize +
  (1|Word) + (1|Subject) + (0+LogFrequency|Subject) + (0+OrthLength|Subject) + (0+PC1|Subject),
  data = beginningReaders)

## End(Not run)
```

---

collin.fnc

*Calculate condition number with intercept included*

---

## Description

Calculates the condition number with the intercept included, following Belsley, Kuh and Welsch (1980).

## Usage

```
collin.fnc(data, colvector)
```

## Arguments

- data A data frame.
- colvector A vector with the column numbers in the data frame for which the collinearity is to be assessed. Only numeric predictors allowed.

**Value**

A list with components

svd	Singular value decomposition
cindex	Condition indices
cnumber	The condition number
pi	The phi matrix

**Author(s)**

F. J. Tweedie

**References**

Belsley, D. A. and Kuh, E. and Welsch, R. E. (1980) *Regression Diagnostics. Identifying Influential Data and Sources of Collinearity*, Wiley Series in Probability and Mathematical Statistics, New York.

**See Also**

[kappa](#)

**Examples**

```
## Not run:
  data(english)
  collin.fnc(english[english$AgeSubj=="young",], 7:29)$cnumber

## End(Not run)
```

---

compare.richness.fnc *Compare Lexical Richness of Two Texts*

---

**Description**

Comparisons of lexical richness between two texts are carried out on the basis of the vocabulary size (number of types) and on the basis of the vocabulary growth rate. Variances of the number of types and of the number of hapax legomena required for the tests are estimated with the help of LNRE models.

**Usage**

```
compare.richness.fnc(text1, text2, digits = 5)
```

**Arguments**

text1	First text in the comparison.
text2	Second text in the comparison.
digits	Number of decimal digits required for the growth rate.

**Details**

The comparison for the vocabulary size is carried out with the test statistic

$$Z = \frac{E[V_1] - E[V_2]}{\sqrt{\sigma(V_1)^2 + \sigma(V_2)^2}}$$

and the comparison of the growth rates with the test statistic

$$Z = \frac{\frac{1}{N_1}E[V_1(1)] - \frac{1}{N_2}E[V_2]}{\sqrt{\frac{1}{N_1^2}\sigma(V_1(1))^2 + \frac{1}{N_2^2}\sigma(V_2(1))^2}}$$

where  $N$  denotes the sample size in tokens,  $V$  the vocabulary size, and  $V(1)$  the number of hapax legomena.

**Value**

A summary listing the Chi-Squared measure of goodness of fit for the LNRE models (available in the zipfR package) used to estimate variances, a table listing tokens, types, hapax legomena and the vocabulary growth rate, and two-tailed tests for differences in the vocabulary sizes and growth rates with Z-score and p-value.

**Note**

It is probably unwise to attempt to apply this function to texts comprising more than 500,000 words.

**Author(s)**

R. Harald Baayen Radboud University Nijmegen and Max Planck Institute for Psycholinguistics, Nijmegen, The Netherlands. baayen@mpi.nl

**References**

Baayen, R. H. (2001) *Word Frequency Distributions*, Kluwer Academic Publishers, Dordrecht.

**Examples**

```
## Not run:
  data(alice, through, oz)
  compare.richness.fnc(tolower(alice), tolower(through[1:length(alice)]))
  compare.richness.fnc(tolower(alice), tolower(oz[1:25942]))

## End(Not run)
```

---

corres-class	<i>Class "corres"</i>
--------------	-----------------------

---

**Description**

A class for correspondence analysis

**Objects from the Class**

Objects can be created by calls of the form `new("corres", ...)`. Correspondence objects can be plotted, summarized, and printed.

**Slots**

`data`: Object of class "list"

**Methods**

No methods defined with class "corres" in the signature.

**Note**

to be expanded

**Author(s)**

R. H. Baayen

**References**

Murtagh

**See Also**

See Also `corres.fnc`.

**Examples**

```
showClass("corres")
```

---

`corres.fnc`*Correspondence Analysis*

---

**Description**

Correspondence analysis for a contingency table.

**Usage**

```
corres.fnc(xtab)
```

**Arguments**

`xtab`            A data frame cross-tabulating frequencies.

**Value**

A correspondence object with summary and plot methods. The summary method lists eigenvalue rates and coordinates, correlations and contributions for Factor 1 and Factor 2. By default, only the first six rows of the factor tables are shown. Full tables are obtained by specifying `header = FALSE` when calling `summary`. For information on higher dimensions, set the option `n` to the desired number (e.g., `n = 3`) within `summary`. See `plot.corres` for documentation of plot options.

**Author(s)**

Extension of the code in Murtagh (2005) by R. Harald Baayen  
Radboud University Nijmegen & Max Planck Institute for Psycholinguistics  
Nijmegen, The Netherlands  
email: baayen@mpi.nl

**References**

F. Murtagh (2005) *Correspondence Analysis and Data Coding with JAVA and R*, Chapman & Hall/CRC, Boca Raton, FL.

**See Also**

See [corsup.fnc](#) for adding supplementary data to a correspondence plot, and [plot.corres](#) for plot options.

**Examples**

```
## Not run:  
data(oldFrench)  
oldFrench.ca = corres.fnc(oldFrench)  
oldFrench.ca  
summary(oldFrench.ca, head = TRUE)
```

```

plot(oldFrench.ca)

# more readable plot
data(oldFrenchMeta)
plot(oldFrench.ca, rlabels = oldFrenchMeta$Genre,
      rcol = as.numeric(oldFrenchMeta$Genre), rcex = 0.5,
      extreme = 0.1, ccol = "blue")

# create subset of proze texts

prose = oldFrench[oldFrenchMeta$Genre=="prose" &
                 !is.na(oldFrenchMeta$Year),]
proseinfo = oldFrenchMeta[oldFrenchMeta$Genre=="prose" &
                          !is.na(oldFrenchMeta$Year),]
proseinfo$Period = as.factor(proseinfo$Year <= 1250)

prose.ca = corres.fnc(prose)
plot(prose.ca, addcol = FALSE,
      rcol = as.numeric(proseinfo$Period) + 1,
      rlabels = proseinfo$Year, rcex = 0.7)

# and add supplementary data for texts with unknown date of composition
proseSup = oldFrench[oldFrenchMeta$Genre == "prose" &
                    is.na(oldFrenchMeta$Year),]
corsup.fnc(prose.ca, bycol = FALSE, supp = proseSup, font = 2,
           cex = 0.8, labels = substr(rownames(proseSup), 1, 4))

## End(Not run)

```

---

corsup.fnc

*Supplementary rows or columns in correspondence analysis*


---

## Description

Corsup calculates supplementary rows or columns for correspondence analysis.

## Usage

```
corsup.fnc(corres, bycol = TRUE, supp, plot = TRUE, font = 3, labels = "",
           cex = 1)
```

## Arguments

corres	A correspondence object.
bycol	A logical value indicating whether supplementary columns (the default) or supplementary rows are required.
supp	Supplementary rows or columns from a data frame with the same structure as the data frame used for the <code>corres</code> object.

plot	A logical value indicating whether supplementary rows or columns should be added to an already existing plot.
font	An integer specifying the font to be used for plotting.
labels	A character vector with row or column names to be used for plotting.
cex	A real specifying the font size required for plotting.

**Value**

If `plot = FALSE`, a matrix with the supplementary coordinates. Otherwise, supplementary rows or columns are added to an already existing plot of a correspondence object.

**Author(s)**

Extension of the code in Murtagh (2005) by R. Harald Baayen  
 Radboud University Nijmegen & Max Planck Institute for Psycholinguistics  
 Nijmegen, The Netherlands  
 email: baayen@mpi.nl

**References**

F. Murtagh (2005) *Correspondence Analysis and Data Coding with JAVA and R*, Chapman & Hall/CRC, Boca Raton, FL.

**See Also**

[corres.fnc](#)

**Examples**

```
## Not run:
data(oldFrench)
data(oldFrenchMeta)
prose = oldFrench[oldFrenchMeta$Genre=="prose" &
  !is.na(oldFrenchMeta$Year),]
proseinfo = oldFrenchMeta[oldFrenchMeta$Genre=="prose" &
  !is.na(oldFrenchMeta$Year),]
proseinfo$Period = as.factor(proseinfo$Year <= 1250)

prose.ca = corres.fnc(prose)
plot(prose.ca, addcol = FALSE,
  rcol = as.numeric(proseinfo$Period) + 1,
  rlabels = proseinfo$Year, rcex = 0.7)

proseSup = oldFrench[oldFrenchMeta$Genre == "prose" &
  is.na(oldFrenchMeta$Year),]
corsup.fnc(prose.ca, bycol = FALSE, supp = proseSup, font = 2,
  cex = 0.8, labels = substr(rownames(proseSup), 1, 4))

## End(Not run)
```

---

danish

*Danish auditory lexical decision*

---

**Description**

Auditory lexical decision latencies for Danish complex words.

**Usage**

data(danish)

**Format**

A data frame with 3326 observations on the following 16 variables.

**Subject** a random-effect factor coding participants in the experiment.

**Word** a random-effect factor coding the words for which auditory lexical decision responses were elicited.

**Affix** a random-effect factor coding the affixes in the words.

**LogRT** the dependent variable, log response latency.

**PC1** first principal component orthogonalizing the four response latencies preceding the current trial in the experiment.

**PC2** second principal component orthogonalizing the four response latencies preceding the current trial in the experiment.

**PrevError** factor with levels CORRECT and ERROR coding whether the preceding trial elicited a correct lexical decision.

**Rank** the trial number in the experiment.

**Sex** factor coding the sex of the participant, with levels F (female) and M (male).

**LogWordFreq** log-transformed word frequency.

**LogAffixFreq** log-transformed affix frequency.

**ResidFamSize** residualized morphological family size (taking out LogWordFreq and LogAffixFreq).

**ResidSemRating** residualized semantic rating (taking out morphological family size).

**LogCUP** log-transformed complex uniqueness point (CUP).

**LogUP** log-transformed uniqueness point (UP).

**LogCUPtoEnd** log of the distance (in msec) from the CUP to the end of the word.

**References**

L. Balling and R. H. Baayen (2008). Morphological effects in auditory word recognition: Evidence from Danish. Submitted to *Language and Cognitive Processes*.

**Examples**

```
## Not run:
data(danish)
require(lme4, quietly=TRUE)

# ---- mixed-effects regression with three random intercepts

danish.lmer = lmer(LogRT ~ PC1 + PC2 + PrevError + Rank +
  ResidSemRating + ResidFamSize + LogWordFreq*LogAffixFreq*Sex +
  poly(LogCUP, 2, raw=TRUE) + LogUP + LogCUPtoEnd +
  (1|Subject) + (1|Word) + (1|Affix), data = danish)

danish.lmerA = lmer(LogRT ~ PC1 + PC2 + PrevError + Rank +
  ResidSemRating + ResidFamSize + LogWordFreq*LogAffixFreq*Sex +
  poly(LogCUP, 2, raw=TRUE) + LogUP + LogCUPtoEnd +
  (1|Subject) + (1|Word) + (1|Affix), data = danish,
  subset=abs(scale(resid(danish.lmer)))<2.5)

mcmc = pvals.fnc(danish.lmerA, nsim=10000, withMCMC=TRUE)
mcmc$fixed[,1:5]

## End(Not run)
```

---

dative

*Dative Alternation*


---

**Description**

Data describing the realization of the dative as NP or PP in the Switchboard corpus and the Treebank Wall Street Journal collection.

**Usage**

```
data(dative)
```

**Format**

A data frame with 3263 observations on the following 15 variables.

*Speaker* a factor coding speaker; available only for the subset of spoken English.

*Modality* a factor with levels spoken, written.

*Verb* a factor with the verbs as levels.

*SemanticClass* a factor with levels a (abstract: 'give it some thought'), c (communication: 'tell, give me your name'), f (future transfer of possession: 'owe, promise'), p (prevention of possession: 'cost, deny'), and t (transfer of possession: 'give an armband, send').

*LengthOfRecipient* a numeric vector coding the number of words comprising the recipient.

*AnimacyOfRec* a factor with levels animate and inanimate for the animacy of the recipient.

DefinOfRec a factor with levels definite and indefinite coding the definiteness of the recipient.

PronomOfRec a factor with levels nonpronominal and pronominal coding the pronominality of the recipient.

LengthOfTheme a numeric vector coding the number of words comprising the theme.

AnimacyOfTheme a factor with levels animate and inanimate coding the animacy of the theme.

DefinOfTheme a factor with levels definite and indefinite coding the definiteness of the theme.

PronomOfTheme a factor with levels nonpronominal and pronominal coding the pronominality of the theme.

RealizationOfRecipient a factor with levels NP and PP coding the realization of the dative.

AccessOfRec a factor with levels accessible, given, and new coding the accessibility of the recipient.

AccessOfTheme a factor with levels accessible, given, and new coding the accessibility of the theme.

## References

Bresnan, J., Cueni, A., Nikitina, T. and Baayen, R. H. (2007) Predicting the dative alternation, in Bouma, G. and Kraemer, I. and Zwarts, J. (eds.), *Cognitive Foundations of Interpretation*, Royal Netherlands Academy of Sciences, 33 pages, in press.

## Examples

```
## Not run:
data(dative)

# analysis with CART tree

library(rpart)

# ---- initial tree

dative.rp = rpart(RealizationOfRecipient ~ .,
  data = dative[ , -c(1, 3)]) # exclude the columns with subjects, verbs
plot(dative.rp, compress = TRUE, branch = 1, margin = 0.1)
text(dative.rp, use.n = TRUE, pretty = 0)

# ---- pruning the initial tree

plotcp(dative.rp)
dative.rp1 = prune(dative.rp, cp = 0.041)
plot(dative.rp1, compress = TRUE, branch = 1, margin = 0.1)
text(dative.rp1, use.n = TRUE, pretty = 0)

# analysis with logistic regression

# ---- logistic regression with the rms package
```

```

library(rms)
dative.dd = datadist(dative)
options(datadist = 'dative.dd')
dative.lrm = lrm(RealizationOfRecipient ~
  AccessOfTheme + AccessOfRec + LengthOfRecipient + AnimacyOfRec +
  AnimacyOfTheme + PronomOfTheme + DefinOfTheme + LengthOfTheme+
  SemanticClass + Modality, data = dative)
anova(dative.lrm)
plot(Predict(dative.lrm))

# ---- mixed-effects logistic regression with the lme4 package

library(lme4)

dative.lmer = lmer(RealizationOfRecipient ~ AccessOfTheme +
  AccessOfRec + LengthOfRecipient + AnimacyOfRec +
  AnimacyOfTheme + PronomOfTheme + DefinOfTheme + LengthOfTheme +
  SemanticClass + Modality + (1|Verb),
  data = dative, family = "binomial")

print(dative.lmer, corr=FALSE)

# multiple comparisons for Accessibility of Theme
library(multcomp)
par(mar=c(5,8,3,1))
AcOfTheme.glt <- glt(dative.lmer, linfct = mcp(AccessOfTheme = "Tukey"))
plot(AcOfTheme.glt)
abline(v=0)
summary(AcOfTheme.glt)

## End(Not run)

```

---

dativeSimplified

*Dative Alternation - simplified data set*


---

### Description

Data describing the realization of the dative as NP or PP in the Switchboard corpus and the Treebank Wall Street Journal collection. Simplified version of the dative data set.

### Usage

```
data(dativeSimplified)
```

### Format

A data frame with 903 observations on the following 5 variables.

Verb a factor with the verbs as levels.

AnimacyOfRec a factor with levels animate and inanimate for the animacy of the recipient.

LengthOfTheme a numeric vector coding the number of words comprising the theme.

AnimacyOfTheme a factor with levels animate and inanimate coding the animacy of the theme.

RealizationOfRec a factor with levels NP and PP coding the realization of the dative.

## References

Bresnan, J., Cueni, A., Nikitina, T. and Baayen, R. H. (2007) Predicting the dative alternation, in Bouma, G. and Kraemer, I. and Zwarts, J. (eds.), *Cognitive Foundations of Interpretation*, Royal Netherlands Academy of Sciences, 33 pages, in press.

## Examples

```
## Not run:  
  data(dative)  
  
## End(Not run)
```

---

degreesOrKnots.fnc      *Extract degree of polynomial or knots for restricted cubic spline*

---

## Description

Extract degree of polynomial or knots for restricted cubic spline from the predictor name

## Usage

```
degreesOrKnots.fnc(name)
```

## Arguments

name                    name of predictor, e.g. poly(X, 2, raw = TRUE)

## Details

attempts to find degrees or knots if present in input name

## Value

Returns an integer for degrees or knots

## Note

not intended for independent use

**Author(s)**

R. H. Baayen

**See Also**

See Also as [plotLMER.fnc](#)

**Examples**

```
## Not run: not intended for independent use
```

---

durationsGe

*Durational measurements on the Dutch prefix ge-*

---

**Description**

Durational measurements on the Dutch prefix *ge-* in the Spoken Dutch Corpus.

**Usage**

```
data(durationsGe)
```

**Format**

A data frame with 428 observations on the following 8 variables.

*Word* a factor with the words as levels.

*Frequency* a numeric vector with the word's absolute frequency in the Spoken Dutch Corpus.

*Speaker* a factor with the speakers as levels.

*Sex* a factor with levels *female* and *male*, this information is missing for one speaker.

*YearOfBirth* a numeric vector with years of birth.

*DurationOfPrefix* a numeric vector with the duration of the prefix *-ont* in seconds.

*SpeechRate* a numeric vector coding speech rate in number of syllables per second.

*NumberSegmentsOnset* a numeric vector for the number of segments in the onset of the stem.

**References**

Pluymaekers, M., Ernestus, M. and Baayen, R. H. (2005) Frequency and acoustic length: the case of derivational affixes in Dutch, *Journal of the Acoustical Society of America*, 118, 2561-2569.

**Examples**

```
## Not run:
data(durationsGe)
durationsGe$Frequency = log(durationsGe$Frequency + 1)
durationsGe$YearOfBirth = durationsGe$YearOfBirth - 1900

durationsGe.lm = lm(DurationOfPrefix ~ Frequency+SpeechRate, data = durationsGe)
summary(durationsGe.lm)

# ---- model criticism

plot(durationsGe.lm)
outliers = c(271, 392, 256, 413, 118, 256)
durationsGe.lm = lm(DurationOfPrefix ~ Frequency + SpeechRate,
  data = durationsGe[-outliers, ])
summary(durationsGe.lm)

## End(Not run)
```

---

durationsOnt

*Durational measurements on the Dutch prefix ont-*


---

**Description**

Durational measurements on the Dutch prefix *ont-* in the Spoken Dutch Corpus.

**Usage**

```
data(durationsOnt)
```

**Format**

A data frame with 102 observations on the following 11 variables.

**Word** a factor with the words as levels.

**Frequency** a numeric vector with the word's logarithmically transformed frequency in the Spoken Dutch Corpus.

**Speaker** a factor with speakers as levels.

**Sex** a factor with levels female and male.

**YearOfBirth** a numeric vector coding year of birth of the speaker - 1900.

**DurationOfPrefix** a numeric vector for the duration of ont- in seconds

**DurationPrefixVowel** a numeric vector for the duration of the vowel in the prefix in seconds.

**DurationPrefixNasal** a numeric vector for the duration of the nasal in the prefix in seconds.

**DurationPrefixPlosive** a numeric vector for the duration of the plosive in the prefix in seconds.

**NumberOfSegmentsOnset** a numeric vector for the number of segments in the onset of the stem.

**PlosivePresent** a factor with levels no and yes for whether the plosive is realized in the signal.

**SpeechRate** a numeric vector coding speech rate in number of syllables per second.

## References

Pluymaekers, M., Ernestus, M. and Baayen, R. H. (2005) Frequency and acoustic length: the case of derivational affixes in Dutch, *Journal of the Acoustical Society of America*, 118, 2561-2569.

## Examples

```

data(durationsOnt)

##### modeling the duration of the prefix

prefix.lm = lm(DurationOfPrefix ~ (YearOfBirth + SpeechRate) * Frequency,
  data = durationsOnt)
summary(prefix.lm)

# ---- model criticism

plot(prefix.lm)
outliers = c(36, 35, 17, 72)
prefix.lm = lm(DurationOfPrefix ~ (YearOfBirth + SpeechRate) * Frequency,
  data = durationsOnt[-outliers,])
summary(prefix.lm)

##### modeling the presence of the /t/

library(rms)
durationsOnt.dd = datadist(durationsOnt)
options(datadist = 'durationsOnt.dd')

plosive.lrm = lrm(PlosivePresent ~ SpeechRate + YearOfBirth,
  data = durationsOnt, x = TRUE, y = TRUE)
plosive.lrm
validate(plosive.lrm, bw = TRUE, B = 200)

##### modeling the duration of the /n/

nasal.lm = lm(DurationPrefixNasal ~ PlosivePresent + Frequency +
  YearOfBirth, data = durationsOnt)
summary(nasal.lm)

# ---- model criticism

plot(nasal.lm)
outliers = c(71, 28, 62, 33)
nasal.lm = lm(DurationPrefixNasal ~ PlosivePresent + Frequency +
  YearOfBirth, data = durationsOnt[-outliers,])
summary(nasal.lm)

```

**Description**

A distance matrix for the conversations of 165 speakers in the Spoken Dutch Corpus. Metadata on the speakers are available in a separate dataset, `dutchSpeakersDistMeta`.

**Usage**

```
data(dutchSpeakersDist)
```

**Format**

A data frame for a 165 by 165 matrix of between-speaker differences.

**Source**

<http://lands.let.kun.nl/cgn/> data collected and analyzed in collaboration with Patrick Juola

**References**

Juola, P. (2003) The time course of language change, *Computers and the Humanities*, 37, 77-96.

Juola, P. and Baayen, R. H. (2005) A Controlled-corpus Experiment in Authorship Identification by Cross-entropy, *Literary and Linguistic Computing*, 20, 59-67.

**Examples**

```
## Not run:
data(dutchSpeakersDist)
dutchSpeakersDist.d = as.dist(dutchSpeakersDist)
dutchSpeakersDist.mds = cmdscale(dutchSpeakersDist.d, k = 3)

data(dutchSpeakersDistMeta)
dat = data.frame(dutchSpeakersDist.mds,
  Sex = dutchSpeakersDistMeta$Sex,
  Year = dutchSpeakersDistMeta$AgeYear,
  EduLevel = dutchSpeakersDistMeta$EduLevel)
dat = dat[!is.na(dat$Year),]

par(mfrow=c(1,2))
plot(dat$Year, dat$X1, xlab="year of birth",
  ylab = "dimension 1", type = "p")
lines(lowess(dat$Year, dat$X1))
boxplot(dat$X3 ~ dat$Sex, ylab = "dimension 3")
par(mfrow=c(1,1))

cor.test(dat$X1, dat$Year, method="sp")
t.test(dat$X3~dat$Sex)

## End(Not run)
```

---

dutchSpeakersDistMeta *Metadata for dutchSpeakersDist*

---

**Description**

Meta-data for the cross-entropy based between-speaker distance matrix dutchSpeakersDist

**Usage**

```
data(dutchSpeakersDistMeta)
```

**Format**

A data frame with 165 observations on the following 6 variables.

Speaker a factor with speakers as levels.

Sex a factor with levels female and male.

AgeYear a numeric vector with the speakers' year of birth.

AgeGroup a factor with levels age18to24, age25to34, age35to44, age45to55, and age56up.

ConversationType a factor with levels femaleOnly maleFemale, maleOnly, and unknown.

EduLevel a factor with levels EduUnknown, high, low mid

**Source**

<http://lands.let.kun.nl/cgn/>

**References**

Juola, P. (2003) The time course of language change, *Computers and the Humanities*, 37, 77-96.

Juola, P. and Baayen, R. H. (2005) A Controlled-corpus Experiment in Authorship Identification by Cross-entropy, *Literary and Linguistic Computing*, 20, 59-67.

**Examples**

```
## Not run:  
  data(dutchSpeakersDistMeta)  
  
## End(Not run)
```

---

 english

*English visual lexical decision and naming latencies*


---

### Description

This data set gives mean visual lexical decision latencies and word naming latencies to 2284 monomorphemic English nouns and verbs, averaged for old and young subjects, with various predictor variables.

### Usage

```
data(english)
```

### Format

A data frame with 4568 observations on the following variables.

`RTlexdec` numeric vector of log RT in visual lexical decision.

`RTnaming` numeric vector of log RT in word naming.

`Familiarity` numeric vector of subjective familiarity ratings.

`Word` a factor with 2284 words.

`AgeSubject` a factor with as levels the age group of the subject: young versus old.

`WordCategory` a factor with as levels the word categories N (noun) and V (verb).

`WrittenFrequency` numeric vector with log frequency in the CELEX lexical database.

`WrittenSpokenFrequencyRatio` numeric vector with the logged ratio of written frequency (CELEX) to spoken frequency (British National Corpus).

`FamilySize` numeric vector with log morphological family size.

`DerivationalEntropy` numeric vector with derivational entropy.

`InflectionalEntropy` numeric vector with inflectional entropy.

`NumberSimplexSynsets` numeric vector with the log-transformed count of synonym sets in WordNet in which the word is listed.

`NumberComplexSynsets` numeric vector with the log-transformed count of synonym sets in WordNet in which the word is listed as part of a compound.

`LengthInLetters` numeric vector with length of the word in letters.

`Ncount` numeric vector with orthographic neighborhood density, defined as the number of lemmas in CELEX with the same length (in letters) at Hamming distance 1.

`MeanBigramFrequency` numeric vector with mean log bigram frequency.

`FrequencyInitialDiphone` numeric vector with log frequency of initial diphone.

`ConspelV` numeric vector with type count of orthographic neighbors.

`ConspelN` numeric vector with token count of orthographic neighbors.

`ConphonV` numeric vector with type count of phonological neighbors.

ConphonN numeric vector with token count of phonological neighbors.  
 ConfriendsV numeric vector with type counts of consistent words.  
 ConfriendsN numeric vector with token counts of consistent words.  
 ConffvV numeric vector with type count of forward inconsistent words  
 ConffvN numeric vector with token count of forward inconsistent words  
 ConfbvV numeric vector with type count of backward inconsistent words  
 ConfbvN numeric vector with token count of backward inconsistent words  
 NounFrequency numeric vector with the frequency of the word used as noun.  
 VerbFrequency numeric vector with the frequency of the word used as verb.  
 CV factor specifying whether the initial phoneme of the word is a consonant (C) or a vowel (V).  
 Obstruent factor specifying whether the initial phoneme of the word is a continuant (cont) or an  
 obstruent (obst).  
 Frication factor specifying whether the initial phoneme has a burst (burst) or frication (frication)  
 for consonant-initial words, and for vowel-initial words whether the vowel is long or short.  
 Voice factor indicating whether the initial phoneme is voiced or voiceless.  
 FrequencyInitialDiphoneWord numeric vector with the log-transformed frequency of the initial  
 diphone given that it is syllable-initial.  
 FrequencyInitialDiphoneSyllable numeric vector with the log-transformed frequency of the  
 initial diphone given that it is word initial.  
 CorrectLexdec numeric vector with the proportion of subjects that accepted the item as a word in  
 lexical decision.

### Source

Balota, D.A., Cortese, M.J. and Pilotti, M. (1999) *Visual lexical decision latencies for 2906 words*. Available at [http://www.artsci.wustl.edu/~dbalota/lexical\\_decision.html](http://www.artsci.wustl.edu/~dbalota/lexical_decision.html).

Spieler, D. H. and Balota, D. A. (1998) *Naming latencies for 2820 words*, <http://www.artsci.wustl.edu/~dbalota/naming.html>.

### References

Balota, D., Cortese, M., Sergent-Marshall, S., Spieler, D. and Yap, M. (2004) Visual word recognition for single-syllable words, *Journal of Experimental Psychology:General*, 133, 283-316.

Baayen, R.H., Feldman, L. and Schreuder, R. (2006) Morphological influences on the recognition of monosyllabic monomorphemic words, *Journal of Memory and Language*, 53, 496-512.

### Examples

```

data(english)

# ---- orthogonalize orthographic consistency measures

items = english[english$AgeSubject == "young",]
items.pca = prcomp(items[, c(18:27)], center = TRUE, scale = TRUE)

```

```

x = as.data.frame(items.pca$rotation[,1:4])
items$PC1 = items.pca$x[,1]
items$PC2 = items.pca$x[,2]
items$PC3 = items.pca$x[,3]
items$PC4 = items.pca$x[,4]
items2 = english[english$AgeSubject != "young", ]
items2$PC1 = items.pca$x[,1]
items2$PC2 = items.pca$x[,2]
items2$PC3 = items.pca$x[,3]
items2$PC4 = items.pca$x[,4]
english = rbind(items, items2)

# ---- add Noun-Verb frequency ratio

english$NVratio = log(english$NounFrequency+1)-log(english$VerbFrequency+1)

# ---- build model with ols() from rms

library(rms)
english.dd = datadist(english)
options(datadist = 'english.dd')

english.ols = ols(RTlexdec ~ Voice + PC1 + MeanBigramFrequency +
  rcs(WrittenFrequency, 5) + rcs(WrittenSpokenFrequencyRatio, 3) +
  NVratio + WordCategory + AgeSubject +
  rcs(FamilySize, 3) + InflectionalEntropy +
  NumberComplexSynsets + rcs(WrittenFrequency, 5) : AgeSubject,
  data = english, x = TRUE, y = TRUE)

# ---- plot partial effects

plot(Predict(english.ols))

# ---- validate the model

validate(english.ols, bw = TRUE, B = 200)

```

---

etymology

*Etymological age and regularity in Dutch*


---

### Description

Estimated etymological age for regular and irregular monomorphemic Dutch verbs, together with other distributional predictors of regularity.

### Usage

```
data(etymology)
```

**Format**

A data frame with 285 observations on the following 14 variables.

Verb a factor with the verbs as levels.

WrittenFrequency a numeric vector of logarithmically transformed frequencies in written Dutch (as available in the CELEX lexical database).

NcountStem a numeric vector for the number of orthographic neighbors.

MeanBigramFrequency a numeric vector for mean log bigram frequency.

InflectionalEntropy a numeric vector for Shannon's entropy calculated for the word's inflectional variants.

Auxiliary a factor with levels hebben, zijn and zijnheb for the verb's auxiliary in the perfect tenses.

Regularity a factor with levels irregular and regular.

LengthInLetters a numeric vector of the word's orthographic length.

Denominative a factor with levels Den and N specifying whether a verb is derived from a noun according to the CELEX lexical database.

FamilySize a numeric vector for the number of types in the word's morphological family.

EtymAge an ordered factor with levels Dutch, DutchGerman, WestGermanic, Germanic and IndoEuropean.

Valency a numeric vector for the verb's valency, estimated by its number of argument structures.

NVratio a numeric vector for the log-transformed ratio of the nominal and verbal frequencies of use.

WrittenSpokenRatio a numeric vector for the log-transformed ratio of the frequencies in written and spoken Dutch.

**References**

Baayen, R. H. and Moscoso del Prado Martin, F. (2005) Semantic density and past-tense formation in three Germanic languages, *Language*, 81, 666-698.

Tabak, W., Schreuder, R. and Baayen, R. H. (2005) Lexical statistics and lexical processing: semantic density, information complexity, sex, and irregularity in Dutch, in Kepser, S. and Reis, M., *Linguistic Evidence - Empirical, Theoretical, and Computational Perspectives*, Berlin: Mouton de Gruyter, pp. 529-555.

**Examples**

```
## Not run:
data(etymology)

# ---- EtymAge should be an ordered factor, set contrasts accordingly

etymology$EtymAge = ordered(etymology$EtymAge, levels = c("Dutch",
"DutchGerman", "WestGermanic", "Germanic", "IndoEuropean"))
options(contrasts=c("contr.treatment", "contr.treatment"))

library(rms)
```

```

etymology.dd = datadist(etymology)
options(datadist = 'etymology.dd')

# ---- EtymAge as additional predictor for regularity

etymology.lrm = lrm(Regularity ~ WrittenFrequency +
  rcs(FamilySize, 3) + NcountStem + InflectionalEntropy +
  Auxiliary + Valency + NVratio + WrittenSpokenRatio + EtymAge,
  data = etymology, x = TRUE, y = TRUE)
anova(etymology.lrm)

# ---- EtymAge as dependent variable

etymology.lrm = lrm(EtymAge ~ WrittenFrequency + NcountStem +
  MeanBigramFrequency + InflectionalEntropy + Auxiliary +
  Regularity + LengthInLetters + Denominative + FamilySize + Valency +
  NVratio + WrittenSpokenRatio, data = etymology, x = TRUE, y = TRUE)

# ---- model simplification

etymology.lrm = lrm(EtymAge ~ NcountStem + Regularity + Denominative,
  data = etymology, x = TRUE, y = TRUE)
validate(etymology.lrm, bw=TRUE, B=200)

# ---- plot partial effects and check assumptions ordinal regression

plot(Predict(etymology.lrm))
plot(etymology.lrm)
resid(etymology.lrm, 'score.binary', pl = TRUE)
plot.xmean.ordinaly(EtymAge ~ NcountStem, data = etymology)

## End(Not run)

```

---

faz

*Frankfurter frequencies*


---

## Description

Frequencies of references to previous years in issues of the Frankfurter Allgemeine Zeitung published in 1994.

## Usage

```
data(faz)
```

## Format

A data frame with 800 observations on the following 2 variables.

**Year** a numeric vector coding years referenced in articles published in 1994.

**Frequency** a numeric vector for the frequencies with which years are referenced.

## References

Pollman, T. and Baayen, R. H. (2001) Computing historical consciousness. A quantitative inquiry into the presence of the past in newspaper texts, *Computers and the Humanities*, 35, 237-253.

## Examples

```
## Not run:
data(faz)
faz$Distance = 1:nrow(faz)

# ---- visualization

plot(log(faz$Distance), log(faz$Frequency + 1),
     xlab = "log Distance", ylab = "log Frequency")
abline(v = log(49), lty=1, col="red") # 1945
abline(v = log(54), lty=1, col="red") # 1940
abline(v = log(76), lty=2, col="blue") # 1918
abline(v = log(80), lty=2, col="blue") # 1914

# ---- breakpoint analysis

deviances = rep(0, nrow(faz)-1)
faz$LogFrequency = log(faz$Frequency + 1)
faz$LogDistance = log(faz$Distance)
for (pos in 1 : (nrow(faz)-1)) { # be patient
  breakpoint = log(pos)
  faz$ShiftedLogDistance = faz$LogDistance - breakpoint
  faz$PastBreakPoint = as.factor(faz$ShiftedLogDistance > 0)
  faz.both = lm(LogFrequency~ShiftedLogDistance:PastBreakPoint, data = faz)
  deviances[pos] = deviance(faz.both)
}
breakpoint = log(which(deviances == min(deviances)))

# ---- refit and plot

faz$ShiftedLogDistance = faz$LogDistance - breakpoint
faz$PastBreakPoint = as.factor(faz$ShiftedLogDistance > 0)
faz.both = lm(LogFrequency ~ ShiftedLogDistance:PastBreakPoint, data = faz)

plot(faz$LogDistance, faz$LogFrequency,
     xlab = "log Distance", ylab = "log Frequency", col = "darkgrey")
lines(faz$LogDistance, fitted(faz.both))

## End(Not run)
```

**Description**

Phonological specifications for onset, nucleus and offset for 1697 Dutch monomorphemic words with a final obstruent. These final obstruents may exhibit a voicing alternation that is traditionally described as syllable-final devoicing: underlying /d/ in /hond/ becomes a /t/ when syllable-final ([hOnt]) and remains a /d/ otherwise ([hOn-den]).

**Usage**

```
data(finalDevoicing)
```

**Format**

A data frame with 1697 observations on the following 9 variables.

Word a factor with the words as levels.

Onset1Type a factor for the first consonant in the onset, with levels None, Obstruent and Sonorant.

Onset2Type a factor for the second consonant in the onset, with levels None, Obstruent and Sonorant.

VowelType a factor describing the vowel with levels iuy, long and short.

ConsonantType a factor for the first consonant in the offset, with levels None, Obstruent and Sonorant.

Obstruent a factor describing place and manner of articulation of the final obstruent, with levels F (/f,v/), P (/p,b/), S (/s,z/), T (/t,d/) and X (/x,g/).

Nsyll a numeric vector for the number of syllables in the word.

Stress a factor with levels A (antepenult), F (final) and P (penult).

Voice a factor with levels voiced and voiceless.

**References**

Ernestus, M. and Baayen, R. H. (2003) Predicting the unpredictable: Interpreting neutralized segments in Dutch, *Language*, 79, 5-38.

**Examples**

```
## Not run:
data(finalDevoicing)
library(rpart)

# ---- CART tree

finalDevoicing.rp = rpart(Voice ~ ., data = finalDevoicing[, -1])
plotcp(finalDevoicing.rp)
finalDevoicing.pruned = prune(finalDevoicing.rp, cp = 0.021)
plot(finalDevoicing.pruned, margin = 0.1, compress = TRUE)
text(finalDevoicing.pruned, use.n = TRUE, pretty = 0, cex=0.8)

# ---- logistic regression
```

```

library(rms)

finalDevoicing.dd = datadist(finalDevoicing)
options(datadist='finalDevoicing.dd')

finalDevoicing.lrm = lrm(Voice ~ VowelType + ConsonantType + Obstruent +
  Nsyll + Stress + Onset1Type + Onset2Type, data = finalDevoicing)
anova(finalDevoicing.lrm)

# ---- model simplification

fastbw(finalDevoicing.lrm)

finalDevoicing.lrm = lrm(Voice ~ VowelType + ConsonantType +
  Obstruent + Nsyll, data = finalDevoicing, x = TRUE, y = TRUE)

plot(Predict(finalDevoicing.lrm))

# ---- model validation

validate(finalDevoicing.lrm, B = 200)

## End(Not run)

```

---

getKnots.fnc

*Extracts knots from variable name*


---

## Description

Extracts knots for predictor specified simply as, e.g., X from column names of model@X or model@frame

## Usage

```
getKnots.fnc(colnms, xlb)
```

## Arguments

colnms	columns of model@X
xlb	simple predictor name

## Details

not intended for independent use

## Value

an integer (number of knots)

**Note**

not intended for independent use

**Author(s)**

R. H. Baayen

**See Also**

See Also as [plotLMER.fnc](#)

**Examples**

```
## Not run: not intended for independent use
```

---

*getMCMCintervals.fnc*    *calculate HPD prediction intervals*

---

**Description**

calculate HPD 95% prediction intervals

**Usage**

```
getMCMCintervals.fnc(fixf, mcmcMatrix, m)
```

**Arguments**

<code>fixf</code>	vector of fixed effects coefficients ( <code>fixef(model.lmer)</code> )
<code>mcmcMatrix</code>	MCMC matrix obtained with <code>mcmcSamp</code> or <code>pvals.fnc</code>
<code>m</code>	model matrix

**Details**

not intended for independent use

**Value**

A matrix with columns "lower" and "upper" and rows corresponding to the values of the predictor to be plotted on the X-axis.

**Note**

not intended for independent use

**Author(s)**

R. H. Baayen

**References**

languageR

**See Also**

See Also as [plotLMER.fnc](#)

**Examples**

```
## Not run: not intended for independent use
```

---

getPos.fnc	<i>determine position for labels for interaction plots</i>
------------	--

---

**Description**

determines the position (in the X and Y vectors) for the adding of text to an interaction plot

**Usage**

```
getPos.fnc(vec, pos)
```

**Arguments**

vec	vector of Y values
pos	can be "beg", "mid", "end"

**Details**

not intended for independent use

**Value**

an integer specifying position in vector for X and Y values in plot

**Note**

not indeded for independent use

**Author(s)**

R. H. Baayen

**See Also**

See Also as [plotLMER.fnc](#)

**Examples**

```
## Not run: not intended for independent use
```

---

getRange.fnc                      *Extracts range of predicted values from list of data frames*

---

**Description**

Extracts range of predicted values from list of data frames

**Usage**

```
getRange.fnc(lst)
```

**Arguments**

lst                      a list with one or more data frames with column names Y and optionally lower and upper.

**Details**

not intended for independent use.

**Value**

value                      a two-element vector specifying the range of values in Y

**Note**

not intended for separate use.

**Author(s)**

R. H. Baayen

**See Also**

See Also as [plotLMER.fnc](#)

**Examples**

```
## Not run:  
  not intended for independent use  
  
## End(Not run)
```

---

getRoot.fnc	<i>extract simple name of predictor from expression with poly</i>
-------------	---

---

**Description**

extract X from expressions such as `poly(X, 3, raw = TRUE)`

**Usage**

```
getRoot.fnc(xlabel)
```

**Arguments**

`xlabel` character string for predictor name

**Details**

not intended for independent use

**Value**

a character string (simple name of predictor)

**Note**

not intended for independent use

**Author(s)**

R. H. Baayen

**See Also**

See Also as [plotLMER.fnc](#)

**Examples**

```
## Not run: not intended for independent use
```

---

growth-class

*Class "growth"*

---

### **Description**

A class for the analysis of word frequency distributions

### **Objects from the Class**

Objects can be created by calls of the form `new("growth", ...)`. Growth objects can be plotted, summarized, and printed.

### **Slots**

data: Object of class "list"

### **Methods**

No methods defined with class "growth" in the signature.

### **Note**

to be expanded

### **Author(s)**

R. H. Baayen

### **References**

R. H. Baayen, 2007

### **See Also**

See Also `growth.fnc`.

### **Examples**

```
showClass("corres")
```

---

growth.fnc                      *Calculate vocabulary growth curve and vocabulary richness measures*

---

### Description

This function calculates, for an increasing sequence of text sizes, the observed number of types, hapax legomena, dis legomena, tris legomena, and selected measures of lexical richness.

### Usage

```
growth.fnc(text = alice, size = 646, nchunks = 40, chunks = 0)
```

### Arguments

text	A vector of strings representing a text.
size	An integer giving the size of a text chunk when the text is to be split into a series of equally-sized text chunks.
nchunks	An integer denoting the number of desired equally-sized text chunks.
chunks	An integer vector denoting the token sizes for which growth measures are required. When chunks is specified, size and nchunks are ignored.

### Value

A growth object with methods for plotting, printing. As running this function on large texts may take some time, a period is printed on the output device for each completed chunk to indicate progress.

The data frame with the actual measures, which can be extracted with `object.name@data$data`, has the following columns.

Chunk	a numeric vector with chunk numbers.
Tokens	a numeric vector with the number of tokens up to and including the current chunk.
Types	a numeric vector with the number of types up to and including the current chunk.
HapaxLegomena	a numeric vector with the corresponding count of hapax legomena.
DisLegomena	a numeric vector with the corresponding count of dis legomena.
TrisLegomena	a numeric vector with the corresponding count of tris legomena.
Yule	a numeric vector with Yule's K.
Zipf	a numeric vector with the slope of Zipf's rank-frequency curve in the double-logarithmic plane.
TypeTokenRatio	a numeric vector with the ratio of types to tokens.
Herdan	a numeric vector with Herdan's C.
Guiraud	a numeric vector with Guiraud's R.
Sichel	a numeric vector with Sichel's S.
Lognormal	a numeric vector with mean log frequency.

**Author(s)**

R. H. Baayen

**References**

R. H. Baayen (2001) *Word Frequency Distributions*, Dordrecht: Kluwer Academic Publishers.  
Tweedie, F. J. & Baayen, R. H. (1998) How variable may a constant be? Measures of lexical richness in perspective, *Computers and the Humanities*, 32, 323-352.

**See Also**

See Also [plot.growth](#), and the zipfR package.

**Examples**

```
## Not run:  
data(alice)  
alice.growth = growth.fnc(alice)  
plot(alice.growth)  
  
## End(Not run)
```

---

growth2vgc.fnc	<i>Conversion of growth object into a vgc object</i>
----------------	--

---

**Description**

This function converts a growth object (as defined in the languageR package) to a vgc object (as defined in the zipfR package).

**Usage**

```
growth2vgc.fnc(growth)
```

**Arguments**

growth            A growth object obtained with growth.fnc().

**Value**

A vgc object as defined in the zipfR library.

**Author(s)**

R. H. Baayen

## References

R. H. Baayen (2001) *Word Frequency Distributions*, Dordrecht: Kluwer Academic Publishers.  
zipfR Website: <URL: <http://purl.org/stefan.evert/zipfR/>>

## See Also

See also [growth.fnc](#) and the zipfR package.

## Examples

```
## Not run:  
library(zipfR)  
  
data(alice)  
alice.growth = growth.fnc(text = alice, size = 648, nchunks = 40)  
alice.vgc = growth2vgc.fnc(alice.growth)  
plot(alice.vgc)  
  
## End(Not run)
```

---

havelaar

*The determiner 'het' in the Dutch novel Max Havelaar*

---

## Description

The frequency of the determiner 'het' in the Dutch novel 'Max Havelaar' by Multatuli (Eduard Douwes Dekker), in 99 consecutive text fragments of 1000 tokens each.

## Usage

```
data(havelaar)
```

## Format

A data frame with 99 observations on the following 2 variables.

Chunk a numeric vector with the indices of the text fragments.

Frequency a numeric vector with the frequencies of the determiner 'het' in the text fragments.

## Source

The text of Max Havelaar was obtained from the Project Gutenberg at [http://www.gutenberg.org/wiki/Main\\_Page](http://www.gutenberg.org/wiki/Main_Page)

**Examples**

```
## Not run:
data(havelaar)

n = 1000 # token size of text fragments
p = mean(havelaar$Frequency / n) # relative frequencies

plot(qbinom(ppoints(99), n, p), sort(havelaar$Frequency),
     xlab = paste("quantiles of (", n, ", ", round(p, 4),
                 ") - binomial", sep=""), ylab = "frequencies")

lambda = mean(havelaar$Frequency)
ks.test(havelaar$Frequency, "ppois", lambda)
ks.test(jitter(havelaar$Frequency), "ppois", lambda)

## End(Not run)
```

---

heid

*Lexical decision latencies for words ending in -heid*


---

**Description**

A simplified version of the primingHeid dataset.

**Usage**

```
data(heid)
```

**Format**

A data frame with 832 observations on the following 4 variables.

Subject a factor with subjects as levels.

Word a factor with words as levels.

RT a numeric vector with logarithmically transformed reaction times in visual lexical decision.

BaseFrequency a numeric vector with the logarithmically transformed frequency of the base adjective of the word with the suffix *-heid*.

**References**

De Vaan, L., Schreuder, R. and Baayen, R. H. (2007) Regular morphologically complex neologisms leave detectable traces in the mental lexicon, *The Mental Lexicon*, 2, in press.

## Examples

```
## Not run:
data(heid)
heid = aggregate(heid$RT, list(heid$Word, heid$BaseFrequency), mean)
colnames(heid) = c("Word", "BaseFrequency", "MeanRT")

## End(Not run)
```

---

herdan.fnc

*Herdan's C*

---

## Description

This function calculates Herdan's constant C.

## Usage

```
herdan.fnc(text, chunks)
```

## Arguments

**text** A vector of strings representing a text.

**chunks** A vector of chunk sizes for which Herdan's C is required. Duplicate chunk sizes are not allowed, and the number of chunks should be at least 2.

## Value

A list with components

**growth** A data frame with token and type counts for the requested chunk sizes.

**C** Herdan's C.

## Author(s)

R. H. Baayen

## References

Herdan, G. (1960) *Type-Token Mathematics*, The Hague: Mouton.

Herdan, G. (1964) *Quantitative Linguistics*, London: Butterworths.

## See Also

See Also [growth.fnc](#).

**Examples**

```
## Not run:
data(alice)
herdan.fnc(alice, cumsum(rep(floor(length(alice)/40), 40)))

## End(Not run)
```

---

imaging

*fMRI Filtered Signal and Priming Scores for Brain-Damaged Patients*

---

**Description**

Filtered fMRI signal at the most significant voxel and average priming scores for brain-damaged patients, in a study addressing the extent to which phonological and semantic processes recruit the same brain areas.

**Usage**

```
data(imaging)
```

**Format**

A data frame with 35 observations on the following 3 variables.

**Condition** a factor with levels *irregulars* (the morphological condition involving priming using inflected forms of irregular English verbs, e.g., 'began'-'begin') and *semantics* (priming with semantically related words such as 'card' and 'paper').

**BehavioralScore** a numeric vector for the average priming scores.

**FilteredSignal** a numeric vector for the intensity of the filtered fMRI signal at the most significant voxel.

**Details**

Location of data points reconstructed from the pixel map of Figure 2b of Tyler et al. 2005.

**Source**

Tyler, L.K., Marslen-Wilson, W.D. and Stamatakis, E.A. (2005) Differentiating lexical form, meaning, and structure in the neural language system, *PNAS*, 102, 8375-8380.

**Examples**

```
## Not run:
data(imaging)

imaging.lm = lm(FilteredSignal~BehavioralScore*Condition, data=imaging)
summary(imaging.lm)
```

```

plot(imaging$BehavioralScore, imaging$FilteredSignal, type = "n",
     xlim = c(-30, 40), ylim = c(0, 80))
semantics = imaging[imaging$Condition == "semantics",]
irregulars = imaging[imaging$Condition == "irregulars",]
points(semantics$BehavioralScore, semantics$FilteredSignal, col = "black")
points(irregulars$BehavioralScore, irregulars$FilteredSignal, col = "darkgrey")
abline(lm(FilteredSignal ~ BehavioralScore, data = semantics), col = 'black')
abline(lm(FilteredSignal ~ BehavioralScore, data = irregulars),
       col = 'darkgrey')

# model criticism

plot(imaging.lm)
outliers = c(1, 19) # given Cook's distance, or perhaps only
outliers = 1       # the outlier in the semantics subset
imaging.lm = lm(FilteredSignal ~ BehavioralScore * Condition,
               data = imaging[-outliers, ])
summary(imaging.lm)

## End(Not run)

```

---

```
implementInteractions.fnc
```

*implement interactions in the model matrix*

---

## Description

given a model matrix with main effects only, add interactions

## Usage

```
implementInteractions.fnc(m)
```

## Arguments

`m` a (model) matrix (rows observations, columns predictors)

## Details

not intended for independent use

## Value

an updated (model) matrix

## Note

not intended for independent use

**Author(s)**

R. H. Baayen

**See Also**

[plotLMER.fnc](#)

**Examples**

```
## Not run: not intended for independent use
```

---

item.fnc

*Function for by-item regression used by simulateRegression.fnc*

---

**Description**

This function carries out a by-item regression for the simulated data sets generated in `simulateRegression.fnc`. It is not designed to be used independently.

**Usage**

```
item.fnc(data)
```

**Arguments**

`data` A data frame as produced by `make.reg.fnc()`.

**Value**

A model fitted with `lm()`.

**Author(s)**

R. H. Baayen

**See Also**

See Also [simulateRegression.fnc](#) and [make.reg.fnc](#).

**Examples**

```
## Not run:
dat = make.reg.fnc()
dat.lm = item.fnc(dat)
summary(dat.lm)

## End(Not run)
```

---

items.quasif.fnc      *By-item anova for simulated data for quasi-F analysis*

---

### Description

By-item anova for simulated data set as created within simulateQuasif.fnc. Not intended for independent use. Depends on the packages MASS, coda and lme4.

### Usage

```
items.quasif.fnc(dat)
```

### Arguments

dat	Simulated data set with Subjects, Item, and SOA treatment, as created within simulateQuasif.fnc, or the quasif dataset.
-----	---

### Value

A list with components

p	p-value of F-test for SOA.
data	the input data.
model	the fitted model.

### Author(s)

R. H. Baayen

### See Also

See also [simulateQuasif.fnc](#).

### Examples

```
## Not run:  
data(quasif)  
items.quasif.fnc(quasif)  
  
## End(Not run)
```

---

lags.fnc	<i>Calculate vector at specified lag</i>
----------	--

---

### Description

This function calculates for a given dependent variable the value of that variable at lag timesteps earlier in the time series of an experiment.

### Usage

```
lags.fnc(dat, time="Trial", group = "Subject", depvar = "RT", lag=1)
```

### Arguments

dat	A data frame with (minimally) a grouping factor, an time index for successive trails/events, and a behavioral measure
group	A grouping factor such as Subject
time	A sequential time index measure such as Trial number in an experimental list
depvar	The dependent variable, usually a chronometric measure such as RT
lag	The lag for which previous values are to be extracted

### Value

A vector with the values of the dependent variable at the specified lag. The by-group mean is substituted for the first lag timestep(s), for which there is/are no preceding value(s) for the dependent variable.

### Author(s)

R. H. Baayen

### See Also

[acf.fnc](#)

### Examples

```
## Not run:
dfr = data.frame(Subject=c(rep("a", 5), rep("b", 5)),
                 Trial = c(rep(1:5,2)),
                 RT = rnorm(10, 500, 40))
dfr$prevRT = lag.fnc(dfr, time="Trial", group="Subject", depvar="RT")
dfr

## End(Not run)
```

---

latinsquare

*Simulated Latin Square data set with subjects and items*

---

### Description

Simulated lexical decision latencies with SOA as treatment, using a Latin Square design with subjects and items, as available in Raaijmakers et al. (1999).

### Usage

```
data(latinsquare)
```

### Format

A data frame with 144 observations on the following 6 variables.

Group a factor with levels G1, G2 and G3, for groups of subjects

Subject a factor with subjects labelled S1, ... S12.

Word a factor with words labelled W1 ... W12.

RT a numeric vector for reaction times.

SOA a factor with levels long, medium, and short.

List a factor with levels L1, L2, and L3 for lists of words.

### Source

Raaijmakers, J.G.W., Schrijnemakers, J.M.C. & Gremmen, F. (1999) How to deal with "The language as fixed effect fallacy": common misconceptions and alternative solutions, *Journal of Memory and Language*, 41, 416-426.

### Examples

```
## Not run:
data(latinsquare)
library(lme4)
latinsquare.with =
  simulateLatinsquare.fnc(latinsquare, nruns = 1000, with = TRUE)
latinsquare.without =
  simulateLatinsquare.fnc(latinsquare, nruns = 1000, with = FALSE)
latinsquare.with$alpha05
latinsquare.without$alpha05

## End(Not run)
```

lexdec

*Lexical decision latencies for 79 English nouns***Description**

Lexical decision latencies elicited from 21 subjects for 79 English concrete nouns, with variables linked to subject or word.

**Usage**

```
data(lexdec)
```

**Format**

A data frame with 1659 observations on the following 28 variables.

**Subject** a factor for the subjects.

**RT** a numeric vector for logarithmically transformed reaction times.

**Trial** a numeric vector for the rank of the trial in the experimental list.

**Sex** a factor with levels F (female) and M (male).

**NativeLanguage** a factor with levels English and Other, distinguishing between native and non-native speakers of English.

**Correct** a factor with levels correct and incorrect coding whether the word was correctly responded to as a word rather than a nonword.

**PrevType** a factor with levels nonword and word coding whether the item presented at the preceding trial was a word or a nonword.

**PrevCorrect** a factor with levels correct and incorrect coding whether the preceding item elicited a correct response.

**Word** a factor with 79 words as levels.

**Frequency** a numeric vector with logarithmically transformed lemma frequencies as available in the CELEX lexical database.

**FamilySize** a numeric vector with the log-transformed count of a word's morphological family members.

**SynsetCount** a numeric vector with the log-transformed count of synonym sets in WordNet in which the word is listed.

**Length** a numeric vector for the word's length in letters.

**Class** a factor for the semantic category of the word's referent, with levels animal and plant.

**FreqSingular** a numeric vector with the frequency in CELEX of the singular form.

**FreqPlural** a numeric vector with the frequency in CELEX of the plural form.

**DerivEntropy** Shannon's entropy calculated over the frequency distribution of a word's family members.

**Complex** a factor coding morphological complexity with levels complex and simplex.

rInfl a numeric vector for the log of the ratio of the singular to the plural frequency.

meanRT a numeric vector for the by-item mean reaction time averaged over subjects.

SubjFreq a numeric vector for the by-item mean subjective frequency estimate averaged over subjects.

meanSize a numeric vector for the by-item mean size rating averaged over subjects.

meanWeight a numeric vector for the by-item mean weight rating averaged over subjects.

BNCw a numeric vector with the logarithmically transformed frequency in the written part of the British National Corpus.

BNCc a numeric vector with the logarithmically transformed frequency in the context-governed part of the British National Corpus.

BNCd a numeric vector with the logarithmically transformed frequency in the demographic part of the British National Corpus.

BNCcRatio a numeric vector with the log of the ratio of the (absolute) frequencies in the context-governed and written parts of the British National Corpus, normalized for the differences in corpus size.

BNCdRatio a numeric vector with the log of the ratio of the (absolute) frequencies in the demographic and written parts of the British National Corpus, normalized for the differences in corpus size.

## Source

Data collected with Jen Hay, University of Canterbury, Christchurch, New Zealand, 2004.

## Examples

```
## Not run:
data(lexdec)
require(lme4, quietly=TRUE)

lexdec.lmer = lmer(RT ~ 1 + Correct + Trial + PrevType * meanWeight +
Frequency + NativeLanguage * Length + (1|Subject) + (1|Word),
data = lexdec)
pvals.fnc(lexdec.lmer)$summary

# random slopes

lexdec.lmerA = lmer(RT ~ 1 + Correct + Trial + PrevType * meanWeight +
Frequency + NativeLanguage * Length + (Trial|Subject) + (1|Word),
data = lexdec)
anova(lexdec.lmer, lexdec.lmerA)

lexdec.lmerB = lmer(RT ~ 1 + Correct + Trial + PrevType * meanWeight +
Frequency + NativeLanguage * Length + (Trial|Subject) +
(Length|Subject) + (1|Word), data = lexdec)
anova(lexdec.lmerA, lexdec.lmerB)

# model criticism
```

```

qqnorm(resid(lexdec.lmerB))

lexdec.lmerC = lmer(RT ~ 1 + Correct + Trial + PrevType * meanWeight +
Frequency + NativeLanguage * Length +
(Trial|Subject) + (Length|Subject) + (1|Word),
data = lexdec[abs(scale(resid(lexdec.lmerB)))<2,])

qqnorm(resid(lexdec.lmerC))
# for models with random correlation parameters, pvals.fnc does not work
# as mcmcSamp() (now in lme4) is currently under reconstruction
# pvals.fnc(lexdec.lmerC)$summary (currently broken)

## End(Not run)

```

lexicalMeasures

*Lexical measures for 2233 English monomorphemic words***Description**

Lexical distributional measures for 2233 English monomorphemic words. This dataset provides a subset of the data available in the dataset english.

**Usage**

```
data(lexicalMeasures)
```

**Format**

A data frame with 2233 observations on the following 24 variables.

Word a factor with 2284 words.

CeLS numeric vector with log-transformed lemma frequency in the CELEX lexical database.

Fdif numeric vector with the logged ratio of written frequency (CELEX) to spoken frequency (British National Corpus).

Vf numeric vector with log morphological family size.

Dent numeric vector with derivational entropy.

Ient numeric vector with inflectional entropy.

NsyS numeric vector with the log-transformed count of synonym sets in WordNet in which the word is listed.

NsyC numeric vector with the log-transformed count of synonym sets in WordNet in which the word is listed as part of a compound.

Len numeric vector with length of the word in letters.

Ncou numeric vector with orthographic neighborhood density.

Bigr numeric vector with mean log bigram frequency.

InBi numeric vector with log frequency of initial diphone.

spelV numeric vector with type count of orthographic neighbors.

spelN numeric vector with token count of orthographic neighbors.

phonV numeric vector with type count of phonological neighbors.

phonN numeric vector with token count of phonological neighbors.

friendsV numeric vector with type counts of consistent words.

friendsN numeric vector with token counts of consistent words.

ffV numeric vector with type count of forward inconsistent words.

ffN numeric vector with token count of forward inconsistent words.

fbV numeric vector with type count of backward inconsistent words.

fbN numeric vector with token count of backward inconsistent words

ffNonzero a numeric vector with the count of forward inconsistent words with nonzero frequency.

NVratio a numeric vector with the logarithmically transformed ratio of the noun and verb frequencies.

## References

Baayen, R.H., Feldman, L. and Schreuder, R. (2006) Morphological influences on the recognition of monosyllabic monomorphemic words, *Journal of Memory and Language*, 53, 496-512.

## Examples

```
## Not run:
data(lexicalMeasures)
data(lexicalMeasuresDist)

library(rms)
library(cluster)
plot(varclus(as.matrix(lexicalMeasures[,-1])))

lexicalMeasures.cor = cor(lexicalMeasures[,-1], method = "spearman")^2
lexicalMeasures.dist = dist(lexicalMeasures.cor)
pltree(diana(lexicalMeasures.dist))

data(lexicalMeasuresClasses)
x = data.frame(measure = rownames(lexicalMeasures.cor),
cluster = cutree(diana(lexicalMeasures.dist), 5),
class = lexicalMeasuresClasses$Class)
x = x[order(x$cluster), ]
x

## End(Not run)
```

---

lexicalMeasuresClasses

*Classification of lexical measures*


---

### Description

A data frame labelling the lexical measures in the dataset lexicalMeasures as measures of form or meaning.

### Usage

```
data(lexicalMeasuresClasses)
```

### Format

A data frame with 23 observations on the following 3 variables.

Variable a factor with as levels the measures:

Bigr Mean Bigram Frequency.

CeIS CELEX Frequency.

Dent Derivational Entropy.

fbN Token Count of Backward Inconsistent Words.

fbV Type Count of Backward Inconsistent Words.

Fdif Ratio of Frequencies in Written and Spoken English.

ffN Token Count of Forward Inconsistent Words.

ffNonzero Type Count of Forward Inconsistent Words with Nonzero Frequency.

ffV Type Count of Forward Inconsistent Words

friendsN Token Count of Consistent Words.

friendsV Type Count of Consistent Words.

Ient Inflectional Entropy

InBi Initial Bigram Frequency

Len Length in Letters

Ncou Orthographic Neighborhood Density

NsyC Number of Complex Synsets

NsyS Number of Simplex Synsets

NVratio Ratio of Noun and Verb Frequencies

phonN Token Count of Phonological Neighbors.

phonV Type Count of Phonological Neighbors.

spe1N Token Count of Orthographic Neighbors.

spe1V Type Count of Orthographic Neighbors.

Vf Morphological Family Size.

Class a factor with levels Form and Meaning.

Explanation a factor with glosses for the variables.

## References

Baayen, R.H., Feldman, L. and Schreuder, R. (2006) Morphological influences on the recognition of monosyllabic monomorphemic words, *Journal of Memory and Language*, 53, 496-512.

## Examples

```
## Not run:
library(cluster)
data(lexicalMeasures)
data(lexicalMeasuresClasses)

lexicalMeasures.cor = cor(lexicalMeasures[, -1], method = "spearman")^2
x = data.frame(measure = rownames(lexicalMeasures.cor),
               cluster = cutree(diana(dist(lexicalMeasures.cor)), 5),
               class = lexicalMeasuresClasses$class)
x = x[order(x$cluster), ]
x

## End(Not run)
```

---

lmerPlotInt.fnc	<i>Plot the interaction of two linear numeric predictors in a model fitted with lmer</i>
-----------------	--

---

## Description

Visualization of an interaction in a model fitted with lmer of two numeric predictors.

## Usage

```
lmerPlotInt.fnc(lmermodel, xname, yname, intxynname, qntls = seq(0, 1, by = 0.1), view = 30, addStdError = FALSE)
```

## Arguments

lmermodel	an lmer model object
xname	name (character string) of first numeric predictor
yname	name (character string) of second numeric predictor
intxynname	name (character string) of the interaction in the lmer summary
qntls	vector of values to be shown for the second numeric predictor, defaults to deciles
view	specifies the viewing parameter theta for the perspective plot
addStdError	add noise with the standard deviation of the residual error in the lmer model to the plot
ndigits	number of digits to show for the second numeric predictor
nlev	number of levels for the contour plot
which	choices are "matplot" (default), "contour", "persp", "image", and "all", in which case a 2 by 2 panel is shown with all four plots

shadow	the amount of shade for the perspective plot
colour	the color used for the perspective plot, defaults to "lightblue"
fun	for matplotlib displays, a function for transforming the predicted response
ylabel	string, to be added to the Y-axis as y label
...	other arguments

**Value**

A plot is shown on the graphics device.

**Warning**

This function should not be used to plot interactions when one of the predictors also has quadratic or higher terms in the model.

**Author(s)**

R. H. Baayen

**Examples**

```
## Not run:
require(lme4, quietly=TRUE)
lexdec.lmer = lmer(RT~BNCw*Frequency+(1|Subject)+(1|Word), data=lexdec)
lmerPlotInt.fnc(lexdec.lmer, "BNCw", "Frequency", "BNCw:Frequency", which="matplot")

## End(Not run)
```

---

make.reg.fnc

*Make a simulated data set with regression design*

---

**Description**

This convenience function creates a regression data set with subjects, items, and three numerical predictors, and optionally an effect of learning or fatigue. This function is called by simulateRegression.fnc, and is not intended for independent use.

**Usage**

```
make.reg.fnc(nsubj = 10, nitem = 20, beta = c(400, 2, 6, 4),
  learn = FALSE, learnRate = 10, stdevItem = 40, stdevSubj = 80,
  stdevError = 50)
```

**Arguments**

nsubj	Number of subjects (random effect) required.
nitem	Number of items (random effect) required.
beta	A numeric vector with four beta weights: one for the intercept and one for each of three predictors.
learn	A logical variable, if TRUE, a learning or fatigue effect will be implemented, as specified by learnRate.
learnRate	A number indicating learning (if negative) or fatigue (if positive).
stdevItem	A number specifying the standard deviation of the Item random effect.
stdevSubj	A number specifying the standard deviation of the Subject random effect.
stdevError	A number specifying the standard deviation of the Residual Error.

**Value**

A data frame with intercept, predictors labelled X, Y and Z, Item, Subject, the simulated random effects for Item and Subject, the residual errors, and the simulated RTs.

**Author(s)**

R. H. Baayen

**See Also**

[simulateRegression.fnc](#)

**Examples**

```
## Not run:
simdat = make.reg.fnc()
library(lme4, keep.source = FALSE)
simdat.lmer = lmer(RT ~ X + Y + Z + (1|Subject) + (1|Item),
  data = simdat)
simdat.lmer

simdat = make.reg.fnc(learn = TRUE)
simdat.lmer = lmer(RT ~ X + Y + Z + Trial + (1|Subject) + (1|Item),
  data = simdat)
simdat.lmer

## End(Not run)
```

---

makeDefaultMatrix.fnc *Create model matrix with main effects only*

---

**Description**

Creates a model matrix with main effects only

**Usage**

```
makeDefaultMatrix.fnc(model, n = 100, conditioningPred = "", conditioningValue = NULL, control = NA)
```

**Arguments**

model	A model fit with lmer
n	integer specifying number of points to be plotted on X-axis
conditioningPred	name of predictor entering into interaction
conditioningValue	vector of values (numeric or factor level names) to be shown for interaction
control	a two-element list (predictor, value) specifying an additional predictor to be fixed to the given value in a partial effect plot. May be useful for hand-made plots for three-way interactions.

**Details**

not intended for independent use

**Value**

a (model) matrix

**Note**

not intended for independent use

**Author(s)**

R. H. Baayen

**See Also**

See Also as [plotLMER.fnc](#)

**Examples**

```
## Not run: not intended for independent use
```

---

makeSplineData.fnc     *generate simulated data set with nonlinear function*

---

**Description**

creates a data set with  $Y \sim 30 + \cos(X)$  for 10 subjects, to compare restricted cubic spline in lmer with the spline of ols.

**Usage**

```
makeSplineData.fnc(intr=0)
```

**Arguments**

`intr`            integer denoting type of data set: with 0 a data set with simple spline is made, with 1 a data set with a parallel interaction, and with 2 a data set with a crossed interaction.

**Details**

Requires rms package to be attached.

**Value**

A data frame with as values:

<code>y</code>	<code>y = 30 + cos(X)</code>
<code>X</code>	ranges from 2 to 8.28
<code>Subject</code>	random-effects factor with 10 levels
<code>Ranef</code>	subjects-specific changes to intercept
<code>Error</code>	by-observation noise
<code>Y</code>	the dependent variable, <code>y+Ranef+Error</code>

**Note**

intended for illustration only

**Author(s)**

R. H. Baayen

**See Also**

See Also as [plotLMER.fnc](#)

**Examples**

```
## Not run:
require("rms", quietly=TRUE, character=TRUE)
dfr = makeSplineData.fnc()
table(dfr$Subject)
xylowess.fnc(Y ~ X | Subject, data = dfr)

dfr.lmer = lmer(Y ~ rcs(X, 5) + (1|Subject), data = dfr)
dfr$fittedLMER = as.vector(dfr.lmer@X %*% fixef(dfr.lmer))

dfr.dd = datadist(dfr)
options(datadist='dfr.dd')
dfr.ols = ols(Y~Subject+rcs(X), data=dfr, x=T, y=T)
dfr$fittedOLS = fitted(dfr.ols)

# we plot the lmer() fit in blue, the ols() fit in red (both adjusted for
# subject S1), and plot the underlying model in green
plot(dfr[dfr$Subject=="S1",]$X, dfr[dfr$Subject=="S1",]$fittedLMER +
  ranef(dfr.lmer)[[1]]["S1",], type="l", col="blue",
  ylim = range(dfr$y + ranef(dfr.lmer)[[1]]["S1",],
  dfr[dfr$Subject == "S1",]$fittedLMER,
  dfr[dfr$Subject == "S1",]$fittedOLS), xlab="X", ylab="Y")
lines(dfr[dfr$Subject=="S1",]$X, dfr[dfr$Subject=="S1",]$fittedOLS, col="red")
lines(dfr[dfr$Subject=="S1",]$X, dfr[dfr$Subject=="S1",]$y+ranef(dfr.lmer)[[1]]["S1",],
  col="green")
legend(2,29,c("30+cos(x)", "lmer (S1)", "ols (S1)"), lty=rep(1,3),
  col=c("green", "blue", "red"))

## End(Not run)
```

---

moby

*Moby Dick*


---

**Description**

The text of H. Melville's 'Moby Dick', with punctuation marks removed.

**Usage**

```
data(alice)
```

**Format**

A character vector with 215994 words.

**Source**

The project Gutenberg at [http://www.gutenberg.org/wiki/Main\\_Page](http://www.gutenberg.org/wiki/Main_Page)

**Examples**

```
## Not run:  
  data(moby)  
  moby[1:2]  
  
## End(Not run)
```

---

mvrnormplot.fnc

*Scatterplot of bivariate standard normal distribution*

---

**Description**

This function produces a scatterplot for a bivariate standard normal distribution with least squares regression line.

**Usage**

```
mvrnormplot.fnc(r, n, limits)
```

**Arguments**

r	The correlation, defaults to 0.9.
n	Number of simulated data points, defaults to 100.
limits	Optional range for the axes

**Value**

A scatterplot with ordinary least squares regression line is shown on the graphics device, with sample estimate of r added at the top of the plot.

**Author(s)**

R. H. Baayen

**See Also**

mvrnorm (MASS package)

**Examples**

```
## Not run:  
mvrnormplot.fnc(r=0.9, n=100)  
  
## End(Not run)
```

---

`nesscg`*Frequency spectrum for -ness in the demographic BNC*

---

**Description**

Frequency (m) and frequency of frequency (Vm) for string types with the suffix *-ness* in the context-governed subcorpus of the British National Corpus sampling spoken British English.

**Usage**

```
data(nesscg)
```

**Format**

A data frame with 37 observations on the following 2 variables.

m a numeric vector with word frequencies.

Vm a numeric vector with the frequencies of word frequencies.

**Source**

The British National Corpus, see <http://www.natcorp.ox.ac.uk/>

**Examples**

```
## Not run:  
data(nesscg)  
library(zipfR)  
nesscg.spc = spc(m=nesscg$m, Vm = nesscg$Vm)  
plot(nesscg.spc)  
  
## End(Not run)
```

---

`nessdemog`*Frequency spectrum for -ness in the context-governed BNC*

---

**Description**

Frequency (m) and frequency of frequency (Vm) for string types with the suffix *-ness* in the demographic subcorpus of the British National Corpus sampling spoken British English.

**Usage**

```
data(nessdemog)
```

**Format**

A data frame with 15 observations on the following 2 variables.

*m* a numeric vector with word frequencies.

*Vm* a numeric vector with the frequencies of word frequencies.

**Source**

The British National Corpus, see <http://www.natcorp.ox.ac.uk/>

**Examples**

```
data(nessdemog)
library(zipfR)
nessdemog.spc = spc(m=nessdemog$m, Vm = nessdemog$Vm)
plot(nessdemog.spc)
```

---

nessw

*Frequency spectrum for -ness in the written BNC*

---

**Description**

Frequency (*m*) and frequency of frequency (*Vm*) for string types with the suffix *-ness* in the sub-corpus of the British National Corpus sampling written British English.

**Usage**

```
data(nessw)
```

**Format**

A data frame with 189 observations on the following 2 variables.

*m* a numeric vector with word frequencies.

*Vm* a numeric vector with the frequencies of word frequencies.

**Source**

The British National Corpus, see <http://www.natcorp.ox.ac.uk/>

**Examples**

```
## Not run:
data(nessw)
library(zipfR)
nessw.spc = spc(m=nessw$m, Vm = nessw$Vm)
plot(nessw.spc)

## End(Not run)
```

---

 oldFrench

*Frequencies of tag trigrams in Old French texts*


---

**Description**

Frequencies of 35 morphosyntactic tag trigrams in 343 Old French texts.

**Usage**

```
data(oldFrench)
```

**Format**

A data frame with the frequencies of 35 tag trigrams (columns) for 343 Old French texts (rows) in the Nouveau Corpus d'Amsterdam. See `oldFrenchMeta` for details on the texts (and manuscript versions).

**Source**

Data from Nouveau Corpus d'Amsterdam, <http://www.uni-stuttgart.de/lingrom/stein/corpus/>.

**References**

Ernestus, M., van Mulken, M. and Baayen, R. H. (2007) De syntax van Oud-Franse ridders en heiligen in ruimte en tijd To appear in *Onze Taal*.

**Examples**

```
data(oldFrench)
data(oldFrenchMeta)

oldFrench.ca = corres.fnc(oldFrench)

plot(oldFrench.ca, rlabels = oldFrenchMeta$Genre,
      rcol = as.numeric(oldFrenchMeta$Genre), rcex = 0.5,
      extreme = 0.1, ccol = "blue")
```

---

 oldFrenchMeta

*Meta data for the oldFrench data*


---

**Description**

Meta data for the `oldFrench` data, a matrix of frequencies for texts (rows) by tag trigrams (columns). The meta data provide information on the texts, manuscript variants, their authors, their region and approximate date of origin, their general topic, and their genre.

**Usage**

```
data(oldFrenchMeta)
```

**Format**

A data frame with 342 observations on the following 7 variables.

`Textlabels` a factor with texts coded as follows:

Abe J. de Meun, Traduction de la premiere epitre de P. Abelard, 1–821  
 Hy11 Anon, La vie de saint Hylaïre  
 Art J. de Meun, L'art de chevalerie  
 Bar Anon, L'histoire de Barlaam et Josaphat  
 Cathy Anon, La vie de sainte Catherine d'Alexandrie  
 Hy12 Anon, La vie de saint Hylaïre  
 Ch1 Chretien de Troyes, Le Chevalier au lion  
 Ch2 Chretien de Troyes, Le chevalier au lion  
 Clari Robert de Clari, La conquete de Constantinople  
 Marie Rutebeuf, Sainte Marie l'Egyptienne  
 Fab4c Anon, Fabliau nr 4 ms C  
 Fab4e Anon, Fabliau nr 4 ms E  
 Fab4f Anon, Fabliau 4f  
 Faba Anon, Fabliaux nrs 1,2,4,23 et 29 du ms A  
 Fabb Anon, Fabliaux nrs 2 et 4 du ms B  
 Fabd.COD Anon, Fabliaux nrs 2 et 4 du ms D  
 Hy13 Anon, La vie de saint Hylaïre  
 Jacobi Pierre de Beauvais, The Liber Sancti Jacobi  
 Louis J. de Joinville, La vie de saint Louis  
 Cathy1 Anon, La passion saynte Katherine  
 Lancelot Anon, Lancelot do Lac, p. 1.1–20.13  
 Merlin1 Merlin, Robert de Boron  
 Marga Anon, La vie de Sainte Marguerite de Wace  
 Martin Anon, Leben und Wunderthaten des heiligen Martin  
 Merlin2 Anon, Merlin, p.1–29 (ms. Huth)  
 RoseA J. de Meun, Le Roman de la Rose  
 Arthur Anon, La mort le roi Artu, par.1–35  
 NimAf Anon, charroi de Nimes, ms. A, fragment  
 NimB1 Anon, Le charroi de Nimes, ms B1  
 NimB2 Anon, Le charroi de Nimes, ms B2  
 Nouvel Jacquemart Gielee, Renart le Nouvel  
 Jehan Anon, La vie de saint Jehan Bouche d'Or  
 Per0 Chretien de Troyes, Perceval  
 perL Chretien de Troyes, Perceval  
 PerQ Chretien de Troyes, Perceval

PerS Chretien de Troyes, Perceval  
 PerU Chretien de Troyes, Perceval  
 Queste Anon, La queste del saint Graal, p.1.1–41.17  
 Rob Anon, Robert le Diable, v.1–808  
 RomB Anon, Le Roman de Renart, br.VI, ms B  
 RomD Anon, Roman de Renard, br.VI, ms D  
 RomL Anon, Roman de Renard, br. VI, ms L  
 RomO Anon, Le Roman de Renart, br. VI, ms O  
 RoseB Guillaume de Lorris, Le roman de la rose  
 Sapi Anon, Sermo de sapientia, dans: Li dialogue Gregoire lo pape  
 Troi Anon, Le roman de Troie en prose, par.1–19  
 Conqueste Josfroi de Vileharduyn, La conquete de Costentinoble  
 YvA Chretien de Troyes, Le chevalier au lion, v.1–1000  
 YvP Chretien de Troyes, Le chevalier au lion, v.1–1000  
 YvS Chretien de Troyes, Le chevalier au lion, v.1–1000  
 YvV Chretien de Troyes, Le chevalier au lion, v 1–1000

Codes a factor with manuscript variants, indicated by extensions to the text codes.

Author a factor with levels Anon, ChretienDeTroyes, GuillaumeDeLorris, Joinville, Meun, NouvelRenart, PierreDeBeauvais, RobertDeBoron, RobertDeClari, RobertLeDiable, Rutebeuf, and Villeharduyn.

Topic a factor with levels Knight, Other, and Saint.

Genre a factor with levels poetry and prose.

Region a factor with levels R1 (Picardie), R2 (Champenois), and R3 (Nievre-Allier).

Year a numeric vector indicating approximate year of origin.

## Source

Data from Nouveau Corpus d'Amsterdam, <http://www.uni-stuttgart.de/lingrom/stein/corpus/>.

## References

Ernestus, M., van Mulken, M. and Baayen, R. H. (2007) De syntax van Oud-Franse ridders en heiligen in ruimte en tijd To appear in *Onze Taal*.

## Examples

```
## Not run:
data(oldFrench)
data(oldFrenchMeta)

oldFrench.ca = corres.fnc(oldFrench)

plot(oldFrench.ca, rlabels = oldFrenchMeta$Genre,
      rcol = as.numeric(oldFrenchMeta$Genre), rcex = 0.5,
      extreme = 0.1, ccol = "blue")

## End(Not run)
```

oz

*The Wonderful Wizard of Oz*

---

**Description**

The text of L. F. Baum's 'The Wonderful Wizard of Oz', with punctuation marks removed.

**Usage**

```
data(alice)
```

**Format**

A character vector with 39513 words.

**Source**

The project Gutenberg at [http://www.gutenberg.org/wiki/Main\\_Page](http://www.gutenberg.org/wiki/Main_Page)

**Examples**

```
## Not run:  
data(oz)  
oz[1:5]  
  
## End(Not run)
```

---

pairscor.fnc*Scatterplot matrix with correlations*

---

**Description**

A matrix of scatterplots is produced with Pearson and Spearman correlations in the lower triangle. By default, smoothers are added to panels in the upper triangle, and histograms are added to the panels on the diagonal.

**Usage**

```
pairscor.fnc(data, hist = TRUE, smooth = TRUE,  
  cex.points = 1, col.points = "darkgrey")
```

**Arguments**

data	a data frame or matrix with numeric vectors.
hist	a logical indicating whether panels on the diagonal should contain a histogram.
smooth	a logical indicating whether panels in the upper triangle should have a smoother added.
cex.points	a number indicating the size of the points in the panels in the upper triangle, available only when smoothers are added.
col.points	a number or string indicating the color of the points in the panels in the upper triangle, available only when smoothers are added.

**Author(s)**

R. Harald Baayen

**See Also**

See Also [pairs](#) and [panel.smooth](#).

**Examples**

```
## Not run:
data(lexicalMeasures)
pairscore.fnc(lexicalMeasures[,c("CeIS", "Vf", "Ient", "NsyS", "Ncou")])

## End(Not run)
```

---

parsePredName.fnc      *parse character string specifying restricted cubic spline*

---

**Description**

parse character string specifying restricted cubic spline into simple predictor name and number of knots

**Usage**

```
parsePredName.fnc(name)
```

**Arguments**

name	character string for predictor, e.g. rcs(X, 3)
------	--

**Details**

not intended for independent use

**Value**

a list with components

baseName        character string denoting simple predictor name (X)

knots            integer specifying number of knots

**Note**

not intended for independent use

**Author(s)**

R. H. Baayen

**See Also**

See Also as [plotLMER.fnc](#)

**Examples**

```
## Not run: not intended for independent use
```

---

periphrasticDo	<i>The development of periphrastic do in English</i>
----------------	--

---

**Description**

The development of periphrastic *do* in English: Ellegard's counts for the use of *do* across four sentence types in 11 consecutive time periods between 1390 and 1710.

**Usage**

```
data(periphrasticDo)
```

**Format**

A data frame with 44 observations on the following 5 variables.

begin a numeric vector with beginnings of the time periods used by Ellegard.

end a numeric vector with ends of these time periods.

type a factor for sentence type, with levels affdecl (affirmative declarative), affquest (affirmative question), negdecl (negative declarative) and negquest (negative question).

do a numeric vector with the count of sentences with *do*.

other a numeric vector with the count of sentences without *do*.

**Source**

Ellegard, A. (1953) *The auxiliary do: The establishment and regulation of its use in English*, Stockholm: Almqvist & Wiksell.

**References**

Vulanovic, R. and Baayen, R. H. (2006) Fitting the development of periphrastic do in all sentence types, in Grzybek, P. and Koehler, R. (eds.), *Festschrift fuer Gabriel Altmann*, Berlin: Walter de Gruyter, p. 679-688.

**Examples**

```
## Not run:
data(periphrasticDo)

# add midpoints of time periods

periphrasticDo$year = periphrasticDo$begin +
  (periphrasticDo$end-periphrasticDo$begin)/2

# and add an indicator variable distinguishing the first three time periods
# from the others

periphrasticDo$Indicator = rep(c(rep(0, 3), rep(1, 8)), 4)

# fit a logistic regression model

periphrasticDo.glm = glm(cbind(do, other) ~
  (year + I(year^2) + I(year^3)) * type + Indicator * type +
  Indicator * year, data = periphrasticDo, family = "binomial")

anova(periphrasticDo.glm, test = "F")

# visualization of data and model predictions

periphrasticDo$predict = predict(periphrasticDo.glm, type = "response")
par(mfrow=c(2, 2))
for (i in 1:nlevels(periphrasticDo$type)) {
  subset = periphrasticDo[periphrasticDo$type ==
    levels(periphrasticDo$type)[i], ]
  plot(subset$year,
    subset$do/(subset$do + subset$other),
    type = "p", ylab = "proportion", xlab = "year",
    ylim = c(0, 1), xlim = c(1400, 1700))
  mtext(levels(periphrasticDo$type)[i], line = 2)
  lines(subset$year, subset$predict, lty = 1)
}
par(mfrow=c(1, 1))

## End(Not run)
```

---

 phylogeny

*Phylogenetic relations between Papuan and Oceanic languages*


---

**Description**

Phylogenetic relations between Papuan and Oceanic languages: 127 grammatical traits (absent/present) for 31 languages.

**Usage**

data(phylogeny)

**Format**

A data frame with 31 observations on the following 127 variables.

Language a factor for 31 languages: Anem, Ata, Bali, Banoni, Bilua, Buin, Gapapaiwa, Kairiru, Kaulong, Kilivila, Kokota, Kol, Kuot, Lavukaleve, Mali, Motuna, Nalik, Nasioi, Rotokas, Roviana, Savosavo, Siar, Sisiqa, Sudest, Sulka, Taiof, Takia, Touo, Tungag, Yabem and Yeli\_Dnye.

Family a factor with levels Oceanic and Papuan.

Frics a numeric vector, 1: presence, 0: absence

PrenasalizedStops a numeric vector, 1: presence, 0: absence

PhonDistBetweenLAndR a numeric vector, 1: presence, 0: absence

PhonVelarFricOrGlide a numeric vector, 1: presence, 0: absence

PhonVoicingContrAmongStops a numeric vector, 1: presence, 0: absence

PhonConsLength a numeric vector, 1: presence, 0: absence

PhonVowelLength a numeric vector, 1: presence, 0: absence

ContrPhonTypesForVowels a numeric vector, 1: presence, 0: absence

PhonStress a numeric vector, 1: presence, 0: absence

WordFinalConss a numeric vector, 1: presence, 0: absence

ConsClusters a numeric vector, 1: presence, 0: absence

DefOrSpecArt a numeric vector, 1: presence, 0: absence

IndefOrNonSpecArt a numeric vector, 1: presence, 0: absence

ArticleNounOrder a numeric vector, 1: presence, 0: absence

NounInitNps a numeric vector, 1: presence, 0: absence

InclExclDist a numeric vector, 1: presence, 0: absence

PronNum a numeric vector, 1: presence, 0: absence

PronRelationship a numeric vector, 1: presence, 0: absence

PronConflation a numeric vector, 1: presence, 0: absence

MoreThan2DegreesDistDem a numeric vector, 1: presence, 0: absence

NonSpkrAnchoredDem a numeric vector, 1: presence, 0: absence  
VerticalityDem a numeric vector, 1: presence, 0: absence  
ClassifiedDem a numeric vector, 1: presence, 0: absence  
NumDeterminedDecl a numeric vector, 1: presence, 0: absence  
GenderDeterminedDecl a numeric vector, 1: presence, 0: absence  
SuppletiveNouns a numeric vector, 1: presence, 0: absence  
SingMarkedNoun a numeric vector, 1: presence, 0: absence  
DualMarkedNoun a numeric vector, 1: presence, 0: absence  
PlMarkedNoun a numeric vector, 1: presence, 0: absence  
OtherNumMarkedNoun a numeric vector, 1: presence, 0: absence  
LimitedDistNumMarking a numeric vector, 1: presence, 0: absence  
NounClassesGenders a numeric vector, 1: presence, 0: absence  
ConcordBeyondNp a numeric vector, 1: presence, 0: absence  
NumeralClassifiers a numeric vector, 1: presence, 0: absence  
PossClassifiers a numeric vector, 1: presence, 0: absence  
PossClasses a numeric vector, 1: presence, 0: absence  
Inalienability a numeric vector, 1: presence, 0: absence  
MultiplePossConstr a numeric vector, 1: presence, 0: absence  
PrefixMarkedPoss a numeric vector, 1: presence, 0: absence  
SuffixMarkedPoss a numeric vector, 1: presence, 0: absence  
MarkedPossr a numeric vector, 1: presence, 0: absence  
MarkedPossessee a numeric vector, 1: presence, 0: absence  
PossrPossdOrder a numeric vector, 1: presence, 0: absence  
DecimalNumerals a numeric vector, 1: presence, 0: absence  
QuinaryNumerals a numeric vector, 1: presence, 0: absence  
CollectiveNouns a numeric vector, 1: presence, 0: absence  
AdjVerbLexOverlap a numeric vector, 1: presence, 0: absence  
AdjAttributionPred a numeric vector, 1: presence, 0: absence  
CoreCaseMarking a numeric vector, 1: presence, 0: absence  
ObliqueCaseMarking a numeric vector, 1: presence, 0: absence  
Prepositions a numeric vector, 1: presence, 0: absence  
Postpositions a numeric vector, 1: presence, 0: absence  
TamPerson a numeric vector, 1: presence, 0: absence  
VerbPrefixesProclitics a numeric vector, 1: presence, 0: absence  
VerbSuffixesEnclitics a numeric vector, 1: presence, 0: absence  
PunctualContinuous a numeric vector, 1: presence, 0: absence  
RealisIrrealis a numeric vector, 1: presence, 0: absence

SSuffix a numeric vector, 1: presence, 0: absence  
SPrefix a numeric vector, 1: presence, 0: absence  
ASuffix a numeric vector, 1: presence, 0: absence  
APrefix a numeric vector, 1: presence, 0: absence  
OSuffix a numeric vector, 1: presence, 0: absence  
OPrefix a numeric vector, 1: presence, 0: absence  
VerbVarTam a numeric vector, 1: presence, 0: absence  
VerbVarVClass a numeric vector, 1: presence, 0: absence  
VerbVarClauseType a numeric vector, 1: presence, 0: absence  
VerbVarPerson a numeric vector, 1: presence, 0: absence  
NumStemAlt a numeric vector, 1: presence, 0: absence  
PersonStemAlt a numeric vector, 1: presence, 0: absence  
SepVerbNumPerson a numeric vector, 1: presence, 0: absence  
Portmanteau3Plus a numeric vector, 1: presence, 0: absence  
DistributedCategory a numeric vector, 1: presence, 0: absence  
NonCore a numeric vector, 1: presence, 0: absence  
RecipientObj a numeric vector, 1: presence, 0: absence  
X3PlacePreds a numeric vector, 1: presence, 0: absence  
VerbNeg a numeric vector, 1: presence, 0: absence  
VerbDirection a numeric vector, 1: presence, 0: absence  
VerbSuppletion a numeric vector, 1: presence, 0: absence  
ConjugationClasses a numeric vector, 1: presence, 0: absence  
TransIntransAlt a numeric vector, 1: presence, 0: absence  
TransitivizingMorph a numeric vector, 1: presence, 0: absence  
IntransitivizingMorph a numeric vector, 1: presence, 0: absence  
ReflexiveMorph a numeric vector, 1: presence, 0: absence  
ReciprocalMorph a numeric vector, 1: presence, 0: absence  
VerbClassifiers a numeric vector, 1: presence, 0: absence  
Copula a numeric vector, 1: presence, 0: absence  
NonVbPreds a numeric vector, 1: presence, 0: absence  
SerialVerbConstr a numeric vector, 1: presence, 0: absence  
Auxiliaries a numeric vector, 1: presence, 0: absence  
VerbCompounds a numeric vector, 1: presence, 0: absence  
VerbAdjunctConstr a numeric vector, 1: presence, 0: absence  
VbIncorporation a numeric vector, 1: presence, 0: absence  
ExistentialVerb a numeric vector, 1: presence, 0: absence  
IrregularGive a numeric vector, 1: presence, 0: absence

ClosedClassOfVb a numeric vector, 1: presence, 0: absence  
 SvIntransClauses a numeric vector, 1: presence, 0: absence  
 VsIntransClauses a numeric vector, 1: presence, 0: absence  
 VInitTransClauses a numeric vector, 1: presence, 0: absence  
 VMedialTransClauses a numeric vector, 1: presence, 0: absence  
 VFinalTransClauses a numeric vector, 1: presence, 0: absence  
 FixedConstituentOrder a numeric vector, 1: presence, 0: absence  
 ClauseFinalNeg a numeric vector, 1: presence, 0: absence  
 ClauseInitNeg a numeric vector, 1: presence, 0: absence  
 ImpVs.DeclNeg a numeric vector, 1: presence, 0: absence  
 VbAndNonVbPredIdentity a numeric vector, 1: presence, 0: absence  
 SOMorphInBasicConstr a numeric vector, 1: presence, 0: absence  
 SAMorphInBasicConstr a numeric vector, 1: presence, 0: absence  
 SOMorphInComplexConstr a numeric vector, 1: presence, 0: absence  
 SAMorphInComplexConstr a numeric vector, 1: presence, 0: absence  
 SynConflationOfSO a numeric vector, 1: presence, 0: absence  
 ControlledUncontrolled a numeric vector, 1: presence, 0: absence  
 ClauseChaining a numeric vector, 1: presence, 0: absence  
 SimultaneousSequential a numeric vector, 1: presence, 0: absence  
 SayInDesidConstr a numeric vector, 1: presence, 0: absence  
 RelativeClauses a numeric vector, 1: presence, 0: absence  
 PurpSubClauses a numeric vector, 1: presence, 0: absence  
 TemporalSubClauses a numeric vector, 1: presence, 0: absence  
 ComplementClauses a numeric vector, 1: presence, 0: absence  
 CausBySerialVerbConstr a numeric vector, 1: presence, 0: absence  
 CausByBoundAffClit a numeric vector, 1: presence, 0: absence  
 CausByConstrInvolvingSay a numeric vector, 1: presence, 0: absence  
 MorphTopicOrFocus a numeric vector, 1: presence, 0: absence  
 TailHeadLinkage a numeric vector, 1: presence, 0: absence  
 VerbRedup a numeric vector, 1: presence, 0: absence  
 NounRedup a numeric vector, 1: presence, 0: absence

### Source

Dunn, M., Terrill, A., Reesink, G., Foley, R. A. and Levinson, S. C. (2005) Structural phylogenetics and the reconstruction of ancient language history, *Science*, 309, 2072-2075.

**Examples**

```

## Not run:
data(phylogeny)
library(ape)

# joint analysis of both language families

phylogeny.dist = dist(phylogeny[,3:ncol(phylogeny)], method = "binary")
phylogeny.dist.tr = nj(phylogeny.dist)
families =
  as.character(phylogeny$Family[as.numeric(phylogeny.dist.tr$tip.label)])
languages =
  as.character(phylogeny$Language[as.numeric(phylogeny.dist.tr$tip.label)])
phylogeny.dist.tr$tip.label = languages
plot(phylogeny.dist.tr, type="u", font = as.numeric(as.factor(families)))

# analysis of subset of Papuan languages

papuan = phylogeny[phylogeny$Family == "Papuan",]
papuan$Language = as.factor(as.character(papuan$Language))
papuan.meta = papuan[,1:2]
papuan.mat = papuan[, 3:ncol(papuan)]
papuan.meta$Geography = c(
  "Bougainville", "Bismarck Archipelago", "Bougainville",
  "Bismarck Archipelago", "Bismarck Archipelago", "Central Solomons",
  "Bougainville", "Louisiade Archipelago", "Bougainville",
  "Bismarck Archipelago", "Bismarck Archipelago",
  "Bismarck Archipelago", "Central Solomons", "Central Solomons",
  "Central Solomons")
papuan.dist = dist(papuan.mat, method = "binary")
papuan.dist.tr = nj(papuan.dist)
fonts =
  as.character(papuan.meta$Geography[as.numeric(
    papuan.dist.tr$tip.label)])
papuan.dist.tr$tip.label =
  as.character(papuan.meta$Language[as.numeric(
    papuan.dist.tr$tip.label)])
plot(papuan.dist.tr, type = "u", font = as.numeric(as.factor(fonts)))

## End(Not run)

```

---

plot.corres

*Plot method for correspondence objects*


---

**Description**

This function defines a plot method for correspondence objects.

**Usage**

```
## S3 method for class 'corres'  
plot(x, main = "", addcol = TRUE, extreme = 0, rcex = 1, rcol = 1,  
rlabels = "", stretch = 1.4, ccex = 1, ccol = 2, clabels = "", ...)
```

**Arguments**

x	A correspondence object as produced by <code>corres</code> .
main	A string used for the main title of the plot.
addcol	A logical, if true, columns are added to the plot.
extreme	If nonzero, defines quantiles that define the extremes such that only data points exceeding these extremes are plotted.
rcex	sets <code>cex</code> graphical parameter for rows.
rcol	sets color for rows.
rlabels	vector of row labels.
stretch	a number defining the degree to which the columns (or rows) should be stretched out for visual presentation.
ccex	sets <code>cex</code> graphical parameter for columns.
ccol	sets color for columns.
clabels	vector of column labels.
...	other parameters to be passed through to plotting functions.

**Value**

A plot on the graphics device.

**Author(s)**

R. H. Baayen

**See Also**

See Also [corres.fnc](#), [link{corsup.fnc}](#).

**Examples**

```
## Not run:  
data(oldFrench)  
data(oldFrenchMeta)  
  
oldFrench.ca = corres.fnc(oldFrench)  
  
plot(oldFrench.ca)  
  
plot(oldFrench.ca, rlabels = oldFrenchMeta$Genre,  
rcol = as.numeric(oldFrenchMeta$Genre), rcex = 0.5,  
extreme = 0.1, ccol = "blue")
```

```
## End(Not run)
```

---

plot.growth	<i>Plot method for growth objects</i>
-------------	---------------------------------------

---

### Description

This function defines the plot method for growth objects.

### Usage

```
## S3 method for class 'growth'  
plot(x, w = "all", ...)
```

### Arguments

x	A growth object.
w	A character string denoting the name of a specific variable to be plotted.
...	other parameters to be passed through to plotting functions.

### Value

A plot shown on the graphics device.

### Author(s)

R. H. Baayen

### See Also

See Also [growth.fnc](#).

### Examples

```
## Not run:  
data(alice)  
alice.growth = growth.fnc(alice)  
plot(alice.growth)  
plot(alice.growth, w = "Yule")  
  
## End(Not run)
```

---

plotAll.fnc                      *create plot or plots for list with data frames for plot or subplots*

---

### Description

given a list with one or more data frames with values for a plot (or subplot), create the actual plots

### Usage

```
plotAll.fnc(reslist, sameYrange = TRUE, ylabel, xlabel = NA, intrName = NA,
  pos = "end", ylimit = NA, addlines=FALSE, cexsize = 0.6, conditioningVals=NA,
  conditioningColors=1, conditioningLines=1, lineColor=1, addToExistingPlot = FALSE, ...)
```

### Arguments

reslist	list with as components either a data frame or a list with data frames, the data frames specify X and Y coordinates and HPD intervals
sameYrange	logical, if TRUE, the ylim for each panel will be chosen to accomodate the range of values across all panels in the plot
ylabel	label to be used for the vertical axis
xlabel	label to be used for the horizontal axis; this option is active only when a single predictor is plotted
intrName	label for the interaction predictor, if present
pos	location of legend values for interaction
ylimit	if specified, overrides sameYrange for ylim
addlines	if TRUE, adds line between levels of same factor(s)
cexsize	character expansion size for information in the plot for interactions, default is 0.6
conditioningVals	vector of names of the levels of the conditioning factor in the interaction (the factor with different lines in the plot)
conditioningColors	vector of names of the colors to be used for the levels of the conditioning factor in the interaction (the factor with different lines in the plot)
conditioningLines	vector of names of the line types to be used for the levels of the conditioning factor in the interaction (the factor with different lines in the plot), by default solid lines
lineColor	name of color to be used for the lines in the plot
addToExistingPlot	if TRUE, the current plot is added to an already existing plot
...	further graphical parameters to be passed down, none are currently implemented

**Details**

Note that `reslist` may contain as elements lists of data frames, these then specify the separate points or lines to be plotted for a given interaction

**Value**

A plot is produced on the graphics device.

**Note**

not intended for independent use

**Author(s)**

R. H. Baayen

**See Also**

See Also as [plotLMER.fnc](#)

**Examples**

```
## Not run: not intended for independent use
```

---

plotLMER.fnc

*plot a mer object*

---

**Description**

Plot partial effects of a (generalized) linear mixed-effects model fit with `lmer`. For gaussian models, 95% highest posterior density credible intervals can be added.

**Usage**

```
plotLMER.fnc(model, xlabel = NA, xlabs = NA, ylabel = NA, ylimit = NA,
  ilabel = NA, fun = NA, pred = NA, control = NA, ranefs = NA, n = 100,
  intr = NA, mcmcMat = NA, lockYlim = TRUE, addlines = FALSE,
  withList = FALSE, cexsize = 0.5, linecolor = 1, addToExistingPlot = FALSE,
  verbose = TRUE, ...)
```

**Arguments**

<code>model</code>	a mer model object
<code>xlabel</code>	label for X-axis (if other than the variable name in the original model formula)
<code>xlabs</code>	character vector with labels for X-axes in multipanel plot (if other than the variable names in the original model formula); if used, <code>xlabel</code> should not be specified

ylabel	label for Y-axis (if other than the variable name of the dependent variable in the original model formula)
ylim	range for vertical axis; if not specified, this range will be chosen such that all data points across all subplots, including HPD intervals, will be accommodated
ilabel	label for the interaction shown in the lower right-hand margin of the plot, overriding the original variable name in the model formula
fun	a function to be applied for transforming the dependent variable, if NA, no transformation is applied; for models with family = "binomial", fun is set to plogis by default; this can be disabled by setting fun=function(x) return(x).
pred	character string with name of predictor; if specified, a single plot will produced for the partial effect of this specific predictor
control	a two-element list list(predictor, val) specifying a predictor the value of which has to be set to val in the partial effect plot(s); the predictor name should be exactly as specified in names(model@fixef). It is up to the user to make sure that name and value make sense, the code here hands full 'control' to the user.
ranefs	a four-element list Group, Level, Predictor, specifying a random-effect Group (e.g. Subject), a level (e.g., S10) and a value (e.g., LogFrequency) for which partial effects have to be calibrated; implemented only for mcmcMat=NA.
n	integer denoting number of points for the plot, chosen at equally spaced intervals across the empirical range of the predictor variable
intr	a list specifying an interaction to be graphed; obligatory arguments are (1) the name of the interaction variable, followed by (2) a vector of values for that variable, followed by (3) the position for interaction labels ("beg", "mid", or "end", or 'NA' if no labels are desired), optionally followed by (4) a list with as first element a vector of colors and as second element a vector of line types. The number of elements in both vectors should match the number of values specified under (2) for the interaction predictor.
mcmcMat	an MCMC matrix as produced by mcmcSamp or pvals.fnc()\$mcmc for the input model; when this matrix is specified, an attempt is made to add MCMC-based HPD intervals to the predicted values in the plot. Works only for R 2.6.2 or higher, and for lme4 0.999 or higher.
lockYlim	logical specifying whether all subplots should have the same range of values for the vertical axis; if TRUE, this range will be chosen to accomodate all fitted values including HDP intervals for all predictors across all plots
addlines	if TRUE, adds line(s) between levels of same factor(s)
withList	logical, if TRUE, a list will be output with all data frames for the subplots
cexsize	character expansion size (cex) for additional information in the plot for interactions
linecolor	color of lines in the plot, by default set to 1 (black)
addToExistingPlot	default FALSE, if set to TRUE, plot will be added to previous plot, but only if pred is specified
verbose	if TRUE (default), effect sizes and default transformations are reported

... further graphical parameters to be passed down; warning: col, pch, lty and cex will often generate an error as they are internally already fully specified for specialized subplots

### Details

When no predictor is specified, a series of plots is produced for the partial effects of each predictor. The graphs are shown for the reference level for factors and are adjusted for the median value for the other numerical predictors in the model. Interactions are not shown. The user should set up the appropriate number of subplots on the graphics device before running plotLMER.fnc().

Instead of showing all predictors jointly, plotLMER.fnc() can also be used to plot the partial effect of a specific predictor. When a specific predictor is specified (with pred = ...), a single plot is produced for that predictor. In this case, the intr argument can be used to specify a single second predictor that enters into an interaction with the selected main predictor.

Polynomials have to be fitted with poly(..., degree, raw=TRUE) and restricted cubic splines with rcs() from the rms package.

When an MCMC object (as produced by the MCMC code in lme4, version 1.0 or higher) is specified (mcmcMat = ...), an attempt is made to plot MCMC HPD intervals for the predicted values. This code is under development. For each MCMC sample, the posterior values of the fixed effects are used to predict the dependent variable for a range of its values, with other predictors in the model held constant (factors at the reference level, covariates at their medians). For each of the predicted 'posterior' values, the HPD interval is obtained, and added to the graph.

### Value

A plot is produced on the graphical device.

### Note

This code needs much more work, including (i) extension to poly with raw=FALSE, and (ii) general clean-up of the code.

### Author(s)

R. H. Baayen

### References

The 'danish' dataset in the example section is contributed by Laura Winther-Balling, see Winther-Balling, L. and Baayen, R. H., Morphological effects in auditory word recognition: Evidence from Danish, Language and Cognitive Processes, in press.

### See Also

See also other utilities in languageR for facilitating work with lmer

## Examples

```

## Not run:

#####
# fitting a cosine with a spline (simulated data)
#####

require("rms", quietly=TRUE, character=TRUE)
require("lme4", quietly=TRUE, character=TRUE)
dfr = makeSplineData.fnc()
table(dfr$Subject)
xylowess.fnc(Y ~ X | Subject, data = dfr)
# the smoother doesn't recognize the cosine function implemented in makeSplineData.fnc()
dev.off()

dfr.lmer = lmer(Y ~ rcs(X, 5) + (1|Subject), data = dfr)
plotLMER.fnc(dfr.lmer)

# comparison with ols from Design package
dfr.dd = datadist(dfr)
options(datadist='dfr.dd')
dfr.ols = ols(Y~Subject+rcs(X), data=dfr, x=T, y=T)
dfr$fittedOLS = fitted(dfr.ols)
dfr$fittedLMER = as.vector(dfr.lmer@X %*% fixef(dfr.lmer))

# we plot the lmer() fit in blue, the ols() fit in red (both adjusted for
# subject S1), and plot the underlying model in green

plot(dfr[dfr$Subject=="S1",]$X, dfr[dfr$Subject=="S1",]$fittedLMER +
  ranef(dfr.lmer)[[1]][ "S1", ], type="l", col="blue",
  ylim = range(dfr$y + ranef(dfr.lmer)[[1]][ "S1", ],
  dfr[dfr$Subject == "S1",]$fittedLMER,
  dfr[dfr$Subject == "S1",]$fittedOLS),
  xlab="X", ylab="Y")
lines(dfr[dfr$Subject=="S1",]$X, dfr[dfr$Subject=="S1",]$fittedOLS,
  col="red")
lines(dfr[dfr$Subject=="S1",]$X, dfr[dfr$Subject=="S1",]$y+
  ranef(dfr.lmer)[[1]][ "S1", ], col="green")
legend(2,29,c("30+cos(x)", "lmer (S1)", "ols (S1)"), lty=rep(1,3),
  col=c("green", "blue", "red"))

#####
# a model with a raw polynomial: the mcmc credible intervals
# will be different each time this example is run; for large
# numbers of samples, the plotting code will become quite slow
#####

bg.lmer = lmer(LogRT ~ PC1+PC2+PC3 + ReadingScore +

```

```

poly(OrthLength, 2, raw=TRUE) + LogFrequency + LogFamilySize +
(1|Word) + (1|Subject)+(0+OrthLength|Subject) +
(0+LogFrequency|Subject), data = beginningReaders)
mcmc = pvals.fnc(bg.lmer, nsim=1000, withMCMC=TRUE)

pars = par()
par(mfrow=c(3,3), mar=c(5,5,1,1))
plotLMER.fnc(bg.lmer, mcmcMat=mcmc$mcmc, fun=exp, ylabel = "RT (ms)")

#####
# a model with an interaction involving numeric predictors
#####

danish.lmer = lmer(LogRT ~ PC1 + PC2 + PrevError + Rank +
ResidSemRating + ResidFamSize + LogWordFreq*LogAffixFreq*Sex +
poly(LogCUP, 2, raw=TRUE) + LogUP + LogCUPtoEnd +
(1|Subject) + (1|Word) + (1|Affix), data = danish)
danish.lmerA = lmer(LogRT ~ PC1 + PC2 + PrevError + Rank +
ResidSemRating + ResidFamSize + LogWordFreq*LogAffixFreq*Sex +
poly(LogCUP, 2, raw=TRUE) + LogUP + LogCUPtoEnd +
(1|Subject) + (1|Word) + (1|Affix), data = danish,
subset=abs(scale(resid(danish.lmer)))<2.5)
mcmc = pvals.fnc(danish.lmerA, nsim=10000, withMCMC=TRUE)
mcmc$fixed[,1:5]

# plot for reference level of Sex
plotLMER.fnc(danish.lmerA, pred = "LogAffixFreq",
intr=list("LogWordFreq", round(quantile(danish$LogWordFreq),3), "beg",
list(c("red", "green", "blue", "yellow", "purple"), rep(1,5))), ylim=c(6.5,7.0),
mcmcMat=mcmc$mcmc, xlabel = "log affix frequency", ylabel = "log RT auditory lexical decision")

# this model has a significant three-way interaction
# for visualization, we can either relevel Sex and refit,
# or make use of the control option. First releveling:

danish$Sex=relevel(danish$Sex, "F")
danish.lmerF = lmer(LogRT ~ PC1 + PC2 + PrevError + Rank +
  ResidSemRating + ResidFamSize + LogWordFreq*LogAffixFreq*Sex +
  poly(LogCUP, 2, raw=TRUE) + LogUP + LogCUPtoEnd +
  (1|Subject) + (1|Word) + (1|Affix), data = danish)
danish$Sex=relevel(danish$Sex, "M")
danish.lmerM = lmer(LogRT ~ PC1 + PC2 + PrevError + Rank +
  ResidSemRating + ResidFamSize + LogWordFreq*LogAffixFreq*Sex +
  poly(LogCUP, 2, raw=TRUE) + LogUP + LogCUPtoEnd +
  (1|Subject) + (1|Word) + (1|Affix), data = danish)

# Next preparing for using the control option:
#
# names(fixef(danish.lmer))[10] # SexM
# unique(danish.lmer@X[,10])   # 1 0

par(mfrow=c(2,2))

```

```

plotLMER.fnc(danish.lmer, pred="LogWordFreq", ylimit=c(6.5,7.0),
intr=list("LogAffixFreq", quantile(danish$LogAffixFreq), "end"),
control=list("SexM", 0))
mtext("females", line=1.5, cex=0.9)

plotLMER.fnc(danish.lmer, pred="LogWordFreq", ylimit=c(6.5,7.0),
intr=list("LogAffixFreq", quantile(danish$LogAffixFreq), "end"),
control=list("SexM", 1))
mtext("males", line=1.5, cex=0.9)

plotLMER.fnc(danish.lmerF, pred="LogWordFreq", ylimit=c(6.5,7.0),
intr=list("LogAffixFreq", quantile(danish$LogAffixFreq), "end"))
mtext("females", line=1.5, cex=0.9)

plotLMER.fnc(danish.lmerM, pred="LogWordFreq", ylimit=c(6.5, 7.0),
intr=list("LogAffixFreq", quantile(danish$LogAffixFreq), "end"))
mtext("males", line=1.5, cex=0.9)

par(mfrow=c(1,1))

#####
# calculating effect sizes, defined as max - min
#####

# effect size for a covariate

dfr = plotLMER.fnc(danish.lmerA, pred = "LogCUP", withList=TRUE)
max(dfr$LogCUP$Y)-min(dfr$LogCUP$Y)

# effect size for a factor

dfr = plotLMER.fnc(danish.lmerA, pred = "PrevError", withList=TRUE)
max(dfr$PrevError$Y)-min(dfr$PrevError$Y)

# effect sizes for the quantiles in an interaction plot

dfr = plotLMER.fnc(danish.lmerA, pred = "LogAffixFreq",
withList=TRUE,
intr=list("LogWordFreq", round(quantile(danish$LogWordFreq),3), "beg"))

unlist(lapply(dfr$LogAffixFreq, FUN=function(X)return(max(X$Y)-min(X$Y))))

#####
# plotting an interaction between two factors
#####

danish$WordFreqFac = danish$LogWordFreq > median(danish$LogWordFreq)
danish.lmer2 = lmer(LogRT ~ WordFreqFac*Sex +
(1|Subject) + (1|Word) + (1|Affix), data = danish)
mcmc = pvals.fnc(danish.lmer2,nsim=1000,withMCMC=TRUE)

```

```

plotLMER.fnc(danish.lmer2, pred = "Sex",
intr=list("WordFreqFac", c("TRUE", "FALSE"), "end",
list(c("red", "blue"), rep(1,2))),
ylimit=c(6.5,7.0), cexsize=1.0, addlines=TRUE, mcmcMat=mcmc$mcmc)

#####
# a generalized linear mixed-effects model
#####

dative.lmer = lmer(RealizationOfRecipient ~
AccessOfTheme + AccessOfRec + LengthOfRecipient + AnimacyOfRec +
AnimacyOfTheme + PronomOfTheme + DefinOfTheme + LengthOfTheme +
SemanticClass + Modality + (1|Verb),
data = dative, family = "binomial")

par(mfrow=c(3,4),mar=c(5,5,1,1))
plotLMER.fnc(dative.lmer, fun=plogis, addlines=TRUE)

# with user-specified labels for the x-axis
par(mfrow=c(3,4),mar=c(5,5,1,1))
plotLMER.fnc(dative.lmer, fun=plogis, addlines=TRUE,
xlab=unlist(strsplit("abcdefghij","")))

par(pars)

## End(Not run)

```

---

plotlogistic.fit.fnc *Plot for goodness of fit of logistic regression*

---

## Description

This function plots observed proportions against mean predicted probabilities. For a good fit, points should be approximately on a straight line.

## Usage

```
plotlogistic.fit.fnc(x, data, method, where, scalesize, ...)
```

## Arguments

x	A logistic regression model fitted with lmer or lrm.
data	The data set to which the model was fitted.
method	Either "cut", in which case the vector of cut-off points supplied by the "where" argument will be used to partition the fitted probabilities, or "shingle", in which a shingle (using equal.count and its defaults) will be used.

where A vector of cut-off points for partitioning the vector of fitted probabilities, by default seq(0, 1, by=0.1).

scalesize A positive real  $\leq 1$ . If not NA (the default), the circles representing data points in the graph are scaled to reflect the number of data points in the underlying data set. The scalesize parameter specifies how large the largest circle will be compared to 1 inch. For counts with large outliers, small values of scalesize are better. See example below.

... other parameters to be passed through to plotting functions.

### Value

A plot is produced on the graphics device. The R-squared value shown above the plot represents the correlation between the X and Y values in the plot. It does NOT represent the R-squared of the lrm or lmer model.

### Author(s)

R. H. Baayen

### Examples

```
## Not run:
data(dative)
library(lme4, keep.source=FALSE)

dative.lrm = lrm(RealizationOfRecipient ~ AccessOfTheme +
  AccessOfRec + LengthOfRecipient + AnimacyOfRec +
  AnimacyOfTheme + PronomOfTheme + DefinOfTheme + LengthOfTheme +
  SemanticClass + Modality, data = dative)

dative.glmm = lmer(RealizationOfRecipient ~ AccessOfTheme +
  AccessOfRec + LengthOfRecipient + AnimacyOfRec +
  AnimacyOfTheme + PronomOfTheme + DefinOfTheme + LengthOfTheme +
  SemanticClass + Modality + (1|Verb),
  data = dative, family = "binomial")

par(mfrow=c(2,2))
plotlogistic.fit.fnc (dative.lrm, dative)
mtext("lrm", 3, 3)
plotlogistic.fit.fnc (dative.glmm, dative)
mtext("lmer", 3, 3)
plotlogistic.fit.fnc (dative.lrm, dative, scalesize=0.2)
mtext("lrm", 3, 3)
plotlogistic.fit.fnc (dative.glmm, dative, method="shingle")
mtext("lmer", 3, 3)
par(mfrow=c(1,1))

## End(Not run)
```

---

```
preparePredictor.fnc    determine X and Y values for a given (sub)plot
```

---

**Description**

this function figures out the X and Y values for a given (sub)plot, including upper and lower 95% HPD intervals

**Usage**

```
preparePredictor.fnc(pred, model, m, ylabel, fun, val, xlabel, mcmc, ranefs, ...)
```

**Arguments**

pred	character string denoting predictor to be plotted on horizontal axis
model	model fit by lmer
m	matrix as produced by makeDefaultMatrix.fnc
ylabel	label for vertical axis (if other than name of dependent variable)
fun	character string denoting transformation function for dependent variable, currently only "plogis" or "exp"
val	value of interacting variable
xlabel	label for horizontal axis
mcmc	MCMC matrix as produced by mcmcsamp or pvals.fnc()\$mcmc
ranefs	a three-element list Group, Level, Predictor, specifying a random-effect Group (e.g. Subject), a level (e.g., S10) and a value (e.g., LogFrequency) for which partial effects have to be calibrated; implemented only for mcmcMat=NA.
...	further graphical parameters, currently not implemented

**Value**

A data frame with values to be plotted, with columns

X	values of predictor
Y	fitted values
Type	logical for whether predictor is factor
Interaction	logical for whether predictor is interacting predictor
Levels	for factors, the factor level names (only present for factors)

**Note**

not intended for independent use

**Author(s)**

R. H. Baayen

**See Also**

See Also as [plotLMER.fnc](#)

**Examples**

```
## Not run: Not intended for independent use.
```

---

 primingHeid

---

*Primed lexical decision latencies for neologisms ending in -heid*


---

**Description**

Primed lexical decision latencies for Dutch neologisms ending in the suffix *-heid*.

**Usage**

```
data(primingHeid)
```

**Format**

A data frame with 832 observations on the following 13 variables.

**Subject** a factor with subjects as levels.

**Word** a factor with words as levels.

**Trial** a numeric vector for the rank of the trial in its experimental list.

**RT** a numeric vector with log-transformed lexical decision latencies.

**Condition** a factor coding the priming treatment, with levels *baseheid* (prime is the base word) and *heid* (the prime is the neologism)

**Rating** a numeric vector for subjective frequency estimates.

**Frequency** a numeric vector for log-transformed frequencies of the whole word.

**BaseFrequency** a numeric vector for the log-transformed frequencies of the base word.

**LengthInLetters** a numeric vector coding orthographic length in letters.

**FamilySize** a numeric vector for the log-transformed count of the word's morphological family.

**NumberOfSynsets** a numeric vector for the number of synonym sets in WordNet in which the base is listed.

**ResponseToPrime** a factor with levels *correct* and *incorrect* for the response to the prime.

**RTtoPrime** a numeric vector for the log-transformed reaction time to the prime.

## References

De Vaan, L., Schreuder, R. and Baayen, R. H. (2007) Regular morphologically complex neologisms leave detectable traces in the mental lexicon, *The Mental Lexicon*, 2, in press.

## Examples

```
## Not run:
data(primingHeid)

library(lme4, keep.source=FALSE)

primingHeid.lmer = lmer(RT ~ RTtoPrime * ResponseToPrime + Condition +
(1|Subject) + (1|Word), data = primingHeid)
pvals.fnc(primingHeid.lmer)$summary

# model criticism

primingHeid.lmer = lmer(RT ~ RTtoPrime * ResponseToPrime + Condition +
(1|Subject) + (1|Word),
data = primingHeid[abs(scale(resid(primingHeid.lmer)))<2.5,])
pvals.fnc(primingHeid.lmer)$summary

## End(Not run)
```

---

primingHeidPrevRT      *Primed lexical decision latencies for neologisms ending in -heid*

---

## Description

Primed lexical decision latencies for Dutch neologisms ending in the suffix *-heid*, with information on RTs to preceding trials added to the data already in `primingHeid`.

## Usage

```
data(primingHeidPrevRT)
```

## Format

A data frame with 832 observations on the following 17 variables.

`Subject` a factor with subjects as levels.

`Word` a factor with words as levels.

`Trial` a numeric vector for the rank of the trial in its experimental list.

`RT` a numeric vector with log-transformed lexical decision latencies.

`Condition` a factor coding the priming treatment, with levels `baseheid` (prime is the base word) and `heid` (the prime is the neologism)

`Rating` a numeric vector for subjective frequency estimates.

Frequency a numeric vector for log-transformed frequencies of the whole word.  
 BaseFrequency a numeric vector for the log-transformed frequencies of the base word.  
 LengthInLetters a numeric vector coding orthographic length in letters.  
 FamilySize a numeric vector for the log-transformed count of the word's morphological family.  
 NumberOfSynsets a numeric vector for the number of synonym sets in WordNet in which the base is listed.  
 ResponseToPrime a factor with levels correct and incorrect for the response to the prime.  
 RTtoPrime a numeric vector for the log-transformed reaction time to the prime.  
 RTmin1 a numeric vector for reaction time in ms to the item preceding the target.  
 RTmin2 a numeric vector for reaction time in ms to the item preceding the target by two trials.  
 RTmin3 a numeric vector for reaction time in ms to the item preceding the target by three trials.  
 RTmin4 a numeric vector for reaction time in ms to the item preceding the target by four trials.

## References

De Vaan, L., Schreuder, R. and Baayen, R. H. (2007) Regular morphologically complex neologisms leave detectable traces in the mental lexicon, *The Mental Lexicon*, 2, in press.

## Examples

```

## Not run:
data(primingHeidPrevRT)

library(lme4, keep.source=FALSE)

primingHeid.lmer = lmer(RT ~ RTtoPrime * ResponseToPrime + Condition +
log(RTmin1) + (1|Subject) + (1|Word), data = primingHeidPrevRT)
pvals.fnc(primingHeid.lmer)$summary

## End(Not run)

```

---

```
print.corres
```

*Print method for correspondence object*

---

## Description

Prints eigenvalues and eigenvalue rates for a correspondence object.

## Usage

```

## S3 method for class 'corres'
print(x, ...)

```

**Arguments**

x                    A correspondence object.  
...                  other parameters to be passed through to plotting functions.

**Value**

Report of eigenvalues and eigenvalue rates.

**Author(s)**

R. H. Baayen

**See Also**

See also [corres.fnc](#).

**Examples**

```
## Not run:  
data(oldFrench)  
oldFrench.ca = corres.fnc(oldFrench)  
oldFrench.ca  
  
## End(Not run)
```

---

print.growth                    *Print method for growth objects.*

---

**Description**

Print method for growth objects.

**Usage**

```
## S3 method for class 'growth'  
print(x, ...)
```

**Arguments**

x                    A growth object, as produced by growth.fnc.  
...                  other parameters to be passed through to plotting functions.

**Value**

The data frame with chunk sizes and associated vocabulary statistics is printed. To access the data frame that is being shown, use `<my.growth.object>@data$data`.

**Author(s)**

R. H. Baayen

**See Also**See also [growth.fnc](#).**Examples**

```
## Not run:
data(alice)
alice.growth = growth.fnc(alice)
alice.growth
# for accessing the printed data frame:
alice.growth@data$data[1:4,]

## End(Not run)
```

pvals.fnc

*Compute p-values and MCMC confidence intervals for mixed models***Description**

This function calculates p-values and HPD intervals for the parameters of models fitted with `lmer`. For the fixed-effects parameters, anticonservative p-values based on the t statistic with the upper bound for the degrees of freedom are added.

**Usage**

```
pvals.fnc(object, nsim = 10000, ndigits = 4, withMCMC = FALSE,
addPlot=TRUE, ...)
```

**Arguments**

<code>object</code>	An <code>lmer</code> or <code>glmer</code> model object fitted with <code>lmer</code> . Currently, MCMC sampling for generalized linear mixed models may not work off the shelf when there is more than one random intercept in the model. The code for random effects structure with correlation parameters is not implemented in <code>mcmcsmamp</code> , this currently blocks calculation of the fixed and random subtables. For evaluation of such models, likelihood ratio tests are recommended.
<code>nsim</code>	An integer denoting the required number of Markov chain Monte Carlo samples.
<code>ndigits</code>	An integer denoting the number of decimal digits in the output.
<code>withMCMC</code>	A logical indicating whether the output of <code>mcmcsmamp</code> should be returned.
<code>addPlot</code>	If TRUE (default), a trellis graph for the posterior distributions of the parameters is shown on the current graphics device.
<code>...</code>	Optional arguments that can be passed down.

**Value**

A list with components

fixed	A data frame with the estimated coefficients, their MCMC mean, the HPD 95 and the probability based on the t distribution with the number of observations minus the number of fixed-effects coefficients as degrees of freedom. This last p-value is anti-conservative, especially for small data sets.
random	A table with the MCMC means and HPD 95 for the standard deviations and correlations of the random effects. The HPD intervals are based on the untransformed values (log for standard deviations, atanh for correlations), the endpoints of the HPD interval are back-transformed to the scale of the input data frame for ease of interpretation.
mcmc	The output of <code>mcmc</code> samp.

**Author(s)**

R. H. Baayen

**See Also**

See also `lmer`, `mcmc`samp, and `HPDinterval`.

**Examples**

```
## Not run:
data(primingHeid)
library(lme4, keep.source=FALSE)

# remove extreme outliers
primingHeid = primingHeid[primingHeid$RT < 7.1,]

# fit mixed-effects model

primingHeid.lmer = lmer(RT ~ RTtoPrime * ResponseToPrime +
  Condition + (1|Subject) + (1|Word), data = primingHeid)

mcmc = pvals.fnc(primingHeid.lmer, nsim=10000, withMCMC=TRUE)

mcmc$fixed
mcmc$random

## End(Not run)
```

---

quasif	<i>Simulated data set with subjects and items requiring quasi-F ratios</i>
--------	--

---

**Description**

Simulated lexical decision latencies with SOA as treatment, traditionally requiring an analysis using quasi-F ratios, as available in Raaijmakers et al. (1999).

**Usage**

```
data(quasif)
```

**Format**

A data frame with 64 observations on the following 4 variables.

Subject a factor coding subjects.

RT a numeric vector for simulated reaction times in lexical decision.

Item a factor coding items.

SOA a factor coding SOA treatment with levels long and short.

**Source**

Raaijmakers, J.G.W., Schrijnemakers, J.M.C. & Gremmen, F. (1999) How to deal with "The language as fixed effect fallacy": common misconceptions and alternative solutions, *Journal of Memory and Language*, 41, 416-426.

**Examples**

```
## Not run:
data(quasif)
items.quasif.fnc(quasif)

## End(Not run)
```

---

quasiF.fnc	<i>Quasi-F test</i>
------------	---------------------

---

**Description**

The textbook Quasi-F test for a design with subjects, items, and a single factorial predictor. Included for educational purposes for this specific design only.

**Usage**

```
quasiF.fnc(ms1, ms2, ms3, ms4, df1, df2, df3, df4)
```

**Arguments**

ms1	Mean squares Factor
ms2	Mean squares Item:Subject
ms3	Mean squares Factor:Subject
ms4	Mean squares Item
df1	Degrees of freedom Factor
df2	Degrees of freedom Item:Subject
df3	Degrees of freedom Factor:Subject
df4	Degrees of freedom Item

**Value**

A list with components

F	Quasi-F value.
df1	degrees of freedom numerator.
df2	degrees of freedom denominator.
p	p-value.

**Author(s)**

R. H. Baayen

**See Also**

See Also as [quasiFsim.fnc](#).

**Examples**

```
data(quasif)

quasif.lm = lm(RT ~ SOA + Item + Subject +
  SOA:Subject + Item:Subject, data = quasif)
quasif.aov = anova(quasif.lm)

quasiF.fnc(quasif.aov["SOA", "Mean Sq"],
  quasif.aov["Item:Subject", "Mean Sq"],
  quasif.aov["SOA:Subject", "Mean Sq"],
  quasif.aov["Item", "Mean Sq"],
  quasif.aov["SOA", "Df"],
  quasif.aov["Item:Subject", "Df"],
  quasif.aov["SOA:Subject", "Df"],
  quasif.aov["Item", "Df"])

# much simpler is
quasiFsim.fnc(quasif)$quasiF
```

---

quasiFsim.fnc	<i>Quasi-F test for specific simple design</i>
---------------	--

---

**Description**

This function carries out a Quasi-F test for data with columns labelled SOA, Subject, Item. This function is called by `simulate.quasif.fnc`, and is not intended for general use.

**Usage**

```
quasiFsim.fnc(dat)
```

**Arguments**

`dat` A data frame with RT (or RTsim), SOA, Subject and Item as predictors.

**Value**

A list with components

<code>p</code>	The p-value of the quasi-F test.
<code>data</code>	The input data.
<code>model</code>	The linear model fitted to the data.
<code>qF</code>	a list with F, df1, df2 and p-value of quasi-F test.

**Author(s)**

R. H. Baayen

**See Also**

See Also [quasiF.fnc](#).

**Examples**

```
data(quasif)
quasiFsim.fnc(quasif)$quasiF
```

---

 ratings

*Ratings for 81 English nouns*


---

**Description**

Subjective frequency ratings, ratings of estimated weight, and ratings of estimated size, averaged over subjects, for 81 concrete English nouns.

**Usage**

```
data(ratings)
```

**Format**

A data frame with 81 observations on the following 14 variables.

Word a factor with words as levels.

Frequency a numeric vector of logarithmically transformed frequencies

FamilySize a numeric vector of logarithmically transformed morphological family sizes.

SynsetCount a numeric vector with logarithmically transformed counts of the number of synonym sets in WordNet in which the word is listed.

Length a numeric vector for the length of the word in letters.

Class a factor with levels animal and plant.

FreqSingular a numeric vector for the frequency of the word in the singular.

FreqPlural a numeric vector with the frequency of the word in the plural.

DerivEntropy a numeric vector with the derivational entropies of the words.

Complex a factor coding morphological complexity with levels complex and simplex.

rInfl a numeric vector coding the log of ratio of singular to plural frequencies.

meanWeightRating a numeric vector for the estimated weight of the word's referent, averaged over subjects.

meanSizeRating a numeric vector for the estimated size of the word's referent, averaged over subjects.

meanFamiliarity a numeric vector with subjective frequency estimates, averaged over subjects.

**Source**

Data collected together with Jen Hay at the University of Canterbury, Christchurch, New Zealand, 2004.

**Examples**

```
## Not run:
data(ratings)

ratings.lm = lm(meanSizeRating ~ meanFamiliarity * Class +
I(meanFamiliarity^2), data = ratings)

ratings$fitted = fitted(ratings.lm)

plot(ratings$meanFamiliarity, ratings$meanSizeRating,
xlab = "mean familiarity", ylab = "mean size rating", type = "n")
text(ratings$meanFamiliarity, ratings$meanSizeRating,
substr(as.character(ratings$Class), 1, 1), col = 'darkgrey')

plants = ratings[ratings$Class == "plant", ]
animals = ratings[ratings$Class == "animal", ]
plants = plants[order(plants$meanFamiliarity),]
animals = animals[order(animals$meanFamiliarity),]

lines(plants$meanFamiliarity, plants$fitted)
lines(animals$meanFamiliarity, animals$fitted)

## End(Not run)
```

---

regularity

*Regular and irregular Dutch verbs*


---

**Description**

Regular and irregular Dutch verbs and selected lexical and distributional properties.

**Usage**

```
data(regularity)
```

**Format**

A data frame with 700 observations on the following 13 variables.

Verb a factor with the verbs as levels.

WrittenFrequency a numeric vector of logarithmically transformed frequencies in written Dutch (as available in the CELEX lexical database).

NcountStem a numeric vector for the number of orthographic neighbors.

VerbalSynsets a numeric vector for the number of verbal synsets in WordNet.

MeanBigramFrequency a numeric vector for mean log bigram frequency.

InflectionalEntropy a numeric vector for Shannon's entropy calculated for the word's inflectional variants.

Auxiliary a factor with levels hebben, zijn and zijnheb for the verb's auxiliary in the perfect tenses.

Regularity a factor with levels irregular and regular.

LengthInLetters a numeric vector of the word's orthographic length.

FamilySize a numeric vector for the number of types in the word's morphological family.

Valency a numeric vector for the verb's valency, estimated by its number of argument structures.

NVratio a numeric vector for the log-transformed ratio of the nominal and verbal frequencies of use.

WrittenSpokenRatio a numeric vector for the log-transformed ratio of the frequencies in written and spoken Dutch.

## References

Baayen, R. H. and Moscoso del Prado Martin, F. (2005) Semantic density and past-tense formation in three Germanic languages, *Language*, 81, 666-698.

Tabak, W., Schreuder, R. and Baayen, R. H. (2005) Lexical statistics and lexical processing: semantic density, information complexity, sex, and irregularity in Dutch, in Kepser, S. and Reis, M., *Linguistic Evidence - Empirical, Theoretical, and Computational Perspectives*, Berlin: Mouton de Gruyter, pp. 529-555.

## Examples

```
## Not run:
data(regularity)

# ---- predicting regularity with a logistic regression model

library(rms)
regularity.dd = datadist(regularity)
options(datadist = 'regularity.dd')

regularity.lrm = lrm(Regularity ~ WrittenFrequency +
  rcs(FamilySize, 3) + NcountStem + InflectionalEntropy +
  Auxiliary + Valency + NVratio + WrittenSpokenRatio,
  data = regularity, x = TRUE, y = TRUE)

anova(regularity.lrm)

# ---- model validation

validate(regularity.lrm, bw = TRUE, B = 200)
pentrace(regularity.lrm, seq(0, 0.8, by = 0.05))
regularity.lrm.pen = update(regularity.lrm, penalty = 0.6)
regularity.lrm.pen

# ---- a plot of the partial effects

plot(Predict(regularity.lrm.pen))
```

```
# predicting regularity with a support vector machine

library(e1071)
regularity$AuxNum = as.numeric(regularity$Auxiliary)
regularity.svm = svm(regularity[, -c(1,8,10)], regularity$Regularity, cross=10)
summary(regularity.svm)

## End(Not run)
```

---

selfPacedReadingHeid *Self-paced reading latencies for Dutch neologisms*

---

## Description

Self-paced reading latencies for Dutch neologisms ending in the suffix *-heid*.

## Usage

```
data(selfPacedReadingHeid)
```

## Format

A data frame with 1280 observations on the following 18 variables.

**Subject** a factor with subjects as levels.

**Word** a factor with words as levels.

**RT** a numeric vector with logarithmically transformed reading latencies.

**RootFrequency** a numeric vector for the logarithmically transformed frequency of the lowest-level base of the neologism (e.g., *lob* in *[[[lob]+ig]+heid]*).

**Condition** a factor for the priming conditions with levels *baseheid* (neologism is preceded 40 trials back by its base word) and *heidheid* (the neologism is preceded 40 trials back by itself).

**Rating** a numeric vector for the word's subjective frequency estimate.

**Frequency** a numeric vector for the neologism's frequency (all zero).

**BaseFrequency** a numeric vector for the base adjective underlying the neologism (e.g., *lobbig* in *[[[lob]+ig]+heid]*).

**LengthInLetters** a numeric vector coding word length in letters.

**FamilySize** a numeric vector for the logarithmically transformed count of a word's morphological family members.

**NumberOfSynsets** a numeric vector for the count of synonym sets in WordNet in which the word is listed.

**RT4WordsBack** a numeric vector for the log-transformed reading latencies four trials back.

**RT3WordsBack** a numeric vector for the log-transformed reading latencies three trials back.

**RT2WordsBack** a numeric vector for the log-transformed reading latencies two trials back.

**RT1WordBack** a numeric vector for the log-transformed reading latencies one trial back.

**RT1WordLater** a numeric vector for the log-transformed reading latencies one trial later.

**RT2WordsLater** a numeric vector for the log-transformed reading latencies two trials later.

**RTtoPrime** a numeric vector for the log-transformed reading latency for the prime.

## References

De Vaan, L., Schreuder, R. and Baayen, R. H. (2007) Regular morphologically complex neologisms leave detectable traces in the mental lexicon, *The Mental Lexicon*, 2, in press.

## Examples

```
## Not run:
data(selfPacedReadingHeid)

# data validation
plot(sort(selfPacedReadingHeid$RT))
selfPacedReadingHeid = selfPacedReadingHeid[selfPacedReadingHeid$RT > 5 &
  selfPacedReadingHeid$RT < 7.2,]

# fitting a mixed-effects model

library(lme4, keep.source = FALSE)
x = selfPacedReadingHeid[,12:15]
x.pr = prcomp(x, center = TRUE, scale = TRUE)
selfPacedReadingHeid$PC1 = x.pr$x[,1]

selfPacedReadingHeid.lmer = lmer(RT ~ RTtoPrime + LengthInLetters +
  PC1 * Condition + (1|Subject) + (1|Word),
  data = selfPacedReadingHeid)
pvals.fnc(selfPacedReadingHeid.lmer)$summary

# model criticism

selfPacedReadingHeid.lmerA = lmer(RT ~ RTtoPrime + LengthInLetters +
  PC1 * Condition + (1|Subject) + (1|Word), data =
  selfPacedReadingHeid[abs(scale(resid(selfPacedReadingHeid.lmer))) < 2.5, ])

qqnorm(resid(selfPacedReadingHeid.lmerA))

pvals.fnc(selfPacedReadingHeid.lmerA)$summary

## End(Not run)
```

---

shadenormal.fnc

*Shade rejection region for normal probability density function*


---

## Description

This function plots the standard normal probability density function and shades the rejection region.

## Usage

```
shadenormal.fnc(qnts = c(0.025, 0.975))
```

**Arguments**

qnts                    A numeric vector with the Z-scores of the boundaries of the lower and upper rejection regions.

**Value**

A plot on the graphics device.

Type `shadenormal.fnc` to see the code. The `polygon()` function used for the shaded areas takes a sequence of X and Y coordinates, connects the corresponding points, and fills the area(s) enclosed with a specified color. To understand the use of `polygon()`, one can best think of making a polygon with a set of pins, a thread, and a board. Outline the polygon by placing the pins on the board at the corners of the polygon. First fasten the thread to one of the pins, then connect the thread to the second pin, from there to the third pin, and so on, until the first pin has been reached. What `polygon()` requires as input is a vector of the X-coordinates of the pins, and a vector of their Y-coordinates. These coordinates should be in exactly the order in which the thread is to be connected from pin to pin.

For shading the left rejection area, we specify the vectors of X and Y coordinates, beginning at the leftmost point of the tail, proceeding to the right edge of the shaded area, then up, and finally to the left and down to the starting point, thereby closing the polygon. The X-coordinates are therefore specified from left to right, and then from right to left. The corresponding Y-coordinates are all the zeros necessary to get from  $-3$  to  $1.96$  (the default, `qnorm(0.025)`), and then the Y-coordinates of the density in reverse order to return to where we began.

**Author(s)**

R. H. Baayen

**Examples**

```
## Not run:  
shadenormal.fnc()  
  
## End(Not run)
```

---

show.growth

*Plot method for growth objects.*

---

**Description**

A print method for growth objects created with `growth.fnc`.

**Usage**

```
## S3 method for class 'growth'  
show(x)
```

**Arguments**

x                    A growth object.

**Value**

Prints growth object. To access the data frame embedded in the growth object, use `<my.growth.object>@data$data`.

**Author(s)**

R. H. Baayen

**See Also**

See Also [growth.fnc](#).

**Examples**

```
## Not run:
data(alice)
alice.growth = growth.fnc(alice, chunks= c(5000, 10000, 15000))
alice.growth

## End(Not run)
```

---

shrinkage

*Data set illustrating shrinkage*

---

**Description**

Simulated data set for illustrating shrinkage.

**Usage**

```
data(shrinkage)
```

**Format**

A data frame with 200 observations on the following 6 variables.

`intercept` a numeric vector for the intercept.

`frequency` a numeric vector for word frequency.

`subject` a factor for subjects with levels S1, S2, ... , S10.

`error` a numeric vector for residuals.

`ranef` a numeric vector for random effect.

`RT` a numeric vector for simulated RTs.

**Examples**

```
## Not run:
data(shrinkage)

library(lme4, keep.source=FALSE)

shrinkage.lmer = lmer(RT ~ frequency + (1|subject), data = shrinkage)
shrinkage.lmList = lmList(RT ~ frequency | subject, data = shrinkage)

# and visualize the difference between random regression
# and mixed-effects regression

mixed = coef(shrinkage.lmer)[[1]]
random = coef(shrinkage.lmList)
subj = unique(shrinkage[,c("subject", "ranef")])
subj = subj[order(subj$subject),]
subj$random = random[,1]
subj$mixed = mixed[,1]
subj = subj[order(subj$random),]
subj$rank = 1:nrow(subj)

par(mfrow=c(1,2))
plot(subj$rank, subj$random, xlab="rank", ylab="RT", ylim=c(200,550), type="n")
text(subj$rank, subj$random, as.character(subj$subject), cex=0.8, col="red")
mtext("random regression", 3, 1)
points(subj$rank, 400+subj$ranef, col="blue")
abline(h=400)
plot(subj$rank, subj$mixed, xlab="rank", ylab="RT", ylim=c(200,550), type="n")
text(subj$rank, subj$mixed, as.character(subj$subject), cex=0.8, col = "red")
mtext("mixed-effects regression", 3, 1)
points(subj$rank, 400+subj$ranef, col="blue")
abline(h=400)
par(mfrow=c(1,1))

## End(Not run)
```

---

```
simulateLatinsquare.fnc
```

*Simulate simple Latin Square data and compare models*

---

**Description**

This function creates a user-specified number of simulated datasets with a Latin Square design, and compares mixed-effects models with the by-subject anova.

**Usage**

```
simulateLatinsquare.fnc(dat, with = TRUE, trial = 0, nruns = 100,
  nsub = NA, nitem = NA, ...)
```

**Arguments**

dat	A data frame with the structure of the data set <code>latinsquare</code> .
with	Logical, if TRUE, effect of SOA built into the data.
trial	A number which, if nonzero, gives the magnitude of a learning or a fatigue effect.
nruns	A number indicating the required number of simulation runs.
nsub	A number for the number of subjects.
nitem	A number for the number of items.
...	other parameters to be passed through to plotting functions.

**Value**

A list with components

alpha05	Description of 'comp1'
alpha01	proportion of runs in which predictors are significant at the 05 significance level.
res	Data frame with simulation results.
with	Logical, TRUE if SOA effect is built into the simulations.

**Author(s)**

R. H. Baayen

**Examples**

```
## Not run:
  data(latinsquare)
  \dontrun{
    library(lme4)
    simulateLatinsquare.fnc(latinsquare, nruns=100)
  }

## End(Not run)
```

---

simulateQuasif.fnc      *Simulate data for quasi-F analysis and compare models*

---

**Description**

This function creates a user-specified number of simulated datasets, and compares mixed-effects models with quasi-F and F1 and F2 analyses. It should be run with the version of R and the version of languageR used by Baayen, Davidson & Bates (2008, JML), as `mcmcscamp` no longer supports models with random correlation parameters.

**Usage**

```
simulateQuasif.fnc(dat, with = TRUE, nruns = 100, nsub = NA, nitem = NA, ...)
```

**Arguments**

dat	Data frame with a data set with as variables Subject, Item and SOA, as in the quasif data set.
with	Logical, if TRUE, an effect of SOA is built into the simulation.
nruns	Integer for the number of simulation runs.
nsub	Integer denoting the number of subjects.
nitem	Integer denoting the number of items.
...	other parameters to be passed through to plotting functions.

**Details**

Model parameters are estimated from the input data set.

For each completed simulation run, a dot is added to the R console.

**Value**

A list with components

alpha05	Description of 'comp1'
alpha01	proportion of runs in which predictors are significant at the 05 significance level.
res	Data frame with simulation results.
with	Logical, TRUE if SOA effect is built into the simulations.

**Author(s)**

R. H. Baayen

**See Also**

See also [subjects.quasif.fnc](#).

**Examples**

```
## Not run:
data(quasif)
library(lme4)

quasif.sim = simulateQuasif.fnc(quasif, nruns = 1000, with = TRUE)
quasif.sim$alpha05

## End(Not run)
```

---

 simulateRegression.fnc

*Simulate regression data and compare models*


---

### Description

This function creates a user-specified number of simulated regression datasets, and compares mixed-effects regression with random regression, by-subject regression, by-item regression, and by-subject plus by-item regression. Optionally, an effect of learning or fatigue can be incorporated.

### Usage

```
simulateRegression.fnc(beta = c(400, 2, 6, 4), nitem = 20, nsubj = 10,
  stdevItem = 40, stdevSubj = 80, stdevError = 50, nruns = 100, learn = FALSE,
  learnRate = 10, ...)
```

### Arguments

beta	A numeric vector with beta weights for the intercept and three numeric predictors.
nitem	A number specifying the number of items.
nsubj	A number specifying the number of subjects.
stdevItem	A number specifying the standard deviation of the Item random effect.
stdevSubj	A number specifying the standard deviation of the Subject random effect.
stdevError	A number specifying the standard deviation of the Residual Error.
nruns	A number specifying the required number of simulated datasets.
learn	A logical that if TRUE, allows an effect of learning or fatigue into the model.
learnRate	A number specifying the learning rate (if negative) or the effect of fatigue (if positive).
...	other parameters to be passed through to plotting functions.

### Value

A list with components

alpha05	proportion of runs in which predictors are significant at the 05 significance level.
alpha01	proportion of runs in which predictors are significant at the 01 significance level.
ranef	mean estimated random effects.

As this may take some time, the index of each completed run is shown on the output device.

### Author(s)

R. H. Baayen

**See Also**

See Also [make.reg.fnc](#).

**Examples**

```
## Not run:  
library(lme4)  
simulateRegression.fnc(beta = c(400, 2, 6, 4), nruns = 5)  
  
\dontrun{simulateRegression.fnc(beta = c(400, 2, 6, 0), nruns = 1000, learn = TRUE)}  
  
## End(Not run)
```

---

sizeRatings

*Size ratings for 81 English concrete nouns*

---

**Description**

Subjective estimates of the size of the referents of 81 English concrete nouns, collected from 38 subjects.

**Usage**

```
data(sizeRatings)
```

**Format**

A data frame with 3078 observations on the following 7 variables.

**Rating** a numeric vector with subjective estimates of the size of the word's referent.

**Subject** a factor with subjects as levels.

**Word** a factor with words as levels.

**Class** a factor with levels animal and plant.

**Naive** a factor with levels naive and notNaive, coding whether the subject new about the purpose of the experiment.

**Language** a factor with levels English and notEnglish coding whether the subject was a native speaker of English.

**MeanFamiliarity** a numeric vector for the by-item mean familiarity ratings.

**Details**

Data collected with Jen Hay, University of Canterbury, Christchurch, New Zealand, 2004.

**Examples**

```
## Not run:
data(sizeRatings)
library(lme4, keep.source = FALSE)
sizeRatings.lmer = lmer(Rating ~ Class * Naive +
  MeanFamiliarity * Language + (1|Subject) + (1|Word),
  data = sizeRatings)
pvals.fnc(sizeRatings.lmer)$summary

## End(Not run)
```

spanish

*Relative frequencies of tag trigrams in selected Spanish texts***Description**

Relative frequencies of the 120 most frequent tag trigrams in 15 texts contributed by 3 authors.

**Usage**

```
data(spanish)
```

**Format**

A data frame with 120 observations on 15 variables documented in spanishMeta.

**References**

Spasova, M. S. (2006) *Las marcas sintacticas de atribucion forense de autoria de textos escritos en espanol*, Masters thesis, Institut Universitari de Linguistica Aplicada, Universitat Pompeu Fabra, Barcelona.

**Examples**

```
## Not run:
data(spanish)
data(spanishMeta)

# principal components analysis

spanish.t = t(spanish)
spanish.pca = prcomp(spanish.t, center = TRUE, scale = TRUE)
spanish.x = data.frame(spanish.pca$x)
spanish.x = spanish.x[order(rownames(spanish.x)), ]

library(lattice)
splom(~spanish.x[ , 1:3], groups = spanishMeta$Author)

# linear discriminant analysis
```

```

library(MASS)
spanish.pca.lda = lda(spanish.x[ , 1:8], spanishMeta$Author)
plot(spanish.pca.lda)

# cross-validation

n = 8
spanish.t = spanish.t[order(rownames(spanish.t)), ]
predictedClasses = rep("", 15)
for (i in 1:15) {
  training = spanish.t[-i,]
  trainingAuthor = spanishMeta[-i,]$Author
  training.pca = prcomp(training, center=TRUE, scale=TRUE)
  training.x = data.frame(training.pca$x)
  training.x = training.x[order(rownames(training.x)), ]
  training.pca.lda = lda(training[ , 1:n], trainingAuthor)
  predictedClasses[i] =
    as.character(predict(training.pca.lda, spanish.t[ , 1:n])$class[i])
}

ncorrect = sum(predictedClasses==as.character(spanishMeta$Author))
ncorrect
sum(dbinom(ncorrect:15, 15, 1/3))

## End(Not run)

```

---

spanishFunctionWords *Relative frequencies of function words in selected Spanish texts*

---

### Description

Relative frequencies of the 120 most frequent function words in 15 texts contributed by 3 authors.

### Usage

```
data(spanishFunctionWords)
```

### Format

A data frame with 120 observations on 15 variables documented in spanishMeta.

### References

Spasova, M. S. (2006) *Las marcas sintacticas de atribucion forense de autoria de textos escritos en espanol*, Masters thesis, Institut Universitari de Linguistica Aplicada, Universitat Pompeu Fabra, Barcelona.

**Examples**

```

## Not run:
data(spanishFunctionWords)
data(spanishMeta)

# principal components analysis

spanishFunctionWords.t = t(spanishFunctionWords)
spanishFunctionWords.t =
  spanishFunctionWords.t[order(rownames(spanishFunctionWords.t)), ]
spanishFunctionWords.pca =
  prcomp(spanishFunctionWords.t, center = TRUE, scale = TRUE)

sdevs = spanishFunctionWords.pca$sdev^2
n = sum(sdevs/sum(sdevs)> 0.05)

# linear discriminant analysis with cross-validation

library(MASS)

predictedClasses = rep("", 15)
for (i in 1:15) {
  training = spanishFunctionWords.t[-i,]
  trainingAuthor = spanishMeta[-i,]$Author
  training.pca = prcomp(training, center = TRUE, scale = TRUE)
  training.x = data.frame(training.pca$x)
  training.x = training.x[order(rownames(training.x)), ]
  training.pca.lda = lda(training[, 1:n], trainingAuthor)
  cl = predict(training.pca.lda, spanishFunctionWords.t[,1:n])$class[i]
  predictedClasses[i] = as.character(cl)
}

ncorrect = sum(predictedClasses==spanishMeta$Author)
sum(dbinom(ncorrect:15, 15, 1/3))

## End(Not run)

```

---

spanishMeta

*Metadata for the spanish and spanishFunctionWords data sets*


---

**Description**

By-text metadata for the spanish and spanishFunctionWords data sets.

**Usage**

```
data(spanishMeta)
```

**Format**

A data frame with 15 observations on the following 6 variables.

Author a factor with levels C, M, and V.

YearOfBirth a numeric vector with year of birth of the author.

TextName a factor with codes for the texts as levels ( X14458g11 ... X14476g11).

PubDate a numeric vector with data of publication of the text.

Nwords a numeric vector with text sizes in tokens.

FullName a factor with author names: Cela, Mendoza and VargasLLosa.

**References**

Spasova, M. S. (2006) *Las marcas sintacticas de atribucion forense de autoria de textos escritos en espanol*, Masters thesis, Institut Universitari de Linguistica Aplicada, Universitat Pompeu Fabra, Barcelona.

**Examples**

```
## Not run:
data(spanishMeta)

## End(Not run)
```

---

spectrum.fnc

*Frequency spectrum from text vector*

---

**Description**

This function creates a frequency spectrum for a text in character vector form.

**Usage**

```
spectrum.fnc(text)
```

**Arguments**

text A character vector containing the words of a text.

**Value**

A data frame with as column variables

frequency Word frequencies.

freqOffreq The frequencies of the word frequencies.

**Author(s)**

R. H. Baayen

**References**

R. H. Baayen (2001) *Word Frequency Distributions*, Dordrecht: Kluwer.

**See Also**

See Also the zipfR package.

**Examples**

```
## Not run:
data(alice)
alice.spectrum = spectrum.fnc(alice)
head(alice.spectrum)
tail(alice.spectrum)

## End(Not run)
```

---

splitplot

*Simulated data set with split plot design*


---

**Description**

Simulated lexical decision latencies with priming as treatment and reaction time in lexical decision as dependent variable.

**Usage**

```
data(splitplot)
```

**Format**

A data frame with 800 observations on the following 11 variables.

`items` A factor with levels w1, w2, ..., w40, coding 40 word items.

`ritems` The by-word random adjustments to the intercept.

`list` A factor with levels listA and listB. The priming effect is counterbalanced for subjects across these two lists, compare `table(splitplot$list, splitplot$subjects)`.

`rlist` The by-list random adjustments to the intercept.

`priming` A treatment factor with levels primed and unprimed.

`fpriming` The priming effect, -30 for the primed and 0 for the unprimed condition.

`subjects` A factor with levels s1, s2, ... s20 coding 20 subjects.

`rsubject` The by-subject random adjustments to the intercept.

error The by-observation noise.  
 int The intercept.  
 RT The reaction time.

### Source

R. H. Baayen, D. J. Davidson and D. M. Bates. Mixed-effects modeling with crossed random effects for subjects and items. Manuscript under revision for *Journal of Memory and Language*.

### Examples

```
## Not run:
data(splitplot)
table(splitplot$list, splitplot$subjects)
library(lme4)
dat.lmer1 = lmer2(RT ~ list*priming+(1+priming|subjects)+(1+list|items),data=dat)
dat.lmer2 = lmer2(RT ~ list*priming+(1+priming|subjects)+(1|items),data=dat)
dat.lmer3 = lmer2(RT ~ list*priming+(1|subjects)+(1|items),data=dat)
dat.lmer4 = lmer2(RT ~ list*priming+(1|subjects),data=dat)
anova(dat.lmer1, dat.lmer2, dat.lmer3, dat.lmer4)
print(dat.lmer3, corr=FALSE)

## End(Not run)
```

---

subjects.latinsquare.fnc

*By-subject analysis of simple Latin Square data sets*

---

### Description

This function is called by simulateLatinsquare.fnc for by-subject analysis of simulated Latin Square datasets. It is not intended for independent use.

### Usage

```
subjects.latinsquare.fnc(dat)
```

### Arguments

dat	A data frame with variables RT or RTsim , SOA, Subject, Item, Group and List, as in the latinsquare data set.
-----	---

### Value

A list with components

p	The p-value of the by-subject anova.
data	The input dataset.
model	The fitted model.

**Author(s)**

R. H. Baayen

**See Also**

See also [simulateLatinsquare.fnc](#).

**Examples**

```
## Not run:  
data(latinsquare)  
subjects.latinsquare.fnc(latinsquare)$p  
  
## End(Not run)
```

---

subjects.quasif.fnc    *By-subject analysis of data sets requiring quasi-F ratios*

---

**Description**

This function is called by `simulateQuasif.fnc` for by-subject analysis of simulated datasets traditionally requiring quasi-F ratios. It is not intended for independent use.

**Usage**

```
subjects.quasif.fnc(dat)
```

**Arguments**

`dat`                    A data frame with variables RT or RTsim , SOA, Subject and Item.

**Value**

A list with components

`p`                      p-value for by-subject F-test.  
`data`                   Data set with aggregated subject means.  
`model`                  Anova table of fitted model.

**Author(s)**

R. H. Baayen

**See Also**

See also [simulateQuasif.fnc](#).

**Examples**

```
## Not run:
  data(quasif)
  subjects.quasif.fnc(quasif)

## End(Not run)
```

---

summary.corres	<i>Summarize a correspondence object</i>
----------------	--

---

**Description**

This function provides a concise summary of a correspondence object.

**Usage**

```
## S3 method for class 'corres'
summary(object, n = 2, returnList = FALSE, head = TRUE, ...)
```

**Arguments**

object	A correspondence object as produced by <code>corres</code> .
n	A number indicating number of dimensions to be summarized.
returnList	Logical, if TRUE, a list is returned with as components the full information on each factor, instead of only the first 6 lines.
head	Logical, if TRUE, first 6 rows of factor summaries are shown.
...	Additional arguments passed on to summaries.

**Value**

A summary with eigenvalue rates, and coordinates, correlations, and contributions for the factors (by default, 2, unless n is set to a higher number).

**Author(s)**

R. H. Baayen

**See Also**

See also [corres.fnc](#).

**Examples**

```
## Not run:  
data(oldFrench)  
oldFrench.ca = corres.fnc(oldFrench)  
oldFrench.ca  
summary(oldFrench.ca)  
  
## End(Not run)
```

---

summary.growth	<i>Summary method for growth objects</i>
----------------	--

---

**Description**

Summary method for vocabulary growth objects created with `growth.fnc`.

**Usage**

```
## S3 method for class 'growth'  
summary(object, ...)
```

**Arguments**

object	A vocabulary growth object.
...	other parameters to be passed through to plotting functions.

**Value**

The growth object is printed. For access to the data frame inside the object, use `<my.growth.object>@data$data`.

**Author(s)**

R. H. Baayen

**See Also**

See also [growth.fnc](#).

---

tail.growth	<i>Show last rows of growth object.</i>
-------------	---

---

**Description**

Prints last rows of growth object.

**Usage**

```
## S3 method for class 'growth'  
tail(x, n = 6, ...)
```

**Arguments**

x	A growth object.
n	An integer specifying the number of lines to be shown.
...	other parameters to be passed through to plotting functions.

**Author(s)**

R. H. Baayen

**See Also**

See Also [growth.fnc](#).

**Examples**

```
## Not run:  
data(alice)  
alice.growth = growth.fnc(alice)  
tail(alice.growth)  
  
## End(Not run)
```

---

text2spc.fnc	<i>Create a frequency spectrum from a text vector</i>
--------------	---

---

**Description**

This functions takes a text in the form of a character vector as input, and outputs a frequency spectrum object as defined in the zipfR package.

**Usage**

```
text2spc.fnc(text)
```

**Arguments**

text            A text in the form of a character vector.

**Value**

A spc spectrum object as defined in the zipfR package.

**Author(s)**

R. H. Baayen

**See Also**

See the documentation for zipfR for spc objects.

**Examples**

```
## Not run:  
library(zipfR)  
data(alice)  
alice.spc = text2spc.fnc(alice)  
plot(alice.spc)  
  
## End(Not run)
```

---

through

*Through the Looking Glass*

---

**Description**

The text of Lewis Carroll's 'Through the Looking Glass', with punctuation marks removed.

**Usage**

```
data(through)
```

**Format**

A character vector with 29560 words.

**Source**

The project Gutenberg at [http://www.gutenberg.org/wiki/Main\\_Page](http://www.gutenberg.org/wiki/Main_Page)

**Examples**

```
## Not run:  
  data(through)  
  through[1:3]  
  
## End(Not run)
```

---

transforming.fnc	<i>transform vector according to specified function</i>
------------------	---

---

**Description**

Apply function fun to input vector y

**Usage**

```
transforming.fnc(y, fun)
```

**Arguments**

y	numerical vector (for dependent variable)
fun	a function, or NA (in which case no transformation is applied)

**Details**

exists only to make code more readable

**Value**

a numerical vector

**Note**

not intended for independent use

**Author(s)**

R. H. Baayen

**See Also**

See Also as [plotLMER.fnc](#)

**Examples**

```
## Not run: not intended for independent use
```

---

twente

*Frequency spectrum for the Twente News Corpus*

---

**Description**

Frequency (m) and frequency of frequency (Vm) for string types in the Twente News Corpus.

**Usage**

```
data(twente)
```

**Format**

A data frame with 4639 observations on the following 2 variables.

m a numeric vector with word frequencies.

Vm a numeric vector with the frequencies of word frequencies.

**Source**

Twente News Corpus.

**Examples**

```
## Not run:  
data(twente)  
library(zipfR)  
twente.spc = spc(m=twente$m, Vm = twente$Vm)  
plot(twente.spc)  
  
## End(Not run)
```

---

variationLijk

*Variation in spoken Dutch in the use of the suffix -lijk*

---

**Description**

This dataset documents variation in the use of the suffix *-lijk*, as realized in 32 words, in spoken Dutch across region (Flanders versus The Netherlands), sex (females versus males) and education (high versus mid).

**Usage**

```
data(variationLijk)
```

**Format**

A data frame with 32 observations on the following 8 variables.

`nlfemaleHigh` a numeric vector with counts for Dutch females with a mid education level.

`nlfemaleMid` a numeric vector counts for Dutch females with a high education level.

`nlmaleHigh` a numeric vector counts for Dutch males with a high education level.

`nlmaleMid` a numeric vector counts for Dutch males with a mid education level.

`vlfemaleHigh` a numeric vector counts for Flemish females with a high education level.

`vlfemaleMid` a numeric vector counts for Flemish females with a mid education level.

`vlmaleHigh` a numeric vector counts for Flemish males with a high education level.

`vlmaleMid` a numeric vector counts for Flemish males with a mid education level.

**References**

Keune, K., Ernestus, M., Van Hout, R. and Baayen, R.H. (2005) Social, geographical, and register variation in Dutch: From written 'mogelijk' to spoken 'mok', *Corpus Linguistics and Linguistic Theory*, 1, 183-223.

**Examples**

```
## Not run:
data(variationLijk)
variationLijk.ca = corres.fnc(variationLijk)
plot(variationLijk.ca, rcex=0.7, ccol="black",
      rcol = rep("blue", nrow(variationLijk)))

## End(Not run)
```

---

 ver

---

*The Dutch prefix ver-: semantic transparency and frequency*


---

**Description**

Semantic transparency (dichotomous) and frequency for 985 words with the Dutch prefix *ver-*.

**Usage**

```
data(ver)
```

**Format**

A data frame with 985 observations on the following 2 variables.

`Frequency` a numeric vector for the words' frequency.

`SemanticClass` a factor with levels `opaque` and `transparent` coding semantic transparency.

## References

Baayen, R. H. and Lieber, R. (1997) Word Frequency Distributions and Lexical Semantics, *Computers and the Humanities*, 30, 281-291.

## Examples

```
## Not run:
data(ver)
ver$Frequency = log(ver$Frequency)

plot(density(ver$Frequency))

# plot separate densities for opaque and transparent words

ver.transp = ver[ver$SemanticClass == "transparent",]$Frequency
ver.opaque = ver[ver$SemanticClass == "opaque", ]$Frequency

ver.transp.d = density(ver.transp)
ver.opaque.d = density(ver.opaque)
xlimit = range(ver.transp.d$x, ver.opaque.d$x)
ylimmit = range(ver.transp.d$y, ver.opaque.d$y)
plot(ver.transp.d, lty = 1, col = "black",
     xlab = "frequency", ylab = "density",
     xlim = xlimit, ylim = ylimmit, main = "")
lines(ver.opaque.d, col = "darkgrey")
legend(6,0.25, lty=rep(1,2), col=c("black", "darkgrey"),
      legend=c("transparent", "opaque"))

# test whether the difference is significant

ks.test(jitter(ver.transp), jitter(ver.opaque))

## End(Not run)
```

---

verbs

*Dative Alternation - simplified data set*

---

## Description

A simplified version of the dative data set, used for expository purposes only.

## Usage

```
data(verbs)
```

**Format**

A data frame with 903 observations on the following 5 variables.

RealizationOfRec a factor with levels NP and PP.

Verb a factor with the verbs as levels.

AnimacyOfRec a factor with levels animate and inanimate.

AnimacyOfTheme a factor with levels animate and inanimate.

LengthOfTheme a numeric vector coding the length in words of the theme.

**References**

Bresnan, J., Cueni, A., Nikitina, T. and Baayen, R. H. (2007) Predicting the dative alternation, in Bouma, G. and Kraemer, I. and Zwarts, J. (eds.), *Cognitive Foundations of Interpretation*, Royal Netherlands Academy of Sciences, 33 pages, in press.

**Examples**

```
data(verbs)
head(verbs)
xtabs( ~ RealizationOfRec + AnimacyOfRec, data = verbs)
barplot(xtabs( ~ RealizationOfRec + AnimacyOfRec, data = verbs),beside=TRUE)
```

---

 warlpiri

*Ergative case marking in Warlpiri*


---

**Description**

This data set documents the use of ergative case marking in the narratives of native speakers of Lajamanu Warlpiri (8 children, 13 adults) describing events in picture books.

**Usage**

```
data(warlpiri)
```

**Format**

A data frame with 347 observations on the following 9 variables.

Speaker a factor with speakers as levels.

Sentence a factor with sentence as levels.

AgeGroup a factor with levels adult and child.

CaseMarking a factor with levels ergative and other.

WordOrder a factor with levels subInitial (subject initial) and subNotInitial (subject not initial).

AnimacyOfSubject a factor with levels animate and inanimate.

OverttnessOfObject a factor with levels notOvert and overt.

AnimacyOfObject a factor with levels animate and inanimate.

Text a factor with levels texta, textb and textc.

## References

O'Shannessy, C. (2006) *Language contact and child bilingual acquisition: Learning a mixed language and Warlpiri in northern Australia*, PhD Thesis, University of Sydney, Australia.

## Examples

```
## Not run:
data(warlpiri)
library(lme4, keep.source = FALSE)
warlpiri.lmer = lmer(CaseMarking ~ WordOrder * AgeGroup +
  AnimacyOfSubject + (1|Text) + (1|Speaker),
  family = "binomial", data = warlpiri, method = "Laplace")
warlpiri.lmer

## End(Not run)
```

---

weightRatings

*Subjective estimates of the weight of the referents of 81 English nouns*

---

## Description

Subjective estimates on a seven-point scale of the weight of the referents of 81 English nouns.

## Usage

```
data(weightRatings)
```

## Format

A data frame with 1620 observations on the following 7 variables.

Subject a factor with subjects as levels.

Rating a numeric vector.

Trial a numeric vector with the weight ratings.

Sex a factor with levels F and M.

Word a factor with words as levels.

Frequency a numeric vector with log-transformed lemma frequencies as available in the CELEX lexical database.

Class a factor with levels animal and plant.

## References

Data collected with Jen Hay, University of Canterbury, Christchurch, New Zealand, 2004.

**Examples**

```
## Not run:
data(weightRatings)
xyloless.fnc(Rating ~ Frequency | Subject, data = weightRatings,
  xlab = "log Frequency", ylab = "Weight Rating")

## End(Not run)
```

---

writtenVariationLijk *Variation in written Dutch in the use of the suffix -lijk*

---

**Description**

This dataset documents variation in the use of the 80 most frequent words ending in the suffix *-lijk* in written Dutch.

**Usage**

```
data(writtenVariationLijk)
```

**Format**

A data frame with 560 observations on the following 5 variables.

**Corpus** a factor with as levels the sampled newspapers: belang (Het Belang van Limburg), gazet (De Gazet van Antwerpen), laatnieu (Het Laatste Nieuws), limburg (De Limburger), nrc (NRC Handelsblad), stand (De Standaard), and tele (De Telegraaf).

**Word** a factor with the 80 most frequent words ending in *-lijk*.

**Count** a numeric vector with token counts in the CONDIV corpus.

**Country** a factor with levels Flanders and Netherlands.

**Register** a factor with levels National, Quality and Regional coding the type of newspaper.

**References**

Keune, K., Ernestus, M., Van Hout, R. and Baayen, R.H. (2005) Social, geographical, and register variation in Dutch: From written 'mogelijk' to spoken 'mok', *Corpus Linguistics and Linguistic Theory*, 1, 183-223.

**Examples**

```
## Not run:
data(writtenVariationLijk)

library(lme4, keep.source=FALSE)

writtenVariationLijk.lmer = lmer(Count ~ Country * Register + (1|Word),
  data = writtenVariationLijk, family = "poisson", method = "Laplace")
```

```
writtenVariationLijk.lmerA = lmer(Count ~ Country * Register + (Country|Word),
  data = writtenVariationLijk, family = "poisson", method = "Laplace")
anova(writtenVariationLijk.lmer, writtenVariationLijk.lmerA)

writtenVariationLijk.lmerA

## End(Not run)
```

---

xylowess.fnc

*Trellis scatterplot with smoothers*

---

### Description

Convenience function for trellis scatterplots with smoothers added.

### Usage

```
xylowess.fnc(fmla, data,
  span = 2/3, symbolcolor = "darkgrey",
  linecolor = "blue", xlabel = "", ylabel = "", ...)
```

### Arguments

fmla	A formula.
data	A dataframe.
span	Span for the smoother.
symbolcolor	Color for plot symbols.
linecolor	Color for smoother.
xlabel	Label for horizontal axis.
ylabel	Label for vertical axis.
...	Arguments to be passed to methods.

### Value

A trellis scatterplot matrix with smoothers is shown on the graphics device.

### Author(s)

R. H. Baayen

### See Also

See also [xyplot](#).

**Examples**

```
## Not run:
data(weightRatings)
xyloless.fnc(Rating ~ Frequency | Subject, data = weightRatings,
  xlab = "log Frequency", ylab = "Weight Rating")

## End(Not run)
```

---

yule.fnc

*Yule's characteristic constant K*

---

**Description**

This function calculates Yule's characteristic constant K given a frequency spectrum.

**Usage**

```
yule.fnc(spect)
```

**Arguments**

spect            A frequency spectrum as generated by `spectrum.fnc`.

**Value**

Yule's characteristic constant K

**Author(s)**

R. H. Baayen

**References**

Yule, G. U. (1944) *The Statistical Study of Literary Vocabulary*, Cambridge: Cambridge University Press.

Baayen, R. H. (2001) *Word Frequency Distributions*, Dordrecht: Kluwer.

**See Also**

See also [spectrum.fnc](#) and [growth.fnc](#).

**Examples**

```
## Not run:
data(alice)
yule.fnc(spectrum.fnc(alice))

## End(Not run)
```

---

zipf.fnc	<i>Zipf's rank frequency distribution</i>
----------	---

---

### Description

This function calculates Zipf's rank-frequency distribution for a text vector, and optionally produces the rank-frequency plot.

### Usage

```
zipf.fnc(text, plot = FALSE)
```

### Arguments

text	A character vector containing a text.
plot	Logical, if TRUE, a rank-frequency plot is shown on the graphics device.

### Value

A data frame with variables

frequency	Word Frequencies, ordered from large to small.
freqOfFreq	Frequencies of word frequencies.
rank	Zipf rank.

### Author(s)

R. H. Baayen

### References

Zipf, G. K. (1935) *The Psycho-Biology of Language*, Boston: Houghton Mifflin.

Zipf, G. K. (1949) *Human Behavior and the Principle of the Least Effort. An Introduction to Human Ecology*, New York: Hafner.

Baayen, R. H. (2001) *Word Frequency Distributions*, Dordrecht: Kluwer.

### See Also

See also [growth.fnc](#).

### Examples

```
## Not run:  
data(alice)  
alice.zipf = zipf.fnc(alice, plot = TRUE)  
head(alice.zipf)  
  
## End(Not run)
```

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