

# Package ‘nlmeU’

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**Version** 0.71.7

**Title** Functions and Data Supporting 'Linear Mixed-Effects Models: A Step-by-Step Approach'

**Description** Provides functions and datasets to support the book by Galecki and Burzykowski (2013), 'Linear Mixed-Effects Models: A Step-by-Step Approach', Springer. Includes functions for power calculations, log-likelihood contributions, and data simulation for linear mixed-effects models.

**Imports** nlme, stats

**Suggests** reshape, WWGbook, lattice, ellipse, roxygen2, testthat

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**License** GPL-2

**URL** <https://github.com/agalecki/nlmeU>

**Depends** R (>= 3.5.0)

**NeedsCompilation** no

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nlmeU-package	<i>nlmeU: Datasets and Utility Functions Enhancing Functionality of 'nlme' Package</i>
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## Description

Provides functions and datasets to support the book by Galecki and Burzykowski (2013), "Linear Mixed-Effects Models Using R: A Step-by-Step Approach", Springer. Package includes functions for power calculations, log-likelihood contributions, and data simulation for linear mixed-effects models.

## Details

This package provides datasets and utility functions to complement the nlme package, including functions like `logLik1`, `Pwr`, and `simulateY`. It also includes datasets such as `armd`, `armd0`, `armd.wide`, `fcats`, `prt.fiber`, `prt.subjects`, and `SIIdata`.

## Author(s)

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## See Also

Useful links:

- <https://github.com/agalecki/nlmeU>

---

armd	<i>armd Data (867 x 8)</i>
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---

### Description

Data from Age-Related Macular Degeneration (ARMD) clinical trial

### Usage

```
armd
```

### Format

A data frame with 867 rows and 8 columns:

**subject** A factor with 234 levels: 1, 2, 3, 4, 6, ..., 240

**treat.f** A factor with 2 levels: Placebo, Active

**visual0** An integer vector with values ranging from 20 to 85

**miss.pat** A factor with 8 levels: ----, ---X, --X-, --XX, -XX-, ..., X-XX

**time.f** A factor with 4 levels: 4wks, 12wks, 24wks, 52wks

**time** A numeric vector with values 4, 12, 24, 52

**visual** An integer vector with values ranging from 3 to 85

**tp** A numeric vector with values 1, 2, 3, 4 corresponding to time points 4, 12, 24, 52, respectively

### Details

The ARMD data arise from a randomized multi-center clinical trial comparing an experimental treatment (interferon-alpha) versus placebo for patients diagnosed with ARMD.

### Source

Pharmacological Therapy for Macular Degeneration Study Group (1997). Interferon alpha-IIA is ineffective for patients with choroidal neovascularization secondary to age-related macular degeneration. Results of a prospective randomized placebo-controlled clinical trial. Archives of Ophthalmology, 115, 865-872.

### See Also

[armd0](#), [armd.wide](#)

### Examples

```
library(nlmeU)
data(armd)
summary(armd)
```

---

armd.wide	<i>armd.wide Data (240 x 10)</i>
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---

### Description

Data from Age-Related Macular Degeneration (ARMD) clinical trial

### Usage

```
armd.wide
```

### Format

A data frame with 240 rows and 10 columns:

**subject** A factor with 240 levels: 1, 2, 3, 4, 5, ..., 240

**lesion** An integer vector with values 1, 2, 3, 4

**line0** An integer vector with values ranging from 5 to 17

**visual0** An integer vector with values of visual acuity measured at baseline ranging from 20 to 85

**visual4** An integer vector with values of visual acuity measured at 4 weeks ranging from 12 to 84

**visual12** An integer vector with values of visual acuity measured at 12 weeks ranging from 3 to 85

**visual24** An integer vector with values of visual acuity measured at 24 weeks ranging from 5 to 85

**visual52** An integer vector with values of visual acuity measured at 52 weeks from 4 to 85

**treat.f** A factor with 2 levels: Placebo, Active

**miss.pat** A factor with 9 levels: ----, ---X, --X-, --XX, -XX-, ..., XXXX

### Details

The ARMD data arise from a randomized multi-center clinical trial comparing an experimental treatment (interferon-alpha) versus placebo for patients diagnosed with ARMD.

### Source

Pharmacological Therapy for Macular Degeneration Study Group (1997). Interferon alpha-IIA is ineffective for patients with choroidal neovascularization secondary to age-related macular degeneration. Results of a prospective randomized placebo-controlled clinical trial. Archives of Ophthalmology, 115, 865-872.

### See Also

[armd](#), [armd0](#)

### Examples

```
data(armd.wide, package = "nlmeU")
summary(armd.wide)
```

---

armd0	<i>armd0 Data (1107 x 8)</i>
-------	------------------------------

---

### Description

Data from Age-Related Macular Degeneration (ARMD) clinical trial

### Usage

```
armd0
```

### Format

A data frame with 1107 rows and 8 columns:

**subject** A factor with 240 levels: 1, 2, 3, 4, 5, ...

**treat.f** A factor with 2 levels: Placebo, Active

**visual0** An integer vector with values from 20 to 85

**miss.pat** A factor with 9 levels: ----, ---X, --X-, --XX, -XX-, ...

**time.f** A factor with 5 levels: Baseline, 4wks, 12wks, 24wks, 52wks

**time** A numeric vector with values from 0 to 52

**visual** An integer vector with values from 3 to 85

**tp** A numeric vector with values from 0 to 4

### Details

The ARMD data arise from a randomized multi-center clinical trial comparing an experimental treatment (interferon-alpha) versus placebo for patients diagnosed with ARMD.

### Source

Pharmacological Therapy for Macular Degeneration Study Group (1997). Interferon alpha-IIA is ineffective for patients with choroidal neovascularization secondary to age-related macular degeneration. Results of a prospective randomized placebo-controlled clinical trial. Archives of Ophthalmology, 115, 865-872.

### See Also

[armd](#), [armd.wide](#)

### Examples

```
data(armd0, package = "nlmeU")
summary(armd0)
```

---

fcats	<i>fcats Data (4851 x 3)</i>
-------	------------------------------

---

### Description

Data from Flemish Community Attainment-Targets (FCAT) Study

### Usage

fcats

### Format

A data frame with 4851 rows and 3 columns:

**target** A factor with 9 levels: T1(4), T2(6), T3(8), T4(5), T5(9), ..., T9(5)

**id** A factor with 539 levels: 1, 2, 3, 4, 5, ..., 539

**scorec** An integer vector with values from 0 to 9

### Details

An educational study, in which elementary school graduates were evaluated with respect to reading comprehension in Dutch. Pupils from randomly selected schools were assessed for a set of nine attainment targets. The dataset is an example of grouped data, for which the grouping factors are crossed.

### Source

Janssen, R., Tuerlinckx, F., Meulders, M., & De Boeck, P. (2000). A hierarchical IRT model for criterion-referenced measurement. *Journal of Educational and Behavioral Statistics*, 25(3), 285.

### Examples

```
data(fcats, package = "nlmeU")
summary(fcats)
```

---

`logLik1`*Calculates contribution of one subject to the log-likelihood*

---

**Description**

This function is generic; method functions can be written to handle specific classes of objects.

**Usage**

```
logLik1(modfit, dt1, dtInit)
```

**Arguments**

<code>modfit</code>	an object representing a model fitted to data using maximum likelihood estimation.
<code>dt1</code>	a data frame with data for one subject, for whom the log-likelihood function is to be evaluated.
<code>dtInit</code>	an optional auxiliary data frame.

**Details**

Calculates contribution of one subject to the log-likelihood

This function is generic; method functions can be written to handle specific classes of objects.

**Value**

Numeric scalar value representing the contribution of a given subject to the overall log-likelihood returned by `logLik`.

**Author(s)**

Andrzej Galecki and Tomasz Burzykowski

**References**

Galecki, A., & Burzykowski, T. (2013). *\*Linear Mixed-Effects Models: A Step-by-Step Approach\**. Springer.

**Examples**

```
library(nlme)
logLik(fm1 <- lme(distance ~ age, data = Orthodont)) # random is ~ age
dt1 <- subset(Orthodont, Subject == "M01")
logLik1(fm1, dt1)
```

---

logLik1.lme	<i>Calculates contribution of one subject to the log-likelihood for an lme object</i>
-------------	---

---

## Description

This is a method for the `logLik1` generic function.

## Usage

```
## S3 method for class 'lme'  
logLik1(modfit, dt1, dtInit)
```

## Arguments

<code>modfit</code>	an <code>lme</code> object representing a model fitted using maximum likelihood.
<code>dt1</code>	a data frame with data for one subject, for whom the log-likelihood function is to be evaluated.
<code>dtInit</code>	an optional auxiliary data frame.

## Details

Calculates the profile likelihood (with beta profiled out) for one subject. Data with one level of grouping only. The correlation component in `modelStruct` is not implemented.

## Value

Numeric scalar value representing the contribution of a given subject to the overall log-likelihood returned by `logLik` applied to an `lme` object defined by the `modfit` argument.

## Author(s)

Andrzej Galecki and Tomasz Burzykowski

## Examples

```
library(nlme)  
data(armd, package = "nlmeU")  
lm3.form <- visual ~ visual0 + time + treat.f  
fm16.5ml <- lme(lm3.form,  
               random = list(subject = pdDiag(~time)),  
               weights = varPower(form = ~time),  
               data = armd, method = "ML") # M16.5  
df1 <- subset(armd, subject = "1") # Panel R20.7  
logLik1(fm16.5ml, df1)
```



---

missPat	<i>Extract pattern of missing data</i>
---------	--

---

### Description

This function compactly presents the pattern of missing data in a given vector, matrix, or data frame.

### Usage

```
missPat(..., symbols = c("X", "-"), collapse = "", missData = FALSE)
```

### Arguments

...	one or more vectors, matrices, or data frames, compatible for column-wise binding.
symbols	vector containing two single characters used to indicate NA and non-NA values. By default, <code>c("X", "-")</code> .
collapse	an optional character string used in the internal call to <code>paste()</code> to separate results. By default, <code>""</code> .
missData	logical. If TRUE, a data frame with the pattern of missing values is saved in the <code>missData</code> attribute of the returned vector. By default, FALSE.

### Value

Character vector with as many elements as the length of vector(s) or number of rows in matrices/data frames in `...`. Attribute `cnames` contains names of vectors/columns/variables. Optional attribute `missData` contains a data frame with the missing pattern.

### Author(s)

Andrzej Galecki and Tomasz Burzykowski

### See Also

[armd.wide](#)

### Examples

```
## Not run:
data(armd.wide, package = "nlmeU")
dtf <- armd.wide[, c("visual12", "visual24", "visual52")]
missPat(dtf, symbols = c("?", "+"))

## End(Not run)
```

---

print.Pwr                      *Print method for Pwr objects*

---

**Description**

Prints the results of power calculations for objects of class Pwr.lme.

**Usage**

```
## S3 method for class 'Pwr'  
print(x, verbose = attr(x, "verbose"), ...)
```

**Arguments**

x	an object of class Pwr.lme.
verbose	logical. If TRUE, additional information is printed. By default, uses the verbose attribute of x.
...	additional arguments passed to the method.

**Value**

Prints the power calculation results and returns NULL invisibly.

**Author(s)**

Andrzej Galecki and Tomasz Burzykowski

---

prt                              *prt Data (2471 x 9)*

---

**Description**

Data from a Progressive Resistance Randomized Trial.

**Usage**

```
prt
```

**Format**

A data frame with 2471 rows and 9 columns:

**id** A factor with 63 levels: 5, 10, 15, 20, 25, ..., 520 (subject id)

**prt.f** A factor with 2 levels: High, Low, i.e., training (intervention) intensity

**age.f** A factor with 2 levels: Young, Old (stratifying variable)

**sex.f** A factor with 2 levels: Female, Male (stratifying variable)

**bmi** A numeric vector with values of BMI at baseline ranging from 18.36 to 32.29

**iso.fo** A numeric vector with values of isometric force ranging from 0.16 to 2.565

**spec.fo** A numeric vector with values of specific force ranging from 80.5 to 290

**occ.f** A factor with 2 levels: Pre, Pos, i.e., pre- and post-intervention

**fiber.f** A factor with 2 levels: Type 1, Type 2, i.e., Type 1 and Type 2 muscle fiber

**Details**

Data frame prt was obtained by merging prt.subjects and prt.fiber.

**Source**

Clafin, D.R., Larkin, L.M., Cederna, P.S., Horowitz, J.F., Alexander, N.B., Cole, N.M., Galecki, A.T., Chen, S., Nyquist, L.V., Carlson, B.M., Faulkner, J.A., & Ashton-Miller, J.A. (2011). Effects of high- and low-velocity resistance training on the contractile properties of skeletal muscle fibers from young and older humans. *Journal of Applied Physiology*, 111, 1021-1030.

**See Also**

[prt.fiber](#), [prt.subjects](#)

**Examples**

```
data(prt, package = "nlmeU")
summary(prt)
```

---

 prt.fiber

*prt.fiber Data (2471 x 5)*

---

**Description**

Data from a Progressive Resistance Randomized Trial.

**Usage**

```
prt.fiber
```

**Format**

A data frame with 2471 rows and 5 columns:

**id** A factor with 63 levels: 5, 10, 15, 20, 25, ..., 520

**iso.fo** A numeric vector with values of isometric force ranging from 0.16 to 2.565

**spec.fo** A numeric vector with values of specific force ranging from 80.5 to 290

**occ.f** A factor with 2 levels: Pre, Pos, i.e., pre- and post-intervention

**fiber.f** A factor with 2 levels: Type 1, Type 2, i.e., Type 1 and Type 2 muscle fiber

**Details**

PRT trial was aimed for devising evidence-based methods for improving and measuring the mobility and muscle power of elderly men and women.

**Source**

Clafin, D.R., Larkin, L.M., Cederna, P.S., Horowitz, J.F., Alexander, N.B., Cole, N.M., Galecki, A.T., Chen, S., Nyquist, L.V., Carlson, B.M., Faulkner, J.A., & Ashton-Miller, J.A. (2011). Effects of high- and low-velocity resistance training on the contractile properties of skeletal muscle fibers from young and older humans. *Journal of Applied Physiology*, 111, 1021-1030.

**See Also**

[prt](#), [prt.subjects](#)

**Examples**

```
data(prt.fiber, package = "nlmeU")
summary(prt.fiber)
```

---

prt.subjects

*prt.subjects Data (63 x 5)*

---

**Description**

Data from a Progressive Resistance Randomized Trial.

**Usage**

```
prt.subjects
```

**Format**

A data frame with 63 rows and 5 columns:

**id** A factor with 63 levels: 5, 10, 15, 20, 25, ...

**prt.f** A factor with 2 levels: High, Low

**age.f** A factor with 2 levels: Young, Old

**sex.f** A factor with 2 levels: Female, Male

**bmi** A numeric vector with values from 18.4 to 32.3

**Details**

The working hypothesis was that a 12-week program of PRT would increase: (a) the power output of the overall musculature associated with movements of the ankles, knees, and hips; (b) the cross-sectional area and the force and power of permeabilized single fibers obtained from the vastus lateralis muscle; and (c) the ability of young and elderly men and women to safely arrest standardized falls. The training consisted of repeated leg extensions by shortening contractions of the leg extensor muscles against a resistance that was increased as the subject trained using a specially designed apparatus.

**Source**

Claffin, D.R., Larkin, L.M., Cederna, P.S., Horowitz, J.F., Alexander, N.B., Cole, N.M., Galecki, A.T., Chen, S., Nyquist, L.V., Carlson, B.M., Faulkner, J.A., & Ashton-Miller, J.A. (2011). Effects of high- and low-velocity resistance training on the contractile properties of skeletal muscle fibers from young and older humans. *Journal of Applied Physiology*, 111, 1021-1030.

**Examples**

```
data(prt.subjects, package = "nlmeU")
summary(prt.subjects)
```

---

Pwr

*Calculates power based on a model fit*

---

**Description**

This function is generic; method functions can be written to handle specific classes of objects.

**Usage**

```
Pwr(object, ...)
```

**Arguments**

**object** an object containing the results returned by a model fitting function (e.g., [lme](#)).

**...** some methods for this generic function may require additional arguments.

**Value**

Numeric scalar value.

**Author(s)**

Andrzej Galecki and Tomasz Burzykowski

**References**

Galecki, A., & Burzykowski, T. (2013). *\*Linear Mixed-Effects Models: A Step-by-Step Approach\**. Springer.

**See Also**

[Pwr.lme](#)

**Examples**

```
## Not run:
library(nlme)
fm1 <- lme(distance ~ age, data = Orthodont)
Pwr(fm1)

## End(Not run)
```

---

Pwr.lme

*Performs power calculations for an lme object*

---

**Description**

This is a method for the [Pwr](#) generic function. It works for the example given in Galecki and Burzykowski (2013) but may require additional testing, especially for post-hoc power analysis.

**Usage**

```
## S3 method for class 'lme'
Pwr(
  object,
  ...,
  type = c("sequential", "marginal"),
  Terms,
  L,
  verbose = FALSE,
  ddf = numeric(0),
  alpha = 0.05,
  altB = NULL,
  tol = 1e-10
)
```

**Arguments**

object	an <a href="#">lme</a> object containing a model fit, which provides information needed for power calculations.
...	some additional arguments may be required.
type	an optional character string specifying the type of sum of squares to be used in F-tests needed for power calculations. Syntax is the same as for <a href="#">anova.lme</a> in the nlme package.
Terms	an optional integer or character vector specifying which terms in the model should be jointly tested to be zero using a Wald F-test. See <a href="#">anova.lme</a> in the nlme package for details.
L	an optional numeric vector or array specifying linear combinations of the coefficients in the model that should be tested to be zero. See <a href="#">anova.lme</a> in the nlme package for details.
verbose	logical. If TRUE, additional information is provided. See <a href="#">anova.lme</a> in the nlme package for details.
ddf	numeric scalar value. Redefines the default number of denominator degrees of freedom.
alpha	numeric scalar value. By default, 0.05.
altB	matrix or vector containing alternative values for beta parameters.
tol	numeric scalar value for numerical tolerance.

**Value**

A data frame inheriting from class `Pwr.lme`.

**Author(s)**

Andrzej Galecki and Tomasz Burzykowski

**References**

Galecki, A., & Burzykowski, T. (2013). *\*Linear Mixed-Effects Models: A Step-by-Step Approach\**. Springer.

**See Also**

[anova.lme](#)

---

`runScript`*Execute scripts from Galecki and Burzykowski (2013)*

---

**Description**

Executes scripts from the book by Galecki and Burzykowski (2013). If called without arguments, it prints a list of available scripts.

**Usage**

```
runScript(  
  script = NA,  
  package = "nlmeU",  
  subdir = "scriptsR4.5.1",  
  echo = TRUE  
)
```

**Arguments**

<code>script</code>	character string containing the name of the script to be executed. By default, NA.
<code>package</code>	character string containing the package name. By default, "nlmeU".
<code>subdir</code>	subdirectory containing scripts. By default, "scriptsR2.15.0".
<code>echo</code>	logical. If TRUE, the script is executed with output printed. Used by <code>source()</code> . By default, TRUE.

**Value**

The script is executed, and results are printed. If `script` is NA, a list of available scripts is printed.

**Author(s)**

Andrzej Galecki and Tomasz Burzykowski

**References**

Galecki, A., & Burzykowski, T. (2013). *\*Linear Mixed-Effects Models: A Step-by-Step Approach\**. Springer.

**Examples**

```
runScript()
```



SIIdata

*SIIdata Data (1190 x 12)***Description**

Data from Study of Instructional Improvement Project

**Usage**

SIIdata

**Format**

A data frame with 1190 rows and 12 columns:

**sex** A factor with 2 levels: M, F, i.e., males and females, respectively

**minority** A factor with 2 levels: Mnr<sub>t</sub>=No, Mnr<sub>t</sub>=Yes. An indicator variable for the minority status

**mathkind** An integer vector with values from 290 to 629. This is pupil's math score in the spring of the kindergarten year

**mathgain** An integer vector with values from -110 to 253. Number represents pupil's gain in the math achievement score from the spring of kindergarten to the spring of first grade

**ses** A numeric vector with values from -1.61 to 3.21. Value represents socioeconomic status

**yearstea** A numeric vector with values from 0 to 40. It is number of years of teacher's experience in teaching in the first grade

**mathknow** A numeric vector with values from -2.5 to 2.61. Number represents teacher's knowledge of the first-grade math contents (higher values indicate a higher knowledge of the contents)

**housepov** A numeric vector containing proportion of households in the neighborhood of the school below the poverty level with values ranging from 0.012 to 0.564

**mathprep** A numeric vector with values from 1 to 6. Contains the number of preparatory courses on the first-grade math contents and methods followed by the teacher

**classid** A factor with 312 levels: 1, 2, 3, 4, 5, ..., 312. Classroom's id

**schoolid** A factor with 107 levels: 1, 2, 3, 4, 5, ..., 107. School's id

**childid** A factor with 1190 levels: 1, 2, 3, 4, 5, ..., 1190. Pupil's id

**Details**

The SII Project was carried out to assess the math achievement scores of first- and third-grade pupils in randomly selected classrooms from a national US sample of elementary schools (Hill et al, 2005). Data were also analyzed in West et al, 2007. The outcome of interest is `mathgain` variable. Data were created based on `classroom` data from `WWGbook` package.

**Source**

Hill, H., Rowan, B., and Ball, D. (2005). Effect of teachers' mathematical knowledge for teaching on student achievement. *American Educational Research Journal*, 42, 371-406. West, B. T., Welch, K. B., and Galecki, A. T. (2007). *Linear Mixed Models: A Practical Guide Using Statistical Software*. Chapman and Hall/CRC.

**Examples**

```
data(SIIdata, package = "nlmeU")
summary(SIIdata)
```

---

 simulateY

---

*Simulates values of the dependent variable based on a model fit*


---

**Description**

This function is generic; method functions can be written to handle specific classes of objects.

**Usage**

```
simulateY(object, nsim = 1, seed = NULL, ..., verbose = FALSE, sigma)
```

**Arguments**

object	an object with a model fit for which the dependent variable is to be simulated.
nsim	number of simulations. By default, nsim = 1.
seed	integer scalar used to initiate the random number generator.
...	some methods for this generic function may require additional arguments.
verbose	logical. If TRUE, basic information about arguments is provided. By default set to FALSE.
sigma	numeric scalar. Allows simulations employing an alternative value of the scale parameter.

**Value**

Numeric matrix with the number of columns determined by the nsim argument.

**Author(s)**

Andrzej Galecki and Tomasz Burzykowski

**Examples**

```
## Not run:
library(nlme)
fm1 <- lme(distance ~ age, data = Orthodont)
simulateY(fm1)

## End(Not run)
```

---

simulateY.lme	<i>Simulates values for an lme object</i>
---------------	---

---

**Description**

This is a method for the [simulateY](#) generic function.

**Usage**

```
## S3 method for class 'lme'
simulateY(
  object,
  nsim = 1,
  seed = as.integer(runif(1, 0, .Machine$integer.max)),
  ...,
  verbose = FALSE,
  sigma
)
```

**Arguments**

object	an <a href="#">lme</a> object with a model fit for which the dependent variable is to be simulated.
nsim	number of simulations. By default, nsim = 1.
seed	integer scalar used to initiate the random number generator.
...	some methods for this generic function may require additional arguments.
verbose	logical. If TRUE, basic information about arguments is provided. By default set to FALSE.
sigma	numeric scalar. Allows simulations employing an alternative value of the scale parameter.

**Details**

Simulates values of the dependent variable for a fitted [lme](#) model. Data with one level of grouping only.

**Value**

Numeric matrix with the number of columns determined by the nsim argument.

**Author(s)**

Andrzej Galecki and Tomasz Burzykowski

**Examples**

```
## Not run:  
library(nlme)  
fm1 <- lme(distance ~ age, data = Orthodont)  
simulateY(fm1)  
  
## End(Not run)
```

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