

# Package ‘MEMSS’

January 20, 2025

**Version** 0.9-3

**Date** 2019-02-07

**Title** Data Sets from Mixed-Effects Models in S

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**Description** Data sets and sample analyses from Pinheiro and Bates,  
``Mixed-effects Models in S and S-PLUS" (Springer, 2000).

**Depends** R(>= 2.12.0), lme4 (>= 0.999375-36)

**Suggests** lattice

**LazyData** yes

**License** GPL (>= 2)

**NeedsCompilation** no

**Repository** CRAN

**Date/Publication** 2019-02-08 05:13:31 UTC

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Alfalfa

*Split-Plot Experiment on Varieties of Alfalfa*

---

### Description

The Alfalfa data frame has 72 rows and 4 columns.

### Format

This data frame contains the following columns:

**Variety** a factor with levels Cossack, Ladak, and Ranger

**Date** a factor with levels None S1 S20 07

**Block** a factor with levels A to F

**Yield** a numeric vector

### Details

These data are described in Snedecor and Cochran (1980) as an example of a split-plot design. The treatment structure used in the experiment was a  $3 \times 4$  full factorial, with three varieties of alfalfa and four dates of third cutting in 1943. The experimental units were arranged into six blocks, each subdivided into four plots. The varieties of alfalfa (*Cossac*, *Ladak*, and *Ranger*) were assigned randomly to the blocks and the dates of third cutting (*None*, *S1*—September 1, *S20*—September 20, and *O7*—October 7) were randomly assigned to the plots. All four dates were used on each block.

### Source

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York. (Appendix A.1)

Snedecor, G. W. and Cochran, W. G. (1980), *Statistical Methods (7th ed)*, Iowa State University Press, Ames, IA

### Examples

```
str(Alfalfa)
(m1 <- lmer(Yield ~ Variety * Date + (1|Block), Alfalfa, verbose = TRUE))
```

---

Assay

*Bioassay on Cell Culture Plate*

---

### Description

The Assay data frame has 60 rows and 4 columns.

### Format

This data frame contains the following columns:

**Block** an factor with levels A and B identifying the block of the well

**sample** a factor with levels a to f identifying the sample corresponding to the well

**dilut** an ordered factor with levels 1 to 5 indicating the dilution applied to the well

**logDens** a numeric vector of the log-optical density

### Details

These data, courtesy of Rich Wolfe and David Lansky from Searle, Inc., come from a bioassay run on a 96-well cell culture plate. The assay is performed using a split-block design. The 8 rows on the plate are labeled A–H from top to bottom and the 12 columns on the plate are labeled 1–12 from left to right. Only the central 60 wells of the plate are used for the bioassay (the intersection of rows B–G and columns 2–11). There are two blocks in the design: Block A contains columns 2–6 and Block B contains columns 7–11. Within each block, six samples are assigned randomly to rows and five (serial) dilutions are assigned randomly to columns. The response variable is the logarithm of the optical density. The cells are treated with a compound that they metabolize to produce the stain. Only live cells can make the stain, so the optical density is a measure of the number of cells that are alive and healthy.

**Source**

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York. (Appendix A.2)

**Examples**

```
str(Assay)
m1 <- lmer(logDens ~ sample * dilut + (1|Block) + (1|Block:sample) +
           (1|Block:dilut), Assay, verbose = TRUE)
print(m1, corr = FALSE)
anova(m1)
m2 <- lmer(logDens ~ sample + dilut + (1|Block) + (1|Block:sample) +
           (1|Block:dilut), Assay, verbose = TRUE)
print(m2, corr = FALSE)
anova(m2)
m3 <- lmer(logDens ~ sample + dilut + (1|Block) + (1|Block:sample),
           Assay, verbose = TRUE)
print(m3, corr = FALSE)
anova(m3)
anova(m2, m3)
```

---

 BodyWeight

*Rat weight over time for different diets*


---

**Description**

The BodyWeight data frame has 176 rows and 4 columns.

**Format**

This data frame contains the following columns:

**weight** a numeric vector giving the body weight of the rat (grams).

**Time** a numeric vector giving the time at which the measurement is made (days).

**Rat** an factor with levels A to P identifying the rat whose weight is measured.

**Diet** a factor with levels a to c indicating the diet that the rat receives.

**Details**

Hand and Crowder (1996) describe data on the body weights of rats measured over 64 days. These data also appear in Table 2.4 of Crowder and Hand (1990). The body weights of the rats (in grams) are measured on day 1 and every seven days thereafter until day 64, with an extra measurement on day 44. The experiment started several weeks before “day 1.” There are three groups of rats, each on a different diet.

**Source**

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York. (Appendix A.3)

Crowder, M. and Hand, D. (1990), *Analysis of Repeated Measures*, Chapman and Hall, London.

Hand, D. and Crowder, M. (1996), *Practical Longitudinal Data Analysis*, Chapman and Hall, London.

**Examples**

```
str(BodyWeight)
```

---

Cefamandole

*Pharmacokinetics of Cefamandole*

---

**Description**

The Cefamandole data frame has 84 rows and 3 columns.

**Format**

This data frame contains the following columns:

**Subject** a factor giving the subject from which the sample was drawn.

**Time** a numeric vector giving the time at which the sample was drawn (minutes post-injection).

**conc** a numeric vector giving the observed plasma concentration of cefamandole (mcg/ml).

**Details**

Davidian and Giltinan (1995, 1.1, p. 2) describe data obtained during a pilot study to investigate the pharmacokinetics of the drug cefamandole. Plasma concentrations of the drug were measured on six healthy volunteers at 14 time points following an intravenous dose of 15 mg/kg body weight of cefamandole.

**Source**

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York. (Appendix A.4)

Davidian, M. and Giltinan, D. M. (1995), *Nonlinear Models for Repeated Measurement Data*, Chapman and Hall, London.

### Examples

```
require(lattice)
str(Cefamandole)
xyplot(conc ~ Time, Cefamandole, groups = Subject, type = c("g", "b"),
       aspect = 'xy', scales = list(y = list(log = 2)),
       auto.key = list(space = "right", lines= TRUE))
xyplot(conc ~ Time|Subject, Cefamandole, type = c("g", "b"),
       index.cond = function(x,y) min(y), aspect = 'xy',
       scales = list(y = list(log = 2)))
#fm1 <- nlsList(SSbiexp, data = Cefamandole)
```

---

CO2

*Carbon Dioxide uptake in grass plants*


---

### Description

The CO2 data frame has 84 rows and 5 columns of data from an experiment on the cold tolerance of the grass species *Echinochloa crus-galli*.

### Usage

CO2

### Format

This data frame contains the following columns:

**Plant** an factor giving a unique identifier for each plant.

**Type** a factor with levels Quebec Mississippi giving the origin of the plant

**Treatment** a factor with levels nonchilled chilled

**conc** a numeric vector of ambient carbon dioxide concentrations (mL/L).

**uptake** a numeric vector of carbon dioxide uptake rates ( $\mu\text{mol}/m^2 \text{ sec}$ ).

### Details

The  $CO_2$  uptake of six plants from Quebec and six plants from Mississippi was measured at several levels of ambient  $CO_2$  concentration. Half the plants of each type were chilled overnight before the experiment was conducted.

### Source

Potvin, C., Lechowicz, M. J. and Tardif, S. (1990) "The statistical analysis of ecophysiological response curves obtained from experiments involving repeated measures", *Ecology*, **71**, 1389–1400.

Pinheiro, J. C. and Bates, D. M. (2000) *Mixed-effects Models in S and S-PLUS*, Springer.

**Examples**

```

require(stats); require(graphics)
coplot(uptake ~ conc | Plant, data = C02, show.given = FALSE, type = "b")
## fit the data for the first plant
fm1 <- nls(uptake ~ SSasym(conc, Asym, lrc, c0),
  data = C02, subset = Plant == 'Qn1')
summary(fm1)
## fit each plant separately
fm1list <- list()
for (pp in levels(C02$Plant)) {
  fm1list[[pp]] <- nls(uptake ~ SSasym(conc, Asym, lrc, c0),
    data = C02, subset = Plant == pp)
}
## check the coefficients by plant
sapply(fm1list, coef)

```

Dialyzer

*High-Flux Hemodialyzer***Description**

The Dialyzer data frame has 140 rows and 5 columns.

**Format**

This data frame contains the following columns:

**Subject** a factor with levels A to T

**QB** a factor with levels 200 and 300 giving the bovine blood flow rate (dL/min).

**pressure** the transmembrane pressure (dmHg).

**rate** the hemodialyzer ultrafiltration rate (mL/hr).

**index** index of observation within subject—1 through 7.

**Details**

Vonesh and Carter (1992) describe data measured on high-flux hemodialyzers to assess their *in vivo* ultrafiltration characteristics. The ultrafiltration rates (in mL/hr) of 20 high-flux dialyzers were measured at seven different transmembrane pressures (in dmHg). The *in vitro* evaluation of the dialyzers used bovine blood at flow rates of either 200~dl/min or 300~dl/min. The data, are also analyzed in Littell, Milliken, Stroup, and Wolfinger (1996).

**Source**

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York. (Appendix A.6)

Vonesh, E. F. and Carter, R. L. (1992), Mixed-effects nonlinear regression for unbalanced repeated measures, *Biometrics*, **48**, 1-18.

Littell, R. C., Milliken, G. A., Stroup, W. W. and Wolfinger, R. D. (1996), *SAS System for Mixed Models*, SAS Institute, Cary, NC.

**Examples**

```
str(Dialyzer)
```

---

 Earthquake
 

---



---

*Earthquake Intensity*


---

**Description**

The Earthquake data frame has 182 rows and 5 columns.

**Format**

This data frame contains the following columns:

**Quake** a factor with levels A to U

**Richter** the intensity of the earthquake on the Richter scale

**distance** the distance from the seismological measuring station to the epicenter of the earthquake (km)

**soil** a factor with levels S (soil) and R (rock) giving the soil condition at the measuring station

**accel** maximum horizontal acceleration observed (g).

**Details**

Measurements recorded at available seismometer locations for 23 large earthquakes in western North America between 1940 and 1980. They were originally given in Joyner and Boore (1981); are mentioned in Brillinger (1987); and are analyzed in Davidian and Giltinan (1995).

**Source**

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York. (Appendix A.8)

Davidian, M. and Giltinan, D. M. (1995), *Nonlinear Models for Repeated Measurement Data*, Chapman and Hall, London.

Joyner and Boor (1981), Peak horizontal acceleration and velocity from strong-motion records including records from the 1979 Imperial Valley, California, earthquake, *Bulletin of the Seismological Society of America*, **71**, 2011-2038.

Brillinger, D. (1987), Comment on a paper by C. R. Rao, *Statistical Science*, **2**, 448-450.

**Examples**

```
str(Earthquake)
```

---

 ergoStool

*Ergometrics experiment with stool types*


---

**Description**

The ergoStool data frame has 36 rows and 3 columns.

**Format**

This data frame contains the following columns:

**effort** a numeric vector giving the effort (Borg scale) required to arise from a stool

**Type** a factor with levels T1, T2, T3, and T4 giving the stool type

**Subject** a factor with levels A to I

**Details**

Devore (2000) cites data from an article in *Ergometrics* (1993, pp. 519-535) on “The Effects of a Pneumatic Stool and a One-Legged Stool on Lower Limb Joint Load and Muscular Activity.”

**Source**

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York. (Appendix A.9)

Devore, J. L. (2000), *Probability and Statistics for Engineering and the Sciences (5th ed)*, Duxbury, Boston, MA.

**Examples**

```
options(show.signif.stars = FALSE)
str(ergoStool)
print(m1 <- lmer(effort ~ Type + (1|Subject), ergoStool), corr = FALSE)
anova(m1)
```

---

 Fatigue

*Cracks caused by metal fatigue*


---

**Description**

The Fatigue data frame has 262 rows and 3 columns.

**Format**

This data frame contains the following columns:

**Path** the test path (or test unit) identifier - a factor with levels A to U.

**cycles** number of test cycles at which the measurement is made (millions of cycles).

**relLength** relative crack length (dimensionless).

## Details

These data are given in Lu and Meeker (1993) where they state “We obtained the data in Table 1 visually from figure 4.5.2 on page 242 of Bogdanoff and Kozin (1985).” The data represent the growth of cracks in metal for 21 test units. An initial notch of length 0.90 inches was made on each unit which then was subjected to several thousand test cycles. After every 10,000 test cycles the crack length was measured. Testing was stopped if the crack length exceeded 1.60 inches, defined as a failure, or at 120,000 cycles.

## Source

Lu, C. Joseph , and Meeker, William Q. (1993), Using degradation measures to estimate a time-to-failure distribution, *Technometrics*, **35**, 161-174

## Examples

```
require(lattice)
str(Fatigue)
xyplot(reLength ~ cycles | Path, Fatigue, type = c("g", "b"),
       aspect = 'xy', xlab = "Number of test cycles (millions)",
       ylab = "Relative crack length (dimensionless)",
       layout = c(7,3))
```

---

Gasoline

*Refinery yield of gasoline*

---

## Description

The Gasoline data frame has 32 rows and 6 columns.

## Format

This data frame contains the following columns:

**yield** a numeric vector giving the percentage of crude oil converted to gasoline after distillation and fractionation

**endpoint** a numeric vector giving the temperature (degrees F) at which all the gasoline is vaporized

**Sample** the inferred crude oil sample number - a factor with levels A to J

**API** a numeric vector giving the crude oil gravity (degrees API)

**vapor** a numeric vector giving the vapor pressure of the crude oil (lbf/in<sup>2</sup>)

**ASTM** a numeric vector giving the crude oil 10% point ASTM—the temperature at which 10% of the crude oil has become vapor.

### Details

Prater (1955) provides data on crude oil properties and gasoline yields. Atkinson (1985) uses these data to illustrate the use of diagnostics in multiple regression analysis. Three of the covariates—API, vapor, and ASTM—measure characteristics of the crude oil used to produce the gasoline. The other covariate — endpoint—is a characteristic of the refining process. Daniel and Wood (1980) notice that the covariates characterizing the crude oil occur in only ten distinct groups and conclude that the data represent responses measured on ten different crude oil samples.

### Source

Prater, N. H. (1955), Estimate gasoline yields from crudes, *Petroleum Refiner*, **35** (5).  
 Atkinson, A. C. (1985), *Plots, Transformations, and Regression*, Oxford Press, New York.  
 Daniel, C. and Wood, F. S. (1980), *Fitting Equations to Data*, Wiley, New York  
 Venables, W. N. and Ripley, B. D. (1999) *Modern Applied Statistics with S-PLUS (3rd ed)*, Springer, New York.

### Examples

```
require(lattice)
str(Gasoline)
xyplot(yield ~ endpoint | Sample, Gasoline, aspect = 'xy',
       main = "Gasoline data", xlab = "Endpoint (degrees F)",
       ylab = "Percentage yield",
       type = c("g", "p", "r"),
       index.cond = function(x,y) coef(lm(y~x))[2],
       layout = c(5,2))
print(m1 <- lmer(yield ~ endpoint + (1|Sample), Gasoline), corr = FALSE)
m2 <- lmer(yield ~ endpoint + (endpoint|Sample), Gasoline, verbose = 1)
print(m2)
Gasoline$endptC <- with(Gasoline, endpoint - mean(endpoint))
m3 <- lmer(yield ~ endpoint + (endptC|Sample), Gasoline, verbose = 1)
print(m3)
xyplot(endptC ~ `(Intercept)`, ranef(m3)[[1]], type = c("g", "p", "r"),
       aspect = 1)
```

---

Glucose

*Glucose levels over time*

---

### Description

The Glucose data frame has 378 rows and 4 columns.

### Format

This data frame contains the following columns:

**Subject** a factor with levels A to F

**Time** a numeric vector

**conc** a numeric vector of glucose levels

**Meal** an ordered factor with levels 2am < 6am < 10am < 2pm < 6pm < 10pm

### Source

Hand, D. and Crowder, M. (1996), *Practical Longitudinal Data Analysis*, Chapman and Hall, London.

### Examples

```
require(lattice)
str(Glucose)
xyplot(conc ~ Time | Meal * Subject, Glucose)
```

---

Glucose2

*Glucose Levels Following Alcohol Ingestion*

---

### Description

The Glucose2 data frame has 196 rows and 4 columns.

### Format

This data frame contains the following columns:

**Subject** a factor with levels A to G

**Date** a factor with levels 1 2 indicating the occasion in which the experiment was conducted.

**Time** a numeric vector giving the time since alcohol ingestion (in min/10).

**glucose** a numeric vector giving the blood glucose level (in mg/dl).

### Details

Hand and Crowder (Table A.14, pp. 180-181, 1996) describe data on the blood glucose levels measured at 14 time points over 5 hours for 7 volunteers who took alcohol at time 0. The same experiment was repeated on a second date with the same subjects but with a dietary additive used for all subjects.

### Source

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York. (Appendix A.10)

Hand, D. and Crowder, M. (1996), *Practical Longitudinal Data Analysis*, Chapman and Hall, London.

### Examples

```
require(lattice)
str(Glucose2)
xyplot(glucose ~ Time | Subject, Glucose2, type = c("g", "b"),
       groups = Date, aspect = 'xy', layout = c(4,2),
       index.cond = function(x,y) max(y))
```

---

Gun

*Methods for firing naval guns*

---

### Description

The Gun data frame has 36 rows and 4 columns.

### Format

This data frame contains the following columns:

**rounds** a numeric vector

**Method** a factor with levels M1 and M2

**Team** an ordered factor with levels T1S < T3S < T2S < T1A < T2A < T3A < T1H < T3H < T2H

**Physique** an ordered factor with levels Slight < Average < Heavy

### Details

Hicks (p.180, 1993) reports data from an experiment on methods for firing naval guns. Gunners of three different physiques (slight, average, and heavy) tested two firing methods. Both methods were tested twice by each of nine teams of three gunners with identical physique. The response was the number of rounds fired per minute.

### Source

Hicks, C. R. (1993), *Fundamental Concepts in the Design of Experiments (4th ed)*, Harcourt Brace, New York.

### Examples

```
str(Gun)
```

---

IGF *Radioimmunoassay of IGF-I Protein*

---

**Description**

The IGF data frame has 237 rows and 3 columns.

**Format**

This data frame contains the following columns:

**Lot** an ordered factor giving the radioactive tracer lot.

**age** a numeric vector giving the age (in days) of the radioactive tracer.

**conc** a numeric vector giving the estimated concentration of IGF-I protein (ng/ml)

**Details**

Davidian and Giltinan (1995) describe data obtained during quality control radioimmunoassays for ten different lots of radioactive tracer used to calibrate the Insulin-like Growth Factor (IGF-I) protein concentration measurements.

**Source**

Davidian, M. and Giltinan, D. M. (1995), *Nonlinear Models for Repeated Measurement Data*, Chapman and Hall, London.

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York. (Appendix A.11)

**Examples**

```
str(IGF)
```

---

Machines *Productivity Scores for Machines and Workers*

---

**Description**

The Machines data frame has 54 rows and 3 columns.

**Format**

This data frame contains the following columns:

**Worker** an ordered factor giving the unique identifier for the worker.

**Machine** a factor with levels A, B, and C identifying the machine brand.

**score** a productivity score.

**Details**

Data on an experiment to compare three brands of machines used in an industrial process are presented in Milliken and Johnson (p. 285, 1992). Six workers were chosen randomly among the employees of a factory to operate each machine three times. The response is an overall productivity score taking into account the number and quality of components produced.

**Source**

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York. (Appendix A.14)

Milliken, G. A. and Johnson, D. E. (1992), *Analysis of Messy Data, Volume I: Designed Experiments*, Chapman and Hall, London.

**Examples**

```
str(Machines)
```

---

MathAchieve

*Mathematics achievement scores*

---

**Description**

The MathAchieve data frame has 7185 rows and 6 columns.

**Format**

This data frame contains the following columns:

**School** an ordered factor identifying the school that the student attends

**Minority** a factor with levels No Yes indicating if the student is a member of a minority racial group.

**Sex** a factor with levels Male Female

**SES** a numeric vector of socio-economic status.

**MathAch** a numeric vector of mathematics achievement scores.

**MEANSES** a numeric vector of the mean SES for the school.

**Details**

Each row in this data frame contains the data for one student.

**Examples**

```
str(MathAchieve)
```

---

MathAchSchool	<i>School demographic data for MathAchieve</i>
---------------	--

---

### Description

The MathAchSchool data frame has 160 rows and 7 columns.

### Format

This data frame contains the following columns:

**School** a factor giving the school on which the measurement is made.

**Size** a numeric vector giving the number of students in the school

**Sector** a factor with levels Public Catholic

**PRACAD** a numeric vector giving the percentage of students on the academic track

**DISCLIM** a numeric vector measuring the discrimination climate

**HIMINTY** a factor with levels 0 1

**MEANSES** a numeric vector giving the mean SES score.

### Details

These variables give the school-level demographic data to accompany the MathAchieve data.

### Examples

```
str(MathAchSchool)
```

---

Meat	<i>Tenderness of meat</i>
------	---------------------------

---

### Description

The Meat data frame has 30 rows and 4 columns.

### Format

This data frame contains the following columns:

**Storage** an ordered factor specifying the storage treatment - 1 (0 days), 2 (1 day), 3 (2 days), 4 (4 days), 5 (9 days), and 6 (18 days)

**score** a numeric vector giving the tenderness score of beef roast.

**Block** an ordered factor identifying the muscle from which the roast was extracted with levels II < V < I < III < IV

**Pair** an ordered factor giving the unique identifier for each pair of beef roasts with levels II-1 < ... < IV-1

**Details**

Cochran and Cox (section 11.51, 1957) describe data from an experiment conducted at Iowa State College (Paul, 1943) to compare the effects of length of cold storage on the tenderness of beef roasts. Six storage periods ranging from 0 to 18 days were used. Thirty roasts were scored by four judges on a scale from 0 to 10, with the score increasing with tenderness. The response was the sum of all four scores. Left and right roasts from the same animal were grouped into pairs, which were further grouped into five blocks, according to the muscle from which they were extracted. Different storage periods were applied to each roast within a pair according to a balanced incomplete block design.

**Source**

Cochran, W. G. and Cox, G. M. (1957), *Experimental Designs*, Wiley, New York.

**Examples**

```
str(Meat)
```

---

Milk	<i>Protein content of cows' milk</i>
------	--------------------------------------

---

**Description**

The Milk data frame has 1337 rows and 4 columns.

**Format**

This data frame contains the following columns:

**protein** a numeric vector giving the protein content of the milk.

**Time** a numeric vector giving the time since calving (weeks).

**Cow** an ordered factor giving a unique identifier for each cow.

**Diet** a factor with levels barley, barley+lupins, and lupins identifying the diet for each cow.

**Details**

Diggle, Liang, and Zeger (1994) describe data on the protein content of cows' milk in the weeks following calving. The cattle are grouped according to whether they are fed a diet with barley alone, with barley and lupins, or with lupins alone.

**Source**

Diggle, Peter J., Liang, Kung-Yee and Zeger, Scott L. (1994), *Analysis of longitudinal data*, Oxford University Press, Oxford.

**Examples**

```
str(Milk)
```

---

Muscle	<i>Contraction of heart muscle sections</i>
--------	---

---

### Description

The Muscle data frame has 60 rows and 3 columns.

### Format

This data frame contains the following columns:

**Strip** an ordered factor indicating the strip of muscle being measured.

**conc** a numeric vector giving the concentration of CaCl<sub>2</sub>

**length** a numeric vector giving the shortening of the heart muscle strip.

### Details

Baumann and Waldvogel (1963) describe data on the shortening of heart muscle strips dipped in a CaCl<sub>2</sub> solution. The muscle strips are taken from the left auricle of a rat's heart.

### Source

Baumann, F. and Waldvogel, F. (1963), La restitution pastsystolique de la contraction de l'oreillette gauche du rat. Effets de divers ions et de l'acetylcholine, *Helvetica Physiologica Acta*, **21**.

### Examples

```
str(Muscle)
```

---

Nitrendipene	<i>Assay of nitrendipene</i>
--------------	------------------------------

---

### Description

The Nitrendipene data frame has 89 rows and 4 columns.

### Format

This data frame contains the following columns:

**activity** a numeric vector

**NIF** a numeric vector

**Tissue** an ordered factor with levels 2 < 1 < 3 < 4

**log.NIF** a numeric vector

**Source**

Bates, D. M. and Watts, D. G. (1988), *Nonlinear Regression Analysis and Its Applications*, Wiley, New York.

**Examples**

```
str(Nitrendipene)
```

---

Oats

*Split-plot Experiment on Varieties of Oats*

---

**Description**

The Oats data frame has 72 rows and 4 columns.

**Format**

This data frame contains the following columns:

**Block** an ordered factor with levels VI < V < III < IV < II < I

**Variety** a factor with levels Golden Rain Marvellous Victory

**nitro** a numeric vector

**yield** a numeric vector

**Details**

These data have been introduced by Yates (1935) as an example of a split-plot design. The treatment structure used in the experiment was a  $3 \times 4$  full factorial, with three varieties of oats and four concentrations of nitrogen. The experimental units were arranged into six blocks, each with three whole-plots subdivided into four subplots. The varieties of oats were assigned randomly to the whole-plots and the concentrations of nitrogen to the subplots. All four concentrations of nitrogen were used on each whole-plot.

**Source**

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York. (Appendix A.15)

Venables, W. N. and Ripley, B. D. (1999) *Modern Applied Statistics with S-PLUS (3rd ed)*, Springer, New York.

**Examples**

```
str(Oats)
```

---

 Orange

*Growth of orange trees*


---

### Description

The Orange data frame has 35 rows and 3 columns of records of the growth of orange trees.

### Usage

Orange

### Format

This data frame contains the following columns:

**Tree** a factor indicating the tree on which the measurement is made.

**age** a numeric vector giving the age of the tree (days since 1968/12/31)

**circumference** a numeric vector of trunk circumferences (mm). This is probably “circumference at breast height”, a standard measurement in forestry.

### Source

Draper, N. R. and Smith, H. (1998), *Applied Regression Analysis (3rd ed)*, Wiley (exercise 24.N).

Pinheiro, J. C. and Bates, D. M. (2000) *Mixed-effects Models in S and S-PLUS*, Springer.

### Examples

```
require(lattice)
xyplot(circumference ~ age, Orange, groups = Tree, type = c("g", "b"),
       auto.key = list(space = "right", lines = TRUE), aspect = "xy",
       xlab = "Age (days since 1968/12/31)", ylab = "Circumference (mm)")
## Not run:
m1 <- nlmer(circumference ~ SSlogis(age, Asym, xmid, scal) ~ Asym|Tree,
           Orange, verbose = TRUE,
           start = c(Asym = 190, xmid = 730, scal = 350))
.Call("mer_optimize", m1, 1L, 1L, PACKAGE = "lme4")
print(m1)
ranef(m1)

## End(Not run)
```

---

Orthodont

*Growth curve data on an orthodontic measurement*

---

### Description

The Orthodont data frame has 108 rows and 4 columns of the change in an orthodontic measurement over time for several young subjects.

### Format

This data frame contains the following columns:

**distance** a numeric vector of distances from the pituitary to the pterygomaxillary fissure (mm). These distances are measured on x-ray images of the skull.

**age** a numeric vector of ages of the subject (yr).

**Subject** an ordered factor indicating the subject on which the measurement was made. The levels are labelled M01 to M16 for the males and F01 to F13 for the females. The ordering is by increasing average distance within sex.

**Sex** a factor with levels Male and Female

### Details

Investigators at the University of North Carolina Dental School followed the growth of 27 children (16 males, 11 females) from age 8 until age 14. Every two years they measured the distance between the pituitary and the pterygomaxillary fissure, two points that are easily identified on x-ray exposures of the side of the head.

### Source

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York. (Appendix A.17)

Potthoff, R. F. and Roy, S. N. (1964), "A generalized multivariate analysis of variance model useful especially for growth curve problems", *Biometrika*, **51**, 313–326.

### Examples

```
str(Orthodont)
```

---

Ovary

*Counts of Ovarian Follicles*

---

### Description

The Ovary data frame has 308 rows and 3 columns.

### Format

This data frame contains the following columns:

**Mare** an ordered factor indicating the mare on which the measurement is made.

**Time** time in the estrus cycle. The data were recorded daily from 3 days before ovulation until 3 days after the next ovulation. The measurement times for each mare are scaled so that the ovulations for each mare occur at times 0 and 1.

**follicles** the number of ovarian follicles greater than 10 mm in diameter.

### Details

Pierson and Ginther (1987) report on a study of the number of large ovarian follicles detected in different mares at several times in their estrus cycles.

### Source

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York. (Appendix A.18)

Pierson, R. A. and Ginther, O. J. (1987), Follicular population dynamics during the estrus cycle of the mare, *Animal Reproduction Science*, **14**, 219-231.

### Examples

```
str(Ovary)
```

---

Oxide

*Variability in Semiconductor Manufacturing*

---

### Description

The Oxide data frame has 72 rows and 5 columns.

**Format**

This data frame contains the following columns:

**Source** a factor with levels 1 and 2

**Lot** a factor giving a unique identifier for each lot.

**Wafer** a factor giving a unique identifier for each wafer within a lot.

**Site** a factor with levels 1, 2, and 3

**Thickness** a numeric vector giving the thickness of the oxide layer.

**Details**

These data are described in Littell et al. (1996, p. 155) as coming “from a passive data collection study in the semiconductor industry where the objective is to estimate the variance components to determine the assignable causes of the observed variability.” The observed response is the thickness of the oxide layer on silicon wafers, measured at three different sites of each of three wafers selected from each of eight lots sampled from the population of lots.

**Source**

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York. (Appendix A.20)

Littell, R. C., Milliken, G. A., Stroup, W. W. and Wolfinger, R. D. (1996), *SAS System for Mixed Models*, SAS Institute, Cary, NC.

**Examples**

```
str(Oxide)
```

---

 PBG

---

*Effect of Phenylbiguanide on Blood Pressure*


---

**Description**

The PBG data frame has 60 rows and 5 columns.

**Format**

This data frame contains the following columns:

**deltaBP** a numeric vector

**dose** a numeric vector

**Run** an ordered factor with levels T5 < T4 < T3 < T2 < T1 < P5 < P3 < P2 < P4 < P1

**Treatment** a factor with levels MDL 72222 P1acebo

**Rabbit** an ordered factor with levels 5 < 3 < 2 < 4 < 1

**Details**

Data on an experiment to examine the effect of a antagonist MDL 72222 on the change in blood pressure experienced with increasing dosage of phenylbiguanide are described in Ludbrook (1994) and analyzed in Venables and Ripley (1999, section 8.8). Each of five rabbits was exposed to increasing doses of phenylbiguanide after having either a placebo or the HD5-antagonist MDL 72222 administered.

**Source**

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York. (Appendix A.21)

Venables, W. N. and Ripley, B. D. (1999) *Modern Applied Statistics with S-PLUS (3rd ed)*, Springer, New York.

Ludbrook, J. (1994), Repeated measurements and multiple comparisons in cardiovascular research, *Cardiovascular Research*, **28**, 303-311.

**Examples**

```
str(PBG)
```

---

Phenobarb

*Phenobarbital Kinetics*

---

**Description**

The Phenobarb data frame has 744 rows and 7 columns.

**Format**

This data frame contains the following columns:

**Subject** an ordered factor identifying the infant.

**Wt** a numeric vector giving the birth weight of the infant (kg).

**Apgar** an ordered factor giving the the 5-minute Apgar score for the infant. This is an indication of health of the newborn infant.

**ApgarInd** a factor indicating whether the 5-minute Apgar score is < 5 or >= 5.

**time** a numeric vector giving the time when the sample is drawn or drug administered (hr).

**dose** a numeric vector giving the dose of drug administered (ug/kg).

**conc** a numeric vector giving the phenobarbital concentration in the serum (ug/L).

**Details**

Data from a pharmacokinetics study of phenobarbital in neonatal infants. During the first few days of life the infants receive multiple doses of phenobarbital for prevention of seizures. At irregular intervals blood samples are drawn and serum phenobarbital concentrations are determined. The data were originally given in Grasela and Donn(1985) and are analyzed in Boeckmann, Sheiner and Beal (1994), in Davidian and Giltinan (1995), and in Littell et al. (1996).

**Source**

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York. (Appendix A.23)

Davidian, M. and Giltinan, D. M. (1995), *Nonlinear Models for Repeated Measurement Data*, Chapman and Hall, London. (section 6.6)

Grasela and Donn (1985), Neonatal population pharmacokinetics of phenobarbital derived from routine clinical data, *Developmental Pharmacology and Therapeutics*, **8**, 374-383.

Boeckmann, A. J., Sheiner, L. B., and Beal, S. L. (1994), *NONMEM Users Guide: Part V*, University of California, San Francisco.

Littell, R. C., Milliken, G. A., Stroup, W. W. and Wolfinger, R. D. (1996), *SAS System for Mixed Models*, SAS Institute, Cary, NC.

**Examples**

```
str(Phenobarb)
```

---

Pixel	<i>X-ray pixel intensities over time</i>
-------	--

---

**Description**

The Pixel data frame has 102 rows and 4 columns of data on the pixel intensities of CT scans of dogs over time

**Format**

This data frame contains the following columns:

**Dog** a factor with levels A to J designating the dog on which the scan was made

**Side** a factor with levels L and R designating the side of the dog being scanned

**day** a numeric vector giving the day post injection of the contrast on which the scan was made

**pixel** a numeric vector of pixel intensities

**Source**

Pinheiro, J. C. and Bates, D. M. (2000) *Mixed-effects Models in S and S-PLUS*, Springer.

**Examples**

```
options(show.signif.stars = FALSE)
str(Pixel)
summary(Pixel)
(fm1 <- lmer(pixel ~ day + I(day^2) + (1|Dog:Side) + (day|Dog), Pixel))
```

---

Quinidine

*Quinidine Kinetics*

---

### Description

The Quinidine data frame has 1471 rows and 14 columns.

### Format

This data frame contains the following columns:

**Subject** a factor identifying the patient on whom the data were collected.

**time** a numeric vector giving the time (hr) at which the drug was administered or the blood sample drawn. This is measured from the time the patient entered the study.

**conc** a numeric vector giving the serum quinidine concentration (mg/L).

**dose** a numeric vector giving the dose of drug administered (mg). Although there were two different forms of quinidine administered, the doses were adjusted for differences in salt content by conversion to milligrams of quinidine base.

**interval** a numeric vector giving the when the drug has been given at regular intervals for a sufficiently long period of time to assume steady state behavior, the interval is recorded.

**Age** a numeric vector giving the age of the subject on entry to the study (yr).

**Height** a numeric vector giving the height of the subject on entry to the study (in.).

**Weight** a numeric vector giving the body weight of the subject (kg).

**Race** a factor with levels *Caucasian*, *Latin*, and *Black* identifying the race of the subject.

**Smoke** a factor with levels *no* and *yes* giving smoking status at the time of the measurement.

**Ethanol** a factor with levels *none*, *current*, *former* giving ethanol (alcohol) abuse status at the time of the measurement.

**Heart** a factor with levels *No/Mild*, *Moderate*, and *Severe* indicating congestive heart failure for the subject.

**Creatinine** an ordered factor with levels  $< 50$   $< = 50$  indicating the creatine clearance (mg/min).

**glyco** a numeric vector giving the alpha-1 acid glycoprotein concentration (mg/dL). Often measured at the same time as the quinidine concentration.

### Details

Verme et al. (1992) analyze routine clinical data on patients receiving the drug quinidine as a treatment for cardiac arrhythmia (atrial fibrillation of ventricular arrhythmias). All patients were receiving oral quinidine doses. At irregular intervals blood samples were drawn and serum concentrations of quinidine were determined. These data are analyzed in several publications, including Davidian and Giltinan (1995, section 9.3).

**Source**

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York. (Appendix A.25)

Davidian, M. and Giltinan, D. M. (1995), *Nonlinear Models for Repeated Measurement Data*, Chapman and Hall, London.

Verme, C. N., Ludden, T. M., Clementi, W. A. and Harris, S. C. (1992), Pharmacokinetics of quinine in male patients: A population analysis, *Clinical Pharmacokinetics*, **22**, 468-480.

**Examples**

```
str(Quinidine)
```

---

 Rail

---

*Evaluation of Stress in Railway Rails*


---

**Description**

The Rail data frame has 18 rows and 2 columns.

**Format**

This data frame contains the following columns:

**Rail** an ordered factor identifying the rail on which the measurement was made.

**travel** a numeric vector giving the travel time for ultrasonic head-waves in the rail (nanoseconds).  
The value given is the original travel time minus 36,100 nanoseconds.

**Details**

Devore (2000, Example 10.10, p. 427) cites data from an article in *Materials Evaluation* on “a study of travel time for a certain type of wave that results from longitudinal stress of rails used for railroad track.”

**Source**

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York. (Appendix A.26)

Devore, J. L. (2000), *Probability and Statistics for Engineering and the Sciences (5th ed)*, Duxbury, Boston, MA.

**Examples**

```
str(Rail)
(fm1 <- lmer(travel ~ 1 | Rail, Rail))
```

---

RatPupWeight	<i>The weight of rat pups</i>
--------------	-------------------------------

---

**Description**

The RatPupWeight data frame has 322 rows and 5 columns.

**Format**

This data frame contains the following columns:

**weight** a numeric vector

**sex** a factor with levels Male Female

**Litter** a factor, the litter number

**Lsize** a numeric vector

**Treatment** an ordered factor with levels Control < Low < High

**Source**

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York.

**Examples**

```
str(RatPupWeight)
```

---

Relaxin	<i>Assay for Relaxin</i>
---------	--------------------------

---

**Description**

The Relaxin data frame has 198 rows and 3 columns.

**Format**

This data frame contains the following columns:

**Run** an ordered factor with levels 5 < 8 < 9 < 3 < 4 < 2 < 7 < 1 < 6

**conc** a numeric vector

**cAMP** a numeric vector

**Source**

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York.

**Examples**

```
str(Relaxin)
```

---

Remifentanil

*Pharmacokinetics of remifentanil*

---

**Description**

The Remifentanil data frame has 2107 rows and 12 columns.

**Format**

This data frame contains the following columns:

**ID** a numeric vector

**Subject** an ordered factor

**Time** a numeric vector

**conc** a numeric vector

**Rate** a numeric vector

**Amt** a numeric vector

**Age** a numeric vector

**Sex** a factor with levels Female Male

**Ht** a numeric vector

**Wt** a numeric vector

**BSA** a numeric vector

**LBM** a numeric vector

**Source**

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York.

**Examples**

```
str(Remifentanil)
```

---

Soybean

*Growth of soybean plants*

---

### Description

The Soybean data frame has 412 rows and 5 columns.

### Format

This data frame contains the following columns:

**Plot** a factor giving a unique identifier for each plot.

**Variety** a factor indicating the variety; Forrest (F) or Plant Introduction #416937 (P).

**Year** a factor indicating the year the plot was planted.

**Time** a numeric vector giving the time the sample was taken (days after planting).

**weight** a numeric vector giving the average leaf weight per plant (g).

### Details

These data are described in Davidian and Giltinan (1995, 1.1.3, p.7) as “Data from an experiment to compare growth patterns of two genotypes of soybeans: Plant Introduction #416937 (P), an experimental strain, and Forrest (F), a commercial variety.”

### Source

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York. (Appendix A.27)

Davidian, M. and Giltinan, D. M. (1995), *Nonlinear Models for Repeated Measurement Data*, Chapman and Hall, London.

### Examples

```
str(Soybean)
#summary(fm1 <- nlsList(SSlogis, data = Soybean))
```

---

Spruce	<i>Growth of Spruce Trees</i>
--------	-------------------------------

---

**Description**

The Spruce data frame has 1027 rows and 4 columns.

**Format**

This data frame contains the following columns:

**Tree** a factor giving a unique identifier for each tree.

**days** a numeric vector giving the number of days since the beginning of the experiment.

**logSize** a numeric vector giving the logarithm of an estimate of the volume of the tree trunk.

**plot** a factor identifying the plot in which the tree was grown.

**Details**

Diggle, Liang, and Zeger (1994, Example 1.3, page 5) describe data on the growth of spruce trees that have been exposed to an ozone-rich atmosphere or to a normal atmosphere.

**Source**

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York. (Appendix A.28)

Diggle, Peter J., Liang, Kung-Yee and Zeger, Scott L. (1994), *Analysis of longitudinal data*, Oxford University Press, Oxford.

**Examples**

```
str(Spruce)
```

---

Tetracycline1	<i>Pharmacokinetics of tetracycline</i>
---------------	---

---

**Description**

The Tetracycline1 data frame has 40 rows and 4 columns.

**Format**

This data frame contains the following columns:

**conc** a numeric vector

**Time** a numeric vector

**Subject** an ordered factor with levels 5 < 3 < 2 < 4 < 1

**Formulation** a factor with levels tetrachel tetracyn

**Source**

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York.

**Examples**

```
str(Tetracycline1)
```

---

Tetracycline2	<i>Pharmacokinetics of tetracycline</i>
---------------	---

---

**Description**

The Tetracycline2 data frame has 40 rows and 4 columns.

**Format**

This data frame contains the following columns:

**conc** a numeric vector

**Time** a numeric vector

**Subject** an ordered factor with levels 4 < 5 < 2 < 1 < 3

**Formulation** a factor with levels Berkmycin tetramycin

**Source**

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York.

**Examples**

```
str(Tetracycline2)
```

---

Theoph	<i>Pharmacokinetics of theophylline</i>
--------	---

---

**Description**

The Theoph data frame has 132 rows and 5 columns of data from an experiment on the pharmacokinetics of theophylline.

**Usage**

Theoph

**Format**

This data frame contains the following columns:

**Subject** a factor with levels A, . . . , L identifying the subject on whom the observation was made.

**Wt** weight of the subject (kg).

**Dose** dose of theophylline administered orally to the subject (mg/kg).

**Time** time since drug administration when the sample was drawn (hr).

**conc** theophylline concentration in the sample (mg/L).

**Details**

Boeckmann, Sheiner and Beal (1994) report data from a study by Dr. Robert Upton of the kinetics of the anti-asthmatic drug theophylline. Twelve subjects were given oral doses of theophylline then serum concentrations were measured at 11 time points over the next 25 hours.

These data are analyzed in Davidian and Giltinan (1995) and Pinheiro and Bates (2000) using a two-compartment open pharmacokinetic model, for which a self-starting model function, `SSfol`, is available.

**Source**

Boeckmann, A. J., Sheiner, L. B. and Beal, S. L. (1994), *NONMEM Users Guide: Part V*, NONMEM Project Group, University of California, San Francisco.

Davidian, M. and Giltinan, D. M. (1995) *Nonlinear Models for Repeated Measurement Data*, Chapman & Hall (section 5.5, p. 145 and section 6.6, p. 176)

Pinheiro, J. C. and Bates, D. M. (2000) *Mixed-effects Models in S and S-PLUS*, Springer (Appendix A.29)

**See Also**

[SSfol](#)

**Examples**

```
require(lattice)
xyplot(conc ~ Time | Subject, Theoph, aspect = 'xy',
       xlab = "Time since drug administration (hr)",
       ylab = "Theophylline concentration (mg/L)")
Theoph.D <- subset(Theoph, Subject == "D")
fm1 <- nls(conc ~ SSfol(Dose, Time, lKe, lKa, lCl),
          data = Theoph.D)
summary(fm1)
plot(conc ~ Time, data = Theoph.D,
     xlab = "Time since drug administration (hr)",
     ylab = "Theophylline concentration (mg/L)",
     main = "Observed concentrations and fitted model",
     sub = "Theophylline data - Subject 4 only",
     las = 1, col = 4)
xvals <- seq(0, par("usr")[2], len = 55)
```

```
lines(xvals, predict(fm1, newdata = list(Time = xvals)),
      col = 4)
```

---

Wafer

*Modeling of Analog MOS Circuits*

---

### Description

The Wafer data frame has 400 rows and 4 columns.

### Format

This data frame contains the following columns:

**Wafer** a factor with levels 1 2 3 4 5 6 7 8 9 10

**Site** a factor with levels 1 2 3 4 5 6 7 8

**voltage** a numeric vector

**current** a numeric vector

### Source

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York.

### Examples

```
str(Wafer)
```

---

Wheat

*Yields by growing conditions*

---

### Description

The Wheat data frame has 48 rows and 4 columns.

### Format

This data frame contains the following columns:

**Tray** an ordered factor with levels 3 < 1 < 2 < 4 < 5 < 6 < 8 < 9 < 7 < 12 < 11 < 10

**Moisture** a numeric vector

**fertilizer** a numeric vector

**DryMatter** a numeric vector

**Source**

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York.

**Examples**

```
str(Wheat)
```

---

Wheat2

*Wheat Yield Trials*

---

**Description**

The Wheat2 data frame has 224 rows and 5 columns.

**Format**

This data frame contains the following columns:

**Block** an ordered factor with levels 4 < 2 < 3 < 1

**variety** a factor with levels ARAPAHOE BRULE BUCKSKIN CENTURA CENTURK78 CHEYENNE CODY COLT GAGE HOMESTEAD KS831374 LANCER LANCOTA NE83404 NE83406 NE83407 NE83432 NE83498 NE83T12 NE84557 NE85556 NE85623 NE86482 NE86501 NE86503 NE86507 NE86509 NE86527 NE86582 NE86606 NE86607 NE86T666 NE87403 NE87408 NE87409 NE87446 NE87451 NE87457 NE87463 NE87499 NE87512 NE87513 NE87522 NE87612 NE87613 NE87615 NE87619 NE87627 NORKAN REDLAND ROUGHRIDER SCOUT66 SIOUXLAND TAM107 TAM200 VONA

**yield** a numeric vector

**latitude** a numeric vector

**longitude** a numeric vector

**Source**

Pinheiro, J. C. and Bates, D. M. (2000), *Mixed-Effects Models in S and S-PLUS*, Springer, New York.

**Examples**

```
str(Wheat2)
```

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