

# Package ‘gastempt’

March 6, 2019

**Version** 0.4.4

**Type** Package

**Title** Analyzing Gastric Emptying from MRI or Scintigraphy

**Description** Fits gastric emptying time series from MRI or scintigraphic measurements using nonlinear mixed-model population fits with 'nlme' and Bayesian methods with Stan; computes derived parameters such as t50 and AUC.

**License** GPL (>= 3)

**LazyData** TRUE

**NeedsCompilation** yes

**URL** <http://github.com/dmenne/gastempt>

**BugReports** <http://github.com/dmenne/gastempt/issues>

**Depends** R (>= 3.4.0)

**Imports** nlme, Rcpp (>= 0.12.19), dplyr, tibble, ggplot2 (>= 3.0.0), rstan (>= 2.18.1), rstantools (>= 1.5.1), assertthat, stringr, methods, shiny

**Suggests** rmarkdown, knitr, covr, testthat

**LinkingTo** StanHeaders (>= 2.18.0), rstan (>= 2.18.1), BH (>= 1.66.0-1), Rcpp (>= 0.12.19), RcppEigen (>= 0.3.3.4.0)

**RoxygenNote** 6.1.1

**VignetteBuilder** knitr

**Encoding** UTF-8

**SystemRequirements** GNU make

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**R topics documented:**

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coef.nlme_gastempt	<i>Extract coefficients from nlme_gastempt result</i>
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**Description**

Extract coefficients from nlme\_gastempt result

**Usage**

```
## S3 method for class 'nlme_gastempt'
coef(object, ...)
```

**Arguments**

object	Result of a call to nlme_gastempt
...	other arguments

**Value**

a data frame with coefficients. See [nlme\\_gastempt](#) for an example.

---

coef.stan_gastempt	<i>Extract coefficients from stan_gastempt result</i>
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**Description**

Extract coefficients from stan\_gastempt result

**Usage**

```
## S3 method for class 'stan_gastempt'
coef(object, ...)
```

**Arguments**

object	Result of a call to stan_gastempt
...	other arguments

**Value**

a data frame with coefficients. See [nlme\\_gastempt](#) for an example.

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gastemptfunc	<i>Functions for gastric emptying analysis</i>
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**Description**

The `linexp` and the power exponential (`powexp`) functions can be used to fit gastric emptying curves.

**Usage**

```
linexp(t, v0 = 1, tempt = NULL, kappa = NULL, pars = NULL)
linexp_slope(t, v0 = 1, tempt = NULL, kappa = NULL, pars = NULL)
linexp_auc(v0 = 1, tempt = NULL, kappa = NULL, pars = NULL)
powexp(t, v0 = 1, tempt = NULL, beta = NULL, pars = NULL)
powexp_slope(t, v0 = 1, tempt = NULL, beta = NULL, pars = NULL)
linexp_log(t, v0 = 1, logtempt = NULL, logkappa = NULL,
  pars = NULL)
powexp_log(t, v0 = 1, logtempt = NULL, logbeta = NULL, pars = NULL)
```

**Arguments**

t	Time after meal or start of scan, in minutes; can be a vector.
v0	Initial volume at t=0.
tempt	Emptying time constant in minutes (scalar).
kappa	Overshoot term for linexp function (scalar).
pars	Default NULL. If not NULL, the other parameters with exception of t are not used and are retrieved as named parameters from the numeric vector pars instead.
beta	Power term for power exponential function (scalar).
logtempt	Logarithm of emptying time constant in minutes (scalar).
logkappa	Logarithm of overshoot term for linexp function (scalar).
logbeta	Logarithm of power term for power exponential function (scalar).

**Details**

The linexp function can have an initial overshoot to model secretion.

$$\text{vol}(t) = v0 * (1 + \text{kappa} * t / \text{tempt}) * \exp(-t / \text{tempt})$$

The powexp function introduced by Elashof et al. is monotonously decreasing but has more freedom to model details in the function tail.

$$\text{vol}(t) = v0 * \exp(-(t / \text{tempt}) ^ \text{beta})$$

The `_slope` functions return the first derivatives of `linexp` and `powexp`. Use the `_log` functions to enforce positive parameters `tempt` and `beta`. Rarely required for gastric emptying curves.

**Value**

Vector of `length(t)` for computed volume.

**Examples**

```
t = seq(0,100, by=5)
kappa = 1.3
tempt = 60
v0 = 400
beta = 3
pars = c(v0 = v0, tempt = tempt, kappa = kappa)
par(mfrow=c(1,3))
plot(t, linexp(t, v0, tempt, kappa), type = "l", ylab = "volume",
     main = "linexp\nkappa = 1.3 and 1.0")
lines(t, linexp(t, v0, tempt, 1), type = "l", col = "green")
# This should give the same plot as above
plot(t, linexp(t, pars = pars), type = "l", ylab = "volume",
     main = "linexp\nkappa = 1.3 and 1.0\nwith vectored parameters")
lines(t, linexp(t, v0, tempt, 1), type = "l", col = "green")
plot(t, powexp(t, v0, tempt, beta), type = "l", ylab = "volume",
     main = "powexp\nbeta = 2 and 1")
lines(t, powexp(t, v0, tempt, 1), type = "l", col = "green")
```

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nlme_gastempt	<i>Simplified population fit of gastric emptying data</i>
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### Description

Compute coefficients  $v_0$ ,  $tempt$  and  $kappa$  of a mixed model fit to a linexp function with one grouping variable

### Usage

```
nlme_gastempt(d, pnlsTol = 0.001, model = linexp, variant = 1)
```

### Arguments

d	A data frame with columns <ul style="list-style-type: none"> <li>record Record descriptor as grouping variable, e.g. patient ID</li> <li>minute Time after meal or start of recording.</li> <li>vol Volume of meal or stomach</li> </ul>
pnlsTol	The value of pnlsTol at the initial iteration. See <a href="#">nlmeControl</a> When the model does not converge, pnlsTol is multiplied by 5 and the iteration repeated until convergence or $pnlsTol \geq 0.5$ . The effective value of pnlsTol is returned in a separate list item. When it is known that a data set converges badly, it is recommended to set the initial pnlsTol to a higher value, but below 0.5, for faster convergence.
model	linexp (default) or powexp
variant	For both models, there are 3 variants <ul style="list-style-type: none"> <li>variant = 1 The most generic version with independent estimates of all three parameters per record (<math>random = v_0 + tempt + kappa \sim 1 \mid record</math>). The most likely to fail for degenerate cases. If this variant converges, use it.</li> <li>variant = 2 Diagonal random effects (<math>random = pdDiag(v_0 + tempt + kappa) \sim 1; groups =</math> Better convergence in critical cases. Note: I never found out why I have to use the groups parameter instead of the <math>\mid</math>; see also p. 380 of Pinheiro/Bates.</li> <li>variant = 3 Since parameters kappa and beta respectively are the most difficult to estimate, these are fixed in this variant (<math>random = v_0 + tempt \sim 1</math>). This variant converges in all reasonable cases, but the estimates of kappa and beta cannot be use for secondary between-group analysis. If you are only interested in <math>t_{50}</math>, you can use this safe version.</li> </ul>

### Value

A list of class nlme\_gastempt with elements coef, summary, plot, pnlsTol, message

- coef is a data frame with columns:
  - record Record descriptor, e.g. patient ID

- v0 Initial volume at t=0
- tempt Emptying time constant
- kappa Parameter kappa for model = linexp
- beta Parameter beta for model = powexp
- t50 Half-time of emptying
- slope\_t50 Slope in t50; typically in units of ml/minute

On error, coef is NULL

- nlme\_result Result of the nlme fit; can be used for addition processing, e.g. to plot residuals or via summary to extract AIC. On error, nlme\_result is NULL.
- plot A ggplot graph of data and prediction. Plot of raw data is returned even when convergence was not achieved.
- pnlsTol Effective value of pnlsTo after convergence or failure.
- message String "Ok" on success, and the error message of nlme on failure.

### Examples

```
suppressWarnings(RNGversion("3.5.0"))
set.seed(4711)
d = simulate_gastempt(n_record = 10, kappa_mean = 0.9, kappa_std = 0.3,
                     model = linexp)$data
fit_d = nlme_gastempt(d)
# fit_d$coef # direct access
coef(fit_d) # better use accessor function
coef(fit_d, signif = 3) # Can also set number of digits
# Avoid ugly ggplot shading (not really needed...)
library(ggplot2)
theme_set(theme_bw() + theme(panel.spacing = grid::unit(0, "lines")))
# fit_d$plot # direct access is possible
plot(fit_d) # better use accessor function
```

---

plot.nlme\_gastempt      *Plot data points and fit curve of an nlme\_gastempt fit*

---

### Description

Plot data points and fit curve of an nlme\_gastempt fit

### Usage

```
## S3 method for class 'nlme_gastempt'
plot(x, ...)
```

### Arguments

x	Result of a call to nlme_gastempt
...	other arguments

**Value**

a ggplot object. Use `print()` if used non-interactively to show the curve

---

`plot.stan_gastempt`      *Plot data points and fit curve of an stan\_gastempt fit*

---

**Description**

Plot data points and fit curve of an stan\_gastempt fit

**Usage**

```
## S3 method for class 'stan_gastempt'  
plot(x, ...)
```

**Arguments**

`x`                      Result of a call to stan\_gastempt  
`...`                    other arguments

**Value**

a ggplot object. Use `print()` if used non-interactively to show the curve

---

`run_shiny`                      *Run shiny app demonstrating fit strategies with simulated data*

---

**Description**

Run shiny app demonstrating fit strategies with simulated data

**Usage**

```
run_shiny()
```

**Value**

Not used, starts shiny app

---

simulate\_gastempt      *Simulate gastric emptying data following a linexp or powexp function*

---

### Description

Simulate gastric emptying data following a linexp or powexp function

### Usage

```
simulate_gastempt(n_records = 10, v0_mean = 400, v0_std = 50,
  tempt_mean = ifelse(identical(model, linexp), 60, 120),
  tempt_std = tempt_mean/3, kappa_mean = 0.7,
  kappa_std = kappa_mean/3, beta_mean = 0.7, beta_std = beta_mean/3,
  noise = 20, student_t_df = NULL, missing = 0, model = linexp,
  seed = NULL, max_minute = NULL)
```

### Arguments

n_records	Number of records
v0_mean, v0_std	Mean and between record standard deviation of initial volume, typically in ml.
tempt_mean, tempt_std	Mean and between record standard deviation of parameter \$t\_empt\$, typically in minutes.
kappa_mean, kappa_std	For linexp only: Mean and between-record standard deviation of overshoot parameter kappa. For values of kappa above 1, curve has an overshoot that can be used to follow volume time series with secretion.
beta_mean, beta_std	For powexp only: Mean and between-record standard deviation of the so called lag parameter.
noise	Standard deviation of normal noise when student_t_df = NULL; scaling of noise when student_t_df >= 2.
student_t_df	When NULL (default), Gaussian noise is added; when >= 2, Student_t distributed noise is added, which generates more realistic outliers. Values from 2 to 5 are useful, when higher values are used the result comes close to that of Gaussian noise. Values below 2 are rounded to 2.
missing	When 0 (default), all curves have the same number of data points. When > 0, this is the fraction of points that were removed randomly to simulate missing points. Maximum value is 0.5.
model	linexp(default) or powexp
seed	optional seed; not set if seed = NULL (default)
max_minute	Maximal time in minutes; if NULL, a sensible default rounded to hours is used

**Value**

A list with 3 elements:

**record** Data frame with columns `record(chr)`, `v0`, `tempt`, `kappa/beta` giving the effective `linexp` or `powexp` parameters for the individual record. `v0` is rounded to nearest integer.

**data** Data frame with columns `record(chr)`, `minute(dbl)`, `vol(dbl)` giving the time series and grouping parameters. `vol` is rounded to nearest integer.

**stan\_data** A list for use as data in Stan-based fits with elements `prior_v0`, `n`, `n_record`, `record`, `minute`, `volume`.

A comment is attached to the return value that can be used as a title

**Examples**

```
suppressWarnings(RNGversion("3.5.0"))
set.seed(4711)
library(ggplot2)
vol_linexp = simulate_gastempt(n_records = 4, noise = 20)
ggplot(vol_linexp$data, aes(x = minute, y = vol)) + geom_point() +
  facet_wrap(~record) + ggtitle("linexp, noise = 0, no missing")

vol_powexp = simulate_gastempt(n_records = 4, missing = 0.2, student_t_df = 2)
ggplot(vol_powexp$data, aes(x = minute, y = vol)) + geom_point() +
  facet_wrap(~record) + ggtitle("powexp, noise = 10 (default), 20% missing,
  Student-t (df = 2) noise")
```

---

 stan\_gastempt

*Fit gastric emptying curves with Stan*


---

**Description**

Fit gastric emptying curves with Stan

**Usage**

```
stan_gastempt(d, model_name = "linexp_gastro_2b", lkj = 2,
  student_df = 5L, init_r = 0.2, chains = 4, ...)
```

**Arguments**

<code>d</code>	<p>A data frame with columns</p> <ul style="list-style-type: none"> <li>• <code>rec</code> Record descriptor as grouping variable, e.g. patient ID</li> <li>• <code>minute</code> Time after meal or start of recording.</li> <li>• <code>vol</code> Volume of meal or stomach</li> </ul>
<code>model_name</code>	<p>Name of predefined model in <code>gastempt/exec</code>. Use <code>stan_model_names()</code> to get a list of available models.</p>

lkj	LKJ prior for kappa/tempt correlation, only required for model linexp_gastro_2b. Values from 1.5 (strong correlation) to 50 (almost independent) are useful. See <a href="http://www.psychstatistics.com/2014/12/27/d-lkj-priors/">http://www.psychstatistics.com/2014/12/27/d-lkj-priors/</a> for examples.
student_df	Student-t degrees of freedom for residual error; default 5. Use 3 for strong outliers; values above 10 are close to gaussian residual distribution.
init_r	for stan, default = 0.2; Stan's own default is 2, which often results in stuck chains.
chains	for stan; default = 4. For debugging, use 1.
...	Additional parameter passed to sampling

### Value

A list of class `stan_gastempt` with elements `coef`, `fit`, `plot`

- `coef` is a data frame with columns:
  - `rec` Record descriptor, e.g. patient ID
  - `v0` Initial volume at t=0
  - `tempt` Emptying time constant
  - `kappa` Parameter kappa for model = linexp
  - `beta` Parameter beta for model = powexp
  - `t50` Half-time of emptying
  - `slope_t50` Slope in t50; typically in units of ml/minute On error, `coef` is NULL
- `fit` Result of class 'stanfit'
- `plot` A ggplot graph of data and prediction. Plot of raw data is returned even when convergence was not achieved.

### Examples

```
## Not run:
dd = simulate_gastempt(n_records = 6, seed = 471)
d = dd$data
ret = stan_gastempt(d)
print(ret$coef)

## End(Not run)
```

---

stan\_model\_names

*Names and descriptions of precompiled Stan models*

---

### Description

By default, line 2 and 3 of comments starting with # or // in Stan file are returned

**Usage**

```
stan_model_names(n_lines = 2, skip = 1, sep = "\n")
```

**Arguments**

n_lines	Number of comment lines to retrieve
skip	Number of lines to skip from beginning of Stan Model file
sep	separator for multiline strings

**Value**

A data frame with model\_name and the first n\_lines comment lines in model as description

---

t50	<i>Compute half-emptying time from nlme parameters</i>
-----	--

---

**Description**

No closed solution known for [linexp](#), we use a Newton approximation.

**Usage**

```
t50(x)
```

**Arguments**

x	Result of a nlme fit, with named components 'tempt, beta, logbeta, kappa, logkappa' depending on model. Function used 'logbeta' when it is present, in 'x', otherwise beta, and similar for logkappa/kappa.
---	---

**Value**

Half-emptying time. Name of evaluated function is returned as attribute fun. Negative of slope is returned as attribute slope.

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