

Package: mpmm (via r-universe)

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Type Package

Title (Animal) Movement Persistence Mixed-Effects Models

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Description Fit covariates as linear mixed-effects to latent, time-varying movement persistence (autocorrelation).

Depends R (>= 3.5.0)

Imports TMB (>= 1.7.19), lme4 (>= 1.1-15), glmmTMB (>= 1.0.0), Matrix (>= 1.2-12), dplyr (>= 1.0.0), tibble (>= 3.0.0), ggplot2, methods, wesanderson, tidyverse

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Repository <https://ianjonsen.r-universe.dev>

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mpmm-package

mpmm

Description

fit movement persistence mixed-effects models to animal location data

Author(s)

Ian Jonsen

References

Jonsen ID, McMahon CR, Patterson TA, et al. (2019) Movement responses to environment: fast inference of variation among southern elephant seals with a mixed effects model. *Ecology*. 100(1):e02566
<https://doi.org/10.1002/ecy.2566>

See Also

mpmm

anova.mpmm

anova tables

Description

perform likelihood ratio tests on 2 or more mpmm fit objects

Usage

```
## S3 method for class 'mpmm'
anova(object, ...)
```

Arguments

object	an mpmm fit object
...	additional mpmm fit objects

`ellie.ice`

foieGras-filtered Southern elephant seal Argos satellite data with environmental covariates (11 individuals)

Description

Example elephant seal Argos tracking data with environmental covariates. Data were sourced from the Integrated Marine Observing System (IMOS) - IMOS is supported by the Australian Government through the National Collaborative Research Infrastructure Strategy and the Super Science Initiative.

Format

.RData

`ellie.ice.short`

foieGras-filtered Southern elephant seal Argos satellite data with environmental covariates (4 individuals)

Description

Example elephant seal Argos tracking data with environmental covariates. Data were sourced from the Integrated Marine Observing System (IMOS) - IMOS is supported by the Australian Government through the National Collaborative Research Infrastructure Strategy and the Super Science Initiative.

Format

.RData

`inner_control`

Control Values for mpmm

Description

`inner_control` allows specification of control parameters for the inner optimization used by `mpmm`

Usage

```
inner_control(...)
```

Arguments

...	control parameters for the inner optimizer
-----	--

Details

See [MakeADFun](#) and [newton](#) for details and available options. Adapted from S. Wotherspoon <https://github.com/SWotherspoon/RWalc/blob/master/R/RWalc.R>

Value

Returns a list with components:

<code>control</code>	list of control parameters for inner optimizer
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See Also

[MakeADFun](#), [newton](#)

Examples

```
fit <- mpmm(~ ice + (ice | id),
            data = ellie.ice.short,
            inner.control = inner_control(tol = 1e-03))
```

`logLik.mpmm`

Extract log-likelihood

Description

extract log-likelihood from an mpmm fit object

Usage

```
## S3 method for class 'mpmm'
logLik(object, ...)
```

Arguments

<code>object</code>	an mpmm model fit object
<code>...</code>	additional arguments to be ignored

mpmm*Move Persistence Mixed-Effects Model*

Description

Fit a move persistence random walk via TMB to a pre-filtered/regularized animal track and estimate gamma as a linear function of covariates

Usage

```
mpmm(
  formula = NA,
  data = NULL,
  map = NULL,
  control = mpmm_control(),
  inner.control = inner_control()
)
```

Arguments

formula	a right-hand-side regression formula (no response variable)
data	a data frame of observations (see details)
map	a named list of parameters as factors that are to be fixed during estimation, e.g., list(rho = factor(NA))
control	a list of control parameters for the outer optimization (see mpmm_control)
inner.control	a list of control parameters for the inner optimization (see MakeADFun and newton)

Details

The input track is given as a data frame where each row is an observed location and columns
'id' individual animal identifier,
'date' observation time (POSIXct,GMT),
'lon' observed longitude,
'lat' observed latitude,
'tid' identifier for tracks if there are more than one track per individual (optional),
'..' named covariates appended to track

Value

a list with components

states	a data frame of estimated states
fitted	a data frame of fitted locations
par	model parameter summary

<code>data</code>	input dataframe
<code>tmb</code>	the tmb object
<code>opt</code>	the object returned by the optimizer

Examples

```
data(ellie.ice.short)
fit <- mpmm(~ ice + (1 | id), data = ellie.ice.short)
summary(fit)
```

`mpmm_control` *Control Values for mpmm.*

Description

`mpmm_control` selects the numerical minimizer, method, associated control parameters, parameter bounds, and likelihood estimation (REML or ML) used by `mpmm`.

Usage

```
mpmm_control(
  optim = c("nlminb", "optim"),
  method = c("BFGS", "L-BFGS-B"),
  lower = NULL,
  upper = NULL,
  REML = FALSE,
  profile = FALSE,
  verbose = 2,
  ...
)
```

Arguments

<code>optim</code>	the numerical optimizer used in the fit
<code>method</code>	optimization method to be used - one of "BFGS" or "L-BFGS-B" for bounded optimization (default lower and upper bounds are used if not specified)
<code>lower</code>	a list of named parameter lower bounds, if NULL then built in defaults are used when <code>method</code> = "L-BFGS-B", otherwise ignored
<code>upper</code>	a list of named parameter upper bounds, if NULL then built in defaults are used when <code>method</code> = "L-BFGS-B", otherwise ignored
<code>REML</code>	logical; whether to use REML (TRUE) or maximum likelihood
<code>profile</code>	logical; option to improve speed and convergence by using REML parameter estimates as initial values for ML optimization
<code>verbose</code>	integer; report progress during minimization: 0 = silent; 1 = optimizer trace; 2 = parameter trace (default)
...	control parameters for the chosen optimizer

Details

The optimizer used to minimize the objective function is selected by the `optim` argument. Additional control parameters specific to the chosen optimizer are specified via the `dots` argument. See `nlminb` and `optim` for available options. Adapted from S. Wotherspoon <https://github.com/SWotherspoon/RWalc/blob/master/R/RWalc.R>

Value

Returns a list with components

<code>optim</code>	the name of the numerical optimizer as a string, "nlminb" or "optim"
<code>method</code>	optimization method to be used
<code>lower</code>	named list of lower parameter bounds
<code>upper</code>	named list of upper parameter bounds
<code>REML</code>	whether REML is to be used in place of ML
<code>profile</code>	whether to enhance convergence robustness
<code>verbose</code>	level of tracing information to be reported
<code>control</code>	list of control parameters for the optimizer

See Also

`nlminb`, `optim`.

Examples

```
fit <- mpmm(~ ice + (ice | id),
data = ellie.ice.short,
control = mpmm_control(
  optim = "nlminb",
  REML = FALSE,
  eval.max = 2000
))
```

`plot.mpmm`

plot

Description

Visualise fixed and random covariate relationships from an mpmm fit object

Usage

```
## S3 method for class 'mpmm'
plot(x, label = FALSE, lwd = c(0.25, 0.75), ...)
```

Arguments

x	an mpmm fit object
label	add id labels to random effects
lwd	a vector of regression line widths (random effect, fixed effects)
...	additional arguments to be ignored

residuals.mpmm *residuals*

Description

Extract one-step-ahead residuals

Usage

```
## S3 method for class 'mpmm'
residuals(
  object,
  method = "oneStepGaussianOffMode",
  trace = FALSE,
  parallel = TRUE,
  ncores = detectCores() - 1,
  ...
)
```

Arguments

object	an mpmm fit object
method	character naming the method to calculate one-step-ahead residuals
trace	logical; print progress to console
parallel	logical; compute in parallel
ncores	integer; number of cores to use (default = total cores detected - 1)
...	additional arguments to be ignored

Details

Wrapper function for modified [oneStepPredict](#) that calculates one-step-ahead residuals, which are residuals that account for temporal correlation in latent states. The modification allows easier parallel computation.

Value

a list with components

res	a tibble with one-step-ahead residuals for longitude and latitude
-----	---

References

Thygesen, U. H., C. M. Albertsen, C. W. Berg, K. Kristensen, and A. Nielsen. 2017. Validation of ecological state space models using the Laplace approximation. Environmental and Ecological Statistics 24:317–339.

Examples

```
## Not run:  
data(ellie.ice)  
fit <- mpmm(~ ice + (1 | id), data = ellie.ice)  
summary(fit)  
residuals(fit)  
  
## End(Not run)
```

summary.mpmm

object summaries

Description

return a summary of an mpmm fit object

Usage

```
## S3 method for class 'mpmm'  
summary(object, ...)
```

Arguments

object	an mpmm fit object
...	additional arguments to be ignored

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