

Package ‘crawl’

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Type Package

Title Fit Continuous-Time Correlated Random Walk Models to Animal Movement Data

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Description Fit continuous-time correlated random walk models with time indexed covariates to animal telemetry data. The model is fit using the Kalman-filter on a state space version of the continuous-time stochastic movement process.

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crawl-package	<i>Fit Continuous-Time Correlated Random Walk Models to Animal Movement Data</i>
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Description

The Correlated RAndom Walk Library (I know it is not an R library, but, "crawp" did not sound as good) of R functions was designed for fitting continuous-time correlated random walk (CTCRW) models with time indexed covariates. The model is fit using the Kalman-Filter on a state space version of the continuous-time stochastic movement process.

Package:	crawl
Type:	Package
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LazyLoad:	yes

Note

This software package is developed and maintained by scientists at the NOAA Fisheries Alaska Fisheries Science Center and should be considered a fundamental research communication. The

recomendations and conclusions presented here are those of the authors and this software should not be construed as official communication by NMFS, NOAA, or the U.S. Dept. of Commerce. In addition, reference to trade names does not imply endorsement by the National Marine Fisheries Service, NOAA. While the best efforts have been made to insure the highest quality, tools such as this are under constant development and are subject to change.

Author(s)

Devin S. Johnson

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References

Johnson, D., J. London, M. -A. Lea, and J. Durban (2008) Continuous-time correlated random walk model for animal telemetry data. *Ecology* 89(5) 1208-1215.

aic.crw

Calculates AIC for all objects of class crwFit listed as arguments

Description

AIC, delta AIC, and Akaike weights for all models listed as arguments.

Usage

```
aic.crw(...)
```

Arguments

... a series of crwFit objects

Details

The function can either be executed with a series of 'crwFit' objects (see [crwMLE](#)) without the '.crwFit' suffix or the function can be called without any arguments and it will search out all 'crwFit' objects in the current workspace and produce the model selection table for all 'crwFit' objects in the workspace. Caution should be used when executing the function in this way. ALL 'crwFit' objects will be included whether or not the same locations are used! For all of the models listed as arguments (or in the workspace), AIC, delta AIC, and Akaike weights will be calculated.

Value

A table, sorted from lowest AIC value to highest.

Author(s)

Devin S. Johnson

argosDiag2Cov	<i>Transform Argos diagnostic data to covariance matrix form</i>
---------------	--

Description

Using this function the user can transform the Argos diagnostic data for location error into a form usable as a covariance matrix to approximate the location error with a bivariate Gaussian distribution. The resulting data.frame should be attached back to the data with cbind to use with the crwMLE function.

Usage

```
argosDiag2Cov(Major, Minor, Orientation)
```

Arguments

Major	A vector containing the major axis information for each observation (na values are ok)
Minor	A vector containing the minor axis information for each observation (na values are ok)
Orientation	A vector containing the angle orientation of the Major axis from North (na values are ok)

Value

A data.frame with the following columns

ln.sd.x	The log standard deviation of the location error in the x coordinate
ln.sd.y	The log standard deviation of the location error in the y coordinate
rho	The correlation of the bivariate location error ellipse

Author(s)

Devin S. Johnson

as.flat	<i>'Flattening' a list-form crwPredict object into a data.frame</i>
---------	---

Description

“Flattens” a list form `crwPredict` object into a flat data.frame.

Usage

```
as.flat(predObj)
```

Arguments

predObj A crwPredict object

Value

a [data.frame](#) version of a crwPredict list with columns for the state standard errors

Author(s)

Devin S. Johnson

See Also

[northernFurSeal](#) for use example

beardedSeals

Bearded Seal Location Data

Description

Bearded Seal Location Data

Format

A data frame with 27,548 observations on 3 bearded seals in Alaska:

deployid Unique animal ID

ptt Hardware ID

instr Hardware type

date_time Time of location

type Location type

quality Argos location quality

latitude Observed latitude

longitude Observed longitude

error_radius Argos error radius

error_semimajor_axis Argos error ellipse major axis length

error_semiminor_axis Argos error ellipse minor axis length

error_ellipse_orientation Argos error ellipse degree orientation

Source

Marine Mammal Laboratory, Alaska Fisheries Science Center, National Marine Fisheries Service, NOAA 7600 Sand Point Way NE Seattle, WA 98115

check_csv	<i>Start a shiny app to check data stored in a .csv file for model fitting with crwMLE function.</i>
-----------	--

Description

Users can start a beta version of Shiny app that allows for data checking and basic location projection.

Usage

```
check_csv()
```

crwMLE	<i>Fit Continuous-Time Correlated Random Walk Models to Animal Telemetry Data</i>
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Description

The function uses the Kalman filter to estimate movement parameters in a state-space version of the continuous-time movement model. Separate models are specified for movement portion and the location error portion. Each model can depend on time indexed covariates. A “haul out” model where movement is allowed to completely stop, as well as, a random drift model can be fit with this function.

Usage

```
crwMLE(mov.model = ~1, err.model = NULL, activity = NULL, drift = FALSE,
       data, coord = c("x", "y"), Time.name, initial.state, theta, fixPar,
       method = "Nelder-Mead", control = NULL, constr = list(lower = -Inf,
       upper = Inf), prior = NULL, need.hess = TRUE, initialSANN = list(maxit =
       200), attempts = 1)
```

Arguments

mov.model	formula object specifying the time indexed covariates for movement parameters.
err.model	A 2-element list of formula objects specifying the time indexed covariates for location error parameters.
activity	formula object giving the covariate for the activity (i.e., stopped or fully moving) portion of the model.
drift	logical indicating whether or not to include a random drift component. For most data this is usually not necessary. See northernFurSeal for an example using a drift model.

<code>data</code>	data.frame object containing telemetry and covariate data. A 'SpatialPointsDataFrame' object from the package 'sp' or an 'sf' object from the 'sf' package with a geometry column of type <code>sfc_POINT</code> . 'spacetime' will also be accepted. Values for coords will be taken from the spatial data set and ignored in the arguments.
<code>coord</code>	A 2-vector of character values giving the names of the "X" and "Y" coordinates in data.
<code>Time.name</code>	character indicating name of the location time column
<code>initial.state</code>	list object containing the initial state of the Kalman filter.
<code>theta</code>	starting values for parameter optimization.
<code>fixPar</code>	Values of parameters which are held fixed to the given value.
<code>method</code>	Optimization method that is passed to <code>optim</code> .
<code>control</code>	Control list which is passed to <code>optim</code> .
<code>constr</code>	Named list with elements <code>lower</code> and <code>upper</code> that are vectors the same length as <code>theta</code> giving the box constraints for the parameters
<code>prior</code>	A function returning the log-density function of the parameter prior distribution. THIS MUST BE A FUNCTION OF ONLY THE FREE PARAMETERS. Any fixed parameters should not be included.
<code>need.hess</code>	A logical value which decides whether or not to evaluate the Hessian for parameter standard errors
<code>initialSANN</code>	Control list for <code>optim</code> when simulated annealing is used for obtaining start values. See details
<code>attempts</code>	The number of times likelihood optimization will be attempted

Details

A full model specification involves 4 components: a movement model, a stopping model, 2 location error models, and a drift indication. The movement model (`mov.model`) specifies how the movement parameters should vary over time. This is a function of specified, time-indexed, covariates. The movement parameters (sigma for velocity variation and beta for velocity autocorrelation) are both modeled with a log link as $\text{par} = \exp(\eta)$, where η is the linear predictor based on the covariates. The `err.model` specification is a list of 2 such models, one for "longitude" and one for "latitude" (in that order) location error. If only one location error model is given, it is used for both coordinates (parameter values as well). If `drift.model` is set to TRUE, then, 2 additional parameters are estimated for the drift process, a drift variance and a beta multiplier. If `polar.coord=TRUE` then the ad-hoc longitude correction factor described by Johnson et al. (2008) (Ecology 89:1208-1215) is used to adjust the variance scale for the longitude model.

The `initial.state` is a list with the following elements (with the exact names):

`a` A vector with initial state values. It has 4 elements (x location at time 1, x velocity at time 1, y location at time 1, y velocity at time 1) for non-drift models and 6 elements for drift models (x location at time 1, x velocity at time 1, x drift velocity at time 1, etc...).

`theta` and `fixPar` are vectors with the appropriate number of parameters. `theta` contains only those parameters which are to be estimated, while `fixPar` contains all parameter values with NA for parameters which are to be estimated.

The data set specified by `data` must contain a numeric or `POSIXct` column which is used as the time index for analysis. The column name is specified by the `Time.name` argument. If a `POSIXct` column is used it is internally converted to a numeric vector with units of hours. The `spacetime` package supports an `STIDF` object that contains slots for both spatial and time series data types. If `data` is of class `STIDF` then the spatial and temporal information are automatically extracted and `polar.coord`, `Time.name` and `coord` are not required. If your data are not compatible with these data structures, it is better to convert it yourself prior to analysis with `crawl`. Also, for stopping models, the stopping covariate must be between 0 and 1 inclusive, with 1 representing complete stop of the animal (no true movement, however, location error can still occur) and 0 represent unhindered movement. The coordinate location should have NA where no location is recorded, but there is a change in the movement covariates.

The CTCRW models can be difficult to provide good initial values for optimization. If `initialSANN` is specified then simulated annealing is used first to obtain starting values for the specified optimization method. If simulated annealing is used first, then the returned `init` list of the `crwFit` object will be a list with the results of the simulated annealing optimization.

Value

A list with the following elements:

<code>par</code>	Parameter maximum likelihood estimates (including fixed parameters)
<code>estPar</code>	MLE without fixed parameters
<code>se</code>	Standard error of MLE
<code>ci</code>	95% confidence intervals for parameters
<code>Cmat</code>	Parameter covariance matrix
<code>loglik</code>	Maximized log-likelihood value
<code>aic</code>	Model AIC value
<code>initial.state</code>	Initial state provided to <code>crwMLE</code> for model fitting
<code>coord</code>	Coordinate names provided for fitting
<code>fixPar</code>	Fixed parameter values provided
<code>convergence</code>	Indicator of convergence (0 = converged)
<code>message</code>	Messages given by <code>optim</code> during parameter optimization
<code>activity</code>	Model provided for stopping variable
<code>drift</code>	Logical value indicating random drift model
<code>mov.model</code>	Model description for movement component
<code>err.model</code>	Model description for location error component
<code>n.par</code>	number of parameters
<code>nms</code>	parameter names
<code>n.mov</code>	number of movement parameters
<code>n.errX</code>	number of location error parameters for "longitude" error model
<code>n.errY</code>	number of location error parameters for "latitude" error model
<code>stop.mf</code>	covariate for stop indication in stopping models

polar.coord	Logical indicating coordinates are polar latitude and longitude
init	Initial values for parameter optimization
data	Original data.frame used to fit the model
lower	The lower parameter bounds
upper	The upper parameter bounds
need.hess	Logical value
runTime	Time used to fit model

Author(s)

Devin S. Johnson, Josh M. London

crwN211	<i>-2 * log-likelihood for CTCRW models</i>
---------	---

Description

This function is designed for primary use within the [crwMLE](#) model fitting function. But, it can be accessed for advanced R and `crawl` users. Uses the state-space parameterization and Kalman filter method presented in Johnson et al. (2008).

Usage

```
crwN211(theta, fixPar, y, noObs, delta, a, P, mov.mf, err.mfX, err.mfY,
        rho = NULL, activity = NULL, n.errX, n.errY, n.mov, driftMod, prior,
        need.hess, constr = list(lower = -Inf, upper = Inf))
```

Arguments

theta	parameter values.
fixPar	values of parameters held fixed (contains NA for theta values).
y	N by 2 matrix of coordinates with the longitude coordinate in the first column.
noObs	vector with 1 for unobserved locations, and 0 for observed locations.
delta	time difference to next location.
a	initial state mean.
P	initial state covariance matrix
mov.mf	Movement covariate data.
err.mfX	longitude error covariate data.
err.mfY	latitude error covariate data.
rho	A vector of known correlation coefficients for the error model, typically used for modern ARGOS data.
activity	Stopping covariate (= 0 if animal is not moving).

n.errX	number or longitude error parameters.
n.errY	number of latitude error parameters.
n.mov	number or movement parameters.
driftMod	Logical. indicates whether a drift model is specified.
prior	Function of theta that returns the log-density of the prior
need.hess	Whether or not the Hessian will need to be calculated from this call
constr	Named list giving the parameter constraints

Details

This function calls compiled C++ code which can be viewed in the `src` directory of the `crawl` source package.

Value

$-2 * \log$ -likelihood value for specified CTCRW model.

Author(s)

Devin S. Johnson

References

Johnson, D., J. London, M. -A. Lea, and J. Durban. 2008. Continuous-time model for animal telemetry data. *Ecology* 89:1208-1215.

See Also

[crwMLE](#)

crwPostIS

Simulate a value from the posterior distribution of a CTCRW model

Description

The `crwPostIS` draws a set of states from the posterior distribution of a fitted CTCRW model. The draw is either conditioned on the fitted parameter values or "full" posterior draw with approximated parameter posterior

Usage

```
crwPostIS(object.sim, fullPost = TRUE, df = Inf, scale = 1,
  thetaSamp = NULL)
```

Arguments

object.sim	A crwSimulator object from crwSimulator .
fullPost	logical. Draw parameter values as well to simulate full posterior
df	degrees of freedom for multivariate t distribution approximation to parameter posterior
scale	Extra scaling factor for t distribution approximation
thetaSamp	If multiple parameter samples are available in object.sim, setting thetaSamp=n will use the nth sample. Defaults to the last.

Details

The crwPostIS draws a posterior sample of the track state matrices. If fullPost was set to TRUE when the object.sim was build in [crwSimulator](#) then a psuedo-posterior draw will be made by first sampling a parameter value from a multivariate t distribution which approximates the marginal posterior distribution of the parameters. The covariance matrix from the fitted model object is used to scale the MVt approximation. In addition, the factor "scale" can be used to further adjust the approximation. Further, the parameter simulations are centered on the fitted values.

To correct for the MVt approximation, the importance sampling weight is also supplied. When calculating averages of track functions for Bayes estimates one should use the importance sampling weights to calculate a weighted average (normalizing first, so the weights sum to 1).

Value

List with the following elements:

alpha.sim.y	A matrix a simulated latitude state values
alpha.sim.x	Matrix of simulated longitude state values
locType	Indicates prediction types with a "p" or observation times with an "o"
Time	Initial state covariance for latitude
loglik	log likelihood of simulated parameter
par	Simulated parameter value
log.isw	non normalized log importance sampling weight

Author(s)

Devin S. Johnson

See Also

See `demo(northernFurSealDemo)` for example.

crwPredict	<i>Predict animal locations and velocities using a fitted CTCRW model and calculate measurement error fit statistics</i>
------------	--

Description

The `crwMEfilter` function uses a fitted model object from `crwMLE` to predict animal locations (with estimated uncertainty) at times in the original data set and supplemented by times in `predTime`. If `speedEst` is set to `TRUE`, then animal log-speed is also estimated. In addition, the measurement error shock detection filter of de Jong and Penzer (1998) is also calculated to provide a measure for outlier detection.

Usage

```
crwPredict(object.crwFit, predTime = NULL, flat = TRUE, ...)
```

Arguments

<code>object.crwFit</code>	A model object from <code>crwMLE</code> .
<code>predTime</code>	vector of additional prediction times (numeric or <code>POSIXct</code>). Alternatively, a character vector specifying a time interval (see <code>Details</code>).
<code>flat</code>	logical. Should the result be returned as a flat <code>data.frame</code> .
<code>...</code>	Additional arguments for testing new features

Details

The requirements for data are the same as those for fitting the model in `crwMLE`.

- ("predTime") `predTime` can be either passed as a separate vector of `POSIXct` or numeric values for additional prediction times beyond the observed times. If the original data were provided as a `POSIXct` type, then `crwPredict` can derive a sequence of regularly spaced prediction times from the original data. This is specified by providing a character string that corresponds to the `by` argument of the `seq.POSIXt` function (e.g. '1 hour', '30 mins'). `crwPredict` will round the first observed time up to the nearest unit (e.g. '1 hour' will round up to the nearest hour, '30 mins' will round up to the nearest minute) and start the sequence from there. The last observation time is truncated down to the nearest unit to specify the end time.

Value

List with the following elements:

<code>originalData</code>	A <code>data.frame</code> with is data merged with <code>predTime</code> .
<code>alpha.hat</code>	Predicted state
<code>Var.hat</code>	array where <code>Var.hat[, , i]</code> is the prediction covariance matrix for <code>alpha.hat[, i]</code> .
<code>fit.test</code>	A <code>data.frame</code> of chi-square fit (<code>df=2</code>) statistics and naive (pointwise) p-values.

If `flat` is set to `TRUE` then a data set is returned with the columns of the original data plus the state estimates, standard errors (`se`), speed estimates, and the fit statistics and naive p-values.

Author(s)

Devin S. Johnson

References

de Jong, P. and Penzer, J. (1998) Diagnosing shocks in time series. *Journal of the American Statistical Association* 93:796-806.

crwPredictPlot	<i>Plot CRW predicted object</i>
----------------	----------------------------------

Description

Creates 2 types of plots of a crwPredict object: a plot of both coordinate axes with prediction intervals and a plot of just observed locations and predicted locations.

Usage

```
crwPredictPlot(object, plotType = "ll", ...)
```

Arguments

object	crwPredict object.
plotType	type of plot has to be one of the following: "map" or "ll" (default).
...	Further arguments passed to plotting commands.

Value

A plot.

Author(s)

Devin S. Johnson and Sebastian Luque

See Also

See `demo(northernFurSealDemo)` for additional examples.

crwSamplePar	<i>Create a weighted importance sample for posterior predictive track simulation.</i>
--------------	---

Description

The `crwSamplePar` function uses a fitted model object from `crwMLE` and a set of prediction times to construct a list from which `crwPostIS` will draw a sample from either the posterior distribution of the state vectors conditional on fitted parameters or a full posterior draw from an importance sample of the parameters.

Usage

```
crwSamplePar(object.sim, method = "IS", size = 1000, df = Inf,
             grid.eps = 1, crit = 2.5, scale = 1, force.quad)
```

Arguments

<code>object.sim</code>	A simulation object from <code>crwSimulator</code> .
<code>method</code>	Method for obtaining weights for movement parameter samples
<code>size</code>	Size of the parameter importance sample
<code>df</code>	Degrees of freedom for the t approximation to the parameter posterior
<code>grid.eps</code>	Grid size for <code>method="quadrature"</code>
<code>crit</code>	Criterion for deciding "significance" of quadrature points (difference in log-likelihood)
<code>scale</code>	Scale multiplier for the covariance matrix of the t approximation
<code>force.quad</code>	A logical indicating whether or not to force the execution of the quadrature method for large parameter vectors.

Details

The `crwSamplePar` function uses the information in a `crwSimulator` object to create a set of weights for importance sample-resampling of parameters in a full posterior sample of parameters and locations using `crwPostIS`. This function is usually called from `crwPostIS`. The average user should have no need to call this function directly.

Value

List with the following elements:

<code>x</code>	Longitude coordinate with NA at prediction times
<code>y</code>	Similar to above for latitude
<code>locType</code>	Indicates prediction types with a "p" or observation times with an "o"
<code>P1.y</code>	Initial state covariance for latitude

P1.x	Initial state covariance for longitude
a1.y	Initial latitude state
a1.x	Initial longitude state
n.errX	number of longitude error model parameters
n.errY	number of latitude error model parameters
delta	vector of time differences
driftMod	Logical. indicates random drift model
stopMod	Logical. Indicated stop model fitted
stop.mf	stop model design matrix
err.mfX	Longitude error model design matrix
err.mfY	Latitude error model design matrix
mov.mf	Movement model design matrix
fixPar	Fixed values for parameters in model fitting
Cmat	Covariance matrix for parameter sampling distribution
Lmat	Cholesky decomposition of Cmat
par	fitted parameter values
N	Total number of locations
loglik	log likelihood of the fitted model
Time	vector of observation times
coord	names of coordinate vectors in original data
Time.name	Name of the observation times vector in the original data
thetaSampList	A list containing a data frame of parameter vectors and their associated probabilities for a resample

Author(s)

Devin S. Johnson

See Also

See `demo(northernFurSealDemo)` for example.

crwSimulator

Construct a posterior simulation object for the CTCRW state vectors

Description

The `crwSimulator` function uses a fitted model object from `crwMLE` and a set of prediction times to construct a list from which `crwPostIS` will draw a sample from either the posterior distribution of the state vectors conditional on fitted parameters or a full posterior draw from an importance sample of the parameters.

Usage

```
crwSimulator(object.crwFit, predTime = NULL, method = "IS", parIS = 1000,
             df = Inf, grid.eps = 1, crit = 2.5, scale = 1, force.quad)
```

Arguments

<code>object.crwFit</code>	A model object from <code>crwMLE</code> .
<code>predTime</code>	vector of additional prediction times.
<code>method</code>	Method for obtaining weights for movement parameter samples
<code>parIS</code>	Size of the parameter importance sample
<code>df</code>	Degrees of freedom for the t approximation to the parameter posterior
<code>grid.eps</code>	Grid size for <code>method="quadrature"</code>
<code>crit</code>	Criterion for deciding "significance" of quadrature points (difference in log-likelihood)
<code>scale</code>	Scale multiplier for the covariance matrix of the t approximation
<code>force.quad</code>	A logical indicating whether or not to force the execution of the quadrature method for large parameter vectors.

Details

The `crwSimulator` function produces a list and preprocesses the necessary components for repeated track simulation from a fitted CTCRW model from `crwMLE`. The `method` argument can be one of "IS" or "quadrature". If `method="IS"` is chosen standard importance sampling will be used to calculate the appropriate weights via t proposal with `df` degrees of freedom. If `df=Inf` (default) then a multivariate normal distribution is used to approximate the parameter posterior. If `method="quadrature"`, then a regular grid over the posterior is used to calculate the weights. The argument `grid.eps` controls the quadrature grid. The arguments are approximately the upper and lower limit in terms of standard deviations of the posterior. The default is `grid.eps`, in units of 1sd. If `object.crwFit` was fitted with `crwArgoFilter`, then the returned list will also include `p.out`, which is the approximate probability that the observation is an outlier.

Value

List with the following elements:

x	Longitude coordinate with NA at prediction times
y	Similar to above for latitude
locType	Indicates prediction types with a "p" or observation times with an "o"
P1.y	Initial state covariance for latitude
P1.x	Initial state covariance for longitude
a1.y	Initial latitude state
a1.x	Initial longitude state
n.errX	number of longitude error model parameters
n.errY	number of latitude error model parameters
delta	vector of time differences
driftMod	Logical. indicates random drift model
stopMod	Logical. Indicated stop model fitted
stop.mf	stop model design matrix
err.mfX	Longitude error model design matrix
err.mfY	Latitude error model design matrix
mov.mf	Movement model design matrix
fixPar	Fixed values for parameters in model fitting
Cmat	Covariance matrix for parameter sampling distribution
Lmat	Cholesky decomposition of Cmat
par	fitted parameter values
N	Total number of locations
loglik	log likelihood of the fitted model
Time	vector of observation times
coord	names of coordinate vectors in original data
Time.name	Name of the observation times vector in the original data
thetaSampList	A list containing a data frame of parameter vectors and their associated probabilities for a resample

Author(s)

Devin S. Johnson

See Also

See `demo(northernFurSealDemo)` for example.

displayPar	<i>Display the order of parameters along with fixed values and starting values</i>
------------	--

Description

This function takes the model specification arguments to the `crwMLE` function and displays a table with the parameter names in the order that `crwMLE` will use during model fitting. This is useful for specifying values for the `fixPar` or `theta` (starting values for free parameters) arguments.

Usage

```
displayPar(mov.model = ~1, err.model = NULL, activity = NULL,
           drift = FALSE, data, Time.name, theta, fixPar, ...)
```

Arguments

<code>mov.model</code>	formula object specifying the time indexed covariates for movement parameters.
<code>err.model</code>	A 2-element list of formula objects specifying the time indexed covariates for location error parameters.
<code>activity</code>	formula object giving the covariate for the stopping portion of the model.
<code>drift</code>	logical indicating whether or not to include a random drift component.
<code>data</code>	data.frame object containing telemetry and covariate data. A <code>SpatialPointsDataFrame</code> object from the package 'sp' will also be accepted.
<code>Time.name</code>	character indicating name of the location time column
<code>theta</code>	starting values for parameter optimization.
<code>fixPar</code>	Values of parameters which are held fixed to the given value.
<code>...</code>	Additional arguments (probably for testing new features.)

Value

A data frame with the following columns

<code>ParNames</code>	The names of the parameters specified by the arguments.
<code>fixPar</code>	The values specified by the <code>fixPar</code> argument for fixed values of the parameters. In model fitting, these values will remain fixed and will not be estimated.
<code>thetaIdx</code>	This column provides the index of each element of the <code>theta</code> argument and to which parameter it corresponds.
<code>thetaStart</code>	If a value is given for the <code>theta</code> argument it will be placed in this column and its elements will correspond to the <code>thetaIdx</code> column.

Author(s)

Devin S. Johnson

See Also

demo(northernFurSealDemo) for example.

expandPred

Expand a time indexed data set with additional prediction times

Description

Expands a covariate data frame (or vector) that has a separate time index by inserting prediction times and duplicating the covariate values for all prediction time between subsequent data times.

Usage

```
expandPred(x, Time = "Time", predTime, time.col = FALSE)
```

Arguments

x	Data to be expanded.
Time	Either a character naming the column which contains original time values, or a numeric vector of original times
predTime	prediction times to expand data
time.col	Logical value indicating whether to attach the new times to the expanded data

Value

data.frame expanded by predTime

Author(s)

Devin S. Johnson

Examples

```
#library(crawl)
origTime <- c(1:10)
x <- cbind(rnorm(10), c(21:30))
predTime <- seq(1,10, by=0.25)
expandPred(x, Time=origTime, predTime, time.col=TRUE)
```

fillCols	<i>Fill missing values in data set (or matrix) columns for which there is a single unique value</i>
----------	---

Description

Looks for columns in a data set that have a single unique non-missing value and fills in all NA with that value

Usage

```
fillCols(data)
```

Arguments

data	data.frame
------	------------

Value

data.frame

Author(s)

Devin S. Johnson

Examples

```
#library(crawl)
data1 <- data.frame(constVals=rep(c(1,NA),5), vals=1:10)
data1[5,2] <- NA
data1
data2 <- fillCols(data1)
data2

mat1 <- matrix(c(rep(c(1,NA),5), 1:10), ncol=2)
mat1[5,2] <- NA
mat1
mat2 <- fillCols(mat1)
mat2
```

fix_path	<i>Project path away from restricted areas</i>
----------	--

Description

Corrects a path so that it does not travel through a restricted area.

Usage

```
fix_path(xy, time, res_raster, trans)
```

Arguments

xy	Coordinate locations for the path. Can be one of the following classes: (1) a two column matrix, (2) 'SpatialPoints' or 'SpatialPointsDataFrame' object from the sp package, (3) 'crwPredict' object from the crwPredict function (4) 'crwIS' object from the crwPostIS function
time	A vector of times associated with xy locations
res_raster	An indicator raster object with cells = 1 if it is 'off-limits' and 0 otherwise.
trans	A transition matrix object from the gdistance package.

Value

Either matrix or 'SpatialPoints' object with path projected around restricted areas

flatten	<i>'Flattening' a list-form crwPredict object into a data.frame</i>
---------	---

Description

"Flattens" a list form `crwPredict` object into a flat `data.frame`.

Usage

```
flatten(predObj)
```

Arguments

predObj	A <code>crwPredict</code> object
---------	----------------------------------

Value

a `data.frame` version of a `crwPredict` list with columns for the state standard errors

Author(s)

Devin S. Johnson

See Also

[northernFurSeal](#) for use example

get_restricted_segments

Find the sections of a path that pass through a restricted area

Description

This function is used to identify sections of a path that pass through restricted areas. the CTCRW model in `crawl` cannot actively steer paths away from restricted areas as it knows nothing of spatial information. So, this function will identify areas that for which the unrestrained path passes through these areas. If the path/points end within the restricted area, those records will be removed. The user can then use this information to adjust the path as desired.

Usage

```
get_restricted_segments(xy, res_raster)
```

Arguments

<code>xy</code>	A <code>SpatialPoints</code> object from the <code>sp</code> package or a 2-column matrix of x and y locations
<code>res_raster</code>	A raster object from the <code>raster</code> package that indicates restricted areas with a 1, else 0 for unrestricted areas.

Value

A `data.frame` with each row associated with each section of the path that crosses a restricted area. The columns provide the start and end row indices of `xy` where the section occurs and the previous and post locations that are in unrestricted space.

Author(s)

Josh M. London (josh.london@noaa.gov)

harborSeal	<i>Harbor seal relocation data set used in Johnson et al. (2008)</i>
------------	--

Description

Harbor seal relocation data set used in Johnson et al. (2008)

Format

A data frame with 7059 observations on the following 5 variables.

Time a numeric vector.

latitude a numeric vector.

longitude a numeric vector.

DryTime a numeric vector.

Argos_loc_class a factor with levels 0 1 2 3 A B.

Author(s)

Devin S. Johnson

Source

Marine Mammal Laboratory, Alaska Fisheries Science Center, National Marine Fisheries Service, NOAA 7600 Sand Point Way NE Seattle, WA 98115

References

Johnson, D., J. London, M. -A. Lea, and J. Durban (2008) Continuous-time random walk model for animal telemetry data. *Ecology* 89:1208-1215.

intToPOSIX	<i>Reverse as.numeric command that is performed on a vector of type POSIXct</i>
------------	---

Description

Takes integer value produced by `as.numeric(x)`, where `x` is a `POSIXct` vector and returns it to a `POSIXct` vector

Usage

```
intToPOSIX(timeVector, tz = "GMT")
```

Arguments

timeVector A vector of integers produced by `as.numeric` applied to a POSIXct vector
 tz Time zone of the vector (see `as.POSIXct`).

Value

POSIXct vector

Note

There is no check that `as.numeric` applied to a POSIX vector produced `timeVector`. So, caution is required in using this function. It was included simply because I have found it useful

Author(s)

Devin S. Johnson

Examples

```
#library(crawl)
timeVector <- as.numeric(Sys.time())
timeVector
intToPOSIX(timeVector, tz="")
```

<code>mergeTrackStop</code>	<i>Merge a location data set with a dry time (or other stopping) covariate</i>
-----------------------------	--

Description

The function merges a location data set with a stopping variable data set.

Usage

```
mergeTrackStop(data, stopData, Time.name = "Time", interp = c("zeros",
  "ma0"), win = 2, constCol)
```

Arguments

data Location data.
 stopData stopping variable data set.
 Time.name character naming time index variable in both data sets
 interp method of interpolation.
 win window for "ma0" interpolation method.
 constCol columns in data for which the user would like to be constant, such as id or sex.

Details

Simply merges the data frames and interpolates based on the chosen method. Both data frames have to use the same name for the time variable. Also contains stopType which = "o" if observed or "p" for interpolated.

The merged data is truncated to the first and last time in the location data set. Missing values in the stopping variable data set can be interpolated by replacing them with zeros (full movement) or first replacing with zeros then using a moving average to smooth the data. Only the missing values are then replace with this smoothed data. This allows a smooth transition to full movement.

Value

Merged data.frame with new column from stopData. Missing values in the stopping variable will be interpolated

Author(s)

Devin S. Johnson

Examples

```
track <- data.frame(TimeVar=sort(runif(20,0,20)), x=1:20, y=20:1)
track
stopData <- data.frame(TimeVar=0:29, stopVar=round(runif(30)))
stopData
mergeTrackStop(track, stopData, Time.name="TimeVar")
```

northernFurSeal

Northern fur seal pup relocation data set used in Johnson et al. (2008)

Description

Northern fur seal pup relocation data set used in Johnson et al. (2008)

Format

A data frame with 795 observations on the following 4 variables:

Time a numeric vector.

Argos_loc_class a factor with levels 0 1 2 3 A.

latitude a numeric vector.

longitude a numeric vector.

Source

Marine Mammal Laboratory, Alaska Fisheries Science Center, National Marine Fisheries Service, NOAA 7600 Sand Point Way NE Seattle, WA 98115

References

Johnson, D., J. London, M. -A. Lea, and J. Durban (2008) Continuous-time random walk model for animal telemetry data. Ecology 89:1208-1215.

tidy_crwFit	<i>tidy-like method for crwFit object</i>
-------------	---

Description

this function mimics the approach taken by broom::tidy to present model output parameters in a tidy, data frame structure.

Usage

```
tidy_crwFit(fit)
```

Arguments

fit	crwFit object from crawl::crwMLE
-----	----------------------------------

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