Package 'ipsecr'

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Description Estimates the density of a spatially distributed animal population sampled with an array of passive detectors, such as traps. Models incorporating distance-dependent detection are fitted by simulation and inverse prediction as proposed by Efford (2004) <doi:10.1111/j.0030-1299.2004.13043.x>.

License GPL (>= 2)

LazyData yes

LazyDataCompression xz

URL https://github.com/MurrayEfford/ipsecr/,

https://www.otago.ac.nz/density/

NeedsCompilation yes

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ipsecr-package

Spatially Explicit Capture–Recapture by Inverse Prediction

Description

Functions to estimate the density and size of a spatially distributed animal population sampled with an array of passive detectors, such as traps. **ipsecr** addresses 'difficult' models that strictly cannot be fitted by maximum likelihood in the larger parent package **secr** (Efford 2022). The classic example concerns data from single-catch traps.

Details

Package:	ipsecr
Type:	Package
Version:	1.1.2
Date:	2022-06-22
License:	GNU General Public License Version 2 or later

Spatially explicit capture–recapture is a set of methods for studying marked animals distributed in space. Data comprise the locations of detectors (described in an object of class 'traps'), and the detection histories of individually marked animals. Individual histories are stored in an object of class 'capthist' that includes the relevant 'traps' object.

Models for population density (animals per hectare) and detection are defined in **ipsecr** using symbolic formula notation (but only constant models are allowed for detection parameters in **ipsecr** 1.1) Density models may include spatial trend. Habitat is distinguished from nonhabitat with an object of class 'mask'.

Models are fitted in **ipsecr** by simulation and inverse prediction (Efford 2004). A model fitted with **ipsecr.fit** is an object of class ipsecr. Generic methods (plot, print, summary, etc.) are provided.

A link at the bottom of each help page takes you to the help index. Several vignettes complement the help pages:

The analyses in **ipsecr** extend those available in the software Density (see www.otago.ac.nz/density/ for the most recent version of Density). Help is available on the 'DENSITY | secr' forum at

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www.phidot.org and the Google group secrgroup. Feedback on the software is also welcome, including suggestions for additional documentation or new features consistent with the overall design.

'Inverse prediction' uses methods from multivariate calibration (Brown 1982). The goal is to estimate population density (D) and the parameters of a detection function (usually g0 or lambda0 and sigma) by 'matching' statistics from proxyfn(capthist) (the target vector) and statistics from simulations of a 2-D population using the postulated detection model. Statistics (see Note) are defined by the proxy function, which should return a vector equal in length to the number of parameters (default np = 3). Simulations of the 2-D population use either internal C++ code or sim.popn.

The simulated population is sampled with internal C++ code, sim.capthist, or a user-specified function. Simulations match the detector type (e.g., 'single' or 'multi') and detector layout specified in traps(capthist), including allowance for varying effort if the layout has a usage attribute.

The simulated 2-D distribution of animals is Poisson by default.

Simulations are usually conducted on a factorial experimental design in parameter space - i.e. at the vertices of a cuboid 'box' centred on the working values of the parameters, plus an optional number of centre points.

A multivariate linear model is fitted to predict each vector of simulated proxies from the known parameter values. The number of simulations at each design point is increased until a specified precision is reached, up to the limit max.nsim.

Once a model with sufficient precision has been obtained, a new working vector of parameter estimates is 'predicted' by inverting the linear model and applying it to the target vector. A working vector is accepted as the final estimate when it lies within the box; this reduces the bias from using a linear approximation to extrapolate a nonlinear function. If the working vector lies outside the box then a new design is centred on value for each parameter in the working vector.

Once a final estimate is accepted, further simulations are conducted to estimate the variancecovariance matrix. These also provide a parametric bootstrap sample to evaluate possible bias. Set var.nsim = 0 to suppress the variance step.

See Efford et al. (2004) for another description of the method, and Efford et al. (2005) for an application.

If not provided, the starting values are determined automatically with the **secr** function makeStart.

Linear measurements are assumed to be in metres and density in animals per hectare (10 000 m²).

If ncores > 1 the **parallel** package is used to create processes on multiple cores (see Parallel for more).

Author(s)

Murray Efford <murray.efford@otago.ac.nz>

References

Brown, P. J. (1982) Multivariate calibration. *Journal of the Royal Statistical Society, Series B* 44, 287–321.

Efford, M. G. (2004) Density estimation in live-trapping studies. Oikos 106, 598-610.

Efford, M. G. (2022) secr: Spatially explicit capture–recapture models. R package version 4.5.4. https://CRAN.R-project.org/package=secr/ Efford, M. G., Borchers D. L. and Byrom, A. E. (2009) Density estimation by spatially explicit capture–recapture: likelihood-based methods. In: D. L. Thompson, E. G. Cooch and M. J. Conroy (eds) *Modeling Demographic Processes in Marked Populations*. Springer. Pp. 255–269.

Efford, M. G., Dawson, D. K. and Robbins C. S. (2004) DENSITY: software for analysing capture-recapture data from passive detector arrays. *Animal Biodiversity and Conservation* **27**, 217–228.

Efford, M. G., Warburton, B., Coleman, M. C. and Barker, R. J. (2005) A field test of two methods for density estimation. *Wildlife Society Bulletin* **33**, 731–738.

Otis, D. L., Burnham, K. P., White, G. C. and Anderson, D. R. (1978) Statistical inference from capture data on closed animal populations. *Wildlife Monographs* **62**.

See Also

proxy.ms ipsecr.fit, secr.fit, capthist, mask

details

Detail Specification for ipsecr.fit

Description

The function ipsecr.fit allows many options. Some of these are used infrequently and have been bundled as a single argument details to simplify the documentation. They are described here.

Detail components

details\$contrasts may be used to specify the coding of factor predictors. The value should be suitable for the 'contrasts.arg' argument of model.matrix. See 'Trend across sessions' in secr-multisession.pdf for an example.

details\$debug is used only for debugging. In ordinary use it should not be changed from the default (FALSE).

details\$distribution specifies the distribution of the number of individuals detected n; this may be conditional on the number in the masked area ("binomial") or unconditional ("poisson"). distribution affects the sampling variance of the estimated density. The default is "poisson". The component 'distribution' may also take a numeric value larger than nrow(capthist), rather than "binomial" or "poisson".

details\$factorial "full" or "fractional" design. Fractional requires the package **FrF2** (Groemping 2014).

details\$FrF2args list of arguments defining a fractional design

details\$ignorenontarget = TRUE causes the function to ignore non-target information (the attribute 'nontarget').

details\$ignoreusage = TRUE causes the function to ignore usage (varying effort) information in the traps component. The default (details\$ignoreusage = FALSE) is to include usage in the model.

details\$popmethod chooses between internal C++ code, the secr function sim.popn, and a userprovided R function with arguments 'mask', 'D' (density per cell of mask) and 'N' (number of individuals to simulate).

Internal

details\$CHmethod chooses between internal C++ code, the **secr** function sim.capthist, and a user-provided R function with arguments 'traps', 'popn', 'detectfn', 'detectpar', and 'noccasions'. Defaults

- boxsize1 scalar or vector of length np for size of design
- boxsize2 as for boxsize1; used from second box onwards
- centre number of centre points in simulation design
- min.nsim minimum number of simulations per point
- max.nsim maximum number of simulations per point
- · dev.max tolerance for precision of points in predictor space
- · var.nsim number of additional simulations to estimate variance-covariance matrix
- min.nbox minimum number of attempts to 'frame' solution
- · max.nbox maximum number of attempts to 'frame' solution
- · max.ntries maximum number of attempts at each simulation

References

Groemping, U. (2014). R Package FrF2 for Creating and Analyzing Fractional Factorial 2-Level Designs. *Journal of Statistical Software*, **56**, 1–56. https://www.jstatsoft.org/article/view/v056i01.

See Also

ipsecr.fit

Internal

Internal Functions

Description

Functions called internally by ipsecr.

Usage

```
proxy.ms(capthist, model = list(D = ~1, NT = ~1), trapdesigndata = NULL, ...)
proxyfn0(capthist, N.estimator = c("n", "null","zippin","jackknife"), ...)
proxyfn1(capthist, N.estimator = c("n", "null","zippin","jackknife"), ...)
simpop(mask, D, N, distribution)
simCH(traps, popn, detectfn, detectpar, NT, noccasions, details = list())
```

Internal

Arguments

capthist	secr capthist object
model	named list of model formulae (see ipsecr.fit)
trapdesigndata	dataframe with one row for each detector and session
	other arguments (not used)
N.estimator	character name of closed-population estimator
mask	secr mask object
D	numeric density in each mask cell
Ν	integer number of animals to simulate
distribution	character 'even' or 'poisson'
traps	detector locations as secr traps object
popn	animal locations as secr popn object
detectfn	integer code for detection function (see detectfn)
detectpar	named list of parameter values
NT	numeric hazard of non-target interference at each detector
noccasions	integer number of sampling occasions
details	list with optional additional named arguments

Details

proxy.ms is the default proxyfn used by ipsecr.fit.

simpop is used by ipsecr.fit for popmethod 'internal'. It is faster and simpler than the secr function sim.popn.

simCH is used by ipsecr.fit for CHmethod 'internal'. It is faster and simpler than the secr function sim.capthist, and optionally simulates non-target interference.

D and NT are matrices with one column per session.

Value

proxy.ms – a numeric vector of length >= 3 corresponding to proxies for multi-session density and non-target interference models.

proxyfn0, proxyfn1 - a numeric vector of length 3 corresponding to proxies for population size, capture probability intercept and scale of detection.

simCH - a single-session capthist object.

References

Efford, M. G. (2004) Density estimation in live-trapping studies. Oikos 106, 598-610.

See Also

ipsecr.fit

ipsecr.fit

Examples

proxy.ms(captdata)

ipsecr.fit

Spatially Explicit Capture–Recapture by Inverse Prediction

Description

Estimate population density by simulation and inverse prediction (Efford 2004; Efford, Dawson & Robbins 2004). A restricted range of SECR models may be fitted.

Usage

```
ipsecr.fit(capthist, proxyfn = proxy.ms, model = list(D ~ 1, g0 ~ 1, sigma ~ 1),
mask = NULL, buffer = 100, detectfn = "HN", binomN = NULL, start = NULL,
link = list(), fixed = list(), timecov = NULL, sessioncov = NULL,
details = list(), verify = TRUE, verbose = TRUE, ncores = NULL,
seed = NULL, ...)
```

Arguments

capthist	secr capthist object including capture data and detector (trap) layout
proxyfn	function to compute proxy from capthist for each coefficient (beta parameter)
model	list with optional components each symbolically defining a linear predictor for one real parameter using formula notation
mask	mask object
buffer	scalar mask buffer radius in metres if mask not specified
detectfn	integer code or character string for shape of detection function $0 =$ halfnormal, 1 = hazard rate etc. – see detectfn
binomN	integer code for distribution of counts (see Details)
start	vector of initial values for beta parameters, or ipsecr object from which they may be derived
link	list with optional components corresponding to 'real' parameters (e.g., 'D', 'g0', 'sigma'), each a character string in {"log", "logit", "identity", "sin"} for the link function of one real parameter
fixed	list with optional components corresponding to real parameters giving the scalar value to which the parameter is to be fixed
timecov	optional dataframe of values of time (occasion-specific) covariate(s). NOT USED
sessioncov	optional dataframe of values of session-specific covariate(s)

ipsecr.fit

details	list of additional settings, to control estimation (see Details)
verify	logical, if TRUE the input data are checked with verify
verbose	logical, if TRUE then messages are output during execution
ncores	integer number of cores to use for parallel processing
seed	either NULL or an integer that will be used in a call to set.seed
	other arguments passed to proxy function

Details

The vignette should be consulted for a full exposition.

details is used for various specialized settings listed below. Some of these are described separately - see details.

Name	Default	Description
boxsize1	0.2	scalar or vector of length np for size of design
boxsize2	0.05	as for boxsize1; used from second box onwards
boxtype	'absolute'	'absolute' or 'relative'
centre	3	number of centre points in simulation design
dev.max	0.002	tolerance for precision of points in predictor space
var.nsim	1000	number of additional simulations to estimate variance-covariance matrix
min.nsim	10	minimum number of simulations per point
max.nsim	2000	maximum number of simulations per point
min.nbox	2	minimum number of attempts to 'frame' solution
max.nbox	5	maximum number of attempts to 'frame' solution
max.ntries	2	maximum number of attempts at each simulation
distribution	'poisson'	'poisson', 'binomial' or 'even'
even	FALSE	'binomial' or 'even' population distribution in main simulations
binomN	0	integer code for distribution of counts (unused)
ignorenontarget	FALSE	override nontarget attribute of capthist
ignoreusage	FALSE	override usage in traps object of capthist
debug	FALSE	stop at arbitrary points in execution (varies)
savecall	TRUE	optionally suppress saving of call
newdetector	NULL	detector type to override detector(traps(capthist))
contrasts	NULL	coding of factor predictors
popmethod	'internal'	'internal' or 'sim.popn' or a user-provided function
CHmethod	'internal'	'internal' or 'sim.capthist' or a user-provided function
factorial	'full'	'full' or 'fractional' design
FrF2args	NULL	arguments for FrF2 when factorial = 'fractional'

Value

An object of class 'ipsecr', a list comprising:

call	the function call
capthist	input
mask	input

ipsecr.fit

detectfn	input
timecov	input
start	input
link	input
fixed	input
model	input
details	input
designD	list of design data for density
design	list of design data for detection parameters
design0	list of design data for detection parameters (naive animal)
parindx	mapping of coefficients (beta parameters) to real parameters
vars	names of covariates in model
betanames	names of coefficients
realnames	names of 'real' parameters
code	integer completion code: 1 successful, 2 target not within final box, 3 exceeded maximum simulations
beta	estimates of coefficients on link scale
beta.vcov	variance-covariance matrix of estimates
designbeta	vertices of final box (design points)
ip.nsim	total number of simulations
var.nsim.OK	number of successful variance simulations
variance.boots	•
	dataframe summarising simulations for variance estimation
version	package version
starttime	time execution started
proctime	processor time (seconds)

(The order and composition of the output list is subject to change).

References

Efford, M. G. (2004) Density estimation in live-trapping studies. Oikos 106, 598-610.

Efford, M. G., Dawson, D. K. and Robbins C. S. (2004) DENSITY: software for analysing capture-recapture data from passive detector arrays. *Animal Biodiversity and Conservation* **27**, 217–228.

See Also

proxy.ms, predict.ipsecr, summary.ipsecr

Examples

```
ipsecrdemo <- ipsecr.fit(captdata, ncores = 1, buffer = 100, detectfn = 14, seed = 1237)
```

ipsecrdemo

Description

Demonstration data from program Density are provided as a capthist object (captdata) ready for input to ipsecr.fit.

The fitted models are objects of class ipsecr formed by

ipsecrdemo <- ipsecr.fit(captdata, ncores = 1, detectfn = 14, seed = 1237)</pre>

Usage

data(ipsecrdemo)

Details

The raw data are 235 fictional captures of 76 animals over 5 occasions in 100 single-catch traps 30 metres apart on a square grid with origin at (365,365).

The fitted model uses a hazard halfnormal detection function and default values of other arguments.

Object Description ipsecrdemo fitted ipsecr model – null

References

Efford, M. G. (2012) *DENSITY 5.0: software for spatially explicit capture–recapture*. Department of Mathematics and Statistics, University of Otago, Dunedin, New Zealand. https://www.otago.ac.nz/density/.

See Also

capthist, read.capthist, secrdemo

Examples

predict(ipsecrdemo)

newdata

Create Default Design Data

Description

Internal function used to generate a dataframe containing design data for the base levels of all predictors in an secrobject.

Usage

S3 method for class 'ipsecr'
makeNewData(object, all.levels = FALSE, ...)

Arguments

object	fitted ipsecr model object
all.levels	logical; if TRUE then all levels of factors are included
	other arguments (not used)

Details

makeNewData is used by predict in lieu of user-specified 'newdata'. There is seldom any need to call the function makeNewData directly.

Value

A dataframe with one row for each session and group, and columns for the predictors used by object\$model.

See Also

predict.ipsecr,ipsecr.fit

Examples

from previously fitted model
makeNewData(ipsecrdemo)

plot.ipsecr

Description

Plot detection functions using estimates of parameters in an ipsecr object.

Usage

```
## S3 method for class 'ipsecr'
plot(x, newdata = NULL, add = FALSE,
    sigmatick = FALSE, rgr = FALSE, limits = FALSE, alpha = 0.05,
    xval = 0:200, ylim = NULL, xlab = NULL, ylab = NULL, ...)
```

Arguments

х	an ipsecr object
newdata	dataframe of data to form estimates
add	logical to add curve(s) to an existing plot
sigmatick	logical; if TRUE the scale parameter sigma is shown by a vertical line
rgr	logical; if TRUE a scaled curve $r.g(r)$ is plotted instead of $g(r)$
limits	logical; if TRUE pointwise confidence limits are drawn
alpha	alpha level for confidence intervals
xval	vector of distances at for which detection to be plotted
ylim	vector length 2 giving limits of y axis
xlab	label for x axis
ylab	label for y axis
	arguments to pass to lines

Details

newdata is usually NULL, in which case one curve is plotted for each session and group. Otherwise, predict.ipsecr is used to form estimates and plot a curve for each row in newdata.

If axis labels are not provided they default to 'Distance (m)' and 'Detection probability' or 'Detection lambda'.

Approximate confidence limits for g(r) are calculated using a numerical first-order delta-method approximation to the standard error at each xval. The distribution of g(r) is assumed to be normal on the logit scale for non-hazard functions (detectfn 0:13). For hazard detection functions (detectfn 14:18) the hazard is assumed (from version 3.1.1) to be distributed normally on the log scale. Limits are back-transformed to the probability scale g(r).

predict.ipsecr

Value

plot.ipsecr invisibly returns a dataframe of the plotted values (or a list of dataframes in the case that newdata has more than one row).

See Also

Detection functions, plot, ipsecr, detectfnplot

Examples

plot (ipsecrdemo, xval = 0:100, ylim = c(0, 0.4))

predict.ipsecr

SECR Model Predictions

Description

Evaluate a spatially explicit capture–recapture model. That is, compute the 'real' parameters corresponding to the 'beta' parameters of a fitted model for arbitrary levels of any variables in the linear predictor.

Usage

```
## S3 method for class 'ipsecr'
predict(object, newdata = NULL, type = c("response", "link"),
    se.fit = TRUE, alpha = 0.05, savenew = FALSE, ...)
```

Arguments

object	ipsecr object output from ipsecr.fit
newdata	optional dataframe of values at which to evaluate model
type	character; type of prediction required. The default ("response") provides esti- mates of the 'real' parameters.
se.fit	logical for whether output should include SE and confidence intervals
alpha	alpha level for confidence intervals
savenew	logical for whether newdata should be saved
	other arguments passed to newdata

Details

The variables in the various linear predictors are described in secr-models.pdf and listed for the particular model in the vars component of object.

Optional newdata should be a dataframe with a column for each of the variables in the model (see 'vars' component of object). If newdata is missing then a dataframe is constructed automatically.

Default newdata are for a naive animal on the first occasion; numeric covariates are set to zero and factor covariates to their base (first) level. The argument 'all.levels' may be passed to newdata; if TRUE then the default newdata includes all factor levels.

realnames may be used to select a subset of parameters.

Standard errors for parameters on the response (real) scale are by the delta method (Lebreton et al. 1992), and confidence intervals are backtransformed from the link scale.

The value of newdata is optionally saved as an attribute.

Value

When se.fit = FALSE, a dataframe identical to newdata except for the addition of one column for each 'real' parameter. Otherwise, a list with one component for each row in newdata. Each component is a dataframe with one row for each 'real' parameter (density, g0, sigma, b) and columns as below

link	link function
estimate	estimate of real parameter
SE.estimate	standard error of the estimate
lcl	lower 100(1–alpha)% confidence limit
ucl	upper 100(1-alpha)% confidence limit

When newdata has only one row, the structure of the list is 'dissolved' and the return value is one data frame.

For detectpar, a list with the estimated values of detection parameters (e.g., g0 and sigma if detectfn = "halfnormal"). In the case of multi-session data the result is a list of lists (one list per session).

Note

predictDsurface should be used for predicting density at many points from a model with spatial variation. This deals automatically with scaling of x- and y-coordinates, and is much is faster than predict.ipsecr. The resulting Dsurface object has its own plot method.

The argument 'scaled' was removed from both predict methods in version 2.10 as the scaleg0 and scalesigma features had been superceded by other parameterisations.

References

Lebreton, J.-D., Burnham, K. P., Clobert, J. and Anderson, D. R. (1992) Modeling survival and testing biological hypotheses using marked animals: a unified approach with case studies. *Ecological Monographs* **62**, 67–118.

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print.ipsecr

See Also

ipsecr.fit, predictDsurface

Examples

predict (ipsecrdemo)

print.ipsecr Print or Summarise ipsecr Object

Description

Print results from fitting a spatially explicit capture-recapture model or generate a list of summary values.

Usage

```
## S3 method for class 'ipsecr'
print(x, newdata = NULL, alpha = 0.05, call = TRUE, ...)
## S3 method for class 'ipsecr'
summary(object, newdata = NULL, alpha = 0.05, ...)
```

Arguments

х	ipsecr object output from ipsecr.fit
object	ipsecr object output from ipsecr.fit
newdata	optional dataframe of values at which to evaluate model
alpha	alpha level
call	logical; if TRUE the call is printed
	other arguments (not used)

Details

Results from print.ipsecr are potentially complex and depend upon the analysis (see below). Optional newdata should be a dataframe with a column for each of the variables in the model. If newdata is missing then a dataframe is constructed automatically. Default newdata are for a naive animal on the first occasion; numeric covariates are set to zero and factor covariates to their base (first) level. Confidence intervals are 100 (1 - alpha) % intervals.

call	the function call (optional)
version,time	ipsecr version, date and time fitting started, and elapsed time
Detector type	'single', 'multi', 'proximity' etc.
Detector number	number of detectors
Average spacing	

x-range	
y-range	
New detector type	as fitted when details\$newdetector specified
N animals	number of distinct animals detected
N detections	number of detections
N occasions	number of sampling occasions
Mask area	
Model	model formula for each 'real' parameter
Fixed (real)	fixed real parameters
Detection fn	detection function type (halfnormal or hazard-rate)
Distribution	spatial model (details\$distribution)
N parameters	number of parameters estimated
Design points	number of vertices and centre points
Simulations per box	total number
Beta parameters	coef of the fitted model, SE and confidence intervals
vcov	variance-covariance matrix of beta parameters
Real parameters	fitted (real) parameters evaluated at base levels of covariates

Value

The summary method constructs a list of outputs similar to those printed by the print method, but somewhat more concise and re-usable:

versiontime	ipsecr version, and date and time fitting started
traps	detector summary
capthist	capthist summary
mask	mask summary
modeldetails	miscellaneous model characteristics
coef	table of fitted coefficients with CI
predicted	predicted values ('real' parameter estimates)

See Also

ipsecr.fit

Examples

load & print previously fitted null (constant parameter) model

print(ipsecrdemo)

summary(ipsecrdemo)

vcov.ipsecr

Description

Variance-covariance matrix of beta or real parameters from fitted ipsecr model.

Usage

```
## S3 method for class 'ipsecr'
vcov(object, realnames = NULL, newdata = NULL,
    byrow = FALSE, ...)
```

Arguments

object	ipsecr object output from the function ipsecr.fit
realnames	vector of character strings for names of 'real' parameters
newdata	dataframe of predictor values
byrow	logical for whether to compute covariances among 'real' parameters for each row of new data, or among rows for each real parameter
	other arguments (not used)

Details

By default, returns the matrix of variances and covariances among the estimated model coefficients (beta parameters).

If realnames and newdata are specified, the result is either a matrix of variances and covariances for each 'real' parameter among the points in predictor-space given by the rows of newdata or among real parameters for each row of newdata. Failure to specify newdata results in a list of variances only.

Value

A matrix containing the variances and covariances among beta parameters on the respective link scales, or a list of among-parameter variance-covariance matrices, one for each row of newdata, or a list of among-row variance-covariance matrices, one for each 'real' parameter.

See Also

vcov, ipsecr.fit, print.ipsecr

Examples

previously fitted ipsecr model
vcov(ipsecrdemo)

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