foieGras an R package for rapid quality control, behavioural estimation and simulation of animal track data

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8 Abstract

9 text...

10 1 | Introduction

The R package foieGras, pronounced "fwah grah", ...

12 **2** | foieGras **Overview**

The workflow for foieGras is deliberately simple, with much of the usual track data processing checks and formatting handled automatically. The main functions are listed in Table 1. When fitting a model, foieGras automatically detects the type of tracking data location quality classes designations that are typical of Argos data and that can be added to the data by the researcher for other types of track data. Based on the location quality classes and other, optional information on observation errors contained in the data, foieGras chooses an appropriate measurement error model for each observation. This capability allows for combinations of different tracking data types, e.g., Argos and GPS, in a single input data frame and to be fit in a single state-space model.

21 **2.1** | Data preparation

Animal tracking data, consisting of a time-series of location coordinates, can be read into R as a data frame using standard functions such as read.csv. The canonical data format for Argos tracks consists of a data frame with 5 columns corresponding to the following named variables:

id (individual id), date (date and time), 1c (location class), 1on (longitude), 1at (latitude). Optionally, an additional 3 columns, smaj (semi-major axis), smin (semi-minor axis), eor (ellipse orientation), providing Argos error ellipse information may be included.

Other types of track data can be accommodated, for example, by including the 1c column where all 1c = "G" for GPS data. In this case, measurement error in the GPS locations is assumed to have a standard deviation of 0.1 x Argos class 3 locations (approximately 30 m). Other types of track data can be considered in a similar manner (see the package vignette for further details).

Table 1: Main functions for the R package foieGras

Function	Description
fit_mpm	Fit a Move Persistence Model to location data
fit_ssm	Fit a State-Space Model to location data
fmap	Plot fitted/predicted locations on a map with or without a defined projection
grab	Extract fitted/predicted/observed locations from a foieGras model, with or without projection information
osar	Estimate One-Step-Ahead Residuals from a foieGras SSM
plot.fG_mpm	Plot move persistence estimates as 1-D or 2-D (along track) time-series
<pre>plot.fG_osar plot.fG_ssm</pre>	Plot One-Step-Ahead Residuals from a foieGras SSM Visualise the fit of a foieGras SSM to data

2.2 | State-space model fitting - fit_ssm

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State-space models are fit using fit_ssm. There are a large number of options that can be set in fit_ssm (see Suppl for details). We focus only the essential options here:

- data the input data structured as described in 2.1
- vmax a maximum threshold speed (ms⁻¹) to help identify potential outlier locations
- model the process model to be used
- time.step the prediction time interval (h)

The function first invokes an automated data processing stage where the following occurs: 1)
data type (Argos Least-Squares, Argos Kalman Filter/Smoother, GPS, or General (e.g., processed light-level geolocations, acoustic telemetry, coded VHF telemetry) is determined; 2) datetimes are converted to POSIXt format, chronological order is ensured, and duplicate datetime records are removed; 3) observations occurring less than min.dt seconds after a prior observation are removed; 4) a speed filter (sda from the trip R package; Sumner et al., 2009) is used to identify potential outlier locations; 5) locations are projected from spherical lon-lat coordinates to planar x,y coordinates in km.

The function then fits a state-space model to the processed data, where the process model (currently, either a continuous-time rw or a continuous-time crw) is specified by the user and the measurement model(s) are selected automatically (see Jonsen et al., 2020 for model details). The model is fit by numerical optimization of the likelihood using either the optim or nlminb R function. The R package TMB, Template Model Builder (Kristensen et al., 2016), is used to compute the gradient function in C++ via reverse-mode auto-differentiation and the Laplace Approximation is used to integrate out the latent states (random effects). Fits to a single versus multiple individuals are handled automatically, with sequential SSM fits occurring in the latter case. No hierarchical or pooled estimation among individuals is currently available.

fit_ssm returns a foieGras fit object (a nested data frame with class fG_ssm). The outer data frame lists the individual id(s), basic convergence information and a list with class ssm. This list contains dense information on the model parameter and state estimates, predictions, processed data, optimizer results, and other diagnostic and contextual information. Users can extract a simple data frame of SSM fitted (location estimates corresponding to the, typically irregular, observation times) or predicted values (locations predicted at regular time.step intervals) using the grab function.

2.3 | Model checking and visualisation

Before using fitted or predicted locations, a model fit should be checked and visualised to confirm
that the model adequately describes the data. In linear regression and a variety of analogous
methods, goodness-of-fit can be assessed by calculating standard residuals such as Pearson or
deviance residuals. There is no simple way to calculate residuals for latent variable models that have
non-finite state-spaces and that may be nonlinear, but they can be computed based on iterative
forecasts of the model (Thygesen et al., 2017). The osar function computes one-step-ahead
(prediction) residuals and uses the oneStepPredict function from the TMB R package to make this
as efficient as possible. A set of residuals are calculated for the x and y values corresponding to
the fitted values from the SSM.

A generic plot method plot.fg_osar provides an easy way to visualisation the prediction residuals.
Time-series plots of the prediction residuals can be used to detect temporal changes in goodnessof-fit. Quantile-quantile plots of residuals against standard normal quantiles can be used to detect
departures from normality. Sample autocorrelation function plots of the residuals are useful for
detecting autocorrelation not accounted for by the model. Assessing residual autocorrelation can
be particularly important as Argos locations, for example, are themselves derived from a time-series
model (Lopez et al., 2015) which can introduce additional autocorrelation in the location errors.

State-space model fits to data can also be visualised by using the generic plot.fG_ssm function on an fG_ssm data frame.

81 2.4 | Behavioural estimation

- 82 2.5 | Simulation
- 83 3 | Examples
- 84 3.x | Extending the behavioural model using mpmm
- 85 4 | Discussion

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94 Author's Contributions

⁹⁵ IDJ developed the R package; IDJ and TAP developed the state-space models and wrote the manuscript.

97 Data Accessibility

- 98 All code mentioned here is provided in the foieGras package for R available on CRAN at https:
- //CRAN.R-project.org/package=foieGras. The development version of the package is available on
- GitHub at https://github.com/ianjonsen/foieGras. Data used in the examples are available at...

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