Introduction

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Welcome!

- Welcome to this online course on animal movement.
- We are Brian and Johannes.
- Who are you? Pair up and introduce your partner

Outline of the course

Day 1:

- Introduction and exploratory data analysis for movement data
 (J)
- Data cleaning (B)

Day 2:

- Quantifying space use of animals with home ranges (B)
- Multiple instances (J)

Day 3:

- Introduction to habitat selection (B)
- Integrated step selection functions 1 (J)

Day 4:

- Integrated step selection functions 2 (B)
- Simulations from fitted iSSF (J)

Day 5:

- Strategies to model multiple animals (J)
- Validation of models for habitat selection (B)
- Time to discuss questions related to your projects.

Some logistics

- The course is scheduled from Monday (13th of March) to Friday (17th of March) from 2pm to 6pm Berlin time.
- We split these 4h block into two chunks, each roughly structured like this:
 - Lecture ~ 45 min
 - R walkthrough ~ 45 min
 - Introduction of exercises ~ 5 min
- A 20 min break between the two chunks.
- Lectures will be held via zoom.
- During the whole workshop we have a slack channel where you ask questions (we will monitor the channel during the course, feel free to ask questions there also outside the course hours).

Analysis of movement data in R

- The statistical software package R has become a widely used tool for data analysis in ecology and evolution and also in movement ecology.
- A typical analysis usually undergoes a few steps (all of which can be performed in R), this was reviewed by Joo et al. 2020.

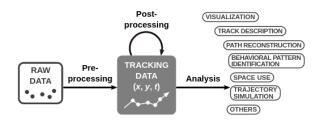


Figure 1: Figure from Joo et al 2020

Packages that we will use

We will mainly use the amt package, but also occasionally other packages for movement analysis.

See also required_packages.R for a list of all packages that we need and to get the latest version of all packages.

Some conventions in R

- %>% or |>: Pipes the output of one function to a next function.
 We will discuss this further later on.
- :: to access a name space form a package.
- use of 'a'.
- means this directory
- .. refers to the parent directory
- data.frame or tibble?

- We often use here::here("path to a file"), when reading in a file.
- The first here calls the function here() from the package here.
- The function here() dynamically creates the absolute path to the project root.

here::here()

[1] "/home/jsigner/git/movement_workshop"

This means, that we save all our data in the root directory data (even though my scripts are in different sub directories).

Brackets ((, [, {)

 round brackets or parentheses (() usually indicate functions or are used in arithmetic calculations.

```
sqrt(3)
[1] 1.732051
or
2 * (3 + 1)
```

Г17 8

• square brackets ([, [[) are used to subset data structures.

letters[1:3]

or

head(iris[["Species"]])

[1] setosa setosa setosa setosa setosa Levels: setosa versicolor virginica • curly brackets or braces ({) are used to form code blocks (e.g., inside a function or control structure).

```
for (i in 1:10) {
   i^2
}
```

Functions

- Functions do something. For example function sqrt() takes the square root for a number.
- It is easy to recognize functions, because they usually have a name (e.g., sqrt) followed by round brackets ().
- Within these round brackets arguments are passed to a function. This arguments can be named or unnamed (as long as they are in the correct order).

Recommended setup

- We would recommend to download the whole repository from GitHub (https://github.com/jmsigner/movement_ workshop_spring_2023)¹.
- Then use the RStudio project (together with RStudio).
- Following these guides, you should have all paths correct.

¹If you are familiar with git, feel free to clone the repository

Geographic data in brief

- Movement data is inherently spatial.
- Thus we will have to deal with tools to work with spatial data (R has a rich set of tools to deal with spatial data; e.g. https://geocompr.robinlovelace.net/).
- We will work frequently with raster data (spatial covariates) and possibly with vector data (i.e., home ranges).
- One of the challenges is to ensure that both tracking data and covariates – have a matching coordinate reference system (CRS).

- The CRS defines the reference system that is being used to explicitly reference a feature in space.
- There are two classes of CRS: geographic (e.g., WGS84) and projected (e.g., UTM) CRS.
- Projected CRS flatten the three dimensional data to the a two-dimensional plane (and introduce some distortion).

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- EPSG codes are four to five digits number that refer to different CRS. See for example www.epsg.io.
- Which CRS is best to use? It depends on the range of the study species. I usually prefer projected CRS, because their units are meters and not degrees.

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Take-home messages 1

- 1. Do not get confused by ::, this is just explicitly calls a function from a package.
- 2. Consider using RStudio projects to have everything in one place.

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- Consider using RStudio projects to have everything in one place.
- 3. Coordinate reference system (CRS) are used to describe where our animals are.
- 4. EPSG-Codes can be used to refer to CRS.

Data

Movement data

- Often times the data we receive is just a time series of coordinates (longitude, latitude and time stamp).
- Depending on the sensors we use, other (meta) information may also be stored (this could include temperature, coordinates in a different [projected] CRS, ...).

```
1 No; CollarD; UTC Date; UTC Time; LMT Date; LMT Time; Origin; SCTS, Date; SCTS, Time; ECEF X [R]; ECE
```

Environmental covariates

- Vector layers (e.g., road networks, rivers, protected areas)
- Raster layers (e.g., land use, remotely sensed data such as NDVI, climatic variables)

Tracks and bursts: the basic building block

- For movement data, we usually read a text file into R (.csv, .txt, ...) as a data frame and then create a analysis-specific object.
- When working with the amt package the function make_track() takes a sequence of coordinates (with or without timestamps) and creates a track. Note, at this point multiple individuals can be mixed and sampling rates can be heterogeneous.

- Bursts can be created from tracks.
- A burst is a sequence of (re)locations from the same individual at equal time intervals (with some tolerance).
- Options to change from tracks to bursts are:
 - Use the function amt::track_resample().

Sampling rates, and resampling

- The function summarize_sampling_rate() takes an track as input and gives a summary of the sampling rate.
- If there are multiple animals present, there is also the function summarize_sampling_rate_many(), which will do the same thing, but for many animals.

- Once a suitable sampling rate is determined, the function track_resample() can be used to take a relocation every predefined time interval (e.g., 30 minutes, 2 hours, ...) within a tolerance.
- The result of track_resample() is again a track with one additional column called burst_.

Take-home messages

- Movement data are often 'just' text files.
- The amt package uses tracks as the basic building block.
- A burst is a track (or a part of a track) with a regular sampling rate.
- Use track_resample() create tracks with equal sampling rates.

If you have gaps and/or different sampling rates, interpolation with continuouse time movement models may be an option.



APPLICATION 🙃 Open Access 🕼 🚯

aniMotum, an R package for animal movement data: Rapid quality control, behavioural estimation and simulation

lan D. Jonsen 🔀, W. James Grecian, Lachlan Phillips, Gemma Carroll, Clive McMahon, Robert G. Harcourt, Mark A. Hindell, Toby A. Patterson

First published: 26 January 2023 | https://doi.org/10.1111/2041-210X.14060

Handling Editor Edward Codling

Movement characteristics (sl_, ta_)

Tracks are still *just* the points as they were collected. If we want to get insights, we have can look at different characteristics of steps (i.e., two consecutive relocations).

This include:

- step length
- turn angle
- speed
- Net squared displacement

Note, unless you take care of different instances or bursts, they are ignored.

Net Squared displacement (NSD)

- The NSD is the squared distance between the first relocation of a track and the every relocation that follows.
- Bunnefeld et al. 2011 described different forms of the NSD that resemble different migratory behaviors.
- The different models can fit to the data (e.g., using nonlinear least square with the function nls() in R).

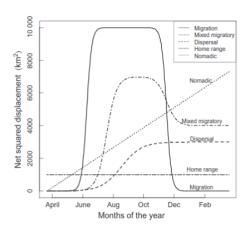


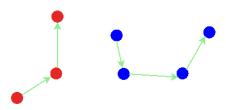
Figure 2: Figure taken from Bunnefeld et al. 2011

Time of the day

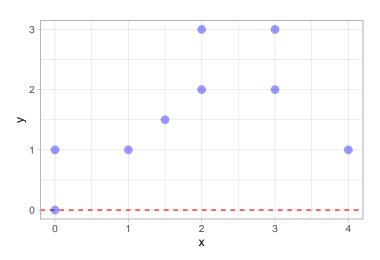
- Time of day can be annotated to steps with the function amt::time_of_day(). This will add an additional column to the data frame of steps tod_end or tod_start depending on the argument when.
- If the data is of sufficient temporal resolution, it is also possible to annotate twilight (dawn and dusk).

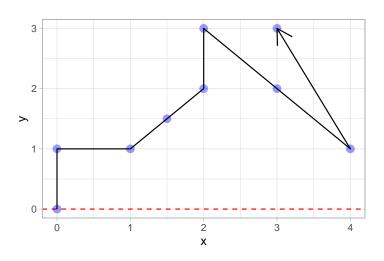
Steps

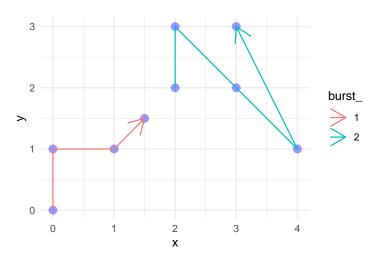
We can start to create a steps-representation.



This can be achieved with the function amt::steps(). If we resampled the data previously, we can even use amt::steps_by_burst().







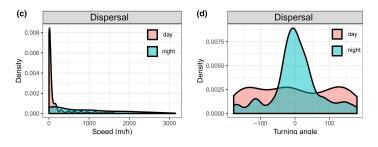
This automatically calculates several step attributes:

- Start and end point
- Step length
- Absolute and relative turn angles
- Duration

This allows already calculate some step characteristics. It becomes even more informative, if we pair this for example with the whether a step was in the night, day or twilight.

Example

Remington Moll observed a (rare) long distance dispersal for White Tail deer³ and looked at the turn angle and step distribution for day and night.



³Moll et. al Ecology and Evolution; https://onlinelibrary.wiley.com/doi/full/10.1002/ece3.7354.

Take-home message

- 1. Movement characteristics (e.g., step length or turn angle) are fundamental to many analyses in discrete time.
- 2. We often compare movement characteristics for different times of day.
- 3. The Net Squared Displacement (NSD) can be used to infer different migratory modes.

Data sets that we use

- We will use several data sets during this course including a data set on fishers, elephants and deer.
- Feel free to use your own data during the exercises, we are happy to help to get it into shape.
- For the R walkthrough we often simulate data. We believe if understand how data is generated it is much easier to understand how a specific method works.

Key resources/publications

Movement ecology

- Bunnefeld, N., Börger, L., van Moorter, B., Rolandsen, C. M., Dettki, H., Solberg, E. J., & Ericsson, G. (2011). A model-driven approach to quantify migration patterns: individual, regional and yearly differences. Journal of Animal Ecology, 80(2), 466-476.
- Bjorneraas et al. 2010: Screening Global Positioning System Location Data for Errors Using Animal Movement Characteristics. https://doi.org/10.1111/j.1937-2817.2010.tb01258.x
- Joo, R., Boone, M. E., Clay, T. A., Patrick, S. C., Clusella-Trullas, S., & Basille, M. (2020). Navigating through the R packages for movement. Journal of Animal Ecology, 89(1), 248-267.

R resource

- Lovlace, R. et al. (2019). Geocomputation with R: https://geocompr.robinlovelace.net/.
- Wickham, H. R for data science: https://r4ds.had.co.nz/