

# wildlifeTG

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**Type** Package

**Title** Time Geograhic Analysis of Wildlife Telemetry Data

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**Description** The package wildlifeTG provides tools for performing time geographic analysis of animal tracking data. The functions provide useful tools for examining wildlife movement from the context of accessibility. This is an alternative view on typical home range estimation procedures. Currently, the package provides functions that facilitate the calculation of the PPA measure of an individuals accessibility space and the jPPA measure of joint accessibility space between two individuals. Please note that the package is still under development.

**License** GPL-3

**Imports** sp, adehabitatLT, rgeos

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wildlifeTG-package

wildlifeTG - Time Geographic Analysis of Wildlife Telemetry Data

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## Description

The package wildlifeTG provides tools for performing time geographic analysis of animal tracking data. The functions provide useful tools for examining wildlife movement from the context of accessibility. This is an alternative view on typical home range estimation procedures. Currently, the package provides functions that facilitate the calculation of the PPA measure of an individual's accessibility space and the jPPA measure of joint accessibility space between two individuals. Please note that the package is still under development.

## Details

wildlifeTG's functions utilize the lttraj objects from the package adehabitat.

## Author(s)

Jed Long

## References

Long, JA, Nelson, TA. (2012) Time geography and wildlife home range delineation. *Journal of Wildlife Management*, 76(2):407-413.

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dyn.ppa

Dynamic PPA Measure of Animal Space Use

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## Description

The function dyn.ppa computes the (dynamic) PPA measure of the accessibility space of an animal. The PPA method can be thought of as an alternative view on the home range; one that explicitly considers the spatial and temporal constraints on movement given known telemetry fixes, and a (dynamic) measure of maximum mobility - termed vmax. The PPA method incorporates dynamic behaviour into the calculation of the vmax parameter used to delineate the original version of the PPA method, but the original method is still an option here.

## Usage

```
dyn.ppa(traj, tol = max(ld(traj)$dt, na.rm = TRUE), dissolve = TRUE,
        ePoints = 360, proj4string = NA, ...)
```

**Arguments**

traj	an object of the class <code>ltraj</code> which contains the time-stamped movement fixes of the first object. Note this object must be a type <code>II ltraj</code> object. For more information on objects of this type see <code>help(ltraj)</code> .
tol	parameter used to filter out those segments where the time between fixes is overly large (often due to irregular sampling or missing fixes); which leads to an overestimation of the activity space via the PPA method. Default is the maximum sampling interval from <code>traj</code> .
dissolve	(logical) whether or not to dissolve output ellipse polygons to create a single output polygon, or keep the individual segment PPA ellipses. Default = <code>TRUE</code> .
ePoints	number of vertices used to construct each PPA ellipse. More points will necessarily provide a more detailed ellipse shape, but will slow computation; default is 360.
proj4string	a string object containing the projection information to be passed included in the output <code>SpatialPolygonsDataFrame</code> object. For more information see the <code>CRS</code> -class in the packages <code>sp</code> and <code>rgdal</code> . Default is <code>NA</code> .
...	additional parameters to be passed to the function <code>dyn.vmax</code> . For example, should include <code>method</code> and/or <code>window</code> parameters, see the documentation for <code>dyn.vmax</code> for more detailed information on what to include here.

**Details**

The function `dyn.ppa` represents an extension to an existing PPA method (Long and Nelson, 2012). Dynamic calculation of the PPA method improves upon the original version by flexibly modelling the `vmax` parameter according to wildlife behaviour. See the function `dyn.vmax` for more information on how to incorporate dynamic behaviour into the `vmax` parameter estimation.

**Value**

This function returns a `SpatialPolygonsDataFrame` representing the dynamic PPA measure of the accessibility space of an individual animal.

**References**

Long, JA, Nelson, TA. (2012) Time geography and wildlife home range delineation. *Journal of Wildlife Management*, 76(2):407-413.

**See Also**

`dyn.vmax`, `di.ppa`

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`dyn.vmax`

*Dynamic Calculation of the Vmax Parameter*

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**Description**

The function `dyn.vmax` computes a dynamic version of the `vmax` parameter for the PPA method. It can be used to incorporate changes in animal movement behaviour into the PPA method calculation to better model that area accessible to an individual animal given the set of known telemetry locations in space and time.

## Usage

```
dyn.vmax(traj, window = "focal", w = 9, class.col = "dt",
  method = "Robson", k = 5, alpha = 0.05, manualVmax = NA,
  vmaxtrunc = NA)
```

## Arguments

traj	an object of the class <code>ltraj</code> which contains the time-stamped movement fixes of the first object. Note this object must be a type <code>II ltraj</code> object. For more information on objects of this type see <code>help(ltraj)</code> .
window	one of 'original', 'focal', 'cumulative', or 'class'; which signifies whether or how to dynamically compute the Vmax parameter. See <b>Details</b> for more information on each of the choices.
method	method for computing the Vmax parameter dynamically; can be one of several options: – "Robson" for the Robson & Whitlock (1964) method, – "RobsonLL" for the R & W (1964) lower $(1 - \alpha) * 100\%$ C.I. limit, – "RobsonUL" for the R & W (1964) upper $(1 - \alpha) * 100\%$ C.I. limit, – "vanderWatt" for the van der Watt (1980) method, – "vanderWattLL" for the van der Watt (1980) lower $(1 - \alpha) * 100\%$ C.I. limit, – "vanderWattUL" for the van der Watt (1980) upper $(1 - \alpha) * 100\%$ C.I. limit.
w	(optional) window size (only used with window = 'focal' or 'cumulative').
class.col	(optional) character indicating the name of the column in the <code>infolocs</code> dataframe of <code>traj</code> containing the categorized behavioural states of the animal (which can be stored as a character or numeric column).
k	(optional) value for the $k$ parameter in the van der watt (1980) method; default is 5.
alpha	(optional) value for the $\alpha$ parameter if using upper or lower C.I. methods; default is 0.05.
manualVmax	(optional) Character name of column in <code>traj</code> storing user input column of vmax values (typically call the column <code>dynVmax</code> ).
vmaxtrunc	(optional) due to irregular sampling intervals, or errors in GPS location, or other effects, the calculation of the vmax parameter through the statistical methods outlined above can be heavily influenced by high outliers. Thus, it may be useful to exclude those segments from calculation of the dynamic Vmax parameter. Default is NA.

## Details

The function `dyn.vmax` represents an intermediary function used to extend and improve upon an existing PPA home range method (Long and Nelson, 2012). Four options are available for computing the vmax parameter dynamically - passed to the `window` option. 1) `original` – if `window = 'original'` the function estimates the original, non-dynamic estimate of vmax, as per Long & Nelson (2012). 1) `focal` – a moving window approach whereby a window of size `w` is moved along the trajectory and vmax computed dynamically within each window and assigned to the central segment.

2) `cumulative` – A moving window of size `w` is again used, only in this case the value is assigned to the end segment. This represents the vmax calculation of the previous `w` segments.

3) `class` – A priori analysis (e.g., obtained via state-space models, or from expert knowledge) is used to identify discrete behavioural states in the telemetry data and these stored in a column which

is then passed into the function.

The class method is the preferred choice, as it allows the use of more sophisticated models for identifying behavioural shifts in telemetry data where we would expect to see clear differences in the vmax parameter based on changing movement behaviour (see Patterson et al. 2008 for a review).

The use of the 'focal' or 'cumulative' methods uses a moving window approach, which is sensitive to edge effects at the initial and ending times of the trajectory. Thus, the dynamic Vmax parameter is only computed for those segments that have a valid window and the dataset is shrunk by  $w-1$  segments.

### Value

This function returns the original traj object with a new column – dynVmax in the infolocs dataframe containing the dynamic vmax parameter for each trajectory segment.

### References

- Long, J.A., Nelson, T.A. 2012. Time geography and wildlife home range delineation. *Journal of Wildlife Management*. 76(2):407-413.
- Robson, D.S., Whitlock, J.H. 1964. Estimation of a truncation point. *Biometrika* 51:33-39.
- van der Watt, P. 1980. A note on estimation bounds of random variables. *Biometrika* 67(3):712-714.

### See Also

dyn.ppa, j.ppa

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j.ppa

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*Joint Potential Path Area (Accessibility Space) of Two Animals*


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### Description

The function j.ppa computes the joint accessibility space between two animals. It can be used to map (as a spatial polygon) the area that could have been jointly accessed by two individual animals in space and time.

### Usage

```
j.ppa(traj1, traj2, t.int = 0.1 *
      as.numeric(names(sort(-table(ld(traj1)$dt)))[1]), tol = max(ld(traj1)$dt,
      na.rm = T), dissolve = TRUE, ePoints = 360, ...)
```

### Arguments

- |       |   |
|-------|---|
| traj1 | an object of the class ltraj which contains the time-stamped movement fixes of the first object. Note this object must be a type II ltraj object. For more information on objects of this type see help(ltraj). |
| traj2 | same as traj1.  |

<code>t.int</code>	(optional) time parameter (in seconds) used to determine the frequency of time slices used to delineate the joint activity space. Default is 1/10th of the mode of the temporal sampling interval from <code>traj1</code> . Smaller values for <code>t.int</code> will result in smoother output polygons.
<code>tol</code>	(optional) parameter used to filter out those segments where the time between fixes is overly large (often due to irregular sampling or missing fixes); which leads to an overestimation of the activity space via the PPA method. Default is the maximum sampling interval from <code>traj1</code> .
<code>dissolve</code>	logical parameter indicating whether (=TRUE; the default) or not (=FALSE) to return a spatially dissolved polygon of the joint activity space.
<code>ePoints</code>	number of vertices used to construct each PPA ellipse. More points will necessarily provide a more detailed ellipse shape, but will slow computation; default is 360.
<code>...</code>	additional parameters to be passed to the function <code>dyn.vmax</code> . For example, should include <code>window</code> and <code>method</code> ; see the documentation for <code>dyn.vmax</code> for more detailed information on what to include here.

## Details

The function `j.ppa` can be used to map areas of potential interaction between two animals. Specifically, this represents a measure of spatial overlap that also considers the temporal sequencing of telemetry points. In this respect it improves significantly over static measures of home range overlap, often used to measure static interaction, and can be considered as a spatial measure of dynamic interaction.

## Value

This function returns a `SpatialPolygonsDataFrame` representing the joint accessibility space between the two animals.

## See Also

`dyn.vmax`, `dyn.ppa`

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`ppaEllipse`

*PPA Ellipse*

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## Description

Internal ellipse calculation function.

## Usage

```
ppaEllipse(x, y, a, b, theta, steps)
```

**Arguments**

x	first coordinate
y	second coordinate
a	semi-major axis
b	semi-minor axis
theta	rotation angle of the ellipse (in radians)
steps	number of segments, from ePoints parameter in <code>dyn.ppa.hr</code>

**Details**

Internal function for calculating ellipses in time geographic analysis.

**Value**

This function returns a polygon ellipse.

**See Also**

`dyn.ppa.hr`

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