

# Package ‘rpostgisLT’

July 14, 2016

**Title** Managing and visualizing animal movement data with PostGIS and R

**Version** 0.0.1

**Description** The project develops the integration of R and PostGIS for managing movement trajectories. The focus is on streamlining the workflow for biologists to store and process animal trajectories in PostGIS and analyze these in R, thus utilizing the strengths of both software. Therefore the main outcome is a new R package rpostgisLT, which will simplify the processing of location datasets into trajectories in PostGIS and provide full integration with the R package adehabitatlt data type lttraj.

**Depends** R (>= 3.3.0)

**License** GPL (>= 3)

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 5.0.1

**Author** Balázs Dukai [aut, cre],  
Mathieu Basille [aut],  
David Bucklin [aut],  
Clément Calenge [ctb]

**Imports** RPostgreSQL, testthat, rpostgis

**Suggests** knitr,  
rmarkdown

**VignetteBuilder** knitr

## R topics documented:

as_pgtraj . . . . .	2
ltraj2pgtraj . . . . .	3
pgtraj2ltraj . . . . .	4

<b>Index</b>	<b>6</b>
--------------	----------

as\_pgtraj

*Converts relocation data into the pgtraj data model.***Description**

This is the core function of the `rpostgisLT` package and it is also used by `ltraj2pgtraj` to import trajectory data into a pgtraj data model. `as_pgtraj` copies the trajectory data which is stored in a database to a traj schema. If the provided schema doesn't exist, it is created on demand. On successful data input, `as_pgtraj` creates a view for each pgtraj, with the views named as `<pgtraj>_params`. The view contains the same step parameters as an `ltraj` object (e.g. `R2n`, `rel.angle`, `dt...`). If the input geometries are projected, their projection is used to create the steps in the schema, otherwise either no projection is used or the function exits.

**Usage**

```
as_pgtraj(conn, schema = "traj", relocation_data = NULL,
          pgtrajs = "pgtraj", animals = "animal", bursts = NULL,
          relocations = NULL, timestamps = NULL, rids = "rid", db = TRUE)
```

**Arguments**

<code>conn</code>	Connection object created with <code>RPostgreSQL</code>
<code>schema</code>	String. Name of the schema that stores or will store the pgtraj data model.
<code>relocation_data</code>	String. Name of the table that stores the relocations, e.g. "public.relocations"
<code>pgtrajs</code>	String. Name of the pgtraj or name of the field that stores the pgtraj names.
<code>animals</code>	String. Name of the animal or name of the field that stores the animal names.
<code>bursts</code>	String. Name of the burst or name of the field that stores the burst names.
<code>relocations</code>	String. Name of the field that contains the relocations in <code>relocation_data</code> .
<code>timestamps</code>	String. Name of the field in <code>relocation_data</code> that contains the timestamps. If <code>NULL</code> , Type I trajectory is assumed.
<code>rids</code>	String. Name of the field in <code>relocation_data</code> that contains the numeric IDs of relocations.
<code>db</code>	Boolean. If <code>TRUE</code> , the relocations are stored in a database table, if <code>FALSE</code> relocations are stored in an R object. It is meant to be used by other functions internally. If you want to import an <code>ltraj</code> from R, use <code>ltraj2pgtraj()</code> .

**Details**

Opening and closing connections have to be done manually by the user. However, the function checks if the provided connection is still valid. Not tested with capital letters for PostgreSQL field names, but it probably won't work. It's a bad practice anyway to force uppercase in PostgreSQL so use lowercase.

**Value**

TRUE on success

**Author(s)**

Balázs Dukai <balazs.dukai@gmail.com>

**References**

<https://cran.r-project.org/web/packages/adehabitatLT/vignettes/adehabitatLT.pdf>

**See Also**

Section on pgtraj data model in the package vignette.

**Examples**

```
## Not run:
as_pgtraj(conn,
  schema = "traj_t4",
  relocation_data = "example_data.relocations_plus",
  pgtrajs = "id",
  animals = "animal",
  bursts = "burst",
  relocations = "geom",
  timestamp = "time",
  rid = "gid")

## End(Not run)
```

---

ltraj2pgtraj

*Export an ltraj object from R into a traj database schema.*

---

**Description**

ltraj2pgtraj creates a new traj schema or uses an existing one and exports an ltraj to the database. Uses as\_pgtraj to insert the values into the traj schema.

**Usage**

```
ltraj2pgtraj(conn, ltraj, schema = "traj", pgtraj = NULL, epsg = NULL,
  comment = NULL)
```

**Arguments**

conn	Connection object created with RPostgreSQL
ltraj	An ltraj object.
schema	String. Name of the schema that stores or will store the pgtraj data model.
pgtraj	String. Name of the new pgtraj. Defaults to the name of the variable that stores the ltraj.
epsg	Numeric. The EPSG code of the Coordinate Reference System of the relocation coordinates in the ltraj. Defaults to 0.
comment	String. A comment that will be stored with the pgtraj in the database.

**Value**

TRUE on success

**Author(s)**

Balázs Dukai <balazs.dukai@gmail.com>

**See Also**

[as\\_pgtraj](#)

**Examples**

```
## Not run: ltraj2pgtraj(conn, ibex, "traj_t2")
```

---

pgtraj2ltraj	<i>Import a pgtraj into an ltraj.</i>
--------------	---------------------------------------

---

**Description**

pgtraj2ltraj imports a single pgtraj from a database into an ltraj object.

**Usage**

```
pgtraj2ltraj(conn, schema = "traj", pgtraj)
```

**Arguments**

conn	Connection object created with RPostgreSQL
schema	String. Name of the schema that stores or will store the pgtraj data model.
pgtraj	String. Name of the pgtraj.

**Value**

an ltraj object

**Author(s)**

Balázs Dukai <balazs.dukai@gmail.com>

**Examples**

```
## Not run: pgtraj2ltraj(conn, "traj_t2", "ibex")
```

# Index

as\_pgtraj, [2](#), [4](#)

ltraj2pgtraj, [2](#), [3](#)

pgtraj2ltraj, [4](#)