

# Package ‘tlocoh’

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**Type** Package

**Title** Constructs homeranges and explores time use patterns from location data

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**Description** Constructs homeranges and explores time use patterns from location data.

**License** GPL-2

**URL** <http://tlocoh.r-forge.r-project.org>

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aoi

*Select an area of interest on a plot using the mouse*

## Description

Allows the user to select an area of interest (box) by clicking on the active plot window

## Usage

```
aoi(draw.poly = TRUE, draw.poly.col = "black", status = TRUE)
```

## Arguments

draw.poly	Whether to draw the box on the plot. T/F.
draw.poly.col	The color to use if drawing the box on the plot. Color value. Ignored if draw.poly=FALSE.
status	Show messages to the user

## Value

A two-column xy data frame containing the upper left point and the lower right point in the first and second rows respectively

## Note

This function requires a plot window to be open

---

aoi2box	<i>Converts a dataframe with upper-left / lower-right coordinates of a box to a closed polygon</i>
---------	--

---

### Description

Converts a dataframe with upper-left / lower-right coordinates of a box to a closed polygon

### Usage

```
aoi2box(aoi)
```

### Arguments

aoi	A two-column two-row data frame or matrix with x in the first column and y in the second column. The upper left point and the lower right point in the first and second rows respectively
-----	---

### Value

Two column dataframe containing the xy coordinates of a closed rectangle that can be plotted using [points](#) or [polygons](#)

### Note

You can use the aoi function to select two coordinates on the active plot window with the mouse. This function facilitates plotting that rectangle.

### See Also

[aoi](#)

### Examples

```
## Not run:
myaoi <- aoi()
polygons(myaoi)

## End(Not run)
```

---

auto.a	<i>Parameters to select a reasonable value of 'a'</i>
--------	---

---

### Description

Returns a list of parameters that other functions accept for the 'a' parameter

### Usage

```
auto.a(meth = "nn", ptp = 0.98, nnn = 2, tct = 1.05)
```

**Arguments**

meth	The name of the method for finding a (default is "nn" method for 'nearest neighbor' method).
ptp	The proportion of total points [0..1]
nnn	The number of nearest neighbors
tct	The temporal continuity threshold, see details.

**Details**

If multiple values are passed for any of the parameters, the resulting data frame will have multiple rows and other parameters will be recycled as needed.

This function does not actually find nearest neighbors. Rather it returns a list of parameters that can be used as input in other functions such as [lxy.nn.add](#).

**Value**

A data frame with columns meth, ptp, nnn, and tct which are the settings that other functions will use to find a value for 'a' should be automatically selected. Columns include:

- *meth* The name of the method for finding a
- *ptp* The proportion of parent points that should have at least a.nn neighbors (default 0.98)
- *nnn* The minimum number of nearest neighbors that each point will get (default 2, so each point can be part of a triangle)
- *tct* A time threshold (expressed as a percentage of the median sampling frequency), such that if a point is temporally disconnected from its 2 nearest temporal neighbors by more than than tct times the median sampling frequency, the point will be excluded from the computation of a. This is to avoid bias from internal time gaps (default 1.05)

**See Also**

[lxy.nn.add](#)

**Examples**

```
## Not run:
lxy <- lxy.nn.add(lxy, ptsh=0.5, a=auto.a())

## End(Not run)
```

---

chop2plot

---

*Divides a long string into separate lines for adding to a plot*


---

**Description**

Breaks a long character object into separate lines at specific characters

**Usage**

```
chop2plot(str, width, char.break = " ", char.break.keep = FALSE, cex = 1)
```

**Arguments**

<code>str</code>	The string to format
<code>width</code>	The maximum width of any single line in inches
<code>char.break</code>	The character where the string should be broken
<code>char.break.keep</code>	Whether to keep the <code>char.break</code> character when the string gets broken. T/F
<code>cex</code>	The character expansion factor that will be used to display the text

**Value**

A two-item character vector containing [1] the formatted character string with `\n` inserted and [2] number of lines

**Note**

This function is designed to break up a potentially long character object that is going to be added to a plot either as a title or text. Line breaks will be inserted so that the string will wrap appropriately at `width` inches when plotted with `cex` character expansion factor.

The character object will be broken at the `char.break` characters such that each line will be  $\leq$  `width` when plotted. Breaks are replaced with carriage returns. If `char.break.keep` is `TRUE`, the break character will be kept, otherwise it will be removed. If no occurrence of the `/codechar.break` is available, the line will be chopped at the last character that fits.

---

cube

---

*Cubes a number*

---

**Description**

Cubes a number

**Usage**

`cube(x)`

**Arguments**

<code>x</code>	a number
----------------	----------

**Value**

$x^3$

---

 cw

*Console Wrap. Formats a string for printing to the console window*


---

**Description**

Takes a potentially long character object and adds carriage returns and blank spaces for better appearance when printing to the terminal window

**Usage**

```
cw(x, final.cr = TRUE, indent = 0, exdent = 0)
```

**Arguments**

x	A character object
final.cr	Whether to add a carriage return at the end. T/F.
indent	First line indent (number of blank spaces)
exdent	Subsequent line(s) indent (number of blank spaces)

**Value**

character object with embedded \n characters

**Note**

This function can prevent long messages from extending beyond the width of the R-console. Works with any function that prints to the console (e.g., cat, print, stop)

---

 findonpath

*findonpath*


---

**Description**

Searches for a file on the system environment path

**Usage**

```
findonpath(fn, status = TRUE)
```

**Arguments**

fn	The filename to search for (without a path)
status	Show messages

**Value**

The full path and name of the found file, or NULL if not found

**Note**

This will return the first found occurrence of file fn, searching 1) the current working directory, 2) the user's R 'home' directory, then 3) the directories on the operating system environment 'path'

---

get.vals	<i>Get x and/or y values from a plot using the mouse</i>
----------	--

---

### Description

Get x and/or y values from a plot using the mouse

### Usage

```
get.vals(axis = c("x", "y")[1], round.to = 1, print = TRUE)
```

### Arguments

axis	Which axes to return values for, character vector
round.to	An integer number that values will be rounded to. For example if round.to=100, values will be rounded to the nearest 100.
print	Whether to print the values in the console, T/F

### Details

This function allows you to get the x and/or y values by clicking on the active plot window. When you're done clicking, right-click and select 'Stop' from the pop-up menu.

### Value

A vector or matrix of the x and/or y values where the mouse was clicked.

---

hist.locoh.lhs	<i>Plot histograms of hull metrics</i>
----------------	--

---

### Description

Plot histograms of hull metrics

### Usage

```
## S3 method for class 'locoh.lhs'
hist(lhs, id = NULL, k = NULL, r = NULL, a = NULL,
     s = NULL, hs.names = NULL, metric = "area",
     include.missing.hulls = TRUE, hmap = NULL, hmap.in.title = TRUE,
     hs.name.in.title = TRUE, title = NULL, title.show = TRUE,
     figs.per.page = NULL, mar = c(3, 3, if (title.show) 2.8 else 0.7, 0.5),
     mgp = c(1.8, 0.5, 0), same.axes.for.all = FALSE, ufat = TRUE,
     breaks = "Sturges", col = "gray80", png.dir = NULL,
     png.dir.make = TRUE, png.width = 800, png.height = png.width,
     png.overwrite = TRUE, png.pointsize = 12 + (png.width - 480)/80,
     title.two.id = FALSE, indicate.missing.hulls.in.axis.lbl = TRUE,
     panel.num = NULL, panel.num.inside.plot = !title.show, ...)
```



**Arguments**

lhs	A <a href="#">LoCoH-hullset</a> object
id	The names of the individual(s) to include in the plot.
k	The k value(s) of the hullset(s) to include in the plot. Numeric vector or comma-delimited character object.
r	The r value(s) of the hullset(s) to include in the plot. Numeric vector or comma-delimited character object.
a	The a value(s) of the hullset(s) to include in the plot. Numeric vector or comma-delimited character object.
s	The s value(s) of the hullset(s) to include in the plot. Numeric vector or comma-delimited character object.
hs.names	The name(s) of saved hullsets to include in the plot
metric	The name(s) of hull metric(s); one histogram will be created for each hull metric.
hmap	A named list of hull metric auxillary parameters, the name of each list element is the name of the variable.
hmap.in.title	Include the hmap value(s) in the plot subtitle. T/F
hs.name.in.title	Include the name of the hullset in the plot subtitle. T/F
title	A title for the plot (over-writes the construction of a title)
title.show	Whether to add a title on the plot. T/F
include.missing.hulls	Whether to include missing hulls on the histogram (using the default value assigned to missing hulls by the hull metrics). T/F
figs.per.page	Number of plots per page
mar	The plot margins. A four item numeric vector
mgp	The distance away from the edge of the plot for the 1) label, 2) tick marks, and 3) axis line. A three-item numeric vector
same.axes.for.all	Whether to use the same range on the axes for all histograms generated (helps to visually see differences)
ufat	User-friendly axis title. T/F
breaks	Number of breaks or a clustering function. See <a href="#">hist</a> .
col	Color value for the bars
png.dir	The directory for a PNG file (filename will be constructed automatically).
png.dir.make	Whether to create png.dir if it doesn't exist. T/F
png.width	The width of the PNG image. Ignored if png.fn is passed.
png.height	The height of the PNG image. Ignored if png.fn is passed.
png.overwrite	Whether to overwrite an existing PNG file if it exists. T/F
title.two.id	Construct a title with the names of two ids (for so.* and to.* association metrics). T/F
indicate.missing.hulls.in.axis.lbl	Specify whether missing hulls are excluded in the axes label, T/F
panel.num	A number or letter to display in the upper left hand corner of the plot when the plot will be used as part of a multi-frame graphic (as in publications). Character

panel.num.inside.plot	Whether to display panel.num inside the plot area itself, as opposed to the title area. Ignored if panel.num is NULL. T/F
png.pointsize	The pointsize (in pixels) for the PNG image (increase to make labels appear larger). Determines the height or width of a character in pixels.
...	Other parameters, including any auxillary parameters to specify particular hull metrics

**Value**

A named list of values of the hull metrics. There is one list element for each metric.

**Note**

To see the names of hull metrics, type `hm.expr()`. Only hull metrics which have been computed can be plotted.

---

hist.locoh.lxy	<i>Plot histograms of properties of a LoCoH-xy object</i>
----------------	---

---

**Description**

Displays histogram(s) of point-to-point step length, velocity, and sampling frequency for a LoCoH-xy object

**Usage**

```
## S3 method for class 'locoh.lxy'
hist(lxy, id = NULL, dt = TRUE, d = TRUE,
     delta.t = TRUE, v = TRUE, figs.per.page = NULL, col = "gray80",
     dt.bins.base = c("secs", "mins", "hours", "days")[4], dt.bins.width = 3600
     * 24 * 7, delta.t.num.sd = NULL, d.tct = 1.2, time.unit = "auto",
     overlay.median = TRUE, breaks = 20)
```

**Arguments**

lxy	A <a href="#">LoCoH-xy</a> object
id	The id value(s) to be plotted
dt	Include a histogram of the number of locations over time) (T/F)
d	Include a histogram of distance travelled per adjacent points (i.e., step length) (T/F)
delta.t	Include a histogram of the time between points (i.e., sampling frequency) (T/F)
v	Include a histogram of point velocity (distance over delta.t) (T/F)
dt.bins.base	Where to start the first bin ((for histogram of number of points over time): "secs", "mins", "hours", "days")
dt.bins.width	The width of the time bins (for histogram of number of points over time). In seconds.
figs.per.page	Number of plots per page

col	The color of the bars
d.tct	temporal connectivity threshold for the distance between two points to be included in the histogram of step length. In other words, delta.t must be $\leq$ median delta.t * d.tct). Ignored if d=F.
delta.t.num.sd	Number of standard deviations for delta.t to be included in the histogram. To omit outliers from appearing in the histogram (which can make the central data more difficult to discern, set delta.t.num.sd to ~2. Ignored if delta.t=F.
time.unit	The unit of time on the x-axis (character). Ignored if delta.t=F.
overlay.median	Plot the median value on the histogram (T/F)
breaks	Argument passed to the <a href="#">hist</a> function, see <a href="#">hist</a>

### Value

A list of frequencies with one element for each of the histograms plotted.

---

hm.expr	<i>Hull metrics expressions</i>
---------	---------------------------------

---

### Description

Returns a list of expression objects that when evaluated in functions will return various hull metrics

### Usage

```
hm.expr(names.only = TRUE, desc = names.only, print = names.only)
```

### Arguments

names.only	Return only the names of hull metrics (not the expressions themselves). T/F.
desc	Include descriptions of the hull metrics in the object returned. T/F.
print	Display the list of hull metrics on the console. T/F.

### Details

Hull metrics are used by many functions. This function returns a list of the expressions to 'pull out' hull metric values from the data structure of a [LoCoH-hullset](#) object and other properties of hull metrics. These objects are used by other T-LoCoH functions for plotting, exporting, creating isopleths, and other things you can do with hull metrics.

Most users will not need to use this function other than to see a list of hull metric names (which you need to know to specify a hull metric in a plotting function for example) and the hull metric descriptions.

This function does not compute any hull metrics, only provide a comprehensive list of all possible metrics. Some hull metrics are computed by default when you create a hullset (see [lxy.lhs](#)). Others metrics must be computed with a special function (e.g., [lhs.ellipses.add](#)). To see which hull metrics exist for a specific LoCoH-hullset, use [summary.locoh.lhs](#).

**Value**

A named list with the following elements:

1. `ufat`. User-friendly axis title
2. `desc`. Description (used to build plot captions)
3. `def`. Definition. If `NULL`, then the definition is the same as `$desc`
4. `ufipt`. User-friendly isopleth plot title. Used when plotting isopleths constructed by sorting hulls on the metric.
5. `nhv`. No hull value (the value to use for parent points that have no hulls, used to highlight the presence of points without hulls in histograms)
6. `iso.dec`. Sort hulls in descending order when used to create isopleths). T/F
7. `expr` Expression object that when evaluated returns the hull metric values
8. `zero2na` Whether 0 values should be colored as NA (i.e., hidden) when classifying hull parent points, T/F
9. `discrete` Whether the metric assumes discrete (integer) values, used in classifying hull parent points T/F
10. `auto.jiggle` The maximum value of a uniform distribution used to apply a random offset to the metric when plotted on scatterplots to better see point density
11. `spao.x` include the metric as a default for the x-axis when the 'auto' option is used in `lhs.plot.scatter.auto`
12. `spao.y` include the metric as a default for the y-axis when the 'auto' option is used in `lhs.plot.scatter.auto`
13. `req.metrics` The underlying metric(s) that need to be saved in the hullset for successful computation. Character vector. This will usually simply be the name of the metric itself, but may be different in the case of derived metrics such as perimeter-area ratio.
14. `req.ap` The name(s) of required auxillary variables that are needed to extract the metric. Character vector.
15. `req.ap.def` A named list of default values of any required auxillary variables
16. `req.ap.subtitle` A named list of expression objects (one for each hmap) that returns a string of hmap values that will be used in the subtitle of plots
17. `req.ap.desc` An named list of expression objects (one for each hmap) that returns a string of hmap values that will be used in the description of plots
18. `fun` The name of the function that creates the hull metric
19. `iso.hm2` Second hull metric used to break ties when creating isopleths (used mostly when the metric returns discrete values)
20. `iso.hm2.dec` Sort second hull metric descending

**See Also**

[lhs.ellipses.add](#), [lhs.visit.add](#), [lhs.revisit.add](#)

---

hulls	<i>Extract hulls from a LoCoH-hullset</i>
-------	---

---

### Description

Extract hulls from a LoCoH-hullset

### Usage

```
hulls(lhs, id = NULL, k = NULL, r = NULL, a = NULL, s = NULL,
      hs.names = NULL)
```

### Arguments

lhs	A LoCoH-hullset object
id	The name(s) of individuals
k	The k value of hullsets
r	The r value of hullsets
a	The a value of hullsets
s	The s value of hullsets
hs.names	The name(s) of saved hullsets

### Details

This function returns a list of SpatialPolygonDataFrame objects that contain hulls. There will be one list element for each hullset in lhs. The data table will contain the hull metrics.

### Value

list of SpatialPolygonDataFrame objects

### See Also

[isopleths](#), [lhs.exp.shp](#), [lhs.exp.csv](#), [Vignette on T-LoCoH data classes](#)

---

iso2raster	<i>Convert isopleths to raster</i>
------------	------------------------------------

---

### Description

Converts isopleths in a SpatialPolygonsDataFrame object to a RasterLayer

### Usage

```
iso2raster(polys, raster = NULL, ext = NULL, dimSize = 100,
           cell.size = NULL, sf.cell.size = 2, ll.round = TRUE, status = TRUE,
           debug = FALSE)
```

**Arguments**

<code>polys</code>	A <code>SpatialPolygonsDataFrame</code> containing isopleths sorted by isopleth level (smallest to largest)
<code>raster</code>	A <code>RasterStack</code> object to be used to set the extent and resolution for the output raster.
<code>ext</code>	An extent object or <code>NULL</code> . Ignored if <code>raster</code> is passed.
<code>dimSize</code>	Numeric value used as the number of cells along the largest dimension of the data is numeric. Ignored if <code>raster</code> is passed.
<code>cell.size</code>	. Ignored if <code>raster</code> is passed.
<code>sf.cell.size</code>	. Ignored if <code>raster</code> is passed.
<code>ll.round</code>	Anchor the lower left coordinate of the raster extent to a multiple of the cell size. Ignored if <code>raster</code> is passed.
<code>status</code>	Show status messages and progress bar
<code>debug</code>	Show debugging info

**Details**

This presumes the SPDF contains isopleths ordered from lowest level to highest. The cell values of the resulting raster will sum up to the largest isopleth leve. In order for the resulting raster to sum to 1, the 100 must be part of the input.

---

<code>isopleths</code>	<i>Extract isopleths from a LoCoH-hullset</i>
------------------------	---

---

**Description**

Extract isopleths from a LoCoH-hullset

**Usage**

```
isopleths(lhs, id = NULL, k = NULL, r = NULL, a = NULL, s = NULL,
          hs.names = NULL, iso.idx = NULL, iso.sort.metric = NULL)
```

**Arguments**

<code>lhs</code>	A LoCoH-hullset object
<code>id</code>	The name(s) of individuals
<code>k</code>	The k value of hullsets
<code>r</code>	The r value of hullsets
<code>a</code>	The a value of hullsets
<code>s</code>	The s value of hullsets
<code>hs.names</code>	The name(s) of saved hullsets
<code>iso.idx</code>	The index(s) of the isopleths to extract. Use <a href="#">summary.locoh.lhs</a> to see the indices of the isopleths.
<code>iso.sort.metric</code>	The name(s) of sort metric(s) for the isopleths that will be extracted. Character. See <a href="#">hm.expr</a> .

**Details**

This function returns a list of `SpatialPolygonDataFrame` objects that contain isopleths. There will be one list element for each isopleth in `lhs`. Note that a `LoCoH-hullset` object can contain multiple hullsets, and a single hullset can have multiple isopleths. Accordingly, the names of elements in the returned list are a combination of a hullset name and an isopleth name.

**Value**

list of `SpatialPolygonDataFrame` objects

**See Also**

[hulls](#), [lhs.exp.shp](#), [lhs.exp.csv](#), [Vignette on T-LoCoH data classes](#)

---

<code>lhs.anv.del</code>	<i>Delete ancillary variables from a LoCoH-hullset</i>
--------------------------	--

---

**Description**

Delete ancillary variables from a `LoCoH-hullset`

**Usage**

```
lhs.anv.del(lhs, anv = NULL, id = NULL, k = NULL, r = NULL, a = NULL,
            s = NULL, hs.names = NULL, status = TRUE)
```

**Arguments**

<code>lhs</code>	A <code>LoCoH-hullset</code> object
<code>anv</code>	The name(s) of ancillary variables to remove
<code>id</code>	The name(s) of individuals to process
<code>k</code>	The <code>k</code> value of hullsets to process
<code>r</code>	The <code>r</code> value of hullsets to process
<code>a</code>	The <code>a</code> value of hullsets to process
<code>s</code>	The <code>s</code> value of hullsets to process
<code>hs.names</code>	The name(s) of hullsets to process
<code>status</code>	Show status messages (T/F)

**Value**

A `LoCoH-hullset` object

**Note**

To see which ancillary variables a hullset contains, use the `summary()` function. Copying ancillary variables is optional when creating a hullset with [lxy.lhs](#).

**See Also**

[summary.locoh.lhs](#)

lhs.dr.add

*Define directional routes***Description**

Defines 'directional routes' by identifying parent points that are 1) temporally contiguous and 2) have hulls in the top n

**Usage**

```
lhs.dr.add(lhs, metric = c("ecc", "par")[2], thresh.val = 0.95,
  thresh.type = c("q", "v")[1], smooth = 1, status = TRUE,
  show.elong.hist = FALSE)
```

**Arguments**

lhs	A LoCoH-hullset object
metric	The name of the hull metric used as a proxy for directionality
thresh.val	The threshold above which a hull is considered part of a directional route
thresh.type	The type of threshold used. If 'q' for quantile, then thresh.val is taken to be a percentile of the full range of the directionality metric values (i.e., $0 < \text{thresh.val} < 1$ ). If 'v', thresh.val is taken to be an actual value of the threshold metric.
smooth	The amount of temporal smoothing applied, expressed as the number of points on either side (temporally) of the parent point whose average of the directionality metric is used for determining if the hull constructed around the parent point should be considered a part of a directional route. For no smoothing, set smooth = 0.
status	Display status messages
show.elong.hist	Whether to display histogram(s) of the distribution of the directionality metric before and after smoothing. T/F

**Value**

A LoCoH-hullset object

**Note**

Directional routes are line segments defined by connecting temporally contiguous hull parent-points that are in the top N percent of hull elongation. Hull elongation is proxied by one of two hull metrics: perimeter:area ratio ('par') or the eccentricity of the bounding ellipse ('ecc'). Hull perimeter:area ratios are automatically computed when a hullset is created; bounding ellipses must be computed separately using [lhs.ellipses.add](#). Once computed, you can plot directional routes by passing dr=TRUE to [plot.locoh.lhs](#).



---

lhs.ellipses.add	<i>Compute hull bounding ellipses</i>
------------------	---------------------------------------

---

### Description

Creates bounding ellipses around each hull and computes the eccentricity as a metric of elongation

### Usage

```
lhs.ellipses.add(lhs, id = NULL, k = NULL, r = NULL, a = NULL,
  s = NULL, hs.names = NULL, status = TRUE, beep = FALSE,
  save.ellipses = TRUE, existing.ellipses = c("overwrite", "abort")[1])
```

### Arguments

lhs	A LoCoH-hullset object
save.ellipses	Save ellipses in the lhs object? (T/F). If False, only the eccentricity values of the ellipses are saved as a hull metric.
existing.ellipses	Whether to overwrite or abort if existing ellipses are saved
id	The name(s) of individuals to analyze
k	The k value of hullsets to analyze
r	The r value of hullsets to analyze
a	The a value of hullsets to analyze
s	The s value of hullsets to analyze
hs.names	The name(s) of saved hullsets to analyze
status	Show status messages (T/F)
beep	Beep when done (T/F)

### Value

A LoCoH-hullset object

### Note

Saving the ellipses is optional. If ellipses are not saved, the eccentricity values will still be computed and saved as a hull metric. The only real reason to save ellipses is if you wish to plot them.

Note creating ellipses is computationally intensive and can take a long time.

lhs.exp.csv

*Export a LoCoH-hullset to csv***Description**

Export hull metrics to a csv file / data frame

**Usage**

```
lhs.exp.csv(lhs, id = NULL, k = NULL, r = NULL, a = NULL, s = NULL,
  hs.names = NULL, csv.save = TRUE, dir = ".", csv.fn.pre = NULL,
  csv.fn.mid = c("none", "short", "long")[2], hm = "all", anv = NULL,
  hsp = NULL, status = TRUE)
```

**Arguments**

lhs	A LoCoH-hullset object
id	The name(s) of individuals to export
k	The k value of hullsets to export
r	The r value of hullsets to export
a	The a value of hullsets to export
s	The s value of hullsets to export
hs.names	The name(s) of saved hullsets to analyze
csv.save	Whether to create csv file(s). If FALSE, a list of data frames will still be returned. T/F
dir	The directory where the csv file(s) will be saved
csv.fn.pre	A prefix for the csv file name(s)
csv.fn.mid	Style of the filename mid-section: none, short, or long
hm	The name of hull metric(s) that will be exported. Character vector or comma separated string.
anv	The name(s) of ancillary variables(s) of the parent point of each hull to be exported. Default is NULL, can also be set to 'all'. Character vector or comma separated string.
hsp	A list containing one hull scatterplot object with regions saved (i.e., object returned by <a href="#">lhs.plot.scatter</a> ), or the index of a hull scatter plot saved in the hullset (see <a href="#">lhs.hsp.add</a> ). If passed, a column for the region number will be included in the attribute table. Applicable only to hulls and hull parent points.
status	Show messages. T/F

**Value**

List (one list element for each hullset) of data frames of hull metrics and possibly ancillary variables

**Note**

This function exports hull metrics only. If you want to export the coordinates of the hulls themselves for further analysis in a GIS, see `/link{lhs.exp.shp}`.

If file is left blank, no csv file will be written to disk but the function will still return a list of data frames (one for each hullset)

If you have created a scatterplot legend with manually defined regions (see tutorial for details), the region number of each hull parent point can be included in the csv file if you pass the hullscatter plot as parameter hsp.

**See Also**

[lhs.exp.shp](#)

---

lhs.exp.mov

---

Create a Quicktime animation from a LoCoH-hullset object

---

**Description**

Create a Quicktime animation from a LoCoH-hullset object

**Usage**

```
lhs.exp.mov(lhs, id = NULL, k = NULL, r = NULL, a = NULL, s = NULL,
  hs.names = NULL, all.ids.at.once = TRUE,
  all.ids.col.unique = all.ids.at.once, all.ids.col = NULL,
  all.ids.legend = c("bottomright", "bottom", "bottomleft", "left", "topleft",
  "top", "topright", "right", "center")[5], all.ids.legend.cex = 0.8,
  dt.start = NULL, dt.end = NULL, frame.method = c("auto", "time",
  "location")[1], frame.rtd = "auto", rtd.center = TRUE,
  clean.multihull.frames = 4, xlim = NULL, ylim = NULL, dt.label = TRUE,
  dt.label.col = "black", dt.label.col.bg = NA, dt.label.x = NULL,
  dt.label.y = NULL, title = NULL, title.show = TRUE, axes.show = TRUE,
  axes.ticks = axes.show, axes.titles = FALSE, mar.map = c(0.7 + (if
  (axes.ticks) 0.9 else 0) + (if (axes.titles) 1.3 else 0), 0.5 + (if
  (axes.ticks) 0.9 else 0) + (if (axes.titles) 1.3 else 0), if (title.show) 2.1
  else 0.5, 0.5), mgp.map = c(0.4 + if (axes.ticks) 1.2 else 0, 0.4, 0),
  col.hull.active = "red", col.hull.alpha = 255, tz.local = NULL,
  tz.local.check = TRUE, col.by.hour.of.day = FALSE,
  col.hod = colorRampPalette(colors()[c(24, 30, 553, 121, 26, 121, 553, 30,
  24)])(24), width = if (screen.test) 7 else 608, height = NULL,
  max.frames = NULL, png.pointsize = 16 + (width - 480)/80,
  screen.test = FALSE, tmp.dir = NULL, tmp.files.delete = TRUE,
  prompt.continue = TRUE, fn.mov = NULL, fn.mov.dir = getwd(),
  fn.mov.exists = c("auto.increment", "overwrite", "stop", "ask")[1],
  duration = NULL, fps = NULL, skip = NULL, ffmpeg = "ffmpeg.exe",
  create.mov = TRUE, info.only = TRUE, shp.csv = NULL, layers = NULL,
  tiff.fn = NULL, tiff.pct = FALSE, tiff.bands = c(4, 3, 2),
  tiff.buff = 500, tiff.fill.plot = TRUE, bg2png = !is.null(layers),
  crop.layers.to.extent = TRUE, date.bar = 0.85, date.bar.bins = 12,
  col.db = "darkblue", cex.axis.db = 0.7, beep = FALSE,
  report.time = TRUE, status = TRUE)
```

**Arguments**

lhs	A <a href="#">LoCoH-hullset</a> object
id	The id value(s) to be on the plot
k	The k value of hullsets to export
r	The r value of hullsets to export
a	The a value of hullsets to export
s	The s value of hullsets to export
hs.names	The name(s) of saved hullsets to export
all.ids.at.once	Display all the individual ids simultaneously. If False, an animation will be created for each id. T/F.
all.ids.col.unique	Whether to use unique colors for each individual when plotting multiple individuals simultaneously, T/F
all.ids.col	A named list of color values; the element names must match the name(s) of the ids
all.ids.legend	Where to place a legend showing the color of each id (for an animation showing the movement of multiple individuals simultaneously): 'bottomright', 'bottom', 'bottomleft', 'left', 'topleft', 'top', 'topright', 'right', or 'center'. May also be NULL, in which case the legend will not be displayed. Ignored if <code>ids.legend.bln=FALSE</code> or <code>all.ids.col.unique = FALSE</code> .
all.ids.legend.cex	The character expansion factor for the id legend. See parameter <code>all.ids.legend</code> above.
dt.start	The starting date-time. An object of class POSIXt or one that can be coerced to POSIXt. If NULL, the earliest date-time in the series will be used.
dt.end	The ending date-time. An object of class POSIXt or one that can be coerced to POSIXt. If NULL, the last date-time in the series will be used.
frame.method	How each frame should be defined temporally, "time" - each frame represents a fixed amount of time, "location" each frame is a point in the series
frame.rtd	The real-time duration of each frame (in seconds). If "auto" (default), the lowest median sampling frequency will be used
xlim	A two-element numeric vector for the range of the x-axis (in map units)
ylim	A two-element numeric vector for the range of the y-axis (in map units)
dt.label	Add a label for the date of the frame
dt.label.col	A color value/name for the date label (ignored if <code>dt.label=FALSE</code> )
dt.label.x	The x-coordinate for the date label (ignored if <code>dt.label=FALSE</code> )
dt.label.y	The y-coordinate for the date label (ignored if <code>dt.label=FALSE</code> )
dt.label.col.bg	The background color for the date label (NA for transparent background, ignored if <code>dt.label=FALSE</code> )
title	A title for the map
title.show	Whether to show the title. T/F.
axes.show	Whether to show the axes (ticks, labels and titles). Can be over-written by <code>axes.ticks</code> and <code>axes.titles</code> . T/F.

<code>axes.ticks</code>	Whether to show the tick marks and labels on the axes. T/F.
<code>axes.titles</code>	Whether to show axes titles. T/F.
<code>mar.map</code>	Margin settings for the map, see <a href="#">par</a>
<code>mgp.map</code>	Locations of the axes elements for the map, see <a href="#">par</a>
<code>tz.local</code>	The name of the time zone of the study area
<code>tz.local.check</code>	Check whether <code>tz.local</code> is a valid timezone name (not implemented) T/F.
<code>col.by.hour.of.day</code>	Whether to color the active point by the hour of day (i.e., dark colors at night, orange for day time locations). T/F
<code>col.hod</code>	A vector of 24 color values used to symbolize the color of the active point by the hour of day. Ignored if <code>col.by.hour.of.day</code> = FALSE.
<code>rtd.center</code>	Whether to center the start and end time of the first frame around the time stamp of the first location, to help ensure that each frame has only one hull in it. T/F.
<code>clean.multihull.frames</code>	A numeric value that determines whether and how much a hull can be time shifted to avoid having two active hulls in the same frame. Pass 0 to disable cleaning multi-hull frames. See details.
<code>col.hull.active</code>	The color of the active hull
<code>col.hull.alpha</code>	A number 0..255 for the alpha value for the hull color (semi-transparency); 0=completely transparent, 255=opaque
<code>width</code>	The width of each frame in pixels (if <code>screen.test=FALSE</code> ) or inches (if <code>screen.test=TRUE</code> ).
<code>height</code>	The height of each frame in pixels (if <code>screen.test=FALSE</code> ) or inches (if <code>screen.test=TRUE</code> ).
<code>max.frames</code>	The maximum number of frames to produce.
<code>png.pointsize</code>	The pointsize (in pixels) for the PNG image, equivalent to the height or width of a character in pixels (increase to make labels appear larger)
<code>screen.test</code>	Create up to three sample frame(s) on the screen (instead of PNG files)
<code>tmp.dir</code>	A directory where temporary PNG files for each frame will be created, character.
<code>tmp.files.delete</code>	Delete the temporary PNG files when done, T/F
<code>prompt.continue</code>	Whether to present a summary of the encoding settings and get user confirmation before continuing, T/F
<code>fn.mov</code>	The path and filename of output Quicktime *.mov file. If NULL a filename will be automatically generated
<code>fn.mov.dir</code>	The directory where the animation will be saved (ignored if a value for <code>fn.mov</code> is passed)
<code>fn.mov.exists</code>	What to do if the animation file already exists: "auto.increment", "overwrite", "stop", or "ask".
<code>duration</code>	The desired duration of the animation (in seconds)
<code>fps</code>	A numeric value for frames per second
<code>skip</code>	Output every nth frame. To include every frame set <code>skip=1</code> . Integer.
<code>ffmpeg</code>	The name of the ffmpeg file. See notes.
<code>create.mov</code>	Whether to actually create the mov file. Set to FALSE preview a few frames without actually encoding them.

info.only	Only return info
shp.csv	The path and filename of a csv file that contains information about shapefiles, including layer names, file, and symbology.
layers	The name(s) of layers in shp.csv to display in the background. Will be displayed using the symbology in shp.csv. Character vector or comma delimited string.
bg2png	Save the plot background elements as a static raster image (to improve speed), ignored if screen.test=TRUE
crop.layers.to.extent	Whether to crop the shapefile layers to the view extent (may speed up drawing time)
date.bar	The height of the lower section of the plot to devote to the time bar, in inches. To hide the time bar completely, set date.bar=0.
date.bar.bins	The target number of bins (tick marks + 1) on the time bar (integer)
col.db	A single color value for the date bar axes / tick labels, character
cex.axis.db	Character expansion factor for the labels on the date bar axis.
beep	Beep when one, T/F
report.time	Show the time taken when done, T/F
status	Show progress bar and status messages
tiff.fn	The path and name of a GeoTIFF file (e.g., satellite image) that will be displayed in the background. See notes.
tiff.pct	Whether or to convert the GeoTIFF to an indexed 256 color RGB image, which may speed up drawing. T/F.
tiff.bands	A vector of three integers corresponding to the bands of the GeoTIFF image that will be mapped to the red, green and blue color guns respectively.
tiff.buff	A numeric buffer distance that the range of the plot will be expanded so the points are not right on the edge of the GeoTIFF.
tiff.fill.plot	Whether to fill the entire plot area with the GeoTIFF. T/F.

### Value

A list with information about each \*.mov file created. Each element of the list is another list with two elements: fn (the full filename) and dim (a two-element numeric vector with the frame width and height). If no \*.mov file(s) were created, returns NULL.

### Note

To create the animation, two and only two of the following parameters must be passed: duration, fps, and skip. The third parameter will be computed. To include every frame, pass fps, set skip=1, and leave duration out.

Larger values for fps will result in the animation running 'faster'. Values between 10 and 20 often work well; beyond 30 fps the eye can't keep up with the motion. Note if you pass values for fps and duration, an appropriate value for skip will be computed but the final duration of the animation may not be exactly equal to duration because only interger values of skip are allowed.

One will normally want to run the function a few times without actually encoding to tweak the frame layout (e.g., where the date label and legend appear). To see what a frame in the output will \*approximately\* look like, set screen.test=TRUE. Once the screen sample looks good, next set max.frames=3, tmp.dir="." (or another folder), create.mov=FALSE, and tmp.files.delete=FALSE.

This will generate a few sample frames in PNG files and not delete them so you can inspect them. Once these look good, create the full animation by setting `create.mov=TRUE`.

If `frame.method` is 'auto', the script will use time-based frames when multiple individuals are being animated simultaneously, and location-based frames otherwise.

`duration` (the duration of the animation in seconds) should not be confused with `frame.rtd` which is the real-time duration of a single frame in seconds (e.g., `frame.rtd=3600` means each frame will represent 1 hour).

Passing a positive value for `clean.multihull.frames` (default value is 4) enables time-shifting hulls if needed to prevent two hulls appearing in one frame and no hulls in the previous or next frame (which produces jerky playback when animated). The maximum amount of time a hull can be shifted is calculated by `frame.rtd / clean.multihull.frames`. Thus for example if `frame.rtd=7200` (two hours), and `clean.multihull.frames=4`, then if a frame has two hulls appearing in it and the parent point of one of those hulls lies within 30 minutes (2 hours / 4) of the beginning or end of the frame, *and* there is no hull in the adjacent frame, that hull will be moved to the earlier / later frame. To disable this effect, set `clean.multihull.frames = 0`.

If `date.bar` is too small or too large, you might get a 'margins too large' error. Try values around 1, or hide the date bar completely by setting `date.bar=0`.

The output animation is encoded in QuickTime format. The Quicktime file is encoded using the 'animation' codec, a lossless format that 'scrubs' well (i.e., you can drag the scroll bar to view frame by frame). This requires installing the open source encoding program `ffmpeg`. `ffmpeg` is a command line program that Linux and Windows users can download from <http://ffmpeg.org/download.html>. Windows users should save the `ffmpeg.exe` file to the working directory or a directory on the path environment variable (e.g., `c:\windows`). Mac users can download `ffmpegX` from <http://ffmpegx.com/download.html> but this has not been tested (pass name to `ffmpeg`).

If `ffmpeg` is not available, you can still use this function to generate the individual frames and then use another utility (e.g., ImageMagick, Quicktime Pro) to combine the frames into a video file. For best results use a 'lossless' compression method in the encoding program. To create the individual frames only, set `tmp.dir="."` (the working directory) and `tmp.files.delete=FALSE`.

If `fn.mov.exists = "auto.increment"`, a two-digit number will be appended to the \*.mov filename to avoid overwriting an existing file

---

lhs.exp.shp

---

Export a LoCoH-hullset to shapefile

---

## Description

Export hulls, hull parent points, and/or isopleths from a `locoh-hullset` object to ESRI's shapefile format

## Usage

```
lhs.exp.shp(lhs, id = NULL, k = NULL, r = NULL, a = NULL, s = NULL,
  hs.names = NULL, hpp = FALSE, hulls = FALSE, iso = FALSE,
  nn = FALSE, ellipses = FALSE, allpts = FALSE, iso.idx = NULL,
  iso.metric = NULL, dir = ".", file.base = "", file.base.auto = TRUE,
  avl.file = NULL, status = TRUE, show.time = TRUE, hm = "all",
  anv = NULL, hsp = NULL, metadata = TRUE)
```

**Arguments**

lhs	A LoCoH-hullset object
id	The name(s) of individuals to export
k	The k value of hullsets to export
r	The r value of hullsets to export
a	The a value of hullsets to export
s	The s value of hullsets to export
hs.names	The name(s) of saved hullsets to export
hulls	Export hulls (as a polygon shapefile). T/F
hpp	Export hull parent points (as a point shapefile). T/F
iso	Export isopleths (as a polygon shapefile). T/F
iso.metric	Hull sort metric(s) for the isopleths that will be exported (acts a filter). Character vector.
iso.idx	Numeric vector of the indices of the isopleths that will be exported (acts a filter)
ellipses	Export ellipses (as a polygon shapefile). T/F
allpts	Export all points (as a point shapefile). T/F
nn	Export nearest neighbors (as a multipoint shapefile). T/F
dir	The directory where the shapefiles will be placed (use "." for the working directory, and "~" for the user directory)
file.base	The base of the file name without any extension (the script will append to this)
file.base.auto	Whether to automatically generate the file name base from the id, s-value, and k-a-r value. Ignored if file.base is passed. T/F
avl.file	The name of an existing avl (ArcView legend) file
status	Show messages. T/F
show.time	Report time for script to complete. T/F
hm	The name of hull metric(s) that will be included in the attribute table. Default is 'all'. Applicable only to hulls and hull parent points. Character vector or comma separated string.
anv	The name of ancillary variables(s) of the parent point that will be included in the attribute table. Default is NULL. Applicable only to hulls and hull parent points. Character vector or comma separated string.
hsp	A list containing one hull scatterplot object with regions saved (i.e., object returned by <a href="#">lhs.plot.scatter</a> ), or the index of a hull scatter plot saved in the hullset (see <a href="#">lhs.hsp.add</a> ). If passed, a column for the region number will be included in the attribute table. Applies only to hulls and hull parent points.
metadata	Export field descriptions in a meta data file (not yet supported)

**Note**

Filenames for the resulting shapefiles are automatically generated, but you can specify the directory with the `dir` argument and the base of the file name with `file.base`. The constructed filenames will consist of a base followed by the name of the hullset and a suffix indicating the type of feature saved (e.g., '.hulls', '.pts', '.iso', etc. ). If a shapefile already exists with the same name, a unique filename will be constructed by appending a two-digit numeric extension.



Most of the time, exporting hull parent points (hpp=T) is the same as exporting all points (allpts=T), however in some cases not all points have enough neighbors to make a hull so they wouldn't be included in exported of hull parent points. Only hull parent points will have the corresponding hull metrics included in the attribute table.

avl.file, if passed, is the name of an ArcView 3.x legend file that contains symbology info. If a value is passed, the script will make a copy of avl.file with the same base name as the shapefile, so that symbology will automatically created in ArcMap. avl.file is presumed to be relative to the working folder.

## See Also

[lhs.exp.csv](#), [hulls](#), [isopleths](#)

---

lhs.filter.anv	<i>Define subsets of hulls based on an ancillary variable</i>
----------------	---

---

## Description

Define subsets of hulls based on an ancillary variable

## Usage

```
lhs.filter.anv(lhs, id = NULL, k = NULL, r = NULL, a = NULL, s = NULL,
  hs.names = NULL, anv.var = NULL, anv.val = NULL, label = NULL,
  col = NULL, status = TRUE)
```

## Arguments

lhs	A <a href="#">LoCoH-hullset</a> object
id	The name(s) of the individual(s) to include. Character vector or comma-delimited character.
k	The k value of hullsets to include
r	The r value of hullsets to include
a	The a value of hullsets to include
s	The s value of hullsets to include
hs.names	The name(s) of saved hullsets to include
anv.var	The name of a single ancillary variable in the LoCoH-hullset object
anv.val	The value(s) of the ancillary variable that define each group
label	Character vector for the labels for each subset. If omitted the value of the ancillary variable will be used
col	Vector of color values (one per group). If omitted colors drawn from a rainbow pallete will be used
status	Show status messages. T/F

## Details

This will return a list that defines subsets of hulls based on the hull parent point's value of an ancillary variable. This can be passed to several functions, including `plot.locoh.lhs` (future), and `lhs.plot.scatter`, to create plots of subsets of hulls.

`anv.val` is a vector of values of `anv.var` that will be used to define the group(s). For a hull to be included in a group, its value of `anv.var` must equal one of the values in `anv.val`. If `anv.val` is omitted, one group will be created for each unique value of `anv.var`. Defining groups based on a range of values (lower and upper limits) is not yet supported, but you could create a new ancillary variable that classifies ranges of values into discrete groups.

Note that this function can only return subsets for *\*one\** hullset. This means that either `lhs` must contain a single hullset, or other parameters (e.g., `id`, `hs.names`, `k`, `r`, and/or `a`) are passed to select one and only one of the hullsets in `lhs`.

## Value

A list that defines subsets of hulls, with one element for each region in `hsp`. Each element is a list with three elements:

**idx** indices of the hull parent points

**label** the label for the subset

**col** the color

---

<code>lhs.filter.hsp</code>	<i>Define subsets of hulls based on the location of the parent point in hull scatterplot space</i>
-----------------------------	--

---

## Description

Define subsets of hulls based on the location of the parent point in hull scatterplot space

## Usage

```
lhs.filter.hsp(lhs, id = NULL, k = NULL, r = NULL, a = NULL, s = NULL,
  hs.names = NULL, hsp = NULL, reg.idx = NULL, label = NULL,
  col = NULL)
```

## Arguments

<code>lhs</code>	A LoCoH-hullset object
<code>id</code>	The name(s) of the individual(s) to include in the plot. Character vector or comma-delimited character.
<code>k</code>	The <code>k</code> value of hullsets to create isopleths for
<code>r</code>	The <code>r</code> value of hullsets to create isopleths for
<code>a</code>	The <code>a</code> value of hullsets to create isopleths for
<code>s</code>	The <code>s</code> value of hullsets to create isopleths for
<code>hs.names</code>	The name(s) of saved hullsets to create isopleths for

<code>hsp</code>	Either the index of a hull scatterplot(s) saved in lhs (use <a href="#">summary.locoh.lhs</a> to which how many hsp objects have been saved), or a list of objects of class <code>locoh.hsp</code> (i.e., the return value of <a href="#">lhs.plot.scatter</a> ).
<code>reg.idx</code>	A numeric vector of the indices of the regions in hsp to include in the filter
<code>label</code>	Character vector of the labels to use for each subset. If not passed, the saved label values in hsp will be used
<code>col</code>	Vector of color values (one per region). If omitted the colors saved in hsp will be used

### Details

This will return a list that defines subsets of hulls grouped according to which manually-digitized region in scatterplot space the hull parent point falls. This can be passed to several functions, including [plot.locoh.lhs](#) (future), and [lhs.plot.scatter](#), to create plots of subsets of hulls.

The scatterplot space, including the manually defined regions, are saved in a `locoh.hsp` object. Typically hsp will be created using the [lhs.plot.scatter](#) function. hsp objects can also be saved in the lhs object (see [lhs.hsp.add](#)).

Note that this function can only return subsets for *\*one\** hullset and *\*one\** hsp object. This means that either lhs must contain a single hullset, or other parameters (e.g., `id`, `hs.names`, `k`, `r`, and/or `a`) are passed to select one and only one of the hullsets in lhs.

### Value

A list that defines subsets of hulls, with one element for each region in hsp. Each element is a list with three elements:

**idx** indices of the hull parent points

**label** the label of the region of hsp

**col** the color of the region in hsp

---

<code>lhs.hsp.add</code>	<i>Adds a hull scatterplot to a LoCoH-hullset object</i>
--------------------------	--

---

### Description

Adds a hull scatterplot to a LoCoH-hullset object

### Usage

```
lhs.hsp.add(lhs, hsp.lst)
```

### Arguments

<code>lhs</code>	A LoCoH-hullset object
<code>hsp.lst</code>	A list of object(s) of class <code>locoh.hsp</code>

### Note

hsp objects are typically created with [lhs.plot.scatter](#) function. They can then be 'saved' in a LoCoH-hullset object with this function.

**Examples**

```
## Display a scatter plot and manually draw four regions on it
## hsp <- lhs.plot.scatter(lhs, x="area", y="par", regions=4)

## Save hull scatter plot as part of the hullset
## lhs <- lhs.hsp.add(lhs, hsp.lst=hsp)
## summary(lhs)
```

---

lhs.hsp.del	<i>Delete a hull scatterplot from a LoCoH-hullset object</i>
-------------	--

---

**Description**

Deletes a saved hull scatterplot from a LoCoH-hullset object

**Usage**

```
lhs.hsp.del(lhs, hsp.idx = "all", status = TRUE)
```

**Arguments**

lhs	A LoCoH-hullset object
hsp.idx	The index(s) of the hullsets to delete. A numeric vector or 'all'
status	Show messages, T/F

**Value**

A LoCoH-hullset object

**Note**

Use the summary() command to see the indices of saved hull scatterplots

---

lhs.iso.add	<i>Compute isopleths</i>
-------------	--------------------------

---

**Description**

Adds isopleth(s) to a LoCoH-hullset object

**Usage**

```
lhs.iso.add(lhs, id = NULL, k = NULL, r = NULL, a = NULL, s = NULL,
  hs.names = NULL, sort.metric = c("auto", hm.expr(names.only = T, desc = F,
  print = F))[1], iso.levels = c(0.1, 0.25, 0.5, 0.75, 0.95),
  iso.method = c("pt.quantiles", "hm.vals")[1], iso.cap.method = c(">=",
  "<=")[1], scale.iso.levels.to.hm.vals = (iso.method == "hm.vals" &&
  max(iso.levels <= 1)), subset.metric = NULL, subset.vals = NULL,
  allow.gpc = TRUE, beep = FALSE, status = TRUE, ...)
```

**Arguments**

lhs	A LoCoH-hullset object
id	The id(s) of the hullsets to create isopleths for
k	The k value of hullsets to create isopleths for
r	The r value of hullsets to create isopleths for
a	The a value of hullsets to create isopleths for
s	The s value of hullsets to create isopleths for
hs.names	The name(s) of saved hullsets to create isopleths for
sort.metric	The name of a hull metric that will be used to sort the hulls prior to merging into isopleths
iso.method	The method used to define isopleths. Default is "pt.quantiles" which defines isopleths as containing a quantile of points (e.g., the 0.1th isopleth contains 10% of the points). Can also be "hm.vals", in which case the isopleth level represents not a proportion of points enclosed but a value of the hull metric (e.g., the 0.1 isopleth contains hulls whose hull metric is $\leq 0.1$ )
iso.levels	A numeric vector of the levels of the isopleths. See details.
iso.cap.method	A character object specifying how isopleths defined by proportions of enclosed points will be 'capped'. Ignored when iso.method="hm.vals". See details.
scale.iso.levels.to.hm.vals	Whether to linearly scale iso.levels (presumed to be 0..1) to the minimum and maximum of the sort.metric value. This is used in conjunction iso.method='hm.vals', to produce isopleths as the aggregation of hulls whose sort.metric value is $\geq i\%$ of the maximum, where i is the isopleth level. T/F
subset.metric	The name of a hull metric that will be used to create subsets of isopleths
subset.vals	A two-column data frame (or matrix) containing the lower and upper bounds of the hull metric in subset.metric, from which subsets of hulls will be extract for separate isopleth construction. Ignored if subset.metric is not provided. Can also be a character string in the form of "equal intervals, n", or "quantiles, n" where n is the number of strata desired (shorthand versions of these commands are "ei,n" and "q,n" where n is a number). The script will compute the break points either spread equally between the minimum and maximum subset metric value (equal interval) or so an equal number of hulls is in each strata
allow.gpc	Allow functions from the gpcplib package if the functions from rgeos fail. See details. T/F
beep	Beep when done. T/F
status	Show status messages. T/F
...	Additional auxillary parameters for the hull sort metric

**Details**

This function creates isopleth(s) for a LoCoH-hullset object. This involves sorting hulls by one of the hull metrics, and then cummulatively unioning them together until a certain level is reached at which point the union is saved as an isopleth. By default, hulls will be sorted according to the area (for k-method) or number of enclosed points (r- and a-methods), producing density isopleths (i.e., utilization distributions).

When iso.method=="pt.quantiles" (the default), isopleths will be defined by the proportion of points they enclose (given in iso.levels). Thus for example the 50% isopleth contains 50% of the

total points. The default isopleth levels are 0.15, 0.25, ..., 0.95. To get an isopleth that encloses all points in the dataset, `iso.levels` must include '1'.

When `iso.method=="hm.vals"`, isopleths will be defined by the value of the hull metrics (provided in `iso.levels`). So if for example `sort.metric="scg.nn.mean"` (average speed of all points identified as nearest neighbors), and `scg.nn.mean` varied from 0.01 to 3.2, the 0.6 isopleth would be the union of all hulls whose average nearest neighbor speed was 0.6 or less. If you would like hulls to be identified by hull metric values, but don't know the range of hull metric values, you can set isopleth levels between 0..1 and set `scale.iso.levels.to.hm.vals=TRUE`.

By default, isopleths defined by quantiles of enclosed points are 'capped' as the smallest number of progressively unioned hulls that enclose a number of points equal to or greater than the isopleth level. For example, if there are 1000 points in the dataset, the 50% isopleth would be the smallest union of hulls that encloses 500 or more points. Alternately, when `iso.cap.method = "<="`, the isopleth will be the largest number of hulls which enclose up to but not more than the isopleth level. The actual number of points enclosed by each isopleth is included in the data table for the isopleth (which you can view by setting `iso.details=TRUE` when calling [summary.locoh.lhs](#)).

The hull metric used for sorting must already have been computed. Several hull metrics (e.g., hull area, number of enclosed points) are 'automatically' computed when a hullset is initially generated (see [lxy.lhs](#)). Other hull metrics must be created separately with functions such as [lhs.ellipses.add](#) and [lhs.visit.add](#). You can use the [summary.locoh.lhs](#) function to see which isopleths have been saved in a LoCoH-hullset object.

Hulls are unioned using the `gUnion` and `gUnaryUnion` functions from the `rgeos` package, which in general is very fast. These functions occasionally fail when lines are too close together or there are other topological conditions. This is not a problem with the data, but an inherent limitation of the algorithms / processing. If `allow.gpc=TRUE`, functions from the `gpclib` package will be used as backup.

## Value

A [LoCoH-hullset](#) object

## See Also

[isopleths](#)

## Examples

```
# Create 0.5 and 0.95 isopleths. By not specifying the sort.metric, density isopleths will be created
# by default, with hulls sorted by area (k-method) or number of enclosed points (r and a method)
# lhs <- lhs.iso.add(lhs, iso.levels=c(0.5, 0.95))

# Compute hull metrics for a 24 hour inter-visit gap, then produce isopleths with hulls sorted by
# the number of separate visits (e.g., visitation)
# lhs <- lhs.visit.add(lhs, ivg=3600*24)
# lhs <- lhs.iso.add(lhs, sort.metric="nsv", ivg=3600*24)
```

---

lhs.iso.del

Delete an isopleth

---

## Description

Delete isopleth(s) from a LoCoH-hullset object

**Usage**

```
lhs.iso.del(lhs, iso.idx = "all")
```

**Arguments**

lhs	A LoCoH-hullset object
iso.idx	The index(s) of isopleths to delete (numeric vector or 'all')

**Value**

A LoCoH-hullset object

**Note**

To see the indices of isopleths, run [summary.locoh.lhs](#)

---

lhs.iso.rast	<i>Convert isopleths to rasters</i>
--------------	-------------------------------------

---

**Description**

Adds rasterized version of isopleth(s) to a LoCoH-hullset object

**Usage**

```
lhs.iso.rast(lhs, id = NULL, k = NULL, r = NULL, a = NULL, s = NULL,
  hs.names = NULL, sort.metric = NULL, iso.method = "pt.quantiles",
  raster = NULL, dimSize = 100, cell.size = NULL, sf.cell.size = 2,
  ll.round = TRUE, status = TRUE)
```

**Arguments**

lhs	A <a href="#">LoCoH-hullset</a> object
id	The id(s) of the hullsets to create isopleths for
k	The k value of hullsets to create isopleths for
r	The r value of hullsets to create isopleths for
a	The a value of hullsets to create isopleths for
s	The s value of hullsets to create isopleths for
hs.names	The name(s) of saved hullsets to create isopleths for
sort.metric	The name(s) of hull metric(s) used to form isopleths that rasters should be created for
iso.method	The method(s) used to define isopleths that will be converted to raster
raster	A RasterLayer object that will be used to set the extent and cell size of the rasterized isopleth
dimSize	The number of cells along the largest dimension of the track. The according raster will be calculated internally. Default is 100. Ignored if raster is passed.
cell.size	The size of each square cell in map units. Ignored if raster is passed.

sf.cell.size	The number of significant figures to use if the cell size has be computed based on dimSize. Default=2. Ignored if raster is passed.
ll.round	Whether to round the lower left coordinate to the lowest multiple of cell.size. Ignored if raster is passed. (T/F)
status	Show status messages. T/F

### Details

This will take exisging isopleths and create raster versions of them.

### Value

A LoCoH-hullset object

### See Also

[lhs.iso.add](#)

### Examples

```
## Not run:
lhs <- lhs.iso.add(lhs)
lhs <- lhs.iso.rast(lhs)

## End(Not run)
```

---

lhs.merge	<i>Merge LoCoH-hullsets</i>
-----------	-----------------------------

---

### Description

Merge LoCoH-hullsets

### Usage

```
lhs.merge(..., check.class = TRUE)
```

### Arguments

...	Two or more LoCoH-hullset objects
check.class	Whether to enforce that all objects merged belong to class locoh.lhs (T/F)

### Value

A LoCoH-hullset object

### Note

Hullsets in different objects can be merged together, but the hullsets themselves remain separate. If duplicate hullsets are detected (e.g., same id, same mode, same parameter value), an error message will appear.



---

lhs.pep.add	<i>Add hull metrics for proportion of enclosed points by each hull for an ancillary variable</i>
-------------	--

---

### Description

Computes proportion of enclosed points of a second set of points

### Usage

```
lhs.pep.add(lhs, pep.var, pep.val = NULL, npep = TRUE, status = TRUE)
```

### Arguments

lhs	A <a href="#">LoCoH-hullset</a> object
pep.var	Name(s) of ancillary variable(s) saved with the LoCoH-hullset.
pep.val	Vector of value(s) for which the percentage of enclosed points will be calculated. If NULL, all unique values of pep.var will be used.
npep	Whether to compute the normalized proportion of enclosed points hull metric (normalized by the proportion of points in the entire dataset), T/F
status	Show messages, T/F

### Value

A LoCoH-hullset object

### Note

The proportion of enclosed points is a hull metric that measures relative association among N individuals when you have movement data for multiple individuals simultaneous over the same time period. The general idea is to create hulls for all of the locations combined (ignoring which location was for each individual), then for each hull look at the proportion of enclosed points for each individual. If the individuals ignored each other, one would expect the proportion of enclosed points for any given hull to be approximately equal to the proportions of each individual in the entire dataset. Deviations from this random mixing null model reflect places where the individuals did not mix evenly (e.g., one dominated).

Do compute pep, all points must have the same id (because you create hulls with the combined dataset) with the original id values saved as an ancillary variable (see [lxy.id.new](#)). Create hulls as you normally would using the 'a' or 'k' method, although it would be recommended to omit time (let s=0) because the time difference between locations means different things for different pairs of points depending on whether they are from the same or different individuals. Pass the name of the ancillary variable that contains the original ids as pep.var and the original id value(s) as pep.val (if pep.val is omitted the pep metric will be computed for all id values found in pep.var. *npep* (normalized proportion of enclosed points) normalizes pep by the proportion of that individual in the entire dataset, such that *npep=1* means the individual was in the hull in the same proportion as it was in the entire dataset. '

### See Also

[lxy.id.new](#)

---

lhs.plot.isoarea	<i>Plot the area of isopleths as a function of the k/a/r parameter</i>
------------------	--

---

## Description

Creates a plot of the area of isopleths in a LoCoH-hullset area

## Usage

```
lhs.plot.isoarea(lhs, id = NULL, k = NULL, r = NULL, a = NULL,
  s = NULL, hs.names = NULL, sort.metric = NULL, iso.idx = NULL,
  figs.per.page = 1, legend = TRUE, title = NULL, title.show = TRUE,
  subtitle = TRUE, mar = c(3.3, 3.2, if (title.show) (if (subtitle) 3.2 else
  2.3) else 0.5, 0.5), mgp = c(2, 0.7, 0), png.fn = NULL, png.dir = NULL,
  png.dir.make = TRUE, png.width = 800, png.height = png.width,
  png.pointsize = 12 + (png.width - 480)/80, png.fn.pre = NULL,
  png.fn.suf = NULL, png.overwrite = TRUE, panel.num = NULL,
  panel.num.inside.plot = !title.show, bg = "white", legend.space = if
  (legend) 0.05 else 0, ...)
```

## Arguments

lhs	A LoCoH-hullset object
id	The id(s) of the individual(s) to include in the plot
k	A k-value for the number of nearest neighbors around each point to include in the plot
r	A r-value for the number of nearest neighbors around each point to include in the plot
a	A a-value for the number of nearest neighbors around each point to include in the plot
s	The s value(s) of nearest neighbor sets to include in the plot. If NULL, all values will be used
hs.names	The name(s) of saved hullsets to analyze
sort.metric	The name(s) of isopleth sort metrics to include in the plot
iso.idx	The indices of isopleths to include in the plot
figs.per.page	Number of plots per page
legend	Whether to include a legend. T/F.
title	The title to be displayed. Character. If NULL a title will be constructed.
title.show	Whether to show the title. T/F.
subtitle	Whether to add a subtitle to the automatically constructed title (when title=NULL, otherwise ignored)
mar	The plot margins. A four item numeric vector
mgp	The distance away from the edge of the plot for the 1) label, 2) tick marks, and 3) axis line. A three-item numeric vector
png.fn	A filename for a PNG file

png.dir	The directory for a PNG file (filename will be constructed automatically). Ignored if png.fn is passed
png.dir.make	Whether to create png.dir if it doesn't exist. T/F
png.width	The width of the PNG image
png.height	The height of the PNG image
png.pointsize	The pointsize (in pixels) for the PNG image, equivalent to the height or width of a character in pixels (increase to make labels appear larger)
png.fn.pre	A prefix that will be used in the construction of the PNG filename. Ignored if png.fn is passed.
png.fn.suf	A suffix that will be used in the construction of the PNG filename. Ignored if png.fn is passed.
png.overwrite	Whether to overwrite an existing PNG file if it exists. T/F.
panel.num	A number or letter to display in the upper left hand corner of the plot when the plot will be used as part of a multi-frame graphic (as in publications). Character
panel.num.inside.plot	Whether to display panel.num inside the plot area itself, as opposed to the title area. Ignored if panel.num is NULL. T/F
bg	Background color
legend.space	The amount of additional space on the lower end of the x-axis to make room for the legend. Expressed as a proportion of the range of the x-axis values
...	Additional parameters that will be passed to the <a href="#">plot</a> function

---

lhs.plot.isoear	<i>Plot the isopleth edge:area ratio as a function of k/a/r</i>
-----------------	---

---

## Description

Creates a plot of the ratio of edge to area for isopleths

## Usage

```
lhs.plot.isoear(lhs, id = NULL, k = NULL, r = NULL, a = NULL,
  s = NULL, hs.names = NULL, sort.metric = NULL, iso.idx = NULL,
  figs.per.page = 1, legend = TRUE, title = NULL, title.show = TRUE,
  subtitle = TRUE, mar = c(3.3, 3.2, if (title.show) (if (subtitle) 3.2 else
  2.3) else 0.5, 0.5), mgp = c(2, 0.7, 0), png.fn = NULL, png.dir = NULL,
  png.dir.make = TRUE, png.width = 800, png.height = png.width,
  png.pointsize = 12 + (png.width - 480)/80, png.fn.pre = NULL,
  png.fn.suf = NULL, png.overwrite = TRUE, panel.num = NULL,
  panel.num.inside.plot = !title.show, bg = "white", legend.space = if
  (legend) 0.05 else 0, ...)
```

**Arguments**

<code>lhs</code>	A LoCoH-hullset object
<code>id</code>	The id(s) of the individual(s) to include in the plot
<code>k</code>	A k-value for the number of nearest neighbors around each point to include in the plot
<code>r</code>	A r-value for the number of nearest neighbors around each point to include in the plot
<code>a</code>	A a-value for the number of nearest neighbors around each point to include in the plot
<code>s</code>	The s value(s) of nearest neighbor sets to include in the plot. If NULL, all values will be used
<code>hs.names</code>	The name(s) of saved hullsets to analyze
<code>sort.metric</code>	The name(s) of isopleth sort metrics to include in the plot
<code>iso.idx</code>	The indices of isopleths to include in the plot
<code>figs.per.page</code>	Number of plots per page
<code>legend</code>	Whether to include a legend. T/F.
<code>title</code>	The title to be displayed. Character. If NULL a title will be constructed.
<code>title.show</code>	Whether to show the title. T/F.
<code>subtitle</code>	Whether to add a subtitle to the automatically constructed title (when <code>title=NULL</code> , otherwise ignored)
<code>mar</code>	The plot margins. A four item numeric vector
<code>mgp</code>	The distance away from the edge of the plot for the 1) label, 2) tick marks, and 3) axis line. A three-item numeric vector
<code>png.fn</code>	A filename for a PNG file
<code>png.dir</code>	The directory for a PNG file (filename will be constructed automatically). Ignored if <code>png.fn</code> is passed
<code>png.dir.make</code>	Whether to create <code>png.dir</code> if it doesn't exist. T/F
<code>png.width</code>	The width of the PNG image
<code>png.height</code>	The height of the PNG image
<code>png.pointsize</code>	The pointsize (in pixels) for the PNG image, equivalent to the height or width of a character in pixels (increase to make labels appear larger)
<code>png.fn.pre</code>	A prefix that will be used in the construction of the PNG filename. Ignored if <code>png.fn</code> is passed.
<code>png.fn.suf</code>	A suffix that will be used in the construction of the PNG filename. Ignored if <code>png.fn</code> is passed.
<code>png.overwrite</code>	Whether to overwrite an existing PNG file if it exists. T/F.
<code>panel.num</code>	A number or letter to display in the upper left hand corner of the plot when the plot will be used as part of a multi-frame graphic (as in publications). Character
<code>panel.num.inside.plot</code>	Whether to display <code>panel.num</code> inside the plot area itself, as opposed to the title area. Ignored if <code>panel.num</code> is NULL. T/F
<code>bg</code>	Background color
<code>legend.space</code>	The amount of additional space on the lower end of the x-axis to make room for the legend. Expressed as a proportion of the range of the x-axis values
<code>...</code>	Additional parameters that will be passed to the <code>plot</code> function

---

lhs.plot.revisit	<i>Plot distribution of revisit times</i>
------------------	---

---

## Description

Plots a histogram of the revisit times a LoCoH-hullset area

## Usage

```
lhs.plot.revisit(lhs, id = NULL, k = NULL, r = NULL, a = NULL,
  s = NULL, hs.names = NULL, ta.min = "auto", ta.max = NULL,
  breaks = 40, xaxis.vals = NULL, ta.min.auto.tau = 4, col = "gray80",
  figs.per.page = 1, title = NULL, title.show = TRUE, subtitle = TRUE,
  mar = c(2.8, 3.2, if (title.show) (if (subtitle) 3.2 else 2.3) else 0.5,
  0.5), mgp = c(1, 0.7, 0), png.fn = NULL, png.dir = NULL,
  png.dir.make = TRUE, png.width = 800, png.height = png.width,
  png.pointsize = 12 + (png.width - 480)/80, png.fn.pre = NULL,
  png.fn.suf = NULL, png.overwrite = TRUE, panel.num = NULL,
  panel.num.inside.plot = !title.show, bg = "white", legend.space = if
  (legend) 0.05 else 0, ...)
```

## Arguments

lhs	A LoCoH-hullset object
id	The id(s) of the individual(s) to include in the plot
k	A k-value for the number of nearest neighbors around each point to include in the plot
r	A r-value for the number of nearest neighbors around each point to include in the plot
a	A a-value for the number of nearest neighbors around each point to include in the plot
s	The s value(s) of nearest neighbor sets to include in the plot. If NULL, all values will be used
hs.names	The name(s) of saved hullsets to plot
ta.min	The minimum time away (in seconds) to include on the histogram , can also be 'auto'
ta.max	The maximum time away (in seconds) to include. If NULL then no upper limit will be imposed
ta.min.auto.tau	The minimum time away to include in 'auto' expressed as the number of median sampling intervals. Ignored if ta.min is not 'auto'.
xaxis.vals	A numeric vector of time-away values (in seconds) that will be labeled on the x-axis.
breaks	A value for 'breaks' that will be passed to the hist function (see help function for breaks)
col	A color value for the histogram bars
bg	A color value for the plot background

<code>figs.per.page</code>	Number of plots per page
<code>legend</code>	Whether to include a legend. T/F.
<code>title</code>	The title to be displayed. Character. If NULL a title will be constructed.
<code>title.show</code>	Whether to show the title. T/F.
<code>subtitle</code>	Whether to add a subtitle to the automatically constructed title (when <code>title=NULL</code> , otherwise ignored)
<code>mar</code>	The plot margins. A four item numeric vector
<code>mgp</code>	The distance away from the edge of the plot for the 1) label, 2) tick marks, and 3) axis line. A three-item numeric vector
<code>png.fn</code>	A filename for a PNG file
<code>png.dir</code>	The directory for a PNG file (filename will be constructed automatically). Ignored if <code>png.fn</code> is passed
<code>png.dir.make</code>	Whether to create <code>png.dir</code> if it doesn't exist. T/F
<code>png.width</code>	The width of the PNG image
<code>png.height</code>	The height of the PNG image
<code>png.pointsize</code>	The pointsize (in pixels) for the PNG image, equivalent to the height or width of a character in pixels (increase to make labels appear larger)
<code>png.fn.pre</code>	A prefix that will be used in the construction of the PNG filename. Ignored if <code>png.fn</code> is passed.
<code>png.fn.suf</code>	A suffix that will be used in the construction of the PNG filename. Ignored if <code>png.fn</code> is passed.
<code>png.overwrite</code>	Whether to overwrite an existing PNG file if it exists. T/F.
<code>panel.num</code>	A number or letter to display in the upper left hand corner of the plot when the plot will be used as part of a multi-frame graphic (as in publications). Character
<code>panel.num.inside.plot</code>	Whether to display <code>panel.num</code> inside the plot area itself, as opposed to the title area. Ignored if <code>panel.num</code> is NULL. T/F
<code>legend.space</code>	The amount of additional space on the lower end of the x-axis to make room for the legend. Expressed as a proportion of the range of the x-axis values
<code>...</code>	Additional parameters that will be passed to the <a href="#">hist</a> function

## Details

This function will plot a histogram of the revisitation times for a hullset. Revisitation time is simply the time between points enclosed in a hull. Basically, all points enclosed by each hull are examined and their time intervals computed. This may help reveal where there are natural temporal cycles in revisitation, for example if you see a spike in revisitation around 24 hours there may be daily revisitation pattern in some of the hulls. You can specify the minimum amount of time-away to show in the histogram (e.g., if enclosed points that are separated by the median sampling interval are of little interest), as well as the maximum time-away period. Note that revisitation metrics do *\*not\** have to be computed for the hullset for the histogram to be computed.

## Value

A list of objects of class histogram (one for each hullset plotted)

## See Also

[lhs.visit.add](#)

---

lhs.plot.scatter	Create scatterplot of hull metrics
------------------	------------------------------------

---

## Description

Multi-purpose scatterplot function for the hull metrics in a LoCoH-hullset

## Usage

```
lhs.plot.scatter(lhs, id = NULL, k = NULL, r = NULL, a = NULL,
  s = NULL, hs.names = NULL, x.axis = NULL, y.axis = NULL,
  limx = NULL, limy = NULL, trans.x = NULL, trans.y = NULL,
  jiggle.x = "auto", jiggle.y = "auto", filter = NULL,
  filter.label.in.subtitle = TRUE, filter.sampsize.in.subtitle = TRUE,
  filter.col.use = TRUE, filter.axes.uniform = TRUE, title = NULL,
  title.show = TRUE, title.hs.name.include = TRUE, title.two.id = FALSE,
  title.axes = TRUE, col = c("gray50", "spiral")[1], bg = NULL,
  mar = c(4, 3.2, if (title.show) (if (title.hs.name.include) 3.9 else 3.2)
  else 0.5, 0.5), mgp = c(2.1, 0.8, 0), lo.save = TRUE,
  lo.margins.set = TRUE, lo.colors.set = TRUE, lo.bg.as.box = FALSE,
  sat.base = NULL, val.base = NULL, hue.offset = NULL,
  center.method = c("bbox", "mean")[2], cex = 0.6, type = c("p", "l",
  "b")[1], ufat = NULL, figs.per.page = 1, add = FALSE, regions = NULL,
  prompt.labels = TRUE, hsp = NULL, hsp.reg.col = TRUE,
  hsp.reg.out = FALSE, hsp.reg.lbl = FALSE, hsp.override = TRUE,
  png.fn = NULL, png.dir = NULL, png.dir.make = TRUE, png.width = 800,
  png.height = png.width, png.pointsize = 12 + (png.width - 480)/80,
  png.fn.pre = NULL, png.fn.mid = NULL, png.fn.suf = NULL,
  png.exists = c("overwrite", "skip", "abort")[1], status = TRUE,
  desc = c(NONE <- 0, BOTTOM <- 1, TOP <- 3)[ifelse(figs.per.page == 1, 2,
  1)], cex.desc = 0.8, col.desc = NULL, panel.num = NULL,
  panel.num.inside.plot = !title.show, hmap = NULL,
  hmap.in.subtitle = TRUE, check.ap.value.in.hmapparams = TRUE, ...)
```

## Arguments

lhs	A LoCoH-hullset object
id	The name(s) of the individual(s) to include in the plot. Character vector or comma-delimited character.
k	The k value(s) of the hullset(s) to include in the plot. Numeric vector or comma-delimited character object.
r	The r value(s) of the hullset(s) to include in the plot. Numeric vector or comma-delimited character object.
a	The a value(s) of the hullset(s) to include in the plot. Numeric vector or comma-delimited character object.
s	The s value(s) of the hullset(s) to include in the plot. Numeric vector or comma-delimited character object.
hs.names	The name(s) of saved hullsets to include in the plot.
x.axis	The name of a hull metric (see <a href="#">hm.expr</a> )

<code>y.axis</code>	The name of a hull metric
<code>limx</code>	The lower and upper limits of the x-axis. Two-element numeric vector.
<code>limy</code>	The lower and upper limits of the y-axis. Two-element numeric vector.
<code>trans.x</code>	The name of a function that will be used to transform the x-axis values. Can be any R function (e.g., "log", "sqrt"), also "square" and "cube"
<code>trans.y</code>	The name of a function that will be used to transform the y-axis values
<code>jiggle.x</code>	"auto" or a numeric value which will be used to add a normally distributed stochastic value (mean 0, sd=jiggle.x) to the x-axis values for better visualization of the number of points in each group (helpful when the x-values take on discrete values)
<code>jiggle.y</code>	"auto" or a numeric value which will be used to add a normally distributed stochastic value (mean 0, sd=jiggle.y) to the y-axis values
<code>title</code>	The title to be displayed. Character. If NULL a title will be constructed.
<code>title.show</code>	Whether to show the title. T/F.
<code>title.hs.name.include</code>	Whether to include the hullset name as part of the title. Ignored if title is passed. T/F.
<code>title.axes</code>	Whether to include the names of the hull metrics on the axes as part of the subtitle. Ignored if title is passed. T/F.
<code>title.two.id</code>	An ad-hoc way to construct a title consisting of the ids of the hullset and hs2
<code>filter</code>	NULL or a list of filter parameters. See details.
<code>filter.label.in.subtitle</code>	Whether to add the filter label to the plot subtitle. Ignored if title is passed or filter is NULL. T/F.
<code>filter.samplesize.in.subtitle</code>	Whether to add the filter sample size to the plot subtitle. Ignored if title is passed or filter is NULL. T/F.
<code>filter.col.use</code>	Whether the points will be displayed with the color(s) saved in the filter (overriding the 'col' parameter). Ignored if filter is NULL. T/F.
<code>filter.axes.uniform</code>	Whether the upper and lower limits of the axes for each plot will be set as the limit for the entire combined dataset. Ignored if filter is NULL or values are passed for limx or limy. T/F.
<code>col</code>	A single color value, or a vector of color values of the same length as the number of hulls, or "spiral" (see additional parameters below). Ignored if filter.col.use=T or something is passed for hsp (in which case col will be extracted from hsp).
<code>bg</code>	Background color
<code>lo.colors.set</code>	Whether to set the background, foreground color on the plot device. T/F.
<code>lo.bg.as.box</code>	Whether to display the background color as a box on the plot (needed by some wrapper function that want the axes and margins to be a different color). This overrides the setting of lo.colors.set. T/F.
<code>mar</code>	The plot margins. A four item numeric vector.
<code>mgp</code>	The distance away from the edge of the plot for the 1) label, 2) tick marks, and 3) axis line. A three-item numeric vector.
<code>lo.save</code>	Whether to save and reset the plot device margin settings (some wrapper functions that call this function don't want device settings reset). T/F.



lo.margins.set	Whether to save and reset the plot device margin settings (some wrapper functions that call this function don't want device settings reset). T/F.
desc	Which side to display automatically generated descriptive text (e.g. caption). 0=none, 1=bottom, 3=top.
cex.desc	The expansion factor for the descriptive text. Numeric value.
col.desc	The color of the descriptive text. Color value.
sat.base	A number between 0 and 1 for the base saturation (how much of the color is present) of the center color (s=0.4 gives pastels), ramping up to 1 (full color) at edges.
val.base	A number between 0 and 1 controlling how much black is in the center color (0 is total black), ramping out to no blackness at the edges
hue.offset	A number between 0 and $2\pi$ for the color wheel rotation (in radians), which controls the color of points directly to the right of the center.
center.method	Determines how the center of the color will be computed. "bbox" = the center of the bounding box of the full range of points; "mean" = the mean of the data plot
cex	Expansion factor for the points on the scatterplot
type	The type of plot: 'p'=points only, 'l'=line, 'b'=both
ufat	Whether to substitute user-friendly axis titles. T/F.
figs.per.page	The number of plots per page
add	Whether to add to the current plot device. T/F.
regions	Determines whether the user will be prompted to draw regions (polygons) on the scatterplot window with the mouse, that will be saved as part of the scatterplot object returned. Values can be 1) a number whereby the user will be prompted to create N regions with randomly assigned colors, or 2) a vector of color values in which case the regions will be assigned those colors.
prompt.labels	Whether the user should be prompted to also enter labels for each of the regions drawn. Ignored if regions = NULL. T/F.
hsp	Either the index(s) of a hull scatterplot(s) saved in lhs (use summary() command to which how many hsp objects have been saved), or a list of objects of class locoh.hsp. When passed, the parameters in hsp will be used to create the scatterplot
hsp.reg.col	Whether to use the colors assigned to the regions in hsp. Ignored if hsp is NULL. T/F.
hsp.reg.out	Whether to display the outline of the regions in hsp. Ignored if hsp is NULL. T/F.
hsp.reg.lbl	Whether to display the labels of the regions in hsp. Ignored if hsp is NULL. T/F.
hsp.override	Whether the parameters in hsp should override other parameters passed. Ignored if hsp is NULL. T/F.
png.fn	The path and name of the PNG file to create (instead of displaying in a plot window).
png.dir	The directory for a PNG file (filename will be constructed automatically). Ignored if png.fn is passed.
png.dir.make	Whether to create png.dir if it doesn't exist. T/F.
png.width	The width of the PNG image. Ignored if png.fn is passed.

png.height	The height of the PNG image. Ignored if png.fn is passed.
png.pointsize	The pointsize (in pixels) for the PNG image (increase to make labels appear larger). Equivalent to the height or width of a character in pixels.
png.fn.pre	A prefix that will be used in the construction of the PNG filename. Ignored if png.fn is passed.
png.fn.mid	A mid-fix that will be used in the construction of the PNG filename. Ignored if png.fn is passed.
png.fn.suf	A suffix that will be used in the construction of the PNG filename. Ignored if png.fn is passed.
png.exists	What to do if a PNG with the same filename already exists: "overwrite", "skip", or "abort"
status	Whether to show messages. T/F.
panel.num	A number or letter to display in the upper left hand corner of the plot, used when the plot will be part of a multi-frame graphic (as in publications). Character.
panel.num.inside.plot	Whether to display panel.num inside the plot area itself (as opposed to the title area). T/F.
hmap	A named list of hull metric auxillary parameters, the name of each list element is the name of the variable.
hmap.in.subtitle	Include the hmap value(s) in the plot subtitle. T/F
check.ap.value.in.hmaparams	Check to make sure that hull metrics have been computed for the hull metric auxillary parameter values passed (either as separate arguments or in hmap)
...	Other parameters, including any auxillary parameters required by certain hull metrics

## Details

This function has two main purposes. 1) To make scatterplot graphics either in a plot window or PNG file 2) To create a list of objects of class "locoh.hsp", which are basically a collection of the parameters that were used to create the scatterplot including any manually drawn regions created by the user (see regions).

Note that hull metrics must already have been computed. Several hull metrics are 'automatically' computed when the hullset is defined (e.g., hull area, number of enclosed points). Other hull metrics must be created separately with functions such as [lhs.ellipses.add](#) and [lhs.visit.add](#). Auxillary parameters required for hull metrics can be passed either as individual parameters (e.g., ivg=3600\*12) or as a list element (e.g., hmap=list(ivg=86400)).

filter, if passed, will create scatterplots for subsets of hulls. For this to work, filter must be a list whose elements are named lists with the following elements: *idx* = a vector of the indices of the hulls (in the hulls SpatialPolygonsDataFrame), *label* = a label for the subset, and *col* = a color value. [lhs.filter.hsp](#) can be used to create a filter list based on manually drawn regions of a hull scatterplot, and [lhs.filter.anv](#) can be used to create a filter based on ranges of values of an ancillary variable. The arguments filter.sampsize.in.subtitle, filter.col.use, and filter.axes.uniform control how the subsets are treated in the scatterplot(s).

**Value**

If `png.fn` or `png.dir` is passed, the plots are exported to PNG file(s) and the function returns a list of file names and image dimensions (in pixels). Otherwise returns a named list whose element(s) are of class *locoh.hsp*. This list of *locoh.hsp* objects can then be 'permanently' saved in the LoCoH-hullset object using `lhs.hsp.add`, and/or fed into other functions such as `plot.locoh.lhs` (to symbolize hull points).

**See Also**

`lhs.plot.scatter.auto`, `hm.expr`, `lhs.hsp.add`, `lhs.filter.hsp`, `lhs.filter.anv`

---

`lhs.plot.scatter.auto` *Generate multiple scatterplots of hull metrics*

---

**Description**

This is a wrapper function for `lhs.plot.scatter` that creates scatterplots for pairs of hull metrics. It can quickly generate several few dozen scatterplots for the purpose of visually looking for novel associations between hull metrics.

**Usage**

```
lhs.plot.scatter.auto(hs, id = NULL, k = NULL, r = NULL, a = NULL,
  s = NULL, hs.names = NULL, x.metrics = c("auto", "all")[1],
  y.metrics = c("auto", "all")[1], metrics.exclude = c("bearing",
    "sgf.nn.mean", "sgf.enc.mean"), exclude.same.pair = TRUE,
  exclude.reverse.pair = TRUE, png.dir = NULL, png.dir.make = TRUE,
  png.width = 800, png.height = png.width, png.pointsize = 12 + (png.width
    - 480)/80, png.fn.pre = NULL, png.fn.mid = NULL, png.fn.suf = NULL,
  png.exists = c("skip", "overwrite")[1], progress.bar = TRUE,
  test.sample = NULL, enumerate.pairs.only = FALSE,
  new.plot.window = TRUE, hmap = NULL, ...)
```

**Arguments**

<code>hs</code>	A LoCoH-hullset object
<code>id</code>	The names of the individual(s) to include in the plot. Character vector or comma-delimited character.
<code>k</code>	The k value(s) of the hullset(s) to include in the plot. Numeric vector or comma-delimited character object.
<code>r</code>	The r value(s) of the hullset(s) to include in the plot. Numeric vector or comma-delimited character object.
<code>a</code>	The a value(s) of the hullset(s) to include in the plot. Numeric vector or comma-delimited character object.
<code>s</code>	The s value(s) of the hullset(s) to include in the plot. Numeric vector or comma-delimited character object.
<code>hs.names</code>	The name(s) of saved hullsets to include in the plot.
<code>x.metrics</code>	The name(s) of hull metric(s) to plot on the x-axis. Can also be "auto" for the most common ones, or "all"

<code>y.metrics</code>	The name(s) of hull metric(s) to plot on the y-axis. Can also be "auto" for the most common ones, or "all"
<code>metrics.exclude</code>	The name(s) of hull metrics to exclude from both the x and y axes
<code>exclude.same.pair</code>	Whether to exclude a hull metric being plotted against itself. T/F
<code>exclude.reverse.pair</code>	Whether to exclude opposite pairs of metrics. T/F
<code>png.dir</code>	The directory for a PNG file (filename will be constructed automatically). Ignored if <code>png.fn</code> is passed.
<code>png.dir.make</code>	Whether to create <code>png.dir</code> if it doesn't exist. T/F.
<code>png.width</code>	The width of the PNG image. Ignored if <code>png.fn</code> is passed.
<code>png.height</code>	The height of the PNG image. Ignored if <code>png.fn</code> is passed.
<code>png.pointsize</code>	The pointsize (in pixels) for the PNG image (increase to make labels appear larger). Equivalent to the height or width of a character in pixels.
<code>png.fn.pre</code>	A prefix that will be used in the construction of the PNG filename
<code>png.fn.mid</code>	A mid-fix that will be used in the construction of the PNG filename
<code>png.fn.suf</code>	A suffix that will be used in the construction of the PNG filename
<code>png.exists</code>	What to do if a PNG with the same filename already exists: "overwrite" or "skip"
<code>progress.bar</code>	Display a progress bar. T/F
<code>test.sample</code>	An optional number of randomly selected pairs of hull metrics to generate for testing purposes. Or NULL.
<code>enumerate.pairs.only</code>	Whether to only return a two-column data frame of the names of hull metrics. T/F
<code>new.plot.window</code>	Whether to start a new plot window and turn recording on. T/F
<code>hmap</code>	A named list of hull metric auxiliary parameters (for hull metrics that involve additional parameters such as an interval gap period). These can also be passed on their own (e.g., <code>ivg=3600*24</code> )
<code>...</code>	Additional parameters that will be passed to the <a href="#">hist</a> function

**Value**

A data frame containing the pairs of hull metrics

**Note**

This function generates a list of pairs of hull metrics, and then calls [lhs.plot.scatter](#) repeatedly to generate the histograms. This is a quick way to visualize the relationships between hull metrics. If `new.plot.window` is TRUE (default), a new plot window will be opened and plot recording turned on so you can flip through the histograms using the PgUp and PgDn keys. You may also pass a value for `png.dir`) which will create a bunch of PNG files which you can then view them as a slideshow.

**See Also**

[lhs.plot.scatter](#), [hm.expr](#)

---

lhs.revisit.add	<i>Add revisitation hull metrics to a LoCoH-hullset object</i>
-----------------	--

---

## Description

Computes revisitation rate based on a minimum and maximum time away period

## Usage

```
lhs.revisit.add(lhs, ta.min = NULL, ta.max = NULL, ta.cuts = NULL,  
  status = TRUE)
```

## Arguments

lhs	A LoCoH-hullset object
ta.min	Minimum value(s) for time away in seconds (numeric vector)
ta.max	Maximum value(s) for time away in seconds (numeric vector)
ta.cuts	A numeric vector of time values in seconds that define the time-away intervals
status	Show status messages. T/F

## Details

ta.min and ta.max define the minimum and maximum period of time (in seconds) which must pass for another occurrence in the hull to be considered a 'revisit'. They should be the same lengths. If ta.max is NULL, no upper bound will be set

ta.cuts is an alternative way of specifying the time-away intervals. The time values in ta.cuts will serve as the values between time away intervals. For example if ta.cuts = c(1000, 4000, 7000, 10000), three time-away intervals will be examined: 1000 to 4000 seconds, 4000 to 7000 seconds, and 7000 to 10000 seconds. One way to get the values for ta.cuts is to plot the distribution of revisit times using [lhs.plot.revisit](#), and then use the [get.vals](#) function to select time values between clusters of revisit times.

## Value

A LoCoH-hullset object

## See Also

[lhs.plot.revisit](#), [get.vals](#)

---

lhs.revisit.del	<i>Delete all revisitation hull metrics in a LoCoH-hullset object</i>
-----------------	---

---

**Description**

Delete all revisitation hull metrics in a LoCoH-hullset object

**Usage**

```
lhs.revisit.del(lhs, status = TRUE)
```

**Arguments**

lhs	A LoCoH-hullset object
status	Display summary, T/F

**Value**

A LoCoH-hullset object

**See Also**

[lhs.revisit.add](#)

---

lhs.save	<i>Save a LoCoH-hullset object to disk</i>
----------	--

---

**Description**

Saves a LoCoH-hullset object to disk, constructing a filename if needed that reflects the contents

**Usage**

```
lhs.save(lhs, file = NULL, dir = ".", suf = NULL, compress = TRUE,
  auto.num.files = TRUE, width = 2, save.as = NULL)
```

**Arguments**

lhs	A LoCoH-hullset object
file	Optional filename. If omitted a sensible filename will be constructed
dir	Directory for the output file (relative to the working directory), ignored if file passed
suf	A suffix that will be used in the construction of the filename, ignored if file is passed
compress	Compress file. T/F
auto.num.files	Use a number as part of the constructed filename. T/F
width	The number of digits of the auto-number token in the filename (ignored if auto.num.files=F)
save.as	The name of the object when saved (default is the same as the original)

**See Also**[lxy.save](#)


---

lhs.select	<i>Select hullsets</i>
------------	------------------------

---

**Description**

Select hullsets

**Usage**

```
lhs.select(lhs, id = NULL, k = NULL, r = NULL, a = NULL, s = NULL,
           hs.names = NULL, hs.idx = NULL, status = TRUE)
```

**Arguments**

lhs	A LoCoH-hullset object
id	The name(s) of individuals to extract
k	The k value of hullsets to extract
r	The r value of hullsets to extract
a	The a value of hullsets to extract
s	The s value of hullsets to extract
hs.names	The name(s) of saved hullsets to extract
hs.idx	The indices of saved hullsets to extract
status	Show status messages (T/F)

**Value**

A LoCoH-hullset object

---

lhs.so.add	<i>Add hull metrics for association analysis</i>
------------	--

---

**Description**

Add hull metrics for association analysis

**Usage**

```
lhs.so.add(lhs, id = "all", hs2.id = "all", tbuff = 0, ivg = NULL,
           test = 0, skip.dups = TRUE, save.hso = TRUE, status = TRUE,
           piFUN = c("pIntersect", "pIntersectSat")[1])
```

## Arguments

<code>lhs</code>	A <a href="#">LoCoH-hullset</a> object
<code>save.hso</code>	Whether to save the hull intersection list, T/F
<code>skip.dups</code>	Skip duplicate hulls (faster)
<code>id</code>	A character vector of the hullset ids to compute metrics for. Can also be 'all'.
<code>hs2.id</code>	A character vector of the hullset ids to use as the comparison hullsets. Can also be 'all'.
<code>tbuff</code>	A temporal overlap threshold (in seconds). See details.
<code>ivg</code>	The intervisit gap period used to collapse intersecting hulls into discrete visits, see details
<code>test</code>	A two-element numeric vector containing the number of hulls in hullset 1 and hullset 2 respectively to identify intersections
<code>status</code>	Show status messages. T/F
<code>piFUN</code>	The function to use to identify which pairs of hulls intersect: 'pIntersect' or 'pIntersectSat'

## Details

This function computes hull metrics for the spatially overlapping hulls from two ids. Typically this would be used when you have hullset from two individuals (i.e., two animals) and you want to see the spatial and temporal patterns of shared space use.

You can impose a temporal overlap requirement as well by passing a value for `tbuff`. Two hulls will be considered spatially overlapping only if their parent points also were recorded within `tbuff` seconds of each other. This essentially produces metrics for spatially and temporally overlapping hulls.

Hullset metrics are computed for each pair of ids. Thus if a hullset has hulls for three unique ids, each hull will have spatial overlap metrics computed for each of the other two hullsets. You can narrow which id(s) to compute metrics for, and which hullset(s) to use as the comparison, with the `id` and `hs2.id` arguments.

Up to three spatial overlap metrics are computed. `so.count` is simply the number of hulls in hullset 2 that overlap. `so.dtmn` is the minimum amount of time (expressed in seconds) that passes between overlapping hulls. This reflects temporal partitioning of shared space - low values of `so.dtmn` suggest the two individuals don't mind being in the same area at the same time. `so.nsv` (number of separate visits) is similar to `so.count`, but collapses overlapping hulls into discrete visits based on an intervisit gap period `ivg`. `so.nsv` is only computed if a value for `ivg` is passed.

`pIntersect` and `pIntersectSat` are two functions that identify which pairs of hulls actually intersect. Neither are terribly fast, but `pIntersect` appears to work faster than `pIntersectSat`.

## Value

A [LoCoH-hullset](#) object



---

lhs.to.add*Compute hull metrics for temporally overlapping hulls*

---

## Description

Compute hull metrics for pairs of temporally overlapping hulls of separate individuals

## Usage

```
lhs.to.add(lhs, id = "all", hs2.id = "all", maxdt = "auto",  
  save.hto = TRUE, status = TRUE)
```

## Arguments

lhs	A <a href="#">LoCoH-hullset</a> object
id	A character vector of the hullset ids to compute metrics for. Can also be 'all'.
hs2.id	A character vector of the hullset ids to use as the comparison hullsets. Can also be 'all'.
maxdt	The maximum difference in time (in seconds) for two hulls to be considered 'overlapping' in time. Can also be 'auto', in which case half of the smallest of the two median sampling intervals will be used.
save.hto	Whether to save the list of hull indices that temporally overlap in the hullset, T/F
status	Show status messages, T/F

## Details

This will compute hull metrics for pairs of hulls from two individuals. This only works for a LoCoH-hullset object that contains hulls from two or more individuals (id's). For each hull for individual A, for example, it will identify the hulls in individual B that temporally overlap, and compute the mean centroid distance (hull metric name = to.mcd). By default *to.mcd* is computed for all pairs of individuals, but you can specify specific pairs by passing values for *id* and *hs2.id*.

This metric is used for association analysis. You can use this metric for example to look for spatial and temporal patterns in how close individuals get to each other. Other association metrics exist for spatially overlapping hulls (see [lhs.so.add](#)).

## Value

A LoCoH-hullset object

## See Also

[lhs.so.add](#), [lhs.merge](#)

lhs.to.stats

*Computes statistics on the centroid distances for time-overlapped hulls***Description**

Computes statistics on the hull-to-hull centroid distance for time-overlapped hulls, with an option to plot and overlay the distribution of centroid distances for random pairs of hulls

**Usage**

```
lhs.to.stats(lhs, id1 = "all", id2 = "all", n = "all", iso.lower = NULL,
  iso.upper = NULL, iso.oz = TRUE, iso.sort.metric = "auto",
  to.comp.hist = TRUE, breaks = 20, to.mcd.outline.only = FALSE,
  lwd.outline = 3, hist.type = c("density", "counts")[1],
  col.to.mcd = "blue", col.h2h.cd = "red", title = NULL,
  title.show = TRUE, title.id.only = FALSE, title.sub.iso.enc = TRUE,
  mar = c(3, 3, if (title.show) 1.5 + (if (title.sub.iso.enc) 1.3 else 0) else
    0.5, 0.5), mgp = c(1.8, 0.5, 0), figs.per.page = 1, panel.num = NULL,
  panel.num.inside.plot = !title.show, png.dir = NULL,
  png.dir.make = TRUE, png.width = 800, png.height = png.width,
  png.overwrite = TRUE, png.pointsize = 12 + (png.width - 480)/80,
  status = TRUE, ...)
```

**Arguments**

lhs	A LoCoH-hullset object
id1	Hullset 1 id value(s). Can also be 'all'.
id2	Hullset 2 id value(s). Can also be 'all'.
n	The number of randomly selected paired hulls to use as the NULL model of no association. Can also be "all"
iso.oz	Apply isopleth filter to hs1 using the isopleths from hs2. T/F.
iso.lower	The lower level isopleth for the isopleth filter
iso.upper	The upper level isopleth for the isopleth filter
iso.sort.metric	The name of a hull metric that was used to sort hulls in the construction of the isopleths to be used as filters. If auto (default) it will pick the default sort metric used for density isopleths (i.e., area for the k-method, and number of enclosed points for the a and r methods)
to.comp.hist	Draw a histogram of the centroid distances of temporally overlapping hulls
breaks	The number of breaks in the histogram (or another valid value for breaks, see <a href="#">hist</a> ).
to.mcd.outline.only	Show the outline only of the histogram of the mean centroid distance for temporally overlapping hulls. T/F.
lwd.outline	The line width of the histogram outline (ignored if to.mcd.outline.only=F).
hist.type	The type of histogram to plot: 'density' or 'counts'.
col.to.mcd	The color of the outline of the histogram of the distribution of the centroid distances of temporally overlapping hulls.

<code>col.h2h.cd</code>	The color of the outline of the histogram of the distribution of the centroid distances for randomly paired hulls.
<code>title</code>	The title to be displayed. Character. If NULL a title will be constructed.
<code>title.show</code>	Whether to show the title. T/F.
<code>title.id.only</code>	Whether to construct the title from the id values only. T/F. Ignored if <code>title</code> is passed or <code>title.show=FALSE</code>
<code>title.sub.iso.enc</code>	Whether to include the isopleth filter information as the second line of the title, T/F.
<code>mar</code>	The plot margins. A four item numeric vector
<code>mgp</code>	The distance away from the edge of the plot for the 1) label, 2) tick marks, and 3) axis line. A three-item numeric vector
<code>figs.per.page</code>	The number of plots per page.
<code>panel.num</code>	A number or letter to display in the upper left hand corner of the plot when the plot will be used as part of a multi-frame graphic (as in publications). Character
<code>panel.num.inside.plot</code>	Whether to display <code>panel.num</code> inside the plot area itself, as opposed to the title area. T/F
<code>png.dir</code>	The directory for a PNG file (filename will be constructed automatically). Ignored if <code>png.fn</code> is passed
<code>png.dir.make</code>	Whether to create <code>png.dir</code> if it doesn't exist. T/F
<code>png.width</code>	The width of the PNG image
<code>png.height</code>	The height of the PNG image
<code>png.overwrite</code>	Whether to overwrite an existing PNG file if it exists. T/F
<code>png.pointsize</code>	The pointsize (in pixels) for the PNG image, equivalent to the height or width of a character in pixels (increase to make labels appear larger)
<code>status</code>	Display status messages. T/F
<code>...</code>	Additional parameters that will be passed to the <a href="#">plot</a> function

## Details

This returns a list object containing the centroid-to-centroid distances of a random selection of hulls from two individuals, which serves as a NULL model of no interaction. It can also plot the histogram of mean-centroid-distance of time-overlapped hulls with the outline of the centroid distance of random-pairs of hulls overlain on top, to visually see how close the distributions match.

It will also compute the Welch Two Sample t-test to see if the distribution for time-overlapped and randomly paired hulls have statistically significant means, and the Two-sample Kolmogorov-Smirnov test which tells you how likely the two distributions are the same.

Note that before you can use this function, the mean-centroid-distance for time-overlapped hulls must be computed using [lhs.to.add](#)

You can apply an isopleth filter by passing values for `iso.lower` and `iso.upper`. These should be the isopleth level (normally between 0 and 1) that the hull parent point must fall in to be included in the analysis. If, for example, you wanted to see whether association in the core area is significantly different from neutral interaction, you would pass `iso.upper=0.5` and leave `iso.lower` NULL. Note that the isopleths with matching isopleth levels must already be present.

**Value**

A list object

**See Also**

[lhs.to.add](#), [lhs.iso.add](#)

---

lhs.visit.add

---

Add time use hull metrics to a LoCoH-hullset object

---

**Description**

Computes visitation and duration hull metrics based on inter-visit gap value(s)

**Usage**

```
lhs.visit.add(lhs, ivg = NULL, status = TRUE)
```

**Arguments**

lhs	A LoCoH-hullset object
ivg	Value(s) for inter-visit gap (in seconds) (numeric vector)
status	Show status messages. T/F.

**Value**

A LoCoH-hullset object

**Note**

The inter-visit gap is the period of time (in seconds) which must pass before another occurrence in the hull can be considered a separate visit. Occurences in the hull are considered a separate visit only if the animal was absent from the hull for a period of time  $\geq$  ivg.

For each ivg value, the function examines each hull and computes the number of separate visits to the hull (visitation, hull metric name = "nsv") and the mean number of locations per visit (visit duration, hull metric name = "mnlv")

---

locoh.lhs	<i>Class for a hullsets</i>
-----------	-----------------------------

---

## Description

Class for a collection of hullsets and associated objects including hull metrics and isopleths

## Details

An object of class `locoh.lhs` is a list (S3) containing one or more sets of hulls (local MCPs) for a set of points. One hull is saved for each point if the method used to create the hulls returned at least two nearest neighbors for each point, otherwise the number of hulls might be less than the number of points. In addition to hulls (saved as a `SpatialPolygonsDataFrame`), LoCoH-hullset object may also contain:

- the original points and their attribute table
- an index of the points a) used to construct the hull and b) enclosed by each hull
- hull metrics
- isopleths (progressive unions of hulls) as both vector (`SpatialPolygonsDataFrame`) and raster (utilization distribution) formats.
- saved parameters for scatterplots of hull metrics.

In code examples, objects of class `locoh.lhs` are often noted by a 'lhs' suffix, and functions that are designed to work on LoCoH-hullset objects typically start with 'lhs'.

For a complete description of the data structure, see the *T-LoCoH Data Classes* vignette [../doc/tlocoh\\_data\\_classes.pdf](#)

## See Also

[lxy.lhs](#), [summary.locoh.lhs](#), [plot.locoh.lhs](#), [lhs.merge](#), [lhs.iso.add](#), [lhs.save](#), [lhs.exp.shp](#)

---

locoh.lxy	<i>Class for location data</i>
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---

## Description

Data class for locations with associated dates, nearest neighbor info, parameters for a random walk null model, and assorted meta-data

## Details

An object of class `locoh.lxy` is a list containing a series of point locations and ancillary variables that go with those locations (e.g., time stamps, point IDs, animal IDs). These items are bundled together and 'ready to go' as inputs into T-LoCoH functions. The benefits of bundling all of the objects related to a set of point locations in a single list object includes:

- cleaning and error checking only has to be done once when the `lxy` object is created (e.g., with [xyt.lxy](#))

- having all of the ancillary variables together simplifies the task of passing parameters to other functions as well as saving/retrieving your work to disk
- the nearest neighbor lookup-table (which can take a long time to compute) can be reused
- locations for multiple individuals can be saved in one object

In code examples, objects of class `locoh.lxy` are often indicated by a 'lxy' suffix, and functions that are designed to work on LoCoH-xy objects typically start with 'lxy'.

For a description of the data structure, see the *T-LoCoH Data Classes* vignette.

### See Also

[xyt.lxy](#), [lxy.proj.add](#), [lxy.repair](#), [lxy.subset](#), [lxy.merge](#), [lxy.exp.csv](#), [lxy.exp.kml](#), [lxy.exp.shp](#), [move.lxy](#)

---

<code>lxy.amin.add</code>	<i>Finds the value of a such that p percent of points are a nearest neighbor for at least one hull</i>
---------------------------	--

---

### Description

Finds the value of a such that p percent of points are a nearest neighbor for at least one hull

### Usage

```
lxy.amin.add(lxy, id = NULL, s = NULL, ptp = 0.98, nnn = 2,
  prec = NULL, max.iter = 20, status = TRUE)
```

### Arguments

<code>lxy</code>	A <a href="#">LoCoH-xy</a> object
<code>id</code>	The id value(s) to analyze. If NULL all ids will be used.
<code>s</code>	Value(s) for the s term in the time-scaled-distance equation for point-to-point distance
<code>ptp</code>	The proportion of total points that should be a nearest neighbor for at least one hull (0..1]
<code>nnn</code>	The minimum number of nearest neighbors each point should have (can pass a vector of several values)
<code>prec</code>	A numeric value in map units to which the value of 'a' will be found. If NULL, will default to one-half of the median step length of the entire dataset
<code>max.iter</code>	The maximum number of iterations to try to get within prec of the minimum value
<code>status</code>	Show messages, T/F

## Details

This function finds the value of 'a' (within a specified threshold prec) such that the proportion ptp of all points will be a nearest neighbor for at least one hull. This value of 'a' is intended to be a reasonable lower bound for a home range that includes as many points as desired, but minimizes areas where the individual was not observed. This assumes that duplicate points will be offset by a random amount when creating hullset(s).

Note that the value of 'a' such that ptp of points are nearest neighbors does not mean that ptp points are enclosed. Points can be enclosed by hulls that are not a nearest neighbor of any hull parent point.

---

lxy.anv.add	<i>Add ancillary variable(s) to a LoCoH-xy object</i>
-------------	---

---

## Description

Add ancillary variable(s) to a LoCoH-xy object

## Usage

```
lxy.anv.add(lxy, anv, anv.desc = NULL, overwrite = FALSE)
```

## Arguments

lxy	A <a href="#">LoCoH-xy</a> object
anv	A vector, named list or data frame with the same number of values as points
anv.desc	A character vector of descriptions
overwrite	Whether to overwrite existing variables with the same names (T/F)

## Value

A [LoCoH-xy](#) object

---

lxy.exp.csv	<i>Export coordinates to a csv file</i>
-------------	---

---

## Description

Exports coordinates and date-stamps from a LoCoH-xy object to a csv file

## Usage

```
lxy.exp.csv(lxy, dir = ".", file = NULL, id = NULL,
  file.overwrite = FALSE, quote = FALSE)
```

**Arguments**

lxy	A LoCoH-xy object
file	Filename (with extension .csv)
id	The id(s) of the locations to export
file.overwrite	Overwrite existing file, T/F
quote	Delimit strings in quotes, T/F

**Value**

A data frame containing the coordinates and attribute values of lxy

**Note**

This function will export the id value (i.e., animal name) point-id value (ptid), coordinates, and date stamp (if exists) of a LoCoH-xy object to a csv file

If file is not passed, a filename will be automatically constructed.

---

lxy.exp.kml	<i>Export a lxy object to kml (Google Earth)</i>
-------------	--

---

**Description**

Export a lxy object to a format that can be animated in Google Earth

**Usage**

```
lxy.exp.kml(lxy, file, id = NULL, skip = 1, overwrite = TRUE,
  compress = TRUE, pt.scale = 0.5, col = NULL, show.path = TRUE,
  path.col = NULL, path.opaque = 80, path.lwd = 3)
```

**Arguments**

lxy	A LoCoH-xy object
file	The name of the kml file to be created (relative to the working folder). The .kml extension will be added if needed.
id	The id(s) of the individual(s) to include in the plot
skip	The skip factor to use to reduce the number of points exported (i.e., every nth point exported where n=skip)
overwrite	Whether to overwrite an existing kml file (T/F)
compress	Whether to compress the kml file and create a kmz file (T/F)
pt.scale	The relative size of the placemark symbols
col	The point colors (one per id)
show.path	Whether to include the point path in the kml file (T/F)
path.col	The path colors (one per id), ignored if show.path=FALSE
path.opaque	The opacity of the path (0=transparent, 255=fully opaque), ignored if show.path=FALSE
path.lwd	The width of the path, ignored if show.path=FALSE



**Value**

The file name of the kml/kmz file

**Note**

This function will export the locations in a [LoCoH-xy](#) object to a kml file that can be opened in Google Earth for animation. The kml format is not optimized for large datasets, but this works reasonably well for moderate datasets (e.g., <1000). To create animations of larger datasets, use a skip value > 1 or see [lxy.exp.mov](#).

Adapted from plotKML package

**See Also**

[lxy.exp.mov](#)

---

lxy.exp.mov

---

Create a Quicktime animation from a LoCoH-xy object

---

**Description**

Create a Quicktime animation from a LoCoH-xy object

**Usage**

```
lxy.exp.mov(lxy, id = NULL, all.ids.at.once = TRUE,
  all.ids.col.unique = all.ids.at.once, all.ids.col = NULL,
  all.ids.legend = c("bottomright", "bottom", "bottomleft", "left", "topleft",
    "top", "topright", "right", "center")[5], all.ids.legend.cex = 0.8,
  dt.start = NULL, dt.end = NULL, frame.method = c("auto", "time",
    "location")[1], frame.rtd = "auto", xlim = NULL, ylim = NULL,
  dt.label = TRUE, dt.label.x = NULL, dt.label.y = NULL,
  dt.label.col = "black", dt.label.bg = "gray90", title = NULL,
  title.show = TRUE, axes.show = TRUE, axes.ticks = axes.show,
  axes.titles = FALSE, mar.map = c(0.7 + (if (axes.ticks) 0.9 else 0) + (if
    (axes.titles) 1.3 else 0), 0.5 + (if (axes.ticks) 0.9 else 0) + (if
    (axes.titles) 1.3 else 0), if (title.show) 2.1 else 0.5, 0.5),
  mgp.map = c(0.4 + if (axes.ticks) 1.2 else 0, 0.4, 0),
  col.xys.active = "red", col.xys.background = "gray80",
  cex.xys.active = 1.5, cex.xys.background = 0.5, tz.local = NULL,
  tz.local.check = TRUE, col.by.hour.of.day = FALSE,
  col.hod = colorRampPalette(colors()[c(24, 30, 553, 121, 26, 121, 553, 30,
    24)])(24), width = if (screen.test) 7 else 608, height = NULL,
  max.frames = NULL, png.pointsize = 16 + (width - 480)/80,
  screen.test = FALSE, tmp.dir = NULL, tmp.files.delete = TRUE,
  prompt.continue = TRUE, fn.mov = NULL, fn.mov.dir = getwd(),
  fn.mov.exists = c("auto.increment", "overwrite", "stop", "ask")[1],
  duration = NULL, fps = NULL, skip = NULL, ffmpeg = "ffmpeg.exe",
  create.mov = TRUE, fmt = c("mov", "mp4")[1], info.only = TRUE,
  shp.csv = NULL, layers = NULL, tiff.fn = NULL, tiff.bands = c(3, 2,
    1), tiff.col = gray(0:255/255), tiff.pct = FALSE, tiff.buff = 0,
```

```
tiff.fill.plot = TRUE, bg2png = !is.null(layers),
crop.layers.to.extent = TRUE, date.bar = 0.85, date.bar.bins = 12,
col.db = "darkblue", cex.axis.db = 0.7, beep = FALSE,
report.time = TRUE, status = TRUE)
```

## Arguments

<code>lxy</code>	A <a href="#">LoCoH-xy</a> object
<code>id</code>	The id value(s) to be on the plot
<code>all.ids.at.once</code>	Display all the individual ids simultaneously. If False, an animation will be created for each id. T/F.
<code>all.ids.col.unique</code>	Whether to use unique colors for each individual when plotting multiple individuals simultaneously, T/F
<code>all.ids.col</code>	A named list of color values; the element names must match the name(s) of the ids
<code>all.ids.legend</code>	Where to place a legend showing the color of each id (for an animation showing the movement of multiple individuals simultaneously): 'bottomright', 'bottom', 'bottomleft', 'left', 'topleft', 'top', 'topright', 'right', or 'center'. May also be NULL, in which case the legend will not be displayed. Ignored if <code>ids.legend.bln=FALSE</code> or <code>all.ids.col.unique = FALSE</code> .
<code>all.ids.legend.cex</code>	The character expansion factor for the id legend. See parameter <code>all.ids.legend</code> above.
<code>dt.start</code>	The starting date-time. An object of class <code>POSIXt</code> or one that can be coerced to <code>POSIXt</code> . If NULL, the earliest date-time in the series will be used.
<code>dt.end</code>	The ending date-time. An object of class <code>POSIXt</code> or one that can be coerced to <code>POSIXt</code> . If NULL, the last date-time in the series will be used.
<code>frame.method</code>	How each frame should be defined temporally, "time" - each frame represents a fixed amount of time, "location" each frame is a point in the series
<code>frame.rtd</code>	The real-time duration of each frame (in seconds). If "auto" (default), the lowest median sampling frequency will be used
<code>xlim</code>	A two-element numeric vector for the range of the x-axis (in map units)
<code>ylim</code>	A two-element numeric vector for the range of the y-axis (in map units)
<code>dt.label</code>	Add a label for the date of the frame
<code>dt.label.x</code>	The x-coordinate for the date label (ignored if <code>dt.label=FALSE</code> )
<code>dt.label.y</code>	The y-coordinate for the date label (ignored if <code>dt.label=FALSE</code> )
<code>dt.label.col</code>	A color value/name for the date label (ignored if <code>dt.label=FALSE</code> )
<code>dt.label.bg</code>	The background color for the date label. Set to NA for a transparent background. (ignored if <code>dt.label=FALSE</code> )
<code>title</code>	A title for the map
<code>title.show</code>	Whether to show the title. T/F.
<code>axes.show</code>	Whether to show the axes (ticks, labels and titles). Can be over-written by <code>axes.ticks</code> and <code>axes.titles</code> . T/F.
<code>axes.ticks</code>	Whether to show the tick marks and labels on the axes. T/F.

<code>axes.titles</code>	Whether to show axes titles. T/F.
<code>mar.map</code>	Margin settings for the map, see <a href="#">par</a>
<code>mgp.map</code>	Locations of the axes elements for the map, see <a href="#">par</a>
<code>col.xys.active</code>	Color of the active point. Ignored if <code>col.by.hour.of.day</code> = TRUE.
<code>col.xys.background</code>	Color of the non-active points. To hide non-active points, set this to NA.
<code>cex.xys.active</code>	The character expansion factor of the active point
<code>cex.xys.background</code>	The character expansion factor of non-active points
<code>tz.local</code>	The name of the time zone of the study area
<code>tz.local.check</code>	Check whether <code>tz.local</code> is a valid timezone name (not implemented) T/F.
<code>col.by.hour.of.day</code>	Whether to color the active point by the hour of day (i.e., dark colors at night, orange for day time locations). T/F
<code>col.hod</code>	A vector of 24 color values used to symbolize the color of the active point by the hour of day. Ignored if <code>col.by.hour.of.day</code> = FALSE.
<code>screen.test</code>	Create up to three sample frame(s) on the screen (instead of PNG files)
<code>width</code>	The width of each frame in pixels (if <code>screen.test</code> =FALSE) or inches (if <code>screen.test</code> =TRUE).
<code>height</code>	The height of each frame in pixels (if <code>screen.test</code> =FALSE) or inches (if <code>screen.test</code> =TRUE).
<code>max.frames</code>	The maximum number of frames to produce.
<code>png.pointsize</code>	The pointsize (in pixels) for the PNG image, equivalent to the height or width of a character in pixels (increase to make labels appear larger)
<code>tmp.dir</code>	A directory where temporary PNG files for each frame will be created, character.
<code>tmp.files.delete</code>	Delete the temporary PNG files when done, T/F
<code>prompt.continue</code>	Whether to present a summary of the encoding settings and get user confirmation before continuing, T/F
<code>fn.mov</code>	The path and filename of output file (either *.mov or *.mp4). If NULL a filename will be automatically generated
<code>fn.mov.dir</code>	The directory where the animation will be saved (ignored if a value for <code>fn.mov</code> is passed)
<code>fn.mov.exists</code>	What to do if the animation file already exists: "auto.increment", "overwrite", "stop", or "ask".
<code>duration</code>	The desired duration of the animation (in seconds)
<code>fps</code>	A numeric value for frames per second
<code>skip</code>	Output every nth frame. To include every frame set <code>skip</code> =1. Integer.
<code>ffmpeg</code>	The name of the ffmpeg file. See notes.
<code>create.mov</code>	Whether to actually create the mov file. Set to FALSE preview a few frames without actually encoding them.
<code>fnt</code>	Video format: 'mov' (Quicktime animation codec) or 'mp4' (h.264)
<code>info.only</code>	Only return info
<code>shp.csv</code>	The path and filename of a csv file that contains information about shapefiles, including layer names, file, and symbology.

<code>layers</code>	The name(s) of layers in <code>shp.csv</code> to display in the background. Will be displayed using the symbology in <code>shp.csv</code> . Character vector or comma delimited string.
<code>tiff.fn</code>	The path and name of a GeoTIFF file (e.g., satellite image) that will be displayed in the background. See notes.
<code>tiff.bands</code>	A vector of three integers corresponding to the bands of the GeoTIFF image that will be mapped to the red, green and blue color guns respectively.
<code>tiff.col</code>	A vector of color values for plotting single-band images in the background. Ignored if using three bands.
<code>tiff.pct</code>	Whether or to convert the GeoTIFF to an indexed 256 color RGB image, which may speed up drawing. T/F.
<code>tiff.buff</code>	A numeric buffer distance in map units that the range of the plot will be expanded so the points are not right on the edge of the GeoTIFF.
<code>tiff.fill.plot</code>	Whether to fill the entire plot area with the GeoTIFF. T/F.
<code>bg2png</code>	Save the plot background elements as a static raster image (to improve speed), ignored if <code>screen.test=TRUE</code>
<code>crop.layers.to.extent</code>	Whether to crop the shapefile layers to the view extent (may speed up drawing time)
<code>date.bar</code>	The height of the lower section of the plot to devote to the time bar, in inches. To hide the time bar completely, set <code>date.bar=0</code> .
<code>date.bar.bins</code>	The target number of bins (tick marks + 1) on the time bar (integer)
<code>col.db</code>	A single color value for the date bar axes / tick labels, character
<code>cex.axis.db</code>	Character expansion factor for the labels on the date bar axis.
<code>beep</code>	Beep when done, T/F
<code>report.time</code>	Show the time taken when done, T/F
<code>status</code>	Show progress bar and status messages

### Value

A list with information about each file created. Each element of the list is another list with two elements: `fn` (the full filename) and `dim` (a two-element numeric vector with the frame width and height). If no animation file(s) were created, returns `NULL`.

### Note

This function creates an animation from a `LoCoH-xy` object. There are three general steps in the workflow: 1) design the frame layout, 2) choose values for the speed and duration of the animation, and 3) create the animation.

One will normally want to run the function a few times without actually encoding to tweak the frame design (e.g., where the date label and legend appear). To see what a frame in the output will approximately look like, set `screen.test=TRUE`. See Appendix VI of the T-LoCoH tutorial for further details.

If you have locations for all hours of the day, you can color the active point by the hour-of-day by setting `col.by.hour.of.day=TRUE`. This can help you visualize daily patterns in movement, assuming that the time-stamps of each point are in local time. Locations at night will appear dark, sunrise and sunset reddish, and mid-day points blue (these colors can be adjusted by passing a different value for `col.hod`).

To create the animation, two and only two of the following parameters must be passed: duration, fps, and skip. The third parameter will be computed based on the other two. To include every frame, pass fps, set skip=1, and leave duration out.

Larger values for fps will result in the animation running 'faster'. Values between 10 and 20 often work well; beyond 30 fps the eye can't keep up with the motion. Note if you pass values for fps and duration, an appropriate value for skip will be computed but the final duration of the animation may not be exactly equal to duration because only integer values of skip are allowed.

tmp.dir="." (or another folder), create.mov=FALSE, and tmp.files.delete=FALSE. This will generate a few sample frames as PNG files but not delete them so you can inspect them using an image viewer. Once these look good, create the full animation by setting create.mov=TRUE and max.frames=NULL.

If frame.method is 'auto', the script will use time-based frames when multiple individuals are being animated simultaneously, and location-based frames otherwise.

duration (the duration of the animation in seconds) should not be confused with frame.rtd which is the real-time duration of a single frame in seconds (e.g., frame.rtd=3600 means each frame will represent 1 hour).

If date.bar is too small or too large, you might get a 'margins too large' error. Try values around 1, or hide the date bar completely by setting date.bar=0.

The output animation is encoded in QuickTime format or mp4. The Quicktime file is encoded using Quicktime's 'animation' codec, a lossless format that 'scrubs' well (i.e., you can drag the scroll bar to view frame by frame). The mp4 settings use the h.264 compression algorithm with a constant rate factor of 20 (good quality). In both cases, encoding requires installing the open source encoding program ffmpeg. ffmpeg is a command line program that Linux and Windows users can download from <http://ffmpeg.org/download.html>. Windows users should save the ffmpeg.exe file to the working directory or a directory on Window's path environment variable (e.g., c:\windows). Mac users can download ffmpegX from <http://ffmpegx.com/download.html> but this has not been tested (pass a value to ffmpeg).

If ffmpeg is not available, you can still use this function to generate the individual frames and then use another utility (e.g., ImageMagick, Quicktime Pro) to combine the frames into a video file. For best results use a 'lossless' compression method in the encoding program. To create the individual frames only for encoding with another utility, set tmp.dir="." (the working directory) and tmp.files.delete=FALSE.

If fn.mov.exists = "auto.increment", a two-digit number will be appended to the filename to avoid overwriting an existing file

---

lxy.exp.shp

---

Export a LoCoH-xy to shapefile

---

## Description

Export a LoCoH-xy object to point shapefile

## Usage

```
lxy.exp.shp(lxy, id = NULL, dir = ".", file.base = "", anv = "all",
  idsSeparate = FALSE, autoName = c("long", "short", "none")[1],
  status = TRUE)
```

**Arguments**

lxy	A <a href="#">LoCoH-xy</a> object
id	The name(s) of individuals to export. If NULL all individuals will be exported
dir	The directory where the shapefiles will be placed (use "." for the working directory, and "~" for the home directory)
file.base	The base of the file name without any extension (the script will append to this)
anv	The name of ancillary variable(s) to be exported in the attribute table. Character vector or comma separated string.
idsSeparate	Whether to export the locations for each individual in separate shapefiles. T/F
autoName	The rule for constructing a filename: short (consisting of the id(s) only), long (id + num points), or none
status	Show messages. T/F

**Details**

Filenames are automatically generated, but the user can specify the directory and a base (`file.base`). If a shapefile already exists, a unique filename will be constructed using two-digit numeric extension. For shorter filenames, pass `autoName="short"`.

Use the `anv` argument to specify which ancillary variables (i.e., attribute fields) should be exported. To not export any ancillary variables, pass an empty string.

Only the individual locations and their ancillary values are exported. The nearest neighbor table is not exported.

**Value**

A character vector of the shapefile(s) created

**See Also**

[lxy.exp.kml](#), [lxy.exp.mov](#)

---

lxy.gridanv.add

---

Add ancillary variable(s) to a LoCoH-xy object from a raster

---

**Description**

Add ancillary variable(s) to a LoCoH-xy object from a raster

**Usage**

```
lxy.gridanv.add(lxy, band = 1, dtfn = NULL, fn = NULL, anv.name = NULL,
  anv.desc = NULL, date.match = c("closest", "before")[1])
```

**Arguments**

lxy	LoCoH-xy object
band	Numeric vector of which band(s) to read
dtfn	A two-column data frame containing a date and filename
fn	Filename of a raster
anv.name	The name(s) of the ancillary variable(s) (one per band)
anv.desc	A character vector of descriptions
date.match	How to match dates when dtfn is used: 'closest' or 'before'. See note.

**Value**

A LoCoH-xy object

**Note**

This function will create new ancillary variables and fill it with the value of the pixel at each point. The image should be in the same coordinate system as the points. Current the script does not check this. Any points that fall beyond the edges of the image will be given 'NA' values. If an ancillary variable with the same name already exists, it will be overwritten.

The image / raster specified by fn be in a format supported by the readGDAL function in the rdgal package. Multi-band formats are support. If the image has multiple bands, you can specify which band(s) to read using the band parameter. A separate ancillary variable will be created for each band. You can provide name(s) for the ancillary variable(s) with the anv.name parameter.

You can also pass a time-series of images, and the function will use the image that is the closest match in time for each location. To do this, pass a two column data frame to the parameter dtfn. The first column of dtfn must be a date (one of the POSIXt date-time classes), and the second column must be a valid raster filename. If each image is a time composite, (e.g., two-week NDVI), the date should either be the mid-point or the beginning date. Two options for time matching are available: closest = select the image whose date is closest in time, and before = pick the closest image that comes before the location.

---

lxy.id.new

---

*Change the id value(s) in a LoCoH-xy object*


---

**Description**

Changes the id values in a LoCoH-xy object, with option to save the 'old' id values as an ancillary variable

**Usage**

```
lxy.id.new(lxy, id.new, save.old.id = NULL, dup.dt = c("check", "offset",
  "ignore")[1], dup.dt.offset = 5, dt.int.round.to = 0.1,
  tau.diff.max = 0.02, status = TRUE)
```

**Arguments**

lxy	A <a href="#">LoCoH-xy</a> object
id.new	New value(s) for the id (i.e., name of the individual(s)). Can be either a single character object 'broken_tooth', or factor or character vector of the same length as the number of points
save.old.id	Whether to save the old ids as an ancillary variable. Can be T/F, or the name of a column in the ancillary variables data frame, see Details
dup.dt	What to do if duplicate dt values are encountered in the merged object, see details
dup.dt.offset	The maximum offset in seconds to offset duplicate dt values, used only if dup.dt = "offset"
dt.int.round.to	The proportion of the median sampling frequency that time intervals will be rounded to when computing the frequency table of sampling intervals (no change is made to the time stamps)
tau.diff.max	The maximum deviation from tau (the median delta.t of the entire dataset), expressed as a proportion of tau, that time difference between two points must fall for the distance between those two points to be included in the calculation of the median step length
status	Show status messages. T/F

**Details**

This function can be used to change the id value for each point. Note that the id value is the name of the individual animal or device, as opposed to ptid which is a unique numeric integer for each point.

tau.diff.max exists to eliminate the inclusion of temporal outliers in the computation of the median step length. The time difference between points must be within tau.diff.max of tau for that pair of points to be included in step length calculation.

One can also use this function to change the id (name of the individual), rebuild the rw.params or table of sampling frequencies

**Value**

A [LoCoH-xy](#) object containing the original xys, dt, ptid and ancillary values for each point. Other items and attributes (including the comment) will be recreated or set to NULL

---

lxy.identify.aoi	<i>Identify the points that fall within a bounding box</i>
------------------	--

---

**Description**

Identify the points that fall within a bounding box

**Usage**

```
lxy.identify.aoi(lxy, aoi)
```



**Arguments**

lxy	A LoCoH-xy object
aoi	Area-of-interest, a two-column xy data frame containing the upper left point and the lower right point in the first and second rows respectively.

**Value**

A dataframe containing the ptid values and indices of the points from lxy that fall within aoi

**Note**

This function can be used to identify specific points on a plot. For example, you see a cluster of points on a locoh-xy plot, and want to know when those points were recorded, and so on. To do this, you can use the [aoi](#) function to define an area of interest box on the plot window, and then use [lxy.identify.aoi](#) to grab the indices and ptid values for that cluster of points. You can then use this information as a parameter in [lxy.subset](#) function to create a locoh-xy object containing just that cluster of points.

**See Also**

[aoi](#), [plot.locoh.lxy](#), [lxy.subset](#)

---

lxy.lhs

---

*Create a LoCoH-hullset object from a LoCoH-xy object*


---

**Description**

Creates a LoCoH-hullset object (class "locoh.lhs") containing one or more sets of hulls.

**Usage**

```
lxy.lhs(lxy, id = NULL, s = 0, a = NULL, r = NULL, k = NULL,
       kmin = 0, anv.copy = TRUE, decimal.places = 1, offset.dups = 1,
       velocity.metrics = TRUE, ud = NULL, iso.add = FALSE,
       iso.levels = c(0.1, 0.25, 0.5, 0.75, 0.95), pbo.style = 3, beep = FALSE,
       status = TRUE, save.hulls = TRUE, save.enc.pts = TRUE)
```

**Arguments**

lxy	A <a href="#">LoCoH-xy</a> object
id	The id value(s) of lxy for which hullsets will be created. If NULL all ids will be used.
s	Value(s) for the s term in the time-scaled-distance equation for point-to-point distance. When s=0 time is excluded and TSD is equivalent to Euclidean distance. Numeric vector or comma-separated string. Can also be 'all'
a	Value(s) for the adaptive method. Numeric vector or comma-separated string. a also be the results of the <a href="#">auto.a</a> function, see notes.
r	Value(s) for the r method. Numeric vector or comma separate list of values.
k	Value(s) for the k method. Numeric vector or comma separate list of values.

kmin	A minimum number of nearest neighbors for each parent point regardless of the method used. Set k=2 to ensure that every point is part of a hull. Integer.
decimal.places	The number of decimal places for the k/r/a parameter value used when constructing the name of a hullset. Note this has no effect on hull creation or the precision of values used, only the number of digits that appear in the hullset name.
offset.dups	A number of map units to randomly offset duplicate points. Set offset.dups=0 to ignore duplicate points
anv.copy	Copy the ancillary variables data frame (if exists), T/F
velocity.metrics	Compute the velocity hull metrics
ud	Deprecated (no longer used). Use iso.add instead
iso.add	Whether to also create density isopleths, T/F
iso.levels	Isopleth levels (see also <a href="#">lhs.iso.add</a> ), numeric vector. Ignored if lcodeiso.add=FALSE.
pbo.style	Progress bar style (see pbapply package)
beep	Beep when done. T/F
status	Show messages. T/F
save.hulls	Whether to save the hulls. T/F
save.enc.pts	Whether to save the enclosed points. T/F

### Value

An object of class [locoh.lhs](#)

### Note

This function creates a [LoCoH-hullset](#) object from a [LoCoH-xy](#) object. Other functions allow you to do things with hullsets, including computing additional hull metrics, constructing isopleths, directional routes, generating scatterplots, exporting, etc. Note that before you use this function, nearest neighbors must have already been identified and saved in the input [LoCoH-xy](#) object (see [lxy.nn.add](#)).

If iso.add=TRUE, after the hulls are created the function will create density isopleths using the default sort order (area for *k-method*, number of enclosed points for the *a* and *r-methods*). You can control which isopleth levels are created using the iso.levels argument.

When working with a large dataset where memory limits may affect performance, you can choose to not save the hulls or enclosed points by setting save.hulls=FALSE and save.enc.pts=FALSE. See the vignette on working with large datasets.

### See Also

[xyt.lxy](#) for creating LoCoH-xy objects [lhs.iso.add](#) for adding isopleths [lhs.ellipses.add](#) for computing bounding ellipses [lhs.visit.add](#) for computing time-use metrics [lhs.plot.scatter](#) for creating scatterplots of hull metrics [plot.locoh.lhs](#) for plotting a LoCoH-hullset [hulls](#) for extracting hulls as a SpatialPolygonsDataFrame [lhs.exp.shp](#) to export hulls as shapefile

---

lxy.merge	<i>Merges LoCoH-xy object</i>
-----------	-------------------------------

---

## Description

Merges LoCoH-xy objects together

## Usage

```
lxy.merge(..., pts.flds = "all", save.ptid = FALSE, new.id = NULL,
  save.old.id = NULL, dt.int.round.to = 0.1, tau.diff.max = 0.02)
```

## Arguments

...	LoCoH-xy objects
pts.flds	A character vector of column names to include in the merged object, can also be "all" or NULL
save.ptid	If duplicate ptid values are encountered in the merged object, whether to save the original ptid values as an ancillary variable, T/F
new.id	A new id value for the merged object
save.old.id	A boolean or the name of a ancillary variable where the old id values will be saved
dt.int.round.to	The proportion of the median sampling frequency that time intervals will be rounded to when computing the frequency table of sampling intervals (no change is made to the time stamps)
tau.diff.max	The maximum deviation from tau (the median delta.t of the entire dataset), expressed as a proportion of tau, that time difference between two points must fall for the distance between those two points to be included in the calculation of the median step length

## Details

The coordinate system of the LoCoH-xy objects being merged must be the same. Also each lxy object must have the same data columns present, e.g., dt, col, ancillary variables, etc. To merge a subset of the attribute fields, use the `pts.flds` parameter.

Because LoCoH-xy object must have unique ptid values for each location, if the merged object contains duplicate ptid values new one will be created. If `save.ptid=TRUE`, the 'old' ptid values will be saved as `ptid.orig`.

The merged object will no have nearest neighbors table and the random walk parameters will be recomputed.

## See Also

[lxy.id.new](#)

---

lxy.nn.add	<i>Identifies nearest neighbors for a LoCoH-xy object</i>
------------	---

---

## Description

Identifies nearest neighbors for a LoCoH-xy object

## Usage

```
lxy.nn.add(lxy, id = NULL, ptid = NULL, k = NULL, r = NULL, a = NULL,
  s = NULL, kmin = 0, ptsh = NULL, ptsh.idx = 1,
  nn.exists = c("append", "replace", "skip")[1], time.term = c("vmax",
  "dif")[1], FNN.algorithm = c("cover_tree", "kd_tree", "VR", "CR")[2],
  dec.places = 1, ra.init.samp.size = 30, ra.init.quant = 0.8,
  tct = 0.05, beep = FALSE, status = TRUE)
```

## Arguments

lxy	A LoCoH-xy object
id	The name(s) of individuals to analyze
k	Value for the fixed-k method (identify k nearest neighbors of each point)
r	Value for the fixed-r method (identify all points within radius r)
a	Value for the a method (identify all points within cumulative distance a). Can also be a data frame of parameters returned by the <a href="#">auto.a</a> function.
s	The value for s for the time-scaled distance metric
ptid	A vector of ptid indicating which points should have nearest neighbors identified (can be used to speed up processing during testing)
kmin	A minimum number of neighbors to identify for each point
ptsh	The desired proportion of time-selected hulls [0..1], may also be 'all' if the s-ptsh map has already been computed. See details.
ptsh.idx	The index of the saved ptsh-s table. See also <a href="#">summary.locoh.lhs</a>
nn.exists	What to do if nearest neighbors have already been saved: "append", "replace", or "skip"
time.term	The space-time transformation to use in the TSD distance metric: 'vmax' for the maximum velocity transformation (default) or 'dif' for the diffusion transformation
FNN.algorithm	The algorithm to be used in the get.knnx() function in the package FNN
dec.places	The number of decimal places that rmax and amax should be rounded to
ra.init.samp.size	How many randomly selected points to use to come up with the first guess for how many neighbors to find for each point to satisfy a (a-method only)
ra.init.quant	The proportion of randomly selected points to use for the a value when coming up with an initial guess at the number of neighbors to find for each point to satisfy a (a-method only)
tct	Temporal continuity threshold used when a= <a href="#">auto.a</a>
beep	Beep when done (T/F)
status	Show status messages (T/F)

## Details

When  $s > 0$ , the 'distance' (as defined by the time-scaled distance metric') between two points is a function of their separation in time as well as space.

If ptsh is provided, the script will find a value of 's' such that the proportion of hulls which are time-selected (e.g, consecutive points in time) is equal to ptsh. In this case, no value for s should be passed. If ptsh="all" and the proportion of time selected hulls has already been computed, those values of 's' will be used.

## Value

A LoCoH-xy object object with saved information about each points' nearest neighbors

## See Also

[summary.locoh.lxy](#) to see what nearest neighbors have already been identified [auto.a.lxy.ptsh.add](#)

## Examples

```
# lxy <- lxy.nn.add(lxy, k=10, s=0.01)
```

---

lxy.plot.freq	<i>Plots of sampling frequency</i>
---------------	------------------------------------

---

## Description

Creates plots of the temporal extent and sampling frequency of a LoCoH-xy object

## Usage

```
lxy.plot.freq(lxy, id = NULL, deltat.by.date = F, samp.by.date = F,
  cp = F, cp.min = 0, cp.max = 1, time.unit = c("auto", "sec", "min",
    "hour", "day", "week", "month", "year")[1], xlim = NULL, ylim = NULL,
  ylim.deltat.by.date = c(0, 0.999), sbd.y.jiggle = 0.1, sbd.vline = NULL,
  sbd.vline.trunc.units.to = c("secs", "mins", "hours", "days")[4],
  cex.pts = 0.3, desc = c(0, 1, 3)[2], cex.desc = 0.8,
  col.desc = "darkgreen", title = NULL, title.show = TRUE,
  subtitle = "auto", mar = c(2.8, 2.8, if (title.show) 2.8 else 0.5, 0.5),
  mgp = c(1.8, 0.5, 0), figs.per.page = NULL, status = TRUE,
  panel.num = NULL, panel.num.inside.plot = !title.show, png.dir = NULL,
  png.dir.make = TRUE, png.width = 800, png.height = png.width,
  png.overwrite = TRUE, png.pointsize = 12 + (png.width - 480)/80, ...)
```

## Arguments

lxy	A LoCoH-xy object
id	The id(s) of the individual(s) to include in the plot
deltat.by.date	Produce a plot of the sampling interval by date. T/F
samp.by.date	Produce a scatterplot showing the number of samples by date. T/F
cp	Produce a plot of the cumulative proportion of sampling frequencies. T/F

cp.min	The minimum sampling frequency (expressed as a proportion of tau) shown on the plot
cp.max	The maximum sampling frequency (expressed as a proportion of tau) shown on the plot
time.unit	A unit of time that will be used on the y-axis when plotting frequency by date
xlim	The range for the x-axis. Can be a two-element vector (min, max), NULL (range will be computed from the data), or "same" (same range will be used for all individuals).
ylim	The range for the y-axis.
ylim.deltat.by.date	A two-element numeric vector with values 0..1. Will interpret these values as the first and last percentiles of delta.t to plot
sbd.y.jiggle	An amount to 'jiggle' the y values to better see point density
sbd.vline	An interval of time (in seconds) for which vertical lines will be included on the sampling interval by date plot, to help see the dates
sbd.vline.trunc.units.to	A character "secs", "mins", "hours" or "days" specifying where to draw the first vertical lines. quantiles of delta.t when setting the range of ylim when deltat.by.date=TRUE
cex.pts	The expansion factor for points on the plot
desc	Which side to display automatically generated descriptive text (e.g. caption). 0=none, 1=bottom, 3=top.
cex.desc	The expansion factor for the descriptive text. Numeric value.
col.desc	The color of the descriptive text. Color value.
mar	The plot margins. A four item numeric vector
mgp	The distance away from the edge of the plot for the 1) label, 2) tick marks, and 3) axis line. A three-item numeric vector
figs.per.page	The number of plots per page.
status	Show status messages. T/F
title	The title to be displayed. Character. If NULL a title will be constructed.
title.show	Whether to show the title. T/F
subtitle	The plot subtitle. If 'auto' a default subtitle will be used, or pass an empty string '' to omit. Character
panel.num	A number or letter to display in the upper left hand corner of the plot when the plot will be used as part of a multi-frame graphic (as in publications). Character
panel.num.inside.plot	Whether to display panel.num inside the plot area itself, as opposed to the title area. T/F
png.dir	The directory for a PNG file (filename will be constructed automatically). Ignored if png.fn is passed
png.dir.make	Whether to create png.dir if it doesn't exist. T/F
png.width	The width of the PNG image
png.height	The height of the PNG image
png.overwrite	Whether to overwrite an existing PNG file if it exists. T/F
png.pointsize	The pointsize (in pixels) for the PNG image, equivalent to the height or width of a character in pixels (increase to make labels appear larger)
...	Additional parameters that will be passed to the <a href="#">plot</a> function

**Value**

A list of lists, one for each plot containing the filename (NULL if no png made), the image dimensions (NULL), descriptive text, and id

**Note**

Plotting the cumulative percentage of sampling frequencies (codecp=TRUE) can help determine a threshold below which pairs of points should be considered to be part of the same 'burst' (see [lxy.thin.bursts](#)). For example, if most points are sampled 1 hour apart, but there are a few bursts of points that were only 2 minutes apart, the cumulative percentage of sampling frequencies will reveal a cluster of sampling intervals around 2 minutes and the majority around 1 hour.

Plotting the sampling frequency over time can help identify gaps in the data as well as frequencies for harmonising sampling frequency

**See Also**

[lxy.thin.bursts](#), [lxy.thin.byfreq](#)

---

lxy.plot.mtdr	<i>Plot distribution of the ratio the maximum theoretical distance ratio for nearest neighbors</i>
---------------	--

---

**Description**

Plot distributions of the ratio of maximum theoretical distance times 's' over TSD for all unique pairs of nearest neighbors

**Usage**

```
lxy.plot.mtdr(lxy, s = NULL, k = NULL, a = NULL, r = NULL,
  type = c("mtd.tsd", "mtd.ed")[1], offset.dups = 1, id = NULL,
  show.samp.size = TRUE, outline = FALSE, desc = c(0, 1, 3)[2],
  cex.desc = 0.8, col.desc = "darkgreen", mar = c(3, 3, if (title.show) 3
  else 0.7, 0.5), mgp = c(1.9, 0.5, 0), figs.per.page = NULL,
  title = NULL, title.show = TRUE, title.obj.name = FALSE,
  panel.num = NULL, panel.num.inside.plot = !title.show,
  no.sci.notation = FALSE, png.dir = NULL, png.dir.make = TRUE,
  png.width = 800, png.height = png.width, png.overwrite = TRUE,
  png.pointsize = 12 + (png.width - 480)/80, status = TRUE, ...)
```

**Arguments**

lxy	A <a href="#">LoCoH-xy</a> object
s	The s value(s) of nearest neighbor sets to include in the plot. If NULL, all values will be used
k	A k-value for the number of nearest neighbors around each point to include in the plot
r	A r-value for the number of nearest neighbors around each point to include in the plot

<code>a</code>	A a-value for the number of nearest neighbors around each point to include in the plot
<code>type</code>	Which ratio to plot: maximum theoretical distance over TSD, or maximum theoretical distance over Euclidean distance
<code>offset.dups</code>	An amount in map units for which pairs of points with the same location but different times will be offset so that a TSD or Euclidean distance can be calculated (see also <a href="#">lxy.lhs</a> )
<code>id</code>	The id(s) of the individual(s) to include in the plot
<code>show.samp.size</code>	Whether to display the sample size of the number of unique pairs of points for each value of s, T/F
<code>outline</code>	Show outliers in the box plots T/F
<code>desc</code>	Which side to display automatically generated descriptive text (e.g. caption). 0=none, 1=bottom, 3=top.
<code>cex.desc</code>	The expansion factor for the descriptive text. Numeric value.
<code>col.desc</code>	The color of the descriptive text. Color value.
<code>mar</code>	The plot margins. A four item numeric vector
<code>mgp</code>	The distance away from the edge of the plot for the 1) label, 2) tick marks, and 3) axis line. A three-item numeric vector
<code>figs.per.page</code>	The number of plots per page.
<code>title</code>	The title to be displayed. Character. If NULL a title will be constructed.
<code>title.show</code>	Whether to show the title. T/F
<code>title.obj.name</code>	Whether to add the name of the lxy object to the plot title (ignored if title is passed). T/F
<code>panel.num</code>	A number or letter to display in the upper left hand corner of the plot when the plot will be used as part of a multi-frame graphic (as in publications). Character
<code>panel.num.inside.plot</code>	Whether to display panel.num inside the plot area itself, as opposed to the title area. T/F
<code>no.sci.notation</code>	Whether to avoid the use of scientific notation on labels on the x-axis. T/F
<code>png.dir</code>	The directory for a PNG file (filename will be constructed automatically). Ignored if png.fn is passed
<code>png.dir.make</code>	Whether to create png.dir if it doesn't exist. T/F
<code>png.width</code>	The width of the PNG image
<code>png.height</code>	The height of the PNG image
<code>png.overwrite</code>	Whether to overwrite an existing PNG file if it exists. T/F
<code>png.pointsize</code>	The pointsize (in pixels) for the PNG image, equivalent to the height or width of a character in pixels (increase to make labels appear larger)
<code>status</code>	Display status messages. T/F
<code>...</code>	Additional parameters that will be passed to the <a href="#">plot</a> function



## Details

This function plots the ratio of the maximum theoretical distance the individual could have traveled (multiplied by 's') over the total TSD distance metric for all nearest neighbors for the specified value of k/a/r for different values of s. This ratio will of course never be greater than one because the maximum theoretical distance is one of the terms in TSD. The purpose of this function is to see the effect of s on the relative contribution of the time term in the TSD metric.

This function requires that nearest neighbors already be computed. To identify nearest neighbors for a range of 's' values, use [lxy.ptsh.add](#) (see examples).

## Value

A list of plots created, with one element per id, and each element consisting of another list with elements for the filename (or NULL), filename dimensions in pixels (or NULL), the MTD ratios (as a list with one element for each value of s), descriptive text, and the k/a/r value

## Examples

```
## Identify a range of 's' values from space-selection to time-selection, and find 10 nearest neighbors for ea
# lxy <- lxy.ptsh(lxy, nn=TRUE)
## Plot the ratio of maximum theoretical distance travel to TSD
# lxy.plot.mtdr(lxy)
```

---

lxy.plot.pt2ctr

*Plot distance to centroid over time*


---

## Description

Produces a plot of the distance of each point to the centroid over time to see 'natural' periodicities in the data

## Usage

```
lxy.plot.pt2ctr(lxy, id = NULL, idx = NULL, vline = NULL,
  vline.trunc.units.to = c("secs", "mins", "hours", "days")[4],
  figs.per.page = NULL, ...)
```

## Arguments

lxy	A <a href="#">LoCoH-xy</a> object
id	The id value(s) to include on the plot
idx	An optional vector of indices of points that the plot will be restricted to. See notes
vline	An interval of time (in seconds) for the vertical grid lines
vline.trunc.units.to	A character value: "secs", "mins", "hours", or "days". Determines where the first vertical grid lines will appear
figs.per.page	The number of figures per page
...	Additional parameters that will be passed to the <a href="#">plot</a> function

**Note**

The purpose of this graph is to help see the 'natural' movement cycles in a dataset, when no *a priori* information exists that helps identify the time scale of interest. In T-locoh, defining the timescale of interest helps to select a value for *s*.

To 'zoom in' to a section of the timeline, pass a value for *idx*. For example, if there are 5000 points in the dataset

If a value is passed for *vline*, the first vertical line will coincide with a multiple of `vline.trunc.units.to`. For example, if `vline=3600*24` (24 hours), and `vline.trunc.units.to="days"`, the first vertical line will be the beginning of a day regardless of when the point series started. This can be helpful for seeing cycles of a specific frequency.

**See Also**

[lxy.plot.sfinder](#)

---

lxy.plot.ptsh

---

*Plots the proportion of time-selected hulls over 's'*


---

**Description**

Plots the proportion of time-selected hulls over 's'

**Usage**

```
lxy.plot.ptsh(lxy, id = NULL, ptsh.idx = NULL, use.nn = FALSE, k = NULL,
  r = NULL, a = NULL, slim = NULL, desc = c(0, 1, 3)[2],
  cex.desc = 0.8, col.desc = "darkgreen", title = NULL,
  title.show = TRUE, legend = c("none", "bottomright", "bottom",
  "bottomleft", "left", "topleft", "top", "topright", "right", "center")[2],
  mar = c(3, 3, if (title.show) 2.8 else 0.7, 0.5), mgp = c(1.8, 0.5, 0),
  figs.per.page = NULL, panel.num = NULL,
  panel.num.inside.plot = !title.show, png.dir = NULL,
  png.dir.make = TRUE, png.width = 800, png.height = png.width,
  png.overwrite = TRUE, png.pointsize = 12 + (png.width - 480)/80, ...)
```

**Arguments**

<i>lxy</i>	A <a href="#">LoCoH-xy</a> object
<i>id</i>	The name(s) of individuals to plot
<i>ptsh.idx</i>	The index number of the saved ptsh table to use (ignored if <code>use.nn=TRUE</code> )
<i>use.nn</i>	Whether to create the plot based on nearest neighbor sets (as opposed to saved ptsh tables, see Details). T/F.
<i>k</i>	Value for the <i>k</i> method if creating the plot based on nearest neighbor tables (ignored if <code>use.nn=FALSE</code> )
<i>r</i>	Value for the <i>r</i> method if creating the plot based on nearest neighbor tables (ignored if <code>use.nn=FALSE</code> )
<i>a</i>	Value for the <i>a</i> method if creating the plot based on nearest neighbor tables (ignored if <code>use.nn=FALSE</code> )

slim	The lower and upper bounds for s, two-element numeric vector
desc	Which side to display automatically generated descriptive text (e.g. caption). 0=none, 1=bottom, 3=top.
cex.desc	The expansion factor for the descriptive text. Numeric value.
col.desc	The color of the descriptive text. Color value.
title	The title to be displayed. Character. If NULL a title will be constructed.
title.show	Whether to show the title. T/F.
legend	A character object specifying where to put the legend (ignored when use.nn=TRUE)
mar	The plot margins. A four item numeric vector
mgp	The distance away from the edge of the plot for the 1) label, 2) tick marks, and 3) axis line. A three-item numeric vector
figs.per.page	The number of plots per page.
panel.num	A number or letter to display in the upper left hand corner of the plot when the plot will be used as part of a multi-frame graphic (as in publications). Character
panel.num.inside.plot	Whether to display panel.num inside the plot area itself, as opposed to the title area. T/F
png.dir	The directory for a PNG file (filename will be constructed automatically).
png.dir.make	Whether to create png.dir if it doesn't exist. T/F
png.width	The width of the PNG image
png.height	The height of the PNG image
png.overwrite	Whether to overwrite an existing PNG file if it exists. T/F
png.pointsize	The pointsize (in pixels) for the PNG image, equivalent to the height or width of a character in pixels (increase to make labels appear larger)
...	Additional parameters that will be passed to the <a href="#">plot</a> function

## Details

This function will plot the proportion of total hulls that are 'time-selected'. Time-selected means all of the nearest neighbors are were sampled sequentially in time. This is one end of the spectrum as far as nearest neighbor identification goes, the other being space-selected (i.e., time has no bearing). The s parameter in the TSD equation determines the degree to which time plays a role in the point-to-point 'distance'. When  $s=0$ , time plays no role in TSD and TSD is equivalent to Euclidean distance. As s increases, time plays a bigger and bigger role until eventually nearest neighbor selection is equivalent to selection points based only on their separation in time.

In order to plot the proportion of time-selected hulls (ptsh), nearest neighbors must have already been identified. This can be done in one of two ways. The [lxy.ptsh.add](#) function will compute ptsh for different values of s using a random sample of points (to save time), automatically picking values of s such that ptsh is close to target values provided by the user. Alternately, one can identify nearest neighbors for different values of s using [lxy.nn.add](#) and then run this function with use.nn=TRUE. The main difference between [lxy.ptsh.add](#) and [lxy.nn.add](#) is that [lxy.ptsh.add](#) finds nearest neighbors for a random sample of points, and doesn't actually save the nearest neighbor information for individual points, whereas [lxy.nn.add](#) identifies and saves nearest neighbor information for each and every point.

## Value

A list of lists, one for each plot containing the filename (NULL if no png made), the image dimensions (or NULL), the descriptive text, the id, and a matrix of the values

**See Also**

[lxy.ptsh.add](#), [lxy.nn.add](#)

**Examples**

```
# data(toni.lxy)
# toni.lxy <- lxy.ptsh.add(toni.lxy)
# lxy.plot.ptsh(toni.lxy)
```

---

lxy.plot.sfinder	<i>Plot distributions of 's' such that the vmax/diffusion term in TSD are equivalent to the actual displacement</i>
------------------	---

---

**Description**

This will find the value of s that will result in the time term of TSD being approximately equal to the distance term

**Usage**

```
lxy.plot.sfinder(lxy, id = NULL, delta.t = "auto", delta.t.auto.n = 8,
  delta.t.err = 0.01, outline = FALSE, desc = c(0, 1, 3)[2],
  cex.desc = 0.8, col.desc = "darkgreen", time.term = c("vmax", "dif")[1],
  mar = c(3, 3, if (title.show) 2.8 else 0.7, 0.5), mgp = c(1.8, 0.5, 0),
  figs.per.page = NULL, title = NULL, title.show = TRUE,
  panel.num = NULL, panel.num.inside.plot = !title.show, png.dir = NULL,
  png.dir.make = TRUE, png.width = 800, png.height = png.width,
  png.overwrite = TRUE, png.pointsize = 12 + (png.width - 480)/80, ...)
```

**Arguments**

lxy	A LoCoH-xy object
id	The id(s) of the individual(s) to include in the plot
delta.t	numeric vector of delta.t values (in seconds) for which a s-values will be computed. Can also be 'auto' in which case delta.t.auto.n delta.t values will be computed ranging from 3 * tau to the 80th percent point-to-point interval for the entire dataset
delta.t.auto.n	The number of delta-ts to use when delta.t="auto"
delta.t.err	The proportion of delta.t within which the interval between two points must fall to be included in the plot
outline	Show outliers in the box plots T/F
desc	Which side to display automatically generated descriptive text (e.g. caption). 0=none, 1=bottom, 3=top.
cex.desc	The expansion factor for the descriptive text. Numeric value.
time.term	The type of time term to use ('vmax' is computed as deltaT * maximum possible velocity, while 'dif' computes Gaussian diffusion distance)
col.desc	The color of the descriptive text. Color value.
mar	The plot margins. A four item numeric vector

mgp	The distance away from the edge of the plot for the 1) label, 2) tick marks, and 3) axis line. A three-item numeric vector
figs.per.page	The number of plots per page.
title	The title to be displayed. Character. If NULL a title will be constructed.
title.show	Whether to show the title. T/F.
panel.num	A number or letter to display in the upper left hand corner of the plot when the plot will be used as part of a multi-frame graphic (as in publications). Character
panel.num.inside.plot	Whether to display panel.num inside the plot area itself, as opposed to the title area. T/F
png.dir	The directory for a PNG file (filename will be constructed automatically). Ignored if png.fn is passed
png.dir.make	Whether to create png.dir if it doesn't exist. T/F
png.width	The width of the PNG image
png.height	The height of the PNG image
png.overwrite	Whether to overwrite an existing PNG file if it exists. T/F
png.pointsize	The pointsize (in pixels) for the PNG image, equivalent to the height or width of a character in pixels (increase to make labels appear larger)
...	Additional parameters that will be passed to the <a href="#">plot</a> function

### Value

A list of lists, one for each plot containing the filename (NULL if no png made), the image dimensions (or NULL), the descriptive text, the id, and a list of the svals

---

lxy.plot.tspan	<i>Plot distributions of the time span of nearest neighbors</i>
----------------	---

---

### Description

Plot distributions of the time span of nearest neighbors as a proportion of the median sampling interval

### Usage

```
lxy.plot.tspan(lxy, s = NULL, k = NULL, a = NULL, r = NULL, id = NULL,
  type = c("hist", "boxplot", "mean")[1], outline = FALSE, breaks = 20,
  desc = c(0, 1, 3)[2], cex.desc = 0.8, col.desc = "darkgreen",
  col.hist = "gray80", mar = c(if (type == "hist") 1.5 else 3, 3, if (type
== "hist") 2 else if (title.show) 3 else 0.5, 0.5), mgp = c(1.9, 0.5, 0),
figs.per.page = NULL, title = NULL, title.show = TRUE,
title.obj.name = FALSE, panel.num = NULL,
panel.num.inside.plot = !title.show, panel.num.cex = 2,
no.sci.notation = FALSE, png.dir = NULL, png.dir.make = TRUE,
png.width = 800, png.height = png.width, png.overwrite = TRUE,
png.pointsize = 12 + (png.width - 480)/80, status = TRUE, ...)
```

**Arguments**

<code>lxy</code>	A <a href="#">LoCoH-xy</a> object
<code>s</code>	The <code>s</code> value(s) of nearest neighbor sets to include in the plot. If NULL, all values will be used
<code>k</code>	A <code>k</code> -value for the number of nearest neighbors around each point to include in the plot
<code>r</code>	A <code>r</code> -value for the number of nearest neighbors around each point to include in the plot
<code>a</code>	A <code>a</code> -value for the number of nearest neighbors around each point to include in the plot
<code>id</code>	The <code>id</code> (s) of the individual(s) to include in the plot
<code>type</code>	The type of plot for the time span: 'hist', 'boxplot', or 'mean'
<code>outline</code>	Show outliers in the box plots T/F
<code>breaks</code>	The breaks parameter for a histogram, see <a href="#">hist</a>
<code>desc</code>	Which side to display automatically generated descriptive text (e.g. caption). 0=none, 1=bottom, 3=top.
<code>cex.desc</code>	The expansion factor for the descriptive text. Numeric value.
<code>col.desc</code>	The color of the descriptive text. Color value.
<code>col.hist</code>	The color of the histogram bars. Color value.
<code>mar</code>	The plot margins. A four item numeric vector
<code>mgp</code>	The distance away from the edge of the plot for the 1) label, 2) tick marks, and 3) axis line. A three-item numeric vector
<code>figs.per.page</code>	The number of plots per page.
<code>title</code>	The title to be displayed. Character. If NULL a title will be constructed.
<code>title.show</code>	Whether to show the title. T/F
<code>title.obj.name</code>	Whether to add the name of the <code>lxy</code> object to the plot title (ignored if <code>title</code> is passed). T/F
<code>panel.num</code>	A number or letter to display in the upper left hand corner of the plot when the plot will be used as part of a multi-frame graphic (as in publications). Character
<code>panel.num.inside.plot</code>	Whether to display <code>panel.num</code> inside the plot area itself, as opposed to the title area. Ignored if <code>panel.num</code> is NULL. T/F
<code>panel.num.cex</code>	The expansion factor for the panel number. Ignored if <code>panel.num</code> is NULL.
<code>no.sci.notation</code>	Whether to avoid the use of scientific notation on labels on the x-axis. T/F
<code>png.dir</code>	The directory for a PNG file (filename will be constructed automatically). Ignored if <code>png.fn</code> is passed
<code>png.dir.make</code>	Whether to create <code>png.dir</code> if it doesn't exist. T/F
<code>png.width</code>	The width of the PNG image
<code>png.height</code>	The height of the PNG image
<code>png.overwrite</code>	Whether to overwrite an existing PNG file if it exists. T/F
<code>png.pointsize</code>	The pointsize (in pixels) for the PNG image, equivalent to the height or width of a character in pixels (increase to make labels appear larger)
<code>status</code>	Show progress bar and status messages. T/F
<code>...</code>	Additional parameters that will be passed to the <a href="#">plot</a> function

**Value**

A list of plots created, with one element per id, and each element consisting of another list with elements for the filename (or NULL), filename dimensions in pixels (or NULL), the timespan values (as a list with one element for each value of s), descriptive text, and the k/a/r value

---

lxy.proj.add	<i>Specify the projection system for a LoCoH-xy object</i>
--------------	--

---

**Description**

Specify the projection system for a LoCoH-xy object

**Usage**

```
lxy.proj.add(lxy, proj4string = CRS(as.character(NA)))
```

**Arguments**

lxy	<a href="#">LoCoH-xy</a> object
proj4string	An object of class "CRS", projection string

**Details**

This function will add projection information to a [LoCoH-xy](#) object. Note this does not reproject coordinates, it only adds or changes the projection information. Any existing projection information will be overwritten.

**Value**

A [LoCoH-xy](#) object

**See Also**

[CRS-class](#) for arguments accepted by the [CRS](#) function.

**Examples**

```
# lxy <- lxy.proj.add(lxy, CRS("+proj=utm +south +zone=34"))
```

---

lxy.ptsh.add	<i>Compute s vs. proportion-time-selected hulls table</i>
--------------	---

---

## Description

Identify the values of  $s$  that result in proportion  $p$  of hulls being time-selected

## Usage

```
lxy.ptsh.add(lxy, id = NULL, k = 10, n = 200, samp.idx = NULL,
  sinit = 0.005, ptsh.target = 1:9/10, ptsh.max = 0.98, ptsh.buf = 0.01,
  max.iter = 15, max.loops = 10, time.term = c("vmax", "dif")[1],
  FNN.algorithm = c("cover_tree", "kd_tree", "VR", "CR")[2], use.nn = FALSE,
  plotme = TRUE, save = TRUE, nn.add = FALSE, use.pb.n = 200,
  ptsh.exists = c("replace", "append")[2], beep = FALSE, status = TRUE)
```

## Arguments

lxy	A <a href="#">LoCoH-xy</a> object
id	The name(s) of individuals to analyze
k	Value for the k-method that will be used to create hulls for the sample points (and then used to find the proportion of time-selected hulls)
n	Number of randomly selected points to use as samples (ignored if samp.idx is passed)
samp.idx	A vector of indices of the points to use as samples
sinit	The initial value of $s$ that will be used to start the iteration to find that value of $s$ that produces a proportion of time selected hulls $\geq$ ptsh.max
ptsh.target	A vector of values 0..1 that will serve as the target <i>ptsh</i> values. For each ptsh target, script will try to find a value of $s$ that generates the targeted proportion of time selected hulls (within ptsh.buf)
ptsh.max	The highest value of ptsh for which a value of $s$ will be computed. It is generally recommended that this be less than 1.
ptsh.buf	The level of accuracy or precision to which the found values of 's' return the targetted ptsh.
max.iter	The maximum number of times the script will double sinit in an effort to find the value of $s$ that produces ptsh.max
max.loops	The maximum number of intermediate values of $s$ the script will try when 'zooming' in on the target ptsh levels
time.term	The space-time transformation to use in the TSD distance metric: 'vmax' for the maximum velocity transformation (default) or 'dif' for the diffusion transformation.
FNN.algorithm	The algorithm to be used in the get.knnx() function in the package FNN
use.nn	Whether to use saved nearest neighbor sets for the ptsh computations. T/F
plotme	Whether to plot the ptsh vs. $s$ curve. T/F
save	Whether to save the ptsh-s table in the LoCoH-xy object. T/F



nn.add	Whether to also identify and save nearest neighbor sets for all points for $k=k$ and the values of $s$ identified. T/F
use.pb.n	The sample size (i.e., $n$ ) above which a progress bar will be displayed
ptsh.exists	What to do if there is already a ptsh table in the LoCoH-xy object: 'replace' or 'append'
beep	Whether to beep when done. T/F
status	Display status messages. T/F

### Details

When  $s=0$ , nearest neighbors are space-selected (i.e., point-to-point distance is merely Euclidean distance). For large values of  $s$ , nearest neighbors are effectively time-selected (e.g., temporally contiguous). One approach to selecting a value of  $s$  between these two ends of the spectrum is to pick a  $s$  value that returns an intermediate proportion of time-selected-hulls representing the desired balance between space and time selection (which in turn is shaped by the research question / objective). This function will find the value(s) of  $s$  that produce a desired proportion of time-selected-hulls using a randomly-selected subset of points. A general rule of thumb is to pick a value of  $s$  that results in a proportion of time-selected hulls from 0.4 to 0.6.

The script iteratively tries a variety of  $s$  values until it finds one that produces the desired proportion of time-selected hulls within a tolerance of `ptsh.buf`. If for example the vector of target values `ptsh.target` includes 0.5, and `ptsh.buf = 0.01`, the script will try to find a value of  $s$  that results in a proportion of time-selected hulls between 0.49 and 0.51.

### See Also

[lxy.nn.add](#)

---

<code>lxy.repair</code>	<i>Repair a LoCoH-xy object</i>
-------------------------	---------------------------------

---

### Description

Recreates a [LoCoH-xy](#) object

### Usage

```
lxy.repair(lxy, fix.dup.ptid = FALSE, dt.int.round.to = 0.1,
  tau.diff.max = 0.02)
```

### Arguments

<code>lxy</code>	A LoCoH-xy object
<code>fix.dup.ptid</code>	Generate new ptid value(s) if there are any duplicates
<code>dt.int.round.to</code>	The proportion of the median sampling frequency that time intervals will be rounded to when computing the frequency table of sampling intervals (no change is made to the time stamps)
<code>tau.diff.max</code>	The maximum deviation from tau (the median $\delta.t$ of the entire dataset), expressed as a proportion of tau, that time difference between two points must fall for the distance between those two points to be included in the calculation of the median step length

### Details

This will return a new lxy object containing the original xys, dt, and ptid from lxy. All other items and attributes (including the comment) will be recreated or set to NULL

Set fix.dup.ptid=TRUE to automatically generate new ptid values if any duplicates are detected. If fix.dup.ptid=FALSE and duplicate ptid values are detected, an error message will result.

tau.diff.max exists to eliminate the inclusion of temporal outliers in the computation of the median step length. The time difference between points must be within tau.diff.max of tau for that pair of points to be included in step length calculation.

### Value

A [LoCoH-xy](#) object

### See Also

[xyt.lxy](#), [lxy.id.new](#), [lxy.merge](#)

---

lxy.reproject	<i>Project or reproject coordinates</i>
---------------	---

---

### Description

Project or reproject coordinates of a [LoCoH-xy](#) object

### Usage

```
lxy.reproject(lxy, proj, dt.int.round.to = 0.1, tau.diff.max = 0.02,
  status = TRUE)
```

### Arguments

lxy	A <a href="#">LoCoH-xy</a> object
proj	Projection object of class <a href="#">CRS-class</a>
dt.int.round.to	The proportion of the median sampling frequency that time intervals will be rounded to when computing the frequency table of sampling intervals (no change is made to the time stamps)
tau.diff.max	The maximum deviation from tau (expressed as a proportion of tau) that a point-to-point time difference must fall within for the point-to-point distance to be included in the calculation of the median step length
status	Show status messages (T/F)

### Details

This function will reproject the locations to a new coordinate system. In order for this to work,

- The coordinate system of the current locations must be recorded. To see the current coordinate system, run [summary.locoh.lxy](#). To add projection information (which you need to get from the source of the data), use [lxy.proj.add](#).

- Argument `proj4string` which is an object of class [CRS-class](#) containing information about the new projection system

Note that not all coordinate systems can be projected into all other coordinate systems. If you get an error message like *'non finite transformation detected'*, the coordinate system you are trying to project into may not be compatible with the existing coordinate system.

### See Also

[lxy.proj.add](#), [summary.locoh.lxy](#), [CRS](#)

### Examples

```
## Not run:
lxy.utm <- lxy.reproject(lxy, CRS("+proj=utm +north +zone=18 +ellps=WGS84"))

## End(Not run)
```

---

<code>lxy.save</code>	<i>Save a LoCoH-xy object to disk</i>
-----------------------	---------------------------------------

---

### Description

Saves a LoCoH-xy object to disk, constructing a file name if needed that reflects the contents

### Usage

```
lxy.save(lxy, file = NULL, save.as = NULL, dir = ".", suf = NULL,
         compress = TRUE, auto.num.files = TRUE, width = 2)
```

### Arguments

<code>lxy</code>	A <a href="#">LoCoH-xy</a> object
<code>file</code>	Optional file name
<code>save.as</code>	The name of the saved object, if NULL the original object name will be used
<code>dir</code>	Directory for the output file (relative to the working directory), ignored if <code>file</code> passed
<code>suf</code>	A suffix that will be used in the construction of the file name, ignored if <code>file</code> is passed
<code>compress</code>	Compress file (T/F)
<code>auto.num.files</code>	Use a numeric token as part of the constructed file name to get a unique file name(T/F)
<code>width</code>	The number of digits in <code>auto.num.files</code>

### See Also

[xyt.lxy](#), [lhs.save](#)

---

lxy.subset	<i>Take a subset of a LoCoH-xy object</i>
------------	---

---

## Description

Returns a subset of a LoCoH-xy object based on the id, ptid, idx, or dates

## Usage

```
lxy.subset(lxy, id = NULL, ptid = NULL, idx = NULL, dt.start = NULL,
           dt.end = NULL, dt.int.round.to = 0.1, tau.diff.max = 0.02)
```

## Arguments

lxy	A <a href="#">LoCoH-xy</a> object
id	The id value(s) to include in the subset
ptid	A vector of ptid values for the subset of points
idx	A vector of indices for the subset of points
dt.start	A starting date for the subset. POSIXct object (or something that can be coerced to POSIXct)
dt.end	An ending date for the subset. POSIXct object (or something that can be coerced to POSIXct)
dt.int.round.to	The proportion of the median sampling frequency that time intervals will be rounded to when computing the frequency table of sampling intervals (no change is made to the time stamps)
tau.diff.max	The maximum deviation from tau (expressed as a proportion of tau) that a point-to-point time difference must fall within for the point-to-point distance to be included in the calculation of the median step length

## Value

A [LoCoH-xy](#) object

## See Also

[xyt.lxy](#), [lxy.repair](#), [lxy.thin.byfreq](#), [lxy.thin.bursts](#), [lxy.merge](#)

---

lxy.thin.bursts	<i>Thins out the 'bursts' in a GPS dataset</i>
-----------------	--

---

## Description

Thin out closely-timed bursts of locations

## Usage

```
lxy.thin.bursts(lxy, id = NULL, thresh = NULL, replace = c("mean",
  "median")[2], info.only = FALSE, dt.int.round.to = 0.1,
  tau.diff.max = 0.02, status = TRUE)
```

## Arguments

lxy	A <a href="#">LoCoH-xy</a> object
id	The id value(s) to be thinned
thresh	The threshold for delta.t below which a pair of points is considered to be part of a burst, expressed either as a proportion of the median sampling frequency (0..1) or an absolute unit of time (in seconds)
replace	The burst replacement method ('mean' or 'median')
info.only	Show information about number of bursts (only)
dt.int.round.to	The proportion of the median sampling frequency that time intervals will be rounded to when computing the frequency table of sampling intervals (no change is made to the time stamps)
tau.diff.max	The maximum deviation from tau (the median delta.t of the entire dataset), expressed as a proportion of tau, that time difference between two points must fall for the distance between those two points to be included in the calculation of the median step length
status	Show status messages. T/F

## Value

a [LoCoH-xy](#) object

## Note

This function processes 'bursts' of locations, where a 'burst' is a series of locations captured close together in time. Each group of points in a burst is replaced with a single point. This of course presumes the burst of locations is an artifact of data collection and not desirable.

Many GPS devices have a feature to save 'bursts' of points close together in time (relative to the dominant sampling frequency) The 'burst' feature should not be confused with point averaging, whereby a GPS device internally averages locations for a period of time (e.g., 2 minutes) but saves a single location.

thresh is a value for the sampling interval for identifying which points should be considered part of a burst. thresh can be a proportion of the median sampling frequency (0..1) or an absolute unit of time (in seconds).

To identify whether there are bursts in a [LoCoH-xy](#) dataset, and the sampling frequency of those bursts (i.e., the value

The two replacement methods are `/codereplace="mean"`, in which case a burst of locations is replaced by a single point consisting of the spatial and temporal average of the burst, or `/codereplace="median"` in which case a burst of locations is replaced by the location at the temporal median of the burst. If `/codereplace="mean"`, then any ancillary variables for the 'new' points will be set to NA.

Because this function deletes points, the nearest neighbor lookup table of the [LoCoH-xy](#) object (if any) will be deleted, and the parameters for random walk model will be recomputed.

### See Also

[lxy.plot.freq](#), [lxy.thin.byfreq](#)

---

<code>lxy.thin.byfreq</code>	<i>Delete locations to harmonize the sampling frequency and time duration</i>
------------------------------	---

---

### Description

Standardize the sampling frequency and duration across individuals in a [LoCoH-xy](#) object by deleting points

### Usage

```
lxy.thin.byfreq(lxy, id = NULL, trim.ends = TRUE, dt.start = NULL,
  dt.end = NULL, byfreq = TRUE, samp.freq = "lcm", lcm.round = 120,
  lcm.max.iter = 300, status = TRUE, dt.int.round.to = 0.1,
  tau.diff.max = 0.02)
```

### Arguments

<code>lxy</code>	A <a href="#">LoCoH-xy</a> object
<code>id</code>	The id value(s) to be harmonized
<code>trim.ends</code>	Truncate points from either end of the timeline to achieve a common time window, T/F
<code>dt.start</code>	The starting date-time that all individual trajectories will be truncated to. If NULL, the first date-time that all points have in common will be used.
<code>dt.end</code>	The end date-time that all individual trajectories will be truncated to. If NULL, the last date-time that all points have in common will be used.
<code>byfreq</code>	Delete points to achieve a common sampling frequency ( <code>samp.freq</code> ), T/F
<code>samp.freq</code>	The common time step for the output (in seconds). Can also be set to "lcm", in which case the least common multiple of the median time step of each individual will be computed
<code>lcm.round</code>	When <code>samp.freq="lcm"</code> , the median time step for each individual will be rounded to the nearest interval of <code>lcm.round</code> (in seconds)
<code>lcm.max.iter</code>	The maximum number of iterations to be used in the algorithm that finds the least common multiple of the median time steps

<code>dt.int.round.to</code>	The proportion of the median sampling frequency that time intervals will be rounded to when computing the frequency table of sampling intervals (no change is made to the time stamps)
<code>tau.diff.max</code>	The maximum deviation from tau (the median <code>delta.t</code> of the entire dataset), expressed as a proportion of tau, that time difference between two points must fall for the distance between those two points to be included in the calculation of the median step length
<code>status</code>	Show messages, T/F

**Value**

a [LoCoH-xy](#) object

**Note**

This function processes a [LoCoH-xy](#) object that contains movement data for several individuals, and removes points such that the output contains a fixed start and end date for each individual, as well as an approximately uniform sampling frequency (time step).

Before using this function, you should clean your data of all abnormally short time intervals (e.g., bursts). See [lxy.thin.bursts](#).

If you know the time interval the data was \*supposed\* to be sampled (e.g., every 20 minutes), you should pass that value for `samp.freq` (expressed in seconds). If `samp.freq="1cm"`, the function will automatically compute the common time step for the individuals by taking the least common multiple of the median time steps of each individual. You can deal with noise by rounding the median sampling interval to the value of `1cm.round` (expressed in seconds).

The function [lxy.plot.freq](#) can help you see the 'actual' sampling intervals in the data (set by `.date=TRUE`).

Because this function deletes points, the nearest-neighbors lookup table of the [LoCoH-xy](#) object (if any) will be deleted.

**See Also**

[xyt.lxy](#), [lxy.plot.freq](#), [lxy.thin.bursts](#)

---

matperim

*matperim*


---

**Description**

matperim

**Usage**

```
matperim(mat)
```

**Arguments**

mat	2-column matrix representing the xy coordinates of a closed polygon or line segment
-----	---

**Value**

The length of the perimeter

---

move.lxy	<i>Converts a Move object to a LoCoH-xy object</i>
----------	--

---

**Description**

Converts a Move or MoveStack object from the move package to a LoCoH-xy object

**Usage**

```
move.lxy(move.obj, use.utm = FALSE, xcoord = NULL, ycoord = NULL,
  proj = NULL, anv.flds = NULL, ptid = NULL, del.dup.xyt = TRUE,
  dup.dt.check = TRUE, dt.int.round.to = 0.1, tau.diff.max = 0.02,
  req.id = TRUE, warn.latlong = TRUE, status = TRUE)
```

**Arguments**

move.obj	A object of class Move or MoveStack.
use.utm	Import UTM coordinates if found <i>and</i> all points fall within the same UTM zone (T/F).
xcoord	The name of a field in the data table for the x-coordinates. If NULL, the default coordinates will be used.
ycoord	The name of a field in the data table for the y-coordinates. If NULL, the default coordinates will be used.
proj	Projection object of class <a href="#">CRS-class</a> . Used <i>*only*</i> to specify the coordinate system when the coordinates being imported are coming from the data table
anv.flds	The name of field(s) in the Move object data table that will be imported as ancillary variables. Can also be 'all'.
ptid	The name of a numeric field in the Move object data table containing unique numeric values for each point. If NULL new point id values will be created.
del.dup.xyt	Whether to delete duplicate rows with the same x, y, dt, and id value (T/F).
dup.dt.check	Whether to check to make sure there are no duplicate date values for the same id.
dt.int.round.to	The proportion of the median sampling frequency that time intervals will be rounded to when computing the frequency table of sampling intervals (no change is made to the time stamps).
tau.diff.max	The maximum deviation from tau (expressed as a proportion of tau) that a point-to-point time difference must fall within for the point-to-point distance to be included in the calculation of the median step length.
req.id	Require a value for id (T/F).
warn.latlong	Show a warning message if it detects coordinates may be in latitude-longitude (T/F).
status	Show status messages (T/F).



**Value**

A [LoCoH-xy](#) object

**Note**

Move and MoveStack are formal classes from the *Move* package for storage of movement data for a single (Move) or multiple (MoveStack) individuals. `move.lxy` will convert these objects to [LoCoH-xy](#) objects. To import the attribute data associated with each point (termed ancillary variables in T-LoCoH), set `anv.flds` to the names of the field(s) you wish to import (or `anv.flds="all"`). Note that the Move and MoveStack classes store a lot more metadata than LoCoH-xy objects, including unused records, which are not supported by the [LoCoH-xy](#) data class and are not imported.

**See Also**

[xyt.lxy](#)

**Examples**

```
require(move)
leroy.move <- move(system.file("extdata", "leroy.csv.gz", package="move"))
class(leroy.move)
leroy.lxy <- move.lxy(leroy.move, anv.flds=c("ground.speed", "heading"))
summary(leroy.lxy)
```

---

mvee	<i>mvee</i>
------	-------------

---

**Description**

Computes the minimum volume enclosing ellipsoid around a set of points using the Khachiyan Algorithm. In two dimensions, this is equivalent to the bounding ellipse

**Usage**

```
mvee(xy = NULL, tolerance = 0.005, plotme = FALSE, max.iter = 500,
      shiftxy = TRUE, no.ellipse.val = 1, checks = TRUE)
```

**Arguments**

<code>xy</code>	a two-column data frame containing x and y coordinates. If NULL then a random sample set of 10 points will be generated
<code>tolerance</code>	a tolerance value
<code>plotme</code>	Plot the points and ellipse (T/F)
<code>max.iter</code>	Maximum number of iterations before giving up. If the script tries this number of iterations but still can't get to the tolerance value, it displays an error message and returns NULL
<code>shiftxy</code>	Apply a shift to the coordinates to make them smaller and speed up the matrix calculations, then reverse the shift to the center point of the resulting ellipsoid (T/F)

no.ellipse.val	Determines what the function returns if a mvee can not be found (e.g., if the points are colinear). The default value 1 means a NULL will be returned. A value of 2 means a list of NAs will be returned which is more useful if these values will be stored in a data frame.
checks	Check xy object for the correct data type and remove any duplicate points. T/F

### Value

A list containing the "center form" matrix equation of the ellipse. i.e. a 2x2 matrix "A" and a 2x1 vector "C" representing the center of the ellipse such that:  $(x - C)' A (x - C) \leq 1$  List elements include A - 2x2 matrix C - 2x1 vector of the center coordinates elps.axes.lngth - 2x1 vector whose elements are one-half the lengths of the major and minor axes (i.e., variables a and b in the standard ellipse equation) alpha - angle of rotation

### Note

Adapted by Andy Lyons from Matlab code by Nima Moshtagh. Copyright (c) 2009, Nima Moshtagh <http://www.mathworks.com/matlabcentral/fileexchange/9542> <http://www.mathworks.com/matlabcentral/fileexchange/9542> <http://stackoverflow.com/questions/1768197/bounding-ellipse>

---

n2z	<i>Convert NULL to a zero</i>
-----	-------------------------------

---

### Description

Convert NULL to a zero

### Usage

n2z(x)

### Arguments

x                      Input object

---

plot.locoh.lhs	<i>Plot a LoCoH-hullset object</i>
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---

### Description

Multi-purpose plotting function for a LoCoH-hullset object

## Usage

```
## S3 method for class 'locoh.lhs'
plot(lhs, id = NULL, k = NULL, r = NULL, a = NULL,
     s = NULL, hs.names = NULL, iso = FALSE, rast = FALSE, hulls = FALSE,
     hpp = FALSE, dr = FALSE, nn = FALSE, ellipses = FALSE,
     allpts = FALSE, ptid = NULL, ptid.highlight = TRUE, add = FALSE,
     aoI = NULL, iso.idx = NULL, iso.sort.metric = NULL, iso.legend = TRUE,
     legend.space = 0.25, dr.metric = NULL, dr.thresh.val = NULL,
     dr.thresh.type = NULL, dr.smooth = NULL, lwd.dr = 2, pch.allpts = 16,
     cex.nn = 2, cex.hpp = 0.6, cex.allpts = 0.5, cex.pp = 2,
     cex.axis = 0.8, cex.legend = 0.8, col.hpp = "gray50",
     col.hpp.na = "yellow", col.hulls.border = "gray50", col.hulls.fill = NA,
     col.ellipses = "red", col.allpts = c("auto", "gray")[1],
     col.nn = "black", col.nn.pp = "blue", col.iso.fill = 1,
     col.iso.border = NA, col.iso.scale = TRUE, col.dr = "red",
     hpp.classify = c("none", "hsp", hm.expr(names.only = TRUE, print = FALSE,
     desc = FALSE))[1], hpp.classify.bins = 10, hpp.classify.chop = 0.01,
     hpp.classify.legend = TRUE, hpp.classify.common.scale.discrete = TRUE,
     col.ramp = c("gray10,gray90", "rainbow")[1], col.ramp.bins = 10,
     hsp = NULL, record = FALSE, figs.per.page = 1, same.axes.4all = NULL,
     ufat = TRUE, ufpt = TRUE, axes.show = TRUE, axes.ticks = axes.show,
     axes.titles = axes.show, title = NULL, title.show = TRUE,
     title.inc = c("title", "hs.name", "id", "kar", "features", "hpp.classify",
     "hmap", "ptid")[if (is.null(title)) 3:4 else 1], subtitle.inc = c("title",
     "hs.name", "id", "kar", "features", "hpp.classify", "hmap", "ptid")[if
     (is.null(title)) 5:8 else 0], mar = c(if (axes.titles || axes.ticks) 3.3
     else 0.5, if (axes.titles || axes.ticks) 3.2 else 0.5, if (title.show) 3.2
     else 0.5, 0.5), mgp = c(2, 0.7, 0), lo.save = TRUE,
     lo.margins.set = TRUE, desc = c(NONE <- 0, BOTTOM <- 1, TOP <-
     3)[ifelse(figs.per.page == 1, 2, 1)], cex.desc = 0.8,
     col.desc = "darkgreen", tiff.fn = NULL, tiff.pct = FALSE,
     tiff.bands = c(4, 3, 2), tiff.col = gray(0:255/255), tiff.buff = 0,
     tiff.fill.plot = TRUE, shp.csv = NULL, layers = NULL, png.fn = NULL,
     png.dir = NULL, png.dir.make = TRUE, png.fn.pre = NULL,
     png.fn.mid = NULL, png.fn.suf = NULL, png.fn.incl.d.hs.name = TRUE,
     png.each.plot.separate = TRUE, png.width = 800, png.height = png.width,
     png.pointsize = 12 + (png.width - 480)/80, png.overwrite = TRUE,
     status = TRUE, panel.num = NULL, panel.num.inside.plot = !title.show,
     hmap = NULL, iso.level = NULL, xlim = NULL, ylim = NULL,
     check.ap.value.in.hmparms = TRUE, ...)
```

## Arguments

lhs	A <a href="#">LoCoH-hullset</a> object
id	The names of the individual(s) to include in the plot.
k	The k value(s) of the hullset(s) to include in the plot. Numeric vector or comma-delimited character object.
r	The r value(s) of the hullset(s) to include in the plot. Numeric vector or comma-delimited character object.
a	The a value(s) of the hullset(s) to include in the plot. Numeric vector or comma-delimited character object.

<code>s</code>	The <code>s</code> value(s) of the hullset(s) to include in the plot. Numeric vector or comma-delimited character object.
<code>hs.names</code>	The name(s) of saved hullsets to include in the plot
<code>iso</code>	Whether to display isopleths. T/F.
<code>rast</code>	Whether to display rasterized isopleths. T/F.
<code>hulls</code>	Whether to display hulls. T/F.
<code>hpp</code>	Whether to display hull parent-points. T/F.
<code>dr</code>	Whether to display directional routes. T/F.
<code>nn</code>	Whether to display nearest neighbors (requires that a value for <code>ptid</code> is passed and nearest neighbors were saved when creating the hullset object). T/F.
<code>ellipses</code>	Whether to display bounding ellipses (requires that ellipses were saved when computing ellipses for the hullset). T/F.
<code>allpts</code>	Whether to display all points. T/F.
<code>ptid</code>	One or more <code>ptid</code> (point id) values. A separate plot will be drawn 'zoomed in' to each point indicated. Can also be "auto", in which case <code>ptid</code> will be selected at random. Used primarily to inspect the hull, nearest neighbors, and/or ellipse for specific hulls.
<code>ptid.highlight</code>	Whether to highlight the point specified by <code>ptid</code> . T/F.
<code>add</code>	Whether to add to the existing plot. T/F.
<code>aoi</code>	An area-of-interest object (e.g., box), used to 'zoom in' to specific parts of the plot. <code>aoi</code> objects may be created by the function <code>aoi()</code> .
<code>iso.idx</code>	The index(s) of the isopleths to plot. Use the <code>summary()</code> function to see the indices of the isopleths.
<code>iso.sort.metric</code>	The sort metric(s) of the isopleths that will be displayed. Character.
<code>iso.legend</code>	Whether to include a legend for the isopleths.
<code>legend.space</code>	An expansion factor for the x-axis that will be used to make room for the legend.
<code>dr.metric</code>	The name of the metric for the directional routes to be displayed (acts to filter on which directional routes are displayed). See <a href="#">lhs.dr.add</a>
<code>dr.thresh.val</code>	The threshold value for the directional routes to be displayed (acts to filter on which directional routes are displayed). See <a href="#">lhs.dr.add</a>
<code>dr.thresh.type</code>	The threshold type for the directional routes to be displayed (acts to filter on which directional routes are displayed). See <a href="#">lhs.dr.add</a>
<code>dr.smooth</code>	The smoothing factor for the directional routes to be displayed (acts to filter on which directional routes are displayed). See <a href="#">lhs.dr.add</a>
<code>lwd.dr</code>	The line width of directional routes.
<code>pch.allpts</code>	The plot character for all points
<code>cex.nn</code>	The expansion factor for nearest neighbor points.
<code>cex.hpp</code>	The expansion factor for hull parent-points.
<code>cex.allpts</code>	The expansion factor for all points.
<code>cex.pp</code>	The expansion factor for the parent point.
<code>cex.axis</code>	The expansion factor for axis labels.
<code>cex.legend</code>	The expansion factor for the legend.

col.hpp	The color of hull parent points. Either a single color value or a vector of color values of the same length as the number of hulls.
col.hpp.na	The color of hull parent points that have no value of the metric specified by hpp.classify. Single color value.
col.hulls.border	The outline color of hulls. Either a single color value or a vector of color values of the same length as the number of hulls.
col.hulls.fill	The fill color of hulls.
col.ellipses	The outline color of bounding ellipses. Either a single color value or a vector of color values of the same length as the number of hulls.
col.allpts	The color of all points. Single color value.
col.nn	The color of nearest neighbors. Single color value.
col.nn.pp	The color of parent-points. Used only when ptid is passed. Single color value.
col.iso.fill	A number corresponding to a preset color ramp that will be used to display the isopleths. 1=red to blue, 2=yellow to red, 3=blue to red, 4=red to yellow. Alternately, a character vector of color values of the same length as the number of isopleth levels.
col.iso.border	Color value of the isopleth border. Color value (use NA for no border).
col.iso.scale	When plotting a subset of isopleths (by ecc or par), whether to scale the isopleth colors to the values of the subset. Not being used.
col.dr	The color of directional routes.
hpp.classify	How to classify hull parent points. May be "none", the name of a hull metric, or "hsp" (hull scatterplot)
hpp.classify.common.scale.discrete	Whether to classify hull parent points using a common scale when there are multiple maps produced. Ignored if hpp.classify is 'none' or 'hsp'. T/F.
hpp.classify.bins	The number of equal-interval bins of the hull metric value to create when classifying hull parent-points by a hull metric.
hpp.classify.chop	The proportion of hull parent-points at the tails of the distribution to remove when calculating the bins.
hpp.classify.legend	Whether to include a legend for the classification of hull parent-points. T/F.
col.ramp	The first and last values of the color ramp to use when classifying hull parent-points by a hull metric. Two-item vector with color values.
col.ramp.bins	The number of bins in the color ramp when classifying hull parent-points by a hull-metric
hsp	The hull scatterplot object to use when displaying hull parent points by a hull scatterplot. May be a hull scatterplot object or the index of a hull scatterplot saved in the hullset.
record	Whether to open a new plot window and turn on recording. T/F.
figs.per.page	The number of plots per page.
same.axes.4all	Whether to use the same axes ranges for all plots. T/F.
ufat	Whether to use user-friendly-axis-titles. T/F.

<code>ufipt</code>	Whether to use user-friendly-isopleth-plot-titles. T/F.
<code>axes.show</code>	Whether to show the axes. T/F.
<code>axes.ticks</code>	Whether to show the tick marks and labels on the axes. T/F.
<code>axes.titles</code>	Whether to show axes titles. T/F.
<code>title</code>	The title to be displayed. Character. If NULL a title will be constructed.
<code>title.show</code>	Whether to show the title. T/F.
<code>title.inc</code>	The element(s) to include in the title (ignored if <code>title</code> is passed)
<code>subtitle.inc</code>	The element(s) to include in the subtitle (ignored if <code>title</code> is passed)
<code>mar</code>	The plot margins. A four item numeric vector.
<code>mgp</code>	The distance away from the edge of the plot for the 1) label, 2) tick marks, and 3) axis line. A three-item numeric vector.
<code>lo.save</code>	Whether to save and reset the plot device margin settings (some wrapper functions that call this function don't want device settings reset). T/F.
<code>lo.margins.set</code>	Whether to save and reset the plot device margin settings (some wrapper functions that call this function don't want device settings reset). T/F.
<code>desc</code>	Which side to display automatically generated descriptive text (e.g. caption). 0=none, 1=bottom, 3=top.
<code>cex.desc</code>	The expansion factor for the descriptive text. Numeric value.
<code>col.desc</code>	The color of the descriptive text. Color value.
<code>tiff.fn</code>	The path and name of a GeoTIFF file (e.g., satellite image) that will be displayed in the background. See notes.
<code>tiff.pct</code>	Whether or to convert the GeoTIFF to an indexed 256 color RGB image, which may speed up drawing. T/F.
<code>tiff.bands</code>	A vector of exactly one (for a single band image) or exactly three integers corresponding to the bands of the GeoTIFF image that will be mapped to the red, green and blue color guns respectively,. Ignored if <code>tiff.fn</code> only contains one band.
<code>tiff.col</code>	A vector of color values for plotting single-band images in the background. Ignored if using three bands.
<code>tiff.buff</code>	A numeric buffer distance that the range of the plot will be expanded so the points are not right on the edge of the GeoTIFF.
<code>tiff.fill.plot</code>	Whether to fill the entire plot area with the GeoTIFF. T/F.
<code>shp.csv</code>	The path and filename of a csv file that contains information about shapefiles, including layer names, file, and symbology.
<code>layers</code>	The name(s) of layers in <code>shp.csv</code> to display in the background. Will be displayed using the symbology in <code>shp.csv</code> . Character vector or comma delimited string
<code>png.fn</code>	The path and name of the PNG file to create (instead of displaying in a plot window)
<code>png.dir</code>	The directory for a PNG file (filename will be constructed automatically). Ignored if <code>png.fn</code> is passed
<code>png.dir.make</code>	Whether to create <code>png.dir</code> if it doesn't exist. T/F
<code>png.fn.pre</code>	A prefix that will be used in the construction of the PNG filename. Ignored if <code>png.fn</code> is passed
<code>png.fn.mid</code>	A mid-fix that will be used in the construction of the PNG filename. Ignored if <code>png.fn</code> is passed

png.fn.suf	A suffix that will be used in the construction of the PNG filename. Ignored if png.fn is passed
png.fn.incl.d.hs.name	Whether to include the hullset name as part of the PNG filename. (T/F). Ignored if png.fn is passed
png.each.plot.separate	Whether to make each plot in a separate plot / PNG. T/F. Ignored if png.fn is passed
png.width	The width of the PNG image. Ignored if png.fn is passed
png.height	The height of the PNG image. Ignored if png.fn is passed
png.pointsize	The pointsize (in pixels) for the PNG image (increase to make labels appear larger). Equivalent to the height or width of a character in pixels.
png.overwrite	Whether to overwrite an existing PNG file if it exists. T/F
status	Whether to show messages. T/F
panel.num	A number or letter to display in the upper left hand corner of the plot when the plot will be used as part of a multi-frame graphic (as in publications). Character
panel.num.inside.plot	Whether to display panel.num inside the plot area itself, as opposed to the title area
hmap	A named list of hull metric auxillary parameters, the name of each list element is the name of the variable
iso.level	The isopleth levels to plot, numeric vector
xlim	The lower and upper limit of the x-axis, two-element numeric vector
ylim	The lower and upper limit of the y-axis, two-element numeric vector
check.ap.value.in.hmparms	Whether to check if hull metrics exist for the auxillary parameters passed. T/F
...	Other parameters, including any auxillary parameters required by certain hull metrics

### Value

If png.fn or png.dir is passed, a list object where each element is a three-element list of the properties of the PNG file(s) created: \$fn, \$dim, \$desc. Otherwise, returns a vector of desc

### Note

This is a multi-purpose plotting function for [LoCoH-hullset](#) objects. You specify which objects in the hullset to include on the plot by passing parameters.

All hullsets have hulls and hull parent points that can be plotted (e.g., `codehulls=TRUE`, `hpp=TRUE`). All hullsets also have the original locations saved and these can be added to the plot by setting `allpts=TRUE`. Depending how the hullset was created (see [lxy.lhs](#)), some hullsets may also have the nearest neighbors for each parent point saved, which can also be plotted (`nn=TRUE`).

Hullsets for which have bounding ellipses ([lhs.ellipses.add](#)) or directional routes ([lhs.dr.add](#)) have been computed can have these elements plotted by setting `ellipses=TRUE`) and `dr=TRUE`) respectively. If isopleths have been constructed ([lhs.iso.add](#)), these can be plotted by setting `iso=TRUE`. If you only want to plot some of the isopleths, you can specify which one(s) to plot with the parameters `iso.idx` (the index(s) of the isopleths to plot, see [summary.locoh.lhs](#)) or `iso.sort.metric`.

The plot title can be manually set with the title parameter. If no value for title is passed, a title will be constructed. You can define what elements will go in the title and subtitle (i.e., the second line of the title) with the title.inc and subtitle.inc parameters. To omit the plot title completely, set title.show=FALSE.

tiff.fn, tiff.pct, tiff.buff, tiff.bands, tiff.col, and tiff.fill.plot control the display of a GeoTIFF image in the plot background. The GeoTIFF image must be georeferenced in the same coordinate system as the locoh-hullset object, and the pixel values must be 'prestretched' for display. tiff.bands controls which bands in the TIFF file will be displayed using the red, green, and blue color guns if using three bands. In a Landsat TM image, for example, the first four bands are blue, green, red, and infrared. To display a TM GeoTIFF image as 'natural colors', you would set tiff.bands=c(3,2,1). If tiff.pct=T, the script will create a indexed 256-color version of the image, which may result in quicker drawing time particularly if several plots are being drawn. If using a single-band, for example a DEM or classified image, the colors are set by tiff.col. tiff.buff can be used to expand the range of values on the x and y axis so that you see a bit of the background image beyond the extent of the points.

---

plot.locoh.lxy

---

*Plot LoCoH-xy object*


---

## Description

Multi-purpose plotting function for a LoCoH-xy object

## Usage

```
## S3 method for class 'locoh.lxy'
plot(lxy, id = NULL, cex = 0.8, show.start = TRUE,
     show.end = TRUE, col = c("auto", "gray80")[1], connect.dots = TRUE,
     overlay = FALSE, status = TRUE, title = NULL, title.show = TRUE,
     axes.show = TRUE, axes.titles = axes.show, axes.ticks = axes.show,
     mar = c(if (axes.titles || axes.ticks) 3.3 else 0.5, if (axes.titles ||
     axes.ticks) 3.2 else 0.5, if (title.show) 3.2 else 0.5, 0.5), mgp = c(2,
     0.7, 0), lo.save = TRUE, panel.num = NULL,
     panel.num.inside.plot = !title.show, png.fn = NULL, png.dir = NULL,
     png.dir.make = TRUE, png.width = 800, png.height = png.width,
     png.overwrite = TRUE, png.pointsize = 12 + (png.width - 480)/80,
     tiff.fn = NULL, tiff.bands = c(3, 2, 1), tiff.col = gray(0:255/255),
     tiff.pct = FALSE, tiff.buff = 0, tiff.fill.plot = TRUE, layers = NULL,
     shp.csv = NULL, xlim = NULL, ylim = NULL, legend = NULL, ...)
```

## Arguments

lxy	A <a href="#">LoCoH-xy</a> object
id	A vector of the id value(s) to plot
cex	Character expansion factor for the points
show.start	Whether to highlight the starting location (if time stamps are present) in green. T/F.
show.end	Whether to highlight the end location (if time stamps are present) in red. T/F.
col	A single value or vector of color values. Can also be 'auto' in which case the colors saved in lxy will be used (which are rainbow by default).



connect.dots	Whether to draw line segments between consecutive locations. T/F.
overlay	Whether to overlay the plots of all individuals in lxy on one pair of axes (map). T/F.
status	Display status messages. T/F.
title	The title to be displayed. Character. If NULL a title will be constructed.
title.show	Whether to show the title. T/F.
axes.show	Whether to show the axes. T/F.
axes.titles	Whether to show axes titles. T/F.
axes.ticks	Whether to show the tick marks and labels on the axes. T/F.
mar	The plot margins. A four item numeric vector
mgp	The distance away from the edge of the plot for the 1) label, 2) tick marks, and 3) axis line. A three-item numeric vector
lo.save	Whether to save and reset the plot device margin settings (some wrapper functions that call this function don't want device settings reset). T/F.
panel.num	A number or letter to display in the upper left hand corner of the plot when the plot will be used as part of a multi-frame graphic (as in publications). Character
panel.num.inside.plot	Whether to display panel.num inside the plot area itself, as opposed to the title area. T/F
png.fn	The path and name of the PNG file to create (instead of displaying in a plot window)
png.dir	The directory for a PNG file (filename will be constructed automatically). Ignored if png.fn is passed
png.dir.make	Whether to create png.dir if it doesn't exist. T/F
png.width	The width of the PNG image
png.height	The height of the PNG image
png.overwrite	Whether to overwrite an existing PNG file if it exists. T/F
png.pointsize	The pointsize (in pixels) for the PNG image, equivalent to the height or width of a character in pixels (increase to make labels appear larger)
tiff.fn	The path and name of a GeoTIFF file (e.g., satellite image) that will be displayed in the background. See notes.
tiff.bands	A vector of three integers corresponding to the bands of the GeoTIFF image that will be mapped to the red, green and blue color guns respectively.
tiff.col	A vector of color values for plotting single-band images in the background. Ignored if using three bands.
tiff.pct	Whether or to convert the GeoTIFF to an indexed 256 color RGB image, which may speed up drawing. T/F.
tiff.buff	A numeric buffer distance in map units that the range of the plot will be expanded so the points are not right on the edge of the GeoTIFF.
tiff.fill.plot	Whether to fill the entire plot area with the GeoTIFF. T/F.
layers	The name(s) of layers in shp.csv to display in the background. Will be displayed using the symbology in shp.csv. Character vector or comma delimited string
shp.csv	The path and filename of a csv file that contains information about shapefiles, including layer names, file, and symbology.

xlim	The lower and upper limit of the x-axis, two-element numeric vector
ylim	The lower and upper limit of the y-axis, two-element numeric vector
legend	One of the following keywords specifying where to put a legend when overlaying the locations of multiple individuals: bottomright, bottom, bottomleft, left, topleft, top, topright, right or center
...	Additional parameters that will be passed to the <a href="#">plot</a> function

### Details

This is a multi-purpose plotting function for a LoCoH-xy object, with several custom arguments useful for plotting a movement trajectory. You may also extract the SpatialPointsDataFrame element of the LoCoH-xy object directly by referencing the pts element (e.g., x\$pts, where x is a LoCoH-xy object).

### See Also

Vignette on T-LoCoH data classes, Tutorial Vignette (for details on displaying GIS layers and raster images in the background)

---

secs.fmt

*Format a number of seconds as minutes, hours or days*

---

### Description

Returns a formatted version of a time value

### Usage

```
secs.fmt(secsVal, round.pos = 1)
```

### Arguments

secsVal	a time value (in seconds)
round.pos	The number of decimal places in the formatted string

### Value

a character string

### Note

This function only works with a single value. To format a multiple time values stored in a vector, use `sapply` or `lapply`

Based on the value of secsVal, the formatted version will be converted to days, hours, minutes, or seconds

---

square	<i>square</i>
--------	---------------

---

**Description**

square

**Usage**

square(x)

**Arguments**

x	Input object
---	--------------

---

strSplitAtChar	<i>Split a character object into multiple lines</i>
----------------	---

---

**Description**

Split a character object at a specific character to wrap to multiple lines for plotting

**Usage**

```
strSplitAtChar(object, size, char = ".", separator = paste("\n", char, sep = ""), ...)
```

**Arguments**

object	Input character vector (should be length 1)
size	The maximum number of characters in one piece
char	The character to split the input string at
separator	The character to use as a separator in the returned object
...	Other arguments (unused)

**Details**

This will take a long character object and split it into pieces at character `char` such that the length of each piece is  $\leq$  `size`. If `separator` is passed, the individual pieces will be concatenated using the `separator` character and returned as a character vector of length 1. Otherwise each piece will be returned as a separate element of a character vector.

---

strTrim	<i>Trim leading and/or trailing blanks from a character object</i>
---------	--

---

**Description**

Trim leading and/or trailing blanks from a character object

**Usage**

```
strTrim(str, side = "both")
```

**Arguments**

str	Input string
side	Which side(s) of the input string to trim: 'left', 'right', or 'both'

---

summary.locoh.lhs	<i>Show summary of a LoCoH-hullset</i>
-------------------	--

---

**Description**

Displays a summary of a LoCoH-hullset object

**Usage**

```
## S3 method for class 'locoh.lhs'
summary(lhs, file = "", id = NULL, k = NULL,
  r = NULL, a = NULL, s = NULL, hs.names = NULL, iso.details = FALSE,
  hsp.details = FALSE, desc = FALSE, compact = FALSE, ...)
```

**Arguments**

lhs	A <a href="#">LoCoH-hullset</a> object
file	A file name
id	The name(s) of individuals to summarize
k	The k value of hullsets to summarize
r	The r value of hullsets to summarize
a	The a value of hullsets to summarize
s	The s value of hullsets to summarize
hs.names	The name(s) of hullsets to summarize
iso.details	Display details of the isopleths. T/F
hsp.details	Display details about saved hull scatterplots. T/F
desc	Display the hullset description. T/F
compact	Use a compact format. T/F
...	Other arguments

---

summary.locoh.lxy	<i>Show a summary of a LoCoH-xy object</i>
-------------------	--

---

**Description**

Prints a summary of a locoh xy object (set of locations)

**Usage**

```
## S3 method for class 'locoh.lxy'
summary(lxy, file = "", dt.int = FALSE,
        round.coords = 1, ptsh = FALSE, ...)
```

**Arguments**

lxy	A <a href="#">LoCoH-xy</a> object
file	A file name where the results will be saved
dt.int	Whether to show a summary of the sampling interval
round.coords	The number of digits to display for the coordinates of the spatial extent
ptsh	Show table of 's' and proportion of time selected hulls values (if available)
...	Other arguments

**See Also**

[lxy.plot.freq](#), [lxy.ptsh.add](#)

---

textContrastColor	<i>Choose a white or black plot background for a particular color of text</i>
-------------------	---

---

**Description**

Choose a white or black plot background for a particular color of text

**Usage**

```
textContrastColor(x, ...)
```

**Arguments**

x	A color value
...	Other arguments (not used)

---

toni	<i>Tracking data for one buffalo in South Africa</i>
------	--

---

### Description

This dataframe contains the locations from a GPS collar fitted to a buffalo in Kruger National Park, South Africa in 2005-06.

### Usage

```
data(toni)
```

### Format

A dataframe with 6371 rows. Columns are id, lat, long, and timestamp (in UTC). Datum presumed to be WGS84.

### Source

MoveBank <http://www.movebank.org>

Name: Kruger African Buffalo, GPS tracking, South Africa

Acknowledgements: Collection of Kruger Park Buffalo data funded by NSF Grant DEB-0090323 to Wayne M. Getz

Principal Investigator Name: Paul Cross

### Examples

```
data(toni)
head(toni)
plot(toni[,2:3])
```

---

tsd.zvals	<i>Space-time transformation term for TSD</i>
-----------	---

---

### Description

Returns the values for the "time axis" of the TSD distance metric

### Usage

```
tsd.zvals(delta.t, sVal, type, d.bar = NULL, tau = NULL, vmax = NULL)
```

**Arguments**

<code>delta.t</code>	is the time difference in seconds
<code>sVal</code>	is the value of S
<code>type</code>	The type of space-time transformation: 'vamax' for the maximum velocity transformation, or 'dif' for diffusion
<code>d.bar</code>	is the median step length (for the entire dataset)
<code>tau</code>	is the median sampling interval (for the entire dataset)
<code>vmax</code>	The maximum velocity parameter (used only when <code>type='vmax'</code> )

---

<code>xyt.lxy</code>	<i>Create a LoCoH-xy object</i>
----------------------	---------------------------------

---

**Description**

Creates a [LoCoH-xy](#) object from a set of location data

**Usage**

```
xyt.lxy(xy, dt = NULL, tz = NULL, id = NULL, ptid = NULL,
        proj4string = CRS(as.character(NA)), anv = NULL, anv.desc = NULL,
        col = NULL, del.dup.xyt = TRUE, dup.dt.check = TRUE,
        show.dup.dt = FALSE, dt.int.round.to = 0.1, tau.diff.max = 0.02,
        req.id = TRUE, warn.latlong = TRUE, status = TRUE)
```

**Arguments**

<code>xy</code>	A two-column matrix or data frame containing the xy coordinates of the points
<code>dt</code>	Optional vector of date-time values (either POSIXct objects or objects that can be coerced to POSIXct)
<code>id</code>	Optional character vector or factor containing the name(s) of the individual(s) of each location.
<code>ptid</code>	Optional integer vector of point id values
<code>proj4string</code>	Projection string object of class <a href="#">CRS-class</a>
<code>anv</code>	Optional ancillary variables for each point (data frame with same number of records as xy)
<code>anv.desc</code>	Optional character vector with descriptions of the ancillary variables (in the same order as they appear in anv)
<code>tz</code>	The name of the time zone that will be assigned if not explicit in dt.
<code>del.dup.xyt</code>	Whether to delete duplicate rows with the same x, y, dt, and id value. (T/F)
<code>dup.dt.check</code>	Whether to check to make sure there are no duplicate date values for the same id
<code>show.dup.dt</code>	Whether to show duplicate time stamps (if found) (T/F)
<code>col</code>	Optional vector of color values (one for each point), or a single color value

<code>dt.int.round.to</code>	The proportion of the median sampling frequency that time intervals will be rounded to when computing the frequency table of sampling intervals (no change is made to the time stamps)
<code>tau.diff.max</code>	The maximum deviation from the median sampling interval ( <code>tau</code> ), expressed as a proportion of the median sampling interval, see details
<code>req.id</code>	Require a value for <code>id</code> (T/F)
<code>warn.latlong</code>	Show a warning message if coordinates appear to be in geographic coordinates (T/F)
<code>status</code>	Show status messages (T/F)

## Details

At a minimum, a [LoCoH-xy](#) object contains a set of points. It can also contain date-time values for each point, the name of the individual(s) associated with each point, and a nearest neighbors lookup table.

Importing date-time values into R is often a painful process. You may want to try exporting your date-time values from your spreadsheet or database as a text field that is formatted in a style that R will recognize as a valid time. To see a date-time format that R will be able to convert to a date-time object (class *POSIXct*), type `Sys.time()` at the console. Once you've noted the date-time format recognized by R, try to format the date field in your spreadsheet or database in a similar manner, then export the values in a text format (e.g., csv). For example in MS Access, you can use the `format()` function as part of a query to format a date field, e.g. `format(PointDate, "yyyy-mm-dd hh:nn:ss")`. In Excel, you can go to Format - Cells and give the date-time cells a custom format `yyyy-mm-dd hh:mm:ss`. See the Appendices of the T-LoCoH tutorial for more examples.

The time zone parameter `tz` is optional. If `tz` is omitted, R will try to get the timezone from `dt`, and if there is no timezone specified set it to UTC. Valid timezone names are to some extent OS specific, see [http://en.wikipedia.org/wiki/List\\_of\\_zoneinfo\\_time\\_zones](http://en.wikipedia.org/wiki/List_of_zoneinfo_time_zones). If `dt` is of class *POSIXct* (which stores the time zone) and a different value for `tz` is passed, a prompt will ask whether date values should be converted.

By default, if the function finds two or more locations with the same time stamp for the same id (individual), an error will be triggered. This usually results from a data processing error (e.g., duplication of a row), or a conversion problem or rounding issue in the time stamp values. To see which records have duplicate time stamps, pass `show.dup.dt=TRUE`. To disable the checking for duplicate time stamps, pass `dup.dt.check=TRUE`. Duplicate locations are allowed, as are duplicate time stamps for different individuals.

`ptid` is an optional vector of numeric id values (i.e., primary key) for each location. When present, point id values of parent points will be saved in the data table of hulls. This enables linking the outputs of `tlocoh` (hulls and hull metrics) with other data and/or other software tools. See also [lhs.exp.csv](#) and [lhs.exp.shp](#). `ptid` should not be confused with `id`, which is a character vector of the name(s) of the individuals in the dataset.

`tau.diff.max` controls which consecutive pairs of points will be used to compute the median step length ( $\bar{d}$ ) and the maximum observed speed ( $v_{\max}$ ). Pairs of points whose sampling interval (i.e., time difference) deviates from the median by more than `tau.diff.max` will be excluded from this computation (presumably because there was a drop-out in the data, and the distance between those pairs of locations does not represent the characteristic movement pattern). However for data that have a wide distribution of sampling intervals (e.g., manually collected radio telemetry locations), this filtering may result in an insufficient number of pairs. In this case, increase `tau.diff.max` or set it to zero to disable filtering completely.

Other variables, including environmental variables or biometric variables can be brought in using the `anv` parameter. These variables can then be used in hull metrics.



**Value**

A object of class [locoh.lxy](#)

**See Also**

[lxy.nn.add](#), [lxy.repair](#), [lxy.subset](#)

**Examples**

```
## Not run:
#Create an unbounded random walk as a sample dataset
n <- 500; stepsize <- 5
theta <- runif(n, min=0, max=2*pi)
start <- c(100,100)
xy <- data.frame(x=start[1]+cumsum(stepsize * cos(theta)), y=start[2]+cumsum(stepsize*sin(theta)))
plot(xy, pch=20, type="b", lty=1)
timestamps <- Sys.time() + 3600 * (1:n-1)

#Combine the xy locations and timestamps into a LoCoH-xy object
lxy <- xyt.lxy(xy=xy, dt=timestamps, id="broken_tooth")
summary(lxy)

## End(Not run)
```

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