

Package ‘PBSmapping’

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Title Mapping Fisheries Data and Spatial Analysis Tools

Author

Jon T. Schnute <Jon.Schnute@dfo-mpo.gc.ca>, Nicholas Boers, Rowan Haigh <Rowan.Haigh@dfo-mpo.gc.ca>, and Ale
ture-Beil.

Maintainer Jon Schnute <Jon.Schnute@dfo-mpo.gc.ca>

Depends R (>= 2.6.0)

Suggests foreign, maptools, deldir

Description This software has evolved from fisheries research conducted at the Pacific Biological Sta-
tion (PBS) in Nanaimo, British Columbia, Canada. It extends the R language to include
two-dimensional plotting features similar to those commonly
available in a Geographic Information System (GIS). Embedded C
code speeds algorithms from computational geometry, such as
finding polygons that contain specified point events or
converting between longitude-latitude and Universal Transverse
Mercator (UTM) coordinates. It includes data for a global
shoreline and other data sets in the public domain. The R
directory ‘../library/PBSmapping/doc’ includes a complete
user’s guide PBSmapping-UG.pdf. To use this package effectively, please consult the guide.

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addBubbles	<i>Add Bubbles to Maps</i>
------------	----------------------------

Description

Add bubbles proportional to some value (e.g., catch, effort) to a map. The function takes x, y and z arguments and optional arguments to draw bubbles of radius proportional to the z variable.

Usage

```
addBubbles(x, y, z, type=c("perceptual","surface","volume"),
           z.max=max(z,na.rm=TRUE), max.size=0.05, symbol.zero="+",
           symbol.fg=rgb(0,0,0,0.6), symbol.bg=rgb(0,0,0,0.3),
           legend.pos="bottomleft", legend.breaks=pretty(range(z),3)[-1],
           show.actual=FALSE, legend.type=c("nested","horiz","vert"),
           legend.title="Abundance", legend.cex=0.8, ...)
```

Arguments

x	X-coordinates compatible with map already plotted.
y	Y-coordinates compatible with map already plotted.
z	values that will determine relative size of bubbles at (X,Y).
type	scaling option for bubbles where "perceptual" emphasises large z-values while "volume" emphasises small z-values. Type "surface" lies in between.
z.max	maximum value for z (default = max(z)); determines the largest bubble; keeps the same legend for different maps.
max.size	maximum size (inches) of the largest bubble.
symbol.zero	symbol to represent zero-value z-values.
symbol.fg	bubble outline (border) colour.

<code>symbol.bg</code>	bubble interior (fill) colour.
<code>legend.pos</code>	position for the legend.
<code>legend.breaks</code>	break values for categorizing the z-values. The automatic method should work if zeroes are present; otherwise you can specify your own break values for the legend.
<code>show.actual</code>	logical: if FALSE, legend values are obtained using <code>pretty()</code> and consequently the largest bubble will be larger than <code>max(z)</code> . If TRUE, the largest bubble in the legend will correspond to <code>max(z)</code> .
<code>legend.type</code>	display format for legend.
<code>legend.title</code>	title for legend.
<code>legend.cex</code>	size of legend text.
<code>...</code>	additional arguments for <code>points</code> function that plots zero-value symbols.

Details

Modified (and for the legend, strongly inspired) from Tanimura et al. (2006) by Denis Chabot to work with `PBSmapping`.

Furthermore, Chabot's modifications make it possible to draw several maps with bubbles that all have the same scale (instead of each bubble plot having a scale that depends on the maximum z-value for that plot). This is done by making `z.max` equals to the largest z-value from all maps that will be plotted.

The user can also add a legend in one of four corners (see `legend::legend`) or at a specific x-y position. If `legend.pos` is NULL, no legend is drawn.

Note

This function has not yet been fully transformed to work with [EventData](#).

Author(s)

Denis Chabot, Institut Maurice-Lamontagne, Mont-Joli QC

References

Tanimura, S., Kuroiwa, C., and Mizota, T. (2006) Proportional symbol mapping in R. *Journal of Statistical Software* **15**(5).

See Also

[addPolys](#), [surveyData](#)

Examples

```
require(PBSmapping)
data(nepacLL,surveyData)
plotMap(nepacLL, xlim=c(-131.8,-127.2), ylim=c(50.5,52.7),
        col="gainsboro",plt=c(.08,.99,.08,.99))
```

```
addBubbles(surveyData$X, surveyData$Y, surveyData$catch,
  symbol.bg=rgb(.9,.5,0,.6), legend.type="nested",
  symbol.zero="+", col="grey")
```

addLabels *Add Labels to an Existing Plot*

Description

Add the label column of data to the existing plot.

Usage

```
addLabels (data, xlim = NULL, ylim = NULL, polyProps = NULL,
  placement = "DATA", polys = NULL, rollup = 3,
  cex = NULL, col = NULL, font = NULL, ...)
```

Arguments

data	EventData or PolyData to add (<i>required</i>).
xlim	range of X-coordinates.
ylim	range of Y-coordinates.
polyProps	PolyData specifying which labels to plot and their properties. par parameters passed as direct arguments supersede these data.
placement	one of "DATA", "CENTROID", "MEAN_RANGE", or "MEAN_XY".
polys	PolySet to use for calculating label placement.
rollup	level of detail at which to process polys, and it should match that in data. 1 = PIDs only, 2 = outer contours only, and 3 = no roll-up.
cex	vector describing character expansion factors (cycled by EID or PID).
col	vector describing colours (cycled by EID or PID).
font	vector describing fonts (cycled by EID or PID).
...	additional par parameters for the text function.

Details

If data is [EventData](#), it must minimally contain the columns EID, X, Y, and label. Since the EID column does not match a column in polys, set placement = "DATA". The function plots each label at its corresponding X/Y coordinate.

If data is [PolyData](#), it must minimally contain the columns PID and label. If it also contains X and Y columns, set placement = "DATA" to plot labels at those coordinates. Otherwise, set placement to one of "CENTROID", "MEAN_RANGE", or "MEAN_XY". When placement != "DATA", supply a [PolySet](#) polys. Using this [PolySet](#), the function calculates a centroid, mean range, or mean X/Y coordinate for each polygon, and then links those [PolyData](#) with data by PID/SID to determine label coordinates.

If data contains both PID and EID columns, the function assumes it is [PolyData](#) and ignores the EID column.

For additional help on the arguments `cex`, `col`, and `font`, please see [par](#).

Value

[EventData](#) or [PolyData](#) with X and Y columns that can subsequently reproduce the labels on the plot. Modify this data frame to tweak label positions.

See Also

[addPoints](#), [calcCentroid](#), [calcMidRange](#), [calcSummary](#), [EventData](#), [plotPoints](#), [PolyData](#).

Examples

```
#--- create sample PolyData to label Vancouver Island
labelData <- data.frame(PID=33, label="Vancouver Island");
#--- load data
if (!is.null(version$language) && (version$language == "R"))
  data(nepacLL)
#--- plot the map
plotMap (nepacLL, xlim=c(-129, -122.6), ylim=c(48, 51.1))
#--- add the labels
addLabels (labelData, placement="CENTROID", polys=nepacLL, col=2)
```

addLines

Add a PolySet to an Existing Plot as Polylines

Description

Add a [PolySet](#) to an existing plot, where each unique (PID, SID) describes a polyline.

Usage

```
addLines (polys, xlim = NULL, ylim = NULL,
          polyProps = NULL, lty = NULL, col = NULL, ...)
```

Arguments

<code>polys</code>	PolySet to add (<i>required</i>).
<code>xlim</code>	range of X-coordinates.
<code>ylim</code>	range of Y-coordinates.
<code>polyProps</code>	PolyData specifying which polylines to plot and their properties. par parameters passed as direct arguments supersede these data.
<code>lty</code>	vector of line types (cycled by PID).
<code>col</code>	vector of colours (cycled by PID).
<code>...</code>	additional par parameters for the lines function.

Details

The plotting routine does not connect the last vertex of each discrete polyline to the first vertex of that polyline. It clips polys to xlim and ylim before plotting.

For additional help on the arguments lty and col, please see [par](#).

Value

[PolyData](#) consisting of the PolyProps used to create the plot.

See Also

[calcLength](#), [clipLines](#), [closePolys](#), [convLP](#), [fixBound](#), [fixPOS](#), [locatePolys](#), [plotLines](#), [thinPolys](#), [thickenPolys](#).

Examples

```
#--- create a PolySet to plot
polys <- data.frame(PID=rep(1,4),POS=1:4,X=c(0,1,1,0),Y=c(0,0,1,1))
polys <- as.PolySet(polys, projection=1)
#--- plot the PolySet
plotLines(polys, xlim=c(-.5,1.5), ylim=c(-.5,1.5), projection=1)
#--- add the PolySet to the plot (in a different style)
addLines(polys, lwd=5, col=3)
```

 addPoints

Add EventData/PolyData to an Existing Plot as Points

Description

Add [EventData/PolyData](#) to an existing plot, where each unique EID describes a point.

Usage

```
addPoints (data, xlim = NULL, ylim = NULL, polyProps = NULL,
           cex = NULL, col = NULL, pch = NULL, ...)
```

Arguments

data	EventData or PolyData to add (<i>required</i>).
xlim	range of X-coordinates.
ylim	range of Y-coordinates.
polyProps	PolyData specifying which points to plot and their properties. par parameters passed as direct arguments supersede these data.
cex	vector describing character expansion factors (cycled by EID or PID).
col	vector describing colours (cycled by EID or PID).
pch	vector describing plotting characters (cycled by EID or PID).
...	additional par parameters for the points function.

Details

This function clips data to `xlim` and `ylim` before plotting. It only adds [PolyData](#) containing X and Y columns.

For additional help on the arguments `cex`, `col`, and `pch`, please see [par](#).

Value

[PolyData](#) consisting of the PolyProps used to create the plot.

See Also

[combineEvents](#), [convDP](#), [findPolys](#), [locateEvents](#), [plotPoints](#).

Examples

```
#--- load the data (if using R)
if (!is.null(version$language) && (version$language == "R")) {
  data(nepacLL)
  data(surveyData)
}
#--- plot a map
plotMap(nepacLL, xlim=c(-136, -125), ylim=c(48, 57))
#--- add events
addPoints(surveyData, col=1:7)
```

 addPolys

Add a PolySet to an Existing Plot as Polygons

Description

Add a [PolySet](#) to an existing plot, where each unique (PID, SID) describes a polygon.

Usage

```
addPolys (polys, xlim = NULL, ylim = NULL, polyProps = NULL,
          border = NULL, lty = NULL, col = NULL, colHoles = NULL,
          density = NA, angle = NULL, ...)
```

Arguments

<code>polys</code>	PolySet to add (<i>required</i>).
<code>xlim</code>	range of X-coordinates.
<code>ylim</code>	range of Y-coordinates.
<code>polyProps</code>	PolyData specifying which polygons to plot and their properties. par parameters passed as direct arguments supersede these data.
<code>border</code>	vector describing edge colours (cycled by PID).

lty	vector describing line types (cycled by PID).
col	vector describing fill colours (cycled by PID).
colHoles	vector describing hole colours (cycled by PID). The default, NULL, should be used in most cases as it renders holes transparent. colHoles is designed solely to eliminate retrace lines when images are converted to PDF format. If colHoles is specified, underlying information (i.e., previously plotted shapes) will be obliterated. If NA is specified, only outer polygons are drawn, consequently filling holes.
density	vector describing shading line densities (lines per inch, cycled by PID).
angle	vector describing shading line angles (degrees, cycled by PID).
...	additional par parameters for the polygon function.

Details

The plotting routine connects the last vertex of each discrete polygon to the first vertex of that polygon. It supports both borders (`border`, `lty`) and fills (`col`, `density`, `angle`). It clips polys to `xlim` and `ylim` before plotting.

For additional help on the arguments `border`, `lty`, `col`, `density`, and `angle`, please see [polygon](#) and [par](#).

Value

[PolyData](#) consisting of the `PolyProps` used to create the plot.

See Also

[addLabels](#), [addStipples](#), [clipPolys](#), [closePolys](#), [fixBound](#), [fixPOS](#), [locatePolys](#), [plotLines](#), [plotMap](#), [plotPoints](#), [plotPolys](#), [thinPolys](#), [thickenPolys](#).

Examples

```
#--- create a PolySet to plot
polys <- data.frame(PID=rep(1,4),POS=1:4,X=c(0,1,1,0),Y=c(0,0,1,1))
polys <- as.PolySet(polys, projection=1)
#--- plot the PolySet
plotPolys(polys,xlim=c(-.5,1.5),ylim=c(-.5,1.5),density=0,projection=1)
#--- add the PolySet to the plot (in a different style)
addPolys(polys, col=3)
```

addStipples

Add Stipples to an Existing Plot

Description

Add stipples to an existing plot.

Usage

```
addStipples (polys, xlim = NULL, ylim = NULL, polyProps = NULL,
            side = 1, density = 1, distance = 4, ...)
```

Arguments

polys	PolySet that provides the stipple boundaries (<i>required</i>).
xlim	range of X-coordinates.
ylim	range of Y-coordinates.
polyProps	PolyData specifying which polygons to stipple and their properties. par parameters passed as direct arguments supersede these data.
side	one of -1, 0, or 1, corresponding to outside, both sides, or inside, respectively.
density	density of points, relative to the default.
distance	distance to offset points, measured as a percentage of the absolute difference in xlim.
...	additional par parameters for the points function.

Details

This function locates stipples based on the [PolySet](#) polys and does not stipple degenerate lines.

Value

[PolyData](#) consisting of the PolyProps used to create the plot.

See Also

[addPoints](#), [addPolys](#), [plotMap](#), [plotPoints](#), [plotPolys](#), [points](#), [PolySet](#).

Examples

```
#--- load the data (if using R)
if (!is.null(version$language) && (version$language == "R")) {
  data(nepacLL)
}
#--- plot a map
plotMap(nepacLL, xlim=c(-128.66, -122.83), ylim=c(48.00, 51.16))
#--- add stippling
addStipples(nepacLL, col=2, pch=19, cex=0.25)
```

appendPolys	<i>Append a Two-Column Matrix to a PolySet</i>
-------------	------------------------------------------------

Description

Append a two-column matrix to a [PolySet](#), assigning PID and possibly SID values automatically or as specified in its arguments.

Usage

```
appendPolys (polys, mat, PID = NULL, SID = NULL, isHole = FALSE)
```

Arguments

polys	existing PolySet ; if NULL, creates a new PolySet (<i>required</i>).
mat	two-column matrix to append (<i>required</i>).
PID	new polygon's PID.
SID	new polygon's SID.
isHole	Boolean value; if TRUE, mat represents a hole.

Details

If polys contains an SID column and the SID argument equals NULL, this function uses the next available SID.

If polys does not contain an SID column and the caller passes an SID argument, all existing polygons will receive an SID of 1. The new polygon's SID will match the SID argument.

If isHole = TRUE, the polygon's POS values will appropriately represent a hole (reverse order of POS).

If (PID, SID) already exists in the [PolySet](#), the function will issue a warning and duplicate those identifiers.

Value

[PolySet](#) containing mat appended to polys. The function retains attributes from polys.

See Also

[addPolys](#), [clipPolys](#), [closePolys](#), [convLP](#), [fixBound](#), [fixPOS](#), [joinPolys](#), [plotMap](#), [plotPolys](#).

Examples

```
#--- create two simple matrices
a <- matrix(data=c(0,0,1,0,1,1,0,1),ncol=2,byrow=TRUE);
b <- matrix(data=c(2,2,3,2,3,3,2,3), ncol=2,byrow=TRUE);
#--- build a PolySet from them
polys <- appendPolys(NULL, a);
polys <- appendPolys(polys, b);
#--- print the result
print (polys);
```

bcBathymetry

Data: Bathymetry Spanning British Columbia's Coast

Description

Bathymetry data spanning British Columbia's coast.

Usage

```
data(bcBathymetry)
```

Format

Three-element list: x = vector of horizontal grid line locations, y = vector of vertical grid line locations, z = (x by y) matrix containing water depths measured in meters. Positive values indicate distance below sea level and negative values above it.

[contour](#) and [contourLines](#) expect data in this format. [convCP](#) converts the output from [contourLines](#) into a [PolySet](#).

Note

In R, the data must be loaded using the [data](#) function.

Source

Bathymetry data acquired from the Scripps Institution of Oceanography at the University of San Diego.

Using their online form, we requested bathymetry data for the complete [nepacLL](#) region. At forty megabytes, the data were not suitable for distribution in our mapping package. Therefore, we reduced the data to the range $-140^\circ \leq x \leq -122^\circ$ and $47^\circ \leq y \leq 61^\circ$.

References

Smith, W.H.F. and Sandwell, D.T. (1997) Global seafloor topography from satellite altimetry and ship depth soundings. *Science* **277**, 1957–1962.

http://topex.ucsd.edu/WWW_html/mar_topo.html

See Also

[contour](#), [contourLines](#), [convCP](#), [nepacLL](#), [nepacLLhigh](#).

calcArea	<i>Calculate the Areas of Polygons</i>
----------	----------------------------------------

Description

Calculate the areas of polygons found in a [PolySet](#).

Usage

```
calcArea (polys, rollup = 3)
```

Arguments

polys	PolySet to use.
rollup	level of detail in the results; 1 = PIDs only, by summing all the polygons with the same PID, 2 = outer contours only, by subtracting holes from their parent, and 3 = no roll-up.

Details

If rollup equals 1, the results contain an area for each unique PID only. When it equals 2, they contain entries for outer contours only. Finally, setting it to 3 prevents roll-up, and they contain areas for each unique (PID, SID).

Outer polygons have positive areas and inner polygons negative areas. When polygons are rolled up, the routine sums the positive and negative areas and consequently accounts for holes.

If the [PolySet](#)'s projection attribute equals "LL", the function projects the [PolySet](#) in UTM first. If the [PolySet](#)'s zone attribute exists, it uses it for the conversion. Otherwise, it computes the mean longitude and uses that value to determine the zone. The longitude range of zone i is $-186 + 6i^\circ < x \leq -180 + 6i^\circ$.

Value

[PolyData](#) with columns PID, SID (*may be missing*), and area. If the projection equals "LL" or "UTM", the units of area are square kilometres.

See Also

[calcCentroid](#), [calcLength](#), [calcMidRange](#), [calcSummary](#), [locatePolys](#).

Examples

```
#--- load the data (if using R)
if (!is.null(version$language) && (version$language == "R"))
  data(nepacLL)
#--- convert LL to UTM so calculation makes sense
attr(nepacLL, "zone") <- 9
nepacUTM <- convUL(nepacLL)
#--- calculate and print the areas
print(calcArea(nepacUTM))
```

 calcCentroid

Calculate the Centroids of Polygons

Description

Calculate the centroids of polygons found in a [PolySet](#).

Usage

```
calcCentroid (polys, rollup = 3)
```

Arguments

polys	PolySet to use.
rollup	level of detail in the results; 1 = PIDs only, 2 = outer contours only, and 3 = no roll-up. When rollup equals 1 and 2, the function appropriately adjusts for polygons with holes.

Details

If rollup equals 1, the results contain a centroid for each unique PID only. When it equals 2, they contain entries for outer contours only. Finally, setting it to 3 prevents roll-up, and they contain a centroid for each unique (PID, SID).

Value

[PolyData](#) with columns PID, SID (*may be missing*), X, and Y.

See Also

[calcArea](#), [calcLength](#), [calcMidRange](#), [calcSummary](#), [locateEvents](#), [locatePolys](#).

Examples

```
#--- load the data (if using R)
if (!is.null(version$language) && (version$language == "R"))
  data(nepacLL)
#--- calculate and print the centroids for several polygons
print(calcCentroid(nepacLL[is.element(nepacLL$PID, c(33, 39, 47)), ]))
```

calcConvexHull	<i>Calculate the Convex Hull for a Set of Points</i>
----------------	------------------------------------------------------

Description

Calculate the convex hull for a set of points.

Usage

```
calcConvexHull (xydata, keepExtra=FALSE)
```

Arguments

xydata a data frame with columns X and Y containing spatial coordinates.
keepExtra logical: if TRUE, retain any additional columns from the input data frame xydata.

Details

This routine uses the function `chull()` in the package `grDevices`. By default, it ignores all columns other than X and Y; however, the user can choose to retain additional columns in xydata by specifying `keepExtra=TRUE`.

Value

[PolySet](#) with columns PID, POS, X, Y, and additional columns in xydata if `keepExtra=TRUE`.

See Also

[addPoints](#), [addPolys](#), [calcArea](#), [calcCentroid](#), [calcMidRange](#), [calcSummary](#), [locateEvents](#), [plotMap](#), [plotPoints](#), [plotPolys](#).

Examples

```
data(surveyData)
#--- plot the convex hull, and then plot the points
plotMap(calcConvexHull(surveyData), col="moccasin")
addPoints(surveyData, col="blue", pch=17, cex=.6);
```

calcLength	<i>Calculate the Length of Polylines</i>
------------	------------------------------------------

Description

Calculate the length of polylines found in a [PolySet](#).

Usage

```
calcLength (polys, rollup = 3, close = FALSE)
```

Arguments

polys	PolySet to use.
rollup	level of detail in the results; 1 = PIDs only, summing the lengths of each SID within each PID, and 3 = no roll-up. Note: rollup 2 has no meaning in this function and, if specified, will be reset to 3.
close	Boolean value; if TRUE, include the distance between each polygon's last and first vertex, if necessary.

Details

If rollup equals 1, the results contain an entry for each unique PID only. Setting it to 3 prevents roll-up, and they contain an entry for each unique (PID, SID).

If the projection attribute equals "LL", this routine uses Great Circle distances to compute the surface length of each polyline. In doing so, the algorithm simplifies Earth to a sphere.

If the projection attribute equals "UTM" or 1, this routine uses Pythagoras' Theorem to calculate lengths.

Value

[PolyData](#) with columns PID, SID (*may be missing*), and length. If projection equals "UTM" or "LL", lengths are in kilometres. Otherwise, lengths are in the same unit as the input [PolySet](#).

See Also

[calcArea](#), [calcCentroid](#), [calcMidRange](#), [calcSummary](#), [locatePolys](#).

Examples

```
#--- load the data (if using R)
if (!is.null(version$language) && (version$language == "R"))
  data(nepacLL)
#--- calculate the perimeter of Vancouver Island
print(calcLength(nepacLL[nepacLL$PID == 33, ]))
```

calcMidRange	<i>Calculate the Midpoint of the X/Y Ranges of Polygons</i>
--------------	-------------------------------------------------------------

Description

Calculate the midpoint of the X/Y ranges of polygons found in a [PolySet](#).

Usage

```
calcMidRange (polys, rollup = 3)
```

Arguments

polys	PolySet to use.
rollup	level of detail in the results; 1 = PIDs only, 2 = outer contours only, and 3 = no roll-up.

Details

If rollup equals 1, the results contain a mean range for each unique PID only. When it equals 2, they contain entries for outer contours only. Finally, setting it to 3 prevents roll-up, and they contain a mean range for each unique (PID, SID).

Value

[PolyData](#) with columns PID, SID (*may be missing*), X, and Y.

See Also

[calcArea](#), [calcCentroid](#), [calcLength](#), [calcSummary](#).

Examples

```
#--- load the data (if using R)
if (!is.null(version$language) && (version$language == "R"))
  data(nepacLL)
#--- calculate and print the centroids for several polygons
print(calcMidRange(nepacLL[is.element(nepacLL$PID, c(33, 39, 47))], ]))
```

 calcSummary

Apply Functions to Polygons in a PolySet

Description

Apply functions to polygons in a [PolySet](#).

Usage

```
calcSummary (polys, rollup = 3, FUN, ...)
```

Arguments

polys	PolySet to use.
rollup	level of detail in the results; 1 = PIDs only, by removing the SID column, and then passing each PID into FUN, 2 = outer contours only, by making hole SIDs equal to their parent's SID, and then passing each (PID, SID) into FUN, and 3 = no roll-up.
FUN	the function to apply; it must accept a vector and return a vector or scalar.
...	optional arguments for FUN.

Details

If rollup equals 1, the results contain an entry for each unique PID only. When it equals 2, they contain entries for outer contours only. Finally, setting it to 3 prevents roll-up, and they contain an entry for each unique (PID, SID).

Value

[PolyData](#) with columns PID, SID (*may be missing*), X, and Y. If FUN returns a vector of length greater than 1 (say n), names the columns X1, X2, ..., X n and Y1, Y2, ..., Y n .

See Also

[calcArea](#), [calcCentroid](#), [calcConvexHull](#), [calcLength](#), [calcMidRange](#), [combineEvents](#), [findPolys](#), [locateEvents](#), [locatePolys](#), [makeGrid](#), [makeProps](#).

Examples

```
#--- load the data (if using R)
if (!is.null(version$language) && (version$language == "R"))
  data(nepacLL)
#--- calculate and print the centroids for several polygons
print(calcSummary(nepacLL[is.element(nepacLL$PID, c(33, 39, 47)), ],
                 rollup = 3, FUN = mean))
```

calcVoronoi

*Calculate the Voronoi (Dirichlet) Tessellation for a Set of Points***Description**

Calculate the Voronoi (Dirichlet) tessellation for a set of points.

Usage

```
calcVoronoi(xydata, xlim = NULL, ylim = NULL, eps = 1e-09, frac = 0.0001)
```

Arguments

xydata	a data frame with columns X and Y containing the points.
xlim	range of X-coordinates; a bounding box for the coordinates.
ylim	range of Y-coordinates; a bounding box for the coordinates.
eps	the value of epsilon used in testing whether a quantity is zero.
frac	used to detect duplicate input points, which meet the condition $ x_1 - x_2 < \text{frac} \times (\text{xmax} - \text{xmin})$ and $ y_1 - y_2 < \text{frac} \times (\text{ymax} - \text{ymin})$.

Details

This routine ignores all columns other than X and Y.

If the user leaves `xlim` and `ylim` unspecified, the function defaults to the range of the data with each extent expanded by ten percent of the range.

This function sets the attribute projection to 1 and the attribute zone to NULL as it assumes this projection in its calculations.

Value

[PolySet](#) with columns PID, POS, X, and Y.

See Also

[addPoints](#), [addPolys](#), [calcArea](#), [calcCentroid](#), [calcConvexHull](#), [calcMidRange](#), [calcSummary](#), [locateEvents](#), [plotMap](#), [plotPoints](#), [plotPolys](#).

Examples

```
#--- create some EventData
events <- as.EventData(data.frame(EID=1:200,
                                X=rnorm(200),
                                Y=rnorm(200)),
                      projection=1)
#--- calculate the Voronoi tessellation
polys <- calcVoronoi(events)
#--- create PolyData to color it based on area
```

```

polyData <- calcArea(polys)
names(polyData)[is.element(names(polyData), "area")] <- "Z"
colSeq <- seq(0.4, 0.95, length=4)
polyData <- makeProps(polyData,
                      breaks=quantile(polyData$Z,c(0,.25,.5,.75,1)),
                      propName="col",
                      propVals=rgb(colSeq, colSeq, colSeq))
#--- plot the tessellation
plotMap(polys, polyProps=polyData)
#--- plot the points
addPoints(events, pch=19)

```

clipLines

Clip a PolySet as PolyLines

Description

Clip a [PolySet](#), where each unique (PID, SID) describes a polyline.

Usage

```
clipLines (polys, xlim, ylim, keepExtra = FALSE)
```

Arguments

polys	PolySet to clip.
xlim	range of X-coordinates.
ylim	range of Y-coordinates.
keepExtra	Boolean value; if TRUE, tries to carry forward any non-standard columns into the result.

Details

For each discrete polyline, the function does not connect vertices 1 and N. It recalculates the POS values for each vertex, saving the old values in a column named oldPOS. For new vertices, it sets oldPOS to NA.

Value

[PolySet](#) containing the input data, with some points added or removed. A new column oldPOS records the original POS value for each vertex.

See Also

[clipPolys](#), [fixBound](#).

Examples

```
#--- create a triangle to clip
polys <- data.frame(PID=rep(1, 3), POS=1:3, X=c(0,1,0), Y=c(0,0.5,1))
#--- clip the triangle in the X direction, and plot the results
plotLines(clipLines(polys, xlim=c(0,.75), ylim=range(polys[, "Y"])))
```

clipPolys

*Clip a PolySet as Polygons***Description**

Clip a [PolySet](#), where each unique (PID, SID) describes a polygon.

Usage

```
clipPolys (polys, xlim, ylim, keepExtra = FALSE)
```

Arguments

polys	PolySet to clip.
xlim	range of X-coordinates.
ylim	range of Y-coordinates.
keepExtra	Boolean value; if TRUE, tries to carry forward any non-standard columns into the result.

Details

For each discrete polygon, the function connects vertices 1 and N. It recalculates the POS values for each vertex, saving the old values in a column named oldPOS. For new vertices, it sets oldPOS to NA.

Value

[PolySet](#) containing the input data, with some points added or removed. A new column oldPOS records the original POS value for each vertex.

See Also

[clipLines](#), [fixBound](#).

Examples

```
#--- create a triangle that will be clipped
polys <- data.frame(PID=rep(1, 3), POS=1:3, X=c(0,1,.5), Y=c(0,0,1))
#--- clip the triangle in the X direction, and plot the results
plotPolys(clipPolys(polys, xlim=c(0,.75), ylim=range(polys[, "Y"])),
          col=2);
```

`closePolys`*Close a PolySet*

Description

Close a [PolySet](#) of polylines to form polygons.

Usage

```
closePolys (polys)
```

Arguments

`polys` [PolySet](#) to close.

Details

Generally, run `fixBound` before this function. The ranges of a [PolySet](#)'s X and Y columns define the boundary. For each discrete polygon, this function determines if the first and last points lie on a boundary. If both points lie on the same boundary, it adds no points. However, if they lie on different boundaries, it may add one or two corners to the polygon.

When the boundaries are adjacent, one corner will be added as follows:

- top boundary + left boundary implies add top-left corner;
- top boundary + right boundary implies add top-right corner;
- bottom boundary + left boundary implies add bottom-left corner;
- bottom boundary + right boundary implies add bottom-right corner.

When the boundaries are opposite, it first adds the corner closest to a starting or ending polygon vertex. This determines a side (left-right or bottom-top) that connects the opposite boundaries. Then, it adds the other corner of that side to close the polygon.

Value

[PolySet](#) identical to `polys`, except for possible additional corner points.

See Also

[fixBound](#), [fixPOS](#).

Examples

```
#--- 4 corners
polys <- data.frame(PID=c(1, 1, 2, 2, 3, 3, 4, 4),
                   POS=c(1, 2, 1, 2, 1, 2, 1, 2),
                   X = c(0, 1, 2, 3, 0, 1, 2, 3),
                   Y = c(1, 0, 0, 1, 2, 3, 3, 2))
plotPolys(closePolys(polys), col=2)
```

```
#--- 2 corners and 1 opposite
polys <- data.frame(PID=c(1, 1, 2, 2, 3, 3, 3),
                   POS=c(1, 2, 1, 2, 1, 2, 3),
                   X = c(0, 1, 0, 1, 5, 6, 1.5),
                   Y = c(1, 0, 2, 3, 0, 1.5, 3))
plotPolys(closePolys(polys), col=2)
```

 combineEvents

Combine Measurements of Events

Description

Combine measurements associated with events that occur in the same polygon.

Usage

```
combineEvents (events, locs, FUN, ..., bdryOK = TRUE)
```

Arguments

events	EventData with at least four columns (EID, X, Y, Z).
locs	LocationSet usually resulting from a call to findPolys .
FUN	a function that produces a scalar from a vector (e.g., mean , sum).
...	optional arguments for FUN.
bdryOK	Boolean value; if TRUE, include boundary points.

Details

This function combines measurements associated with events that occur in the same polygon. Each event (EID) has a corresponding measurement Z. The locs data frame (usually output from [findPolys](#)) places events within polygons. Thus, each polygon (PID, SID) determines a set of events within it, and a corresponding vector of measurements Z_v. The function returns FUN(Z_v), a summary of measurements within each polygon.

Value

[PolyData](#) with columns PID, SID (*if in* locs), and Z.

See Also

[findCells](#), [findPolys](#), [locateEvents](#), [locatePolys](#), [makeGrid](#), [makeProps](#).

Examples

```
#--- create an EventData data frame: let each event have Z = 1
events <- data.frame(EID=1:10, X=1:10, Y=1:10, Z=rep(1, 10))
#--- example output from findPolys where 1 event occurred in the first
#--- polygon, 3 in the second, and 6 in the third
locs <- data.frame(EID=1:10, PID=c(rep(1, 1), rep(2, 3), rep(3, 6)),
                  Bdry=rep(0, 10))
#--- sum the Z column of the events in each polygon, and print the result
print(combineEvents(events=events, locs=locs, FUN=sum))
```

combinePolys

Combine Several Polygons into a Single Polygon

Description

Combine several polygons into a single polygon by modifying the PID and SID indices.

Usage

```
combinePolys (polys)
```

Arguments

polys [PolySet](#) with one or more polygons, each with possibly several components/holes.

Details

This function accepts a [PolySet](#) containing one or more polygons (PIDs), each with one or more components or holes (SIDs). The SID column need not exist in the input. The function combines these polygons into a single polygon by simply renumbering the PID and SID indices. The resulting [PolySet](#) contains a single PID (with the value 1) and uses the SID value to differentiate between polygons, their components, and holes.

Value

[PolySet](#), possibly with the addition of an SID column if it did not already exist. The function may also reorder columns such that PID, SID, POS, X and Y appear first, in that order.

See Also

[dividePolys](#)

convCP	<i>Convert Contour Lines into a PolySet</i>
--------	---------------------------------------------

Description

Convert output from [contourLines](#) into a [PolySet](#).

Usage

```
convCP (data, projection = NULL, zone = NULL)
```

Arguments

data	contour line data, often from the contourLines function.
projection	optional projection attribute to add to the PolySet.
zone	optional zone attribute to add to the PolySet.

Details

data contains a list as described below. The [contourLines](#) function create a list suitable for the data argument.

A three-element list describes each contour. The named elements in this list include the scalar level, the vector x, and the vector y. Vectors x and y must have equal lengths. A higher-level list (data) contains one or more of these contours lists.

Value

A list with two named elements [PolySet](#) and [PolyData](#). The [PolySet](#) element contains a [PolySet](#) representation of the contour lines. The [PolyData](#) element links each contour line (PID, SID) with a level.

See Also

[contour](#), [contourLines](#), [convLP](#), [makeTopography](#).

Examples

```
#--- create sample data for the contourLines() function
x <- seq(-0.5, 0.8, length=50);
y <- x;
z <- outer(x, y, FUN = function(x,y) { sin(2*pi*(x^2+y^2))^2; } );
data <- contourLines(x, y, z, levels=c(0.2, 0.8));
#--- pass that sample data into convCP()
result <- convCP(data);
#--- plot the result
plotLines(result$PolySet, projection=1);
print(result$PolyData);
```

`convDP`*Convert EventData/PolyData into a PolySet*

Description

Convert [EventData/PolyData](#) into a [PolySet](#).

Usage

```
convDP (data, xColumns, yColumns)
```

Arguments

<code>data</code>	PolyData or EventData .
<code>xColumns</code>	vector of X-column names.
<code>yColumns</code>	vector of Y-column names.

Details

This function expects data to contain several X- and Y-columns. For example, consider data with columns `x1`, `y1`, `x2`, and `y2`. Suppose `xColumns = c("x1", "x2")` and `yColumns = c("y1", "y2")`. The result will contain `nrow(data)` polygons. Each one will have two vertices, `(x1, y1)` and `(x2, y2)` and POS values 1 and 2, respectively. If data includes an SID column, so will the result.

If data contains an EID and not a PID column, the function uses the EIDs as PIDs.

If data contains both PID and EID columns, the function assumes it is [PolyData](#) and ignores the EID column.

Value

[PolySet](#) with the same PIDs as those given in data. If data has an SID column, the result will include it.

See Also

[addPoints](#), [plotPoints](#).

Examples

```
#--- create sample PolyData
polyData <- data.frame(PID=c(1, 2, 3),
                      x1=c(1, 3, 5), y1=c(1, 3, 2),
                      x2=c(1, 4, 5), y2=c(2, 4, 1),
                      x3=c(2, 4, 6), y3=c(2, 3, 1));

#--- print PolyData
print(polyData);
```

```

#--- make a PolySet from PolyData
polys <- convDP(polyData,
                xColumns=c("x1", "x2", "x3"),
                yColumns=c("y1", "y2", "y3"));
#--- print and plot the PolySet
print(polys);
plotLines(polys, xlim=c(0, 7), ylim=c(0, 5), col=2);

```

convLP

Convert Polylines into a Polygon

Description

Convert two polylines into a polygon.

Usage

```
convLP (polyA, polyB, reverse = TRUE)
```

Arguments

polyA	PolySet containing a polyline.
polyB	PolySet containing a polyline.
reverse	Boolean value; if TRUE, reverse polyB's vertices.

Details

The resulting [PolySet](#) contains all the vertices from polyA in their original order. If `reverse = TRUE`, this function appends the vertices from polyB in the reverse order (`nrow(polyB):1`). Otherwise, it appends them in their original order. The PID column equals the PID of polyA. No SID column appears in the result. The resulting polygon is an exterior boundary.

Value

[PolySet](#) with a single PID that is the same as polyA. The result contains all the vertices in polyA and polyB. It has the same projection and zone attributes as those in the input PolySets. If an input PolySet's attributes equal NULL, the function uses the other PolySet's. If the PolySet attributes conflict, the result's attribute equals NULL.

See Also

[addLines](#), [appendPolys](#), [closePolys](#), [convCP](#), [joinPolys](#), [plotLines](#).

Examples

```
#--- create two polylines
polyline1 <- data.frame(PID=rep(1, 2), POS=1:2, X=c(1, 4), Y=c(1, 4));
polyline2 <- data.frame(PID=rep(1, 2), POS=1:2, X=c(2, 5), Y=c(1, 4));

#--- create two plots to demonstrate the effect of 'reverse'
par(mfrow=c(2, 1))
plotPolys(convLP(polyline1, polyline2, reverse = TRUE), col=2);
plotPolys(convLP(polyline1, polyline2, reverse = FALSE), col=3);
```

convUL

*Convert Coordinates between UTM and Lon/Lat***Description**

Convert coordinates between UTM and Lon/Lat.

Usage

```
convUL (xydata, km=TRUE)
```

Arguments

xydata	data frame with columns X and Y.
km	use UTM coordinates in kilometres if true, or metres if false.

Details

xydata must possess a projection attribute that identifies the current projection. If the data frame contains UTM coordinates, it must also have a zone attribute equal to a number between 1 and 60 (inclusive). If it contains longitude/latitude coordinates and the zone attribute is missing, the function computes the mean longitude and uses that value to determine the zone. The longitude range of zone i is $-186 + 6i^\circ < x \leq -180 + 6i^\circ$.

This function converts the X and Y columns of xydata from "LL" to "UTM" or vice-versa. After the conversion, it adjusts the data frame's attributes accordingly.

Value

A data frame identical to xydata, except that the X and Y columns contain the results of the conversion and the projection attribute matches the new projection.

See Also

[closePolys](#), [fixBound](#).

Examples

```
#--- load the data
data(nepacLL, package="PBSmapping")
#--- set the zone attribute
attr(nepacLL, "zone") <- 9
#--- convert and plot the result
nepacUTM <- convUL(nepacLL)
plotMap(nepacUTM)
```

dividePolys

Divide a Single Polygon into Several Polygons

Description

Divide a single polygon (with several outer-contour components) into several polygons, a polygon for each outer contour, by modifying the PID and SID indices.

Usage

```
dividePolys (polys)
```

Arguments

polys [PolySet](#) with one or more polygons, each with possibly several components/holes.

Details

Given the input [PolySet](#), this function renumbers the PID and SID indices so that each outer contour has a unique PID and is followed by all of its holes, identifying them with SIDs greater than one.

Value

[PolySet](#), possibly with the addition of an SID column if it did not already exist. The function may also reorder columns such that PID, SID, POS, X and Y appear first, in that order.

See Also

[combinePolys](#).

 EventData

EventData Objects

Description

PBS Mapping functions that expect EventData will accept properly formatted data frames in their place (see 'Details').

`as.EventData` attempts to coerce a data frame to an object with class `EventData`.

`is.EventData` returns `TRUE` if its argument is of class `EventData`.

Usage

```
as.EventData(x, projection = NULL, zone = NULL)
is.EventData(x, fullValidation = TRUE)
```

Arguments

<code>x</code>	data frame to be coerced or tested.
<code>projection</code>	optional projection attribute to add to <code>EventData</code> , possibly overwriting an existing attribute.
<code>zone</code>	optional zone attribute to add to <code>EventData</code> , possibly overwriting an existing attribute.
<code>fullValidation</code>	Boolean value; if <code>TRUE</code> , fully test <code>x</code> .

Details

We define `EventData` as a data frame with at least three fields named (EID, X, Y). Conceptually, an `EventData` object describes events that take place at specific points (X, Y) in two-dimensional space. Additional fields specify measurements associated with these events. For example, in a fishery context `EventData` could describe fishing events associated with trawl tows, based on the fields:

- EID - fishing event (tow) identification number;
- X, Y - fishing location;
- Duration - length of time for the tow;
- Depth - average depth of the tow;
- Catch - biomass captured.

Like `PolyData`, `EventData` can have attributes `projection` and `zone`, which may be absent. Inserting the string "EventData" as the class attribute's first element alters the behaviour of some functions, including `print` (if `PBSprint` is `TRUE`) and `summary`.

Value

The `as.EventData` method returns an object with classes "EventData" and "data.frame", in that order.

See Also

[LocationSet](#), [PolyData](#), [PolySet](#).

extractPolyData	<i>Extract PolyData from a PolySet</i>
-----------------	----------------------------------------

Description

Extract [PolyData](#) from a [PolySet](#). Columns for the [PolyData](#) include those other than PID, SID, POS, oldPOS, X, and Y.

Usage

```
extractPolyData (polys)
```

Arguments

polys [PolySet](#) to use.

Details

This function identifies the [PolySet](#)'s extra columns and determines if those columns contain unique values for each (PID, SID). Where they do, the (PID, SID) will appear in the [PolyData](#) output with that unique value. Where they do not, the extra column will contain NAs for that (PID, SID).

Value

[PolyData](#) with columns PID, SID, and any extra columns.

See Also

[makeProps](#), [PolyData](#), [PolySet](#).

Examples

```
#--- create a PolySet with an extra column
polys <- data.frame(PID = c(rep(1, 10), rep(2, 10)),
                   POS = c(1:10, 1:10),
                   X = c(rep(1, 10), rep(1, 10)),
                   Y = c(rep(1, 10), rep(1, 10)),
                   colour = (c(rep("green", 10), rep("red", 10))));
#--- extract the PolyData
print(extractPolyData(polys))
```

findCells

*Find the Grid Cells that Contain Events***Description**

Find the grid cells in a [PolySet](#) that contain events specified in [EventData](#). Similar to [findPolys](#), except this function requires a [PolySet](#) resulting from [makeGrid](#). This restriction allows this function to calculate the result with greater efficiency.

Usage

```
findCells (events, polys)
```

Arguments

events [EventData](#) to use.
polys [PolySet](#) to use.

Details

The resulting data frame, a [LocationSet](#), contains the columns EID, PID, SID (*if in polys*), and Bdry, where an event (EID) occurs in a polygon (PID, SID). The Boolean variable Bdry indicates whether an event lies on a polygon's edge. Note that if an event lies properly outside of all the polygons, then a record with (EID, PID, SID) does not occur in the output. It may happen, however, that an event occurs in multiple polygons (i.e., on two or more boundaries). Thus, the same EID can occur more than once in the output.

Value

[LocationSet](#) that links events with polygons.

See Also

[combineEvents](#), [findPolys](#), [locateEvents](#), [locatePolys](#), [LocationSet](#), [makeGrid](#).

Examples

```
#--- create some EventData: points in a diagonal line
events <- data.frame(EID=1:11, X=seq(0, 2, length=11),
                    Y=seq(0, 2, length=11))
events <- as.EventData(events, projection=1);
#--- create a PolySet (a grid)
polys <- makeGrid (x=seq(0, 2, by=0.50), y=seq(0, 2, by=0.50),
                  projection=1)
#--- show a picture
plotPolys(polys, xlim=range(polys$X)+c(-0.1, 0.1),
          ylim=range(polys$Y)+c(-0.1, 0.1), projection=1)
addPoints(events, col=2);
#--- run findCells and print the results
```

```

fc <- findCells(events, polys)
fc <- fc[order(fc$EID, fc$PID, fc$SID), ]
fc$label <- paste(fc$PID, fc$SID, sep=", ")
print (fc)
#--- add labels to the graph
addLabels(as.PolyData(fc[!duplicated(paste(fc$PID,fc$SID)), ],
                    projection=1),
          placement="CENTROID", polys=as.PolySet(polys, projection=1),
          col=4)

```

findPolys

Find the Polygons that Contain Events

Description

Find the polygons in a [PolySet](#) that contain events specified in [EventData](#).

Usage

```
findPolys (events, polys, maxRows = 1e+05)
```

Arguments

events	EventData to use.
polys	PolySet to use.
maxRows	estimated maximum number of rows in the output LocationSet .

Details

The resulting data frame, a [LocationSet](#), contains the columns EID, PID, SID (*if in polys*), and Bdry, where an event (EID) occurs in a polygon (PID, SID) and SID does not correspond to an inner boundary. The Boolean variable Bdry indicates whether an event lies on a polygon's edge. Note that if an event lies properly outside of all the polygons, then a record with (EID, PID, SID) does not occur in the output. It may happen, however, that an event occurs in multiple polygons. Thus, the same EID can occur more than once in the output.

Value

[LocationSet](#) that links events with polygons.

See Also

[combineEvents](#), [findCells](#), [locateEvents](#), [locatePolys](#), [LocationSet](#), [makeGrid](#).

Examples

```
#--- create some EventData: a column of points at X = 0.5
events <- data.frame(EID=1:10, X=.5, Y=seq(0, 2, length=10))
events <- as.EventData(events, projection=1);
#--- create a PolySet: two squares with the second above the first
polys <- data.frame(PID=c(rep(1, 4), rep(2, 4)), POS=c(1:4, 1:4),
                    X=c(0, 1, 1, 0, 0, 1, 1, 0),
                    Y=c(0, 0, 1, 1, 1, 1, 2, 2))
polys <- as.PolySet(polys, projection=1);
#--- show a picture
plotPolys(polys, xlim=range(polys$X)+c(-0.1, 0.1),
          ylim=range(polys$Y)+c(-0.1, 0.1), projection=1);
addPoints(events, col=2);
#--- run findPolys and print the results
print(findPolys(events, polys))
```

 fixBound

Fix the Boundary Points of a PolySet

Description

The ranges of a [PolySet](#)'s X and Y columns define its boundary. This function fixes a [PolySet](#)'s vertices by moving vertices near a boundary to the actual boundary.

Usage

```
fixBound (polys, tol)
```

Arguments

polys	PolySet to fix.
tol	vector (length 1 or 2) specifying a percentage of the ranges to use in defining <i>near</i> to a boundary. If tol has two elements, the first specifies the tolerance for the x-axis and the second the y-axis. If it has only one element, the function uses the same tolerance for both axes.

Details

When moving vertices to a boundary, the function moves them strictly horizontally or vertically, as appropriate.

Value

[PolySet](#) identical to the input, except for possible changes in the X and Y columns.

See Also

[closePolys](#), [fixPOS](#), [isConvex](#), [isIntersecting](#), [PolySet](#).

Examples

```

#--- set up a long horizontal and long vertical line to extend the plot's
#--- limits, and then try fixing the bounds of a line in the top-left
#--- corner and a line in the bottom-right corner
polys <- data.frame(PID=c(1, 1, 2, 2, 3, 3, 4, 4),
                   POS=c(1, 2, 1, 2, 1, 2, 1, 2),
                   X = c(0, 10, 5, 5, 0.1, 4.9, 5.1, 9.9),
                   Y = c(5, 5, 0, 10, 5.1, 9.9, 0.1, 4.9))
polys <- fixBound(polys, tol=0.0100001)
plotLines(polys)

```

fixPOS

*Fix the POS Column of a PolySet***Description**

Fix the POS column of a [PolySet](#) by recalculating it using sequential integers.

Usage

```
fixPOS (polys, exteriorCCW = NA)
```

Arguments

polys	PolySet to fix.
exteriorCCW	Boolean value; if TRUE, orders exterior polygon vertices in a counter-clockwise direction. If FALSE, orders them in a clockwise direction. If NA, maintains their original order.

Details

This function recalculates the POS values of each (PID, SID) as either 1 to N or N to 1, depending on the order of POS (ascending or descending) in the input data. POS values in the input must be properly ordered (ascending or descending), but they may contain fractional values. For example, POS = 2.5 might correspond to a point manually added between POS = 2 and POS = 3. If exteriorCCW = NA, all other columns remain unchanged. Otherwise, it orders the X and Y columns according to exteriorCCW.

Value

[PolySet](#) with the same columns as the input, except for possible changes to the POS, X, and Y columns.

See Also

[closePolys](#), [fixBound](#), [isConvex](#), [isIntersecting](#), [PolySet](#).

Examples

```
### create a PolySet with broken POS numbering
polys <- data.frame(PID = c(rep(1, 10), rep(2, 10)),
                   POS = c(seq(2, 10, length = 10), seq(10, 2, length = 10)),
                   X = c(rep(1, 10), rep(1, 10)),
                   Y = c(rep(1, 10), rep(1, 10)))

### fix the POS numbering
polys <- fixPOS(polys)
### print the results
print(polys)
```

importEvents

Import EventData from a Text File

Description

Import a text file and convert into EventData.

Usage

```
importEvents(EventData, projection=NULL, zone=NULL)
```

Arguments

EventData	filename of EventData text file.
projection	optional projection attribute to add to EventData.
zone	optional zone attribute to add to EventData.

Value

An imported EventData.

See Also

[importPolys](#), [importLocs](#), [importGSHHS](#), [importShapefile](#)

`importGSHHS`*Import Data from a GSHHS Database*

Description

Import data from a GSHHS database and convert data into a PolySet with a PolyData attribute.

Usage

```
importGSHHS(gshhsDB, xlim, ylim, level=1, n=0, xoff=-360)
```

Arguments

<code>gshhsDB</code>	filename of binary GSHHS database. If no file is specified, look for <code>gshhs_f.b</code> in the root of the PBSmapping library directory.
<code>xlim</code>	range of X-coordinates to clip. Range should match the transform <code>xoff</code> .
<code>ylim</code>	range of Y-coordinates to clip.
<code>level</code>	import polygons of level 1 (land), 2 (lakes on land), 3 (islands in lakes), 4 (ponds on islands). Multiple layers are specified in a vector.
<code>n</code>	retain polygons of at least <code>n</code> vertices.
<code>xoff</code>	transform the X-coordinates by specified amount.

Details

This routine requires a binary GSHHS (Global Self-consistent, Hierarchical, High-resolution Shoreline) database file. The GSHHS database has been released in the public domain and may be downloaded from <http://www.soest.hawaii.edu/wessel/gshhs/gshhs.html>. At the time of writing, the most recent database is `gshhs_1.5.zip`. This version uses binary packing of the header information.

This routine will return a PolySet object directly to R.

The PolyData attribute contains four fields: 1) PID which maps poly data to a polygon in the PolySet, 2) LEVEL to indicate the type of polygon (1=land, 2=lake, 3=island, 4=pond), 3) SOURCE to indicate the source of the data (1=WVS, not 1=CIA (WDBII)), and 4) GREENWICH (1 if poly crosses greenwich, 0 otherwise).

Value

A PolySet with a PolyData attribute.

See Also

[importEvents](#), [importLocs](#), [importPolys](#), [importShapefile](#)

Examples

```
## Not run:
#function to properly plot the 4 levels in order working
#from 1 (land) to 4 (ponds on islands in lakes on land)
plotInOrder <- function(polys, ...)
{
  myCols = c("gold1", "steelblue2", "gold1", "steelblue2")
  projection <- attr(polys, "projection")
  if (is.null(projection))
    stop("projection attribute is null")
  Pdata <- attr(polys, "PolyData")
  if (is.null(Pdata))
    stop("PolyData attribute is null")

  plotMap(NULL, xlim=c(min(polys$X), max(polys$X)),
          ylim=c(min(polys$Y), max(polys$Y)), projection=projection, ...)

  #extract all polygons of a certain level
  #(could probably be done more efficiently)
  for(i in 1:4) {
    PIDs <- Pdata[Pdata$LEVEL==i,]$PID
    if (length(PIDs)==0) next
    tmp <- NULL
    for (p in PIDs) {
      if (is.null(tmp))
        tmp <- polys[(polys$PID==p),]
      else
        tmp <- rbind(tmp, polys[(polys$PID==p),])
    }
    attr(tmp, "PolyData") <- Pdata
    addPolys(tmp, col=myCols[i])
  }
}

# You will have to download a copy of 'gshhs_f.b'
# from http://www.soest.hawaii.edu/wessel/gshhs/gshhs.html
# and place it in the PBSmapping library root.

# clip out Manitoulin Island area which includes all four levels
x <- importGSHHS(xlim=c(-84, -81), ylim=c(45.3, 46.5), level=1:4)

#plot map data and a label
plotInOrder(x, main="Manitoulin Island, Ontario, Canada")
text(-82.08, 45.706, "Manitoulin Isl")

## End(Not run)
```

Description

Import a text file and convert into a LocationSet.

Usage

```
importLocs(LocationSet)
```

Arguments

LocationSet filename of LocationSet text file.

Value

An imported LocationSet.

See Also

[importPolys](#), [importEvents](#), [importGSHHS](#), [importShapefile](#)

importPolys

Import PolySet from a text file

Description

Import a text file and convert into a PolySet with optional PolyData attribute.

Usage

```
importPolys(PolySet, PolyData=NULL, projection=NULL, zone=NULL)
```

Arguments

PolySet filename of PolySet text file.
PolyData optional filename of PolyData text file.
projection optional projection attribute to add to EventData.
zone optional zone attribute to add to EventData.

Value

An imported PolySet with optional PolyData attribute.

See Also

[importEvents](#), [importLocs](#), [importGSHHS](#), [importShapefile](#)

importShapefile	<i>Import an ESRI Shapefile</i>
-----------------	---------------------------------

Description

Import an ESRI shapefile (.shp) into either a [PolySet](#) or [EventData](#).

Usage

```
importShapefile (fn, readDBF = TRUE, projection = NULL, zone = NULL)
```

Arguments

fn	file name of the shapefile to import; specifying the extension is optional.
readDBF	Boolean value; if TRUE, it also imports the .dbf (a database containing the feature attributes) associated with the shapefile.
projection	optional projection attribute to override the internally derived value.
zone	optional zone attribute to override the default value of NULL.

Details

This routine imports an ESRI shapefile (.shp) into either a [PolySet](#) or [EventData](#), depending on the type of shapefile. It supports types 1 (Point), 3 (PolyLine), and 5 (Polygon) and imports type 1 into [EventData](#) and types 3 and 5 into a [PolySet](#). In addition to the shapefile (.shp), it requires the related index file (.shx).

If a database containing feature attributes (.dbf) exists, it also imports this database by default. For [EventData](#), it binds the database columns to the [EventData](#) object. For a [PolySet](#), it saves the database in a [PolyData](#) object and attaches that object to the [PolySet](#) in an attribute named "PolyData".

If a .prj file exists, this information is attached as an attribute. If the first 3 characters are 'GEO', then a geographic projection is assumed and projection="LL". If the first 4 characters are 'PROJ', and 'UTM' occurs elsewhere in the string, then the Universal Transverse Mercator projection is assumed and projection="UTM". Otherwise, projection=1.

If an .xml file exists, this information is attached as an attribute.

Value

For points, [EventData](#) with columns EID, X, and Y, possibly with other columns from the attribute database. For polylines and polygons, a [PolySet](#) with columns PID, SID, POS, X, Y and attribute projection. Other attributes that may or may not be attached: parent.child (boolean vector from original input), shpType (numeric shape type: 1, 3, or 5), prj (projection information from .prj file, xml (metadata from an .xml file), [PolyData](#) (data from the attribute database .dbf), and zone (UTM zone).

See Also

[importGSHHS](#), [importEvents](#), [importLocs](#), [importPolys](#)

isConvex

Determine Whether Polygons are Convex

Description

Determine whether polygons found in a [PolySet](#) are convex.

Usage

```
isConvex (polys)
```

Arguments

polys [PolySet](#) to use.

Details

Convex polygons do not self-intersect. In a convex polygon, only the first and last vertices may share the same coordinates (i.e., the polygons are optionally closed).

The function does not give special consideration to holes. It returns a value for each unique (PID, SID), regardless of whether a contour represents a hole.

Value

[PolyData](#) with columns PID, SID (*may be missing*), and convex. Column convex contains Boolean values.

See Also

[isIntersecting](#), [PolySet](#).

Examples

```
#--- load the data (if using R)
if (!is.null(version$language) && (version$language == "R"))
  data(nepacLL);
#--- calculate then print the polygons that are convex
p <- isConvex(nepacLL);
#--- nepacLL actually contains no convex polygons
print(p[p$convex, ]);
```

isIntersecting	<i>Determine Whether Polygons are Self-Intersecting</i>
----------------	---------------------------------------------------------

Description

Determine whether polygons found in a [PolySet](#) are self-intersecting.

Usage

```
isIntersecting (polys, numericResult = FALSE)
```

Arguments

`polys` [PolySet](#) to use.
`numericResult` Boolean value; if TRUE, returns the number of intersections.

Details

When `numericResult = TRUE`, this function counts intersections as the algorithm processes them. It counts certain types (i.e., those involving vertices and those where an edge retraces over an edge) more than once.

The function does not give special consideration to holes. It returns a value for each unique (PID, SID), regardless of whether a contour represents a hole.

Value

[PolyData](#) with columns PID, SID (*may be missing*), and `intersecting`. If `numericResult` is TRUE, `intersecting` contains the number of intersections. Otherwise, it contains a Boolean value.

See Also

[isConvex](#), [PolySet](#).

Examples

```
#--- load the data (if using R)
if (!is.null(version$language) && (version$language == "R"))
  data(nepacLL);
#--- calculate then print the polygons that are self-intersecting
p <- isIntersecting(nepacLL, numericResult = FALSE);
print(p[p$intersecting, ];
```

 joinPolys

 Join One or Two PolySets using a Logic Operation

Description

Join one or two [PolySets](#) using a logic operation.

Usage

```
joinPolys(polysA, polysB=NULL, operation="INT", maxVert=1e+05)
```

Arguments

polysA	PolySet to join.
polysB	optional second PolySet with which to join.
operation	one of "DIFF", "INT", "UNION", or "XOR", representing difference, intersection, union, and exclusive-or, respectively.
maxVert	estimated maximum number of vertices in the output PolySet .

Details

This function interfaces with the General Polygon Clipper library (<http://www.cs.man.ac.uk/aig/staff/alan/software/>) produced by Alan Murta at the University of Manchester. Consequently, we adopt some of his terminology in the details below.

Murta (2004) defines a *generic polygon* (or *polygon set*) as zero or more disjoint boundaries of arbitrary configuration. He relates a *boundary* to a contour, where each may be convex, concave or self-intersecting. In a [PolySet](#), the polygons associated with each unique PID loosely correspond to a generic polygon, as they can represent both inner and outer boundaries. Our use of the term *generic polygon* includes the restrictions imposed by a [PolySet](#). For example, the polygons for a given PID cannot be arranged arbitrarily.

If polysB is NULL, this function sequentially applies the operation between the generic polygons in polysA. For example, suppose polysA contains three generic polygons (A, B, C). The function outputs the [PolySet](#) containing ((A op B) op C).

If polysB is not NULL, this function applies operation between each generic polygon in polysA and each one in polysB. For example, suppose polysA contains two generic polygons (A, B) and polysB contains two generic polygons (C, D). The function's output is the concatenation of A op C, B op C, A op D, B op D, with PIDs 1 to 4, respectively. Generally there are n times m comparisons, where n = number of polygons in polysA and m = number of polygons in polysB. If polysB contains only one generic polygon, the function maintains the PIDs from polysA. It also maintains them when polysA contains only one generic polygon and the operation is difference. Otherwise, if polysA contains only one generic polygon, it maintains the PIDs from polysB.

Value

If polysB is NULL, the resulting **PolySet** contains a single generic polygon (one PID), possibly with several components (SIDs). The function recalculates the PID and SID columns.

If polysB is not NULL, the resulting **PolySet** contains one or more generic polygons (PIDs), each with possibly several components (SIDs). The function recalculates the SID column, and depending on the input, it may recalculate the PID column.

References

Murta, A. (2004) *A General Polygon Clipping Library*. Accessed: July 29, 2004.
<http://www.cs.man.ac.uk/aig/staff/alan/software/gpc.html>

See Also

[addPolys](#), [appendPolys](#), [clipPolys](#), [closePolys](#), [fixBound](#), [fixPOS](#), [locatePolys](#), [plotMap](#), [plotPoints](#), [thickenPolys](#), [thinPolys](#).

Examples

```
#--- load the data (if using R)
if (!is.null(version$language) && (version$language == "R"))
  data(nepacLL)
#--- create a triangle to use in clipping
polysB <- data.frame(PID=rep(1, 3), POS=1:3,
  X=c(-127.5, -124.5, -125.6), Y = c(49.2, 50.3, 48.6))
#--- intersect nepacLL with the single polygon, and plot the result
plotMap(joinPolys(nepacLL, polysB), col=5)
#--- add nepacLL in a different line type to emphasize the intersection
addPolys(nepacLL, border=2, lty=8, density=0)
```

 locateEvents

Locate Events on the Current Plot

Description

Locate events on the current plot (using the [locator](#) function).

Usage

```
locateEvents (EID, n = 512, type = "p", ...)
```

Arguments

EID	vector of event IDs (<i>optional</i>).
n	maximum number of events to locate.
type	one of "n", "p", "l", or "o". If "p" or "o", then the points are plotted; if "l" or "o", then the points are joined by lines.
...	additional par parameters for the locator function.

Details

This function allows its user to define events with mouse clicks on the current plot via the `locator` function. The arguments `n` and `type` are the usual parameters of the `locator` function. If `EID` is not missing, then `n = length(EID)`.

On exit from `locator`, suppose the user defined m events. If `EID` was missing, then the output data frame will contain m events. However, if `EID` exists, then the output data frame will contain `length(EID)` events, and both `X` and `Y` will be `NA` for events `EID[(m+1):n]`. The `na.omit` function can remove rows with `NAs`.

Value

`EventData` with columns `EID`, `X`, and `Y`, and projection attribute equal to the map's projection. The function does not set the zone attribute.

See Also

[addPoints](#), [combineEvents](#), [convDP](#), [EventData](#), [findCells](#), [findPolys](#), [plotPoints](#).

Examples

```
#--- define five events on the current plot, numbering them 10 to 14
## Not run: events <- locateEvents(EID = 10:14)
```

locatePolys

Locate Polygons on the Current Plot

Description

Locate polygons on the current plot (using the `locator` function).

Usage

```
locatePolys (pdata, n = 512, type = "o", ...)
```

Arguments

<code>pdata</code>	<code>PolyData</code> (<i>optional</i>) with columns <code>PID</code> and <code>SID</code> (<i>optional</i>), with two more optional columns <code>n</code> and <code>type</code> .
<code>n</code>	maximum number of points to locate.
<code>type</code>	one of "n", "p", "l", or "o". If "p" or "o", then the points are plotted; if "l" or "o", then the points are joined by lines.
<code>...</code>	additional <code>par</code> parameters for the <code>locator</code> function.

Details

This function allows its user to define polygons with mouse clicks on the current plot via the [locator](#) function. The arguments `n` and `type` are the usual parameters for the [locator](#) function, but the user can specify them for each individual (PID, SID) in a `pdata` object.

If a `pdata` object exists, the function ignores columns other than PID, SID, `n`, and `type`. If `pdata` includes `n`, then an outer boundary has `n > 0` and an inner boundary has `n < 0`.

On exit from [locator](#), suppose the user defined m vertices for a given polygon. For that polygon, the X and Y columns will contain NAs where $POS = (m+1):n$ for outer-boundaries and $POS = (|n|-m):1$ for inner-boundaries. The [na.omit](#) function can remove rows with NAs.

If a `pdata` object does not exist, the output contains only one polygon with a PID equal to 1. One inner-boundary polygon (POS goes from `n` to 1) can be generated by supplying a negative `n`.

If `type = "o"` or `type = "l"`, the function draws a line connecting the last and first vertices.

Value

[PolySet](#) with projection attribute equal to the map's projection. The function does not set the zone attribute.

See Also

[addPolys](#), [appendPolys](#), [clipPolys](#), [closePolys](#), [findCells](#), [findPolys](#), [fixPOS](#), [joinPolys](#), [plotMap](#), [plotPolys](#), [thickenPolys](#), [thinPolys](#).

Examples

```
#--- define one polygon with up to 5 vertices on the current plot
## Not run: polys <- locatePolys(n = 5)
```

LocationSet

LocationSet Objects

Description

PBS Mapping functions that expect `LocationSet`'s will accept properly formatted data frames in their place (see 'Details').

`as.LocationSet` attempts to coerce a data frame to an object with class `LocationSet`.

`is.LocationSet` returns TRUE if its argument is of class `LocationSet`.

Usage

```
as.LocationSet(x)
is.LocationSet(x, fullValidation = TRUE)
```

Arguments

`x` data frame to be coerced or tested.
`fullValidation` Boolean value; if TRUE, fully test `x`.

Details

A [PolySet](#) can define regional boundaries for drawing a map, and [EventData](#) can give event points on the map. Which events occur in which regions? Our function [findPolys](#) resolves this problem. The output lies in a [LocationSet](#), a data frame with three or four columns (EID, PID, SID, Bdry), where SID may be missing. One row in a [LocationSet](#) means that the event EID occurs in the polygon (PID, SID). The boundary (Bdry) field specifies whether (Bdry=T) or not (Bdry=F) the event lies on the polygon boundary. If SID refers to an inner polygon boundary, then EID occurs in (PID, SID) only if Bdry=T. An event may occur in multiple polygons. Thus, the same EID can occur in multiple records. If an EID does not fall in any (PID, SID), or if it falls within a hole, it does not occur in the output [LocationSet](#). Inserting the string "LocationSet" as the first element of a [LocationSet](#)'s class attribute alters the behaviour of some functions, including [print](#) (if [PBSprint](#) is TRUE) and [summary](#).

Value

The `as.LocationSet` method returns an object with classes "LocationSet" and "data.frame", in that order.

See Also

[EventData](#), [PolyData](#), [PolySet](#).

makeGrid	<i>Make a Grid of Polygons</i>
----------	--------------------------------

Description

Make a grid of polygons, using PIDs and SIDs according to the input arguments.

Usage

```
makeGrid(x,y,byrow=TRUE,addSID=TRUE,projection=NULL,zone=NULL)
```

Arguments

`x` vector of X-coordinates (of length m).
`y` vector of Y-coordinates (of length n).
`byrow` Boolean value; if TRUE, increment PID along X.
`addSID` Boolean value; if TRUE, include an SID column in the resulting [PolySet](#).
`projection` optional projection attribute to add to the [PolySet](#).
`zone` optional zone attribute to add to the [PolySet](#).

Details

This function makes a grid of polygons, labeling them according to byrow and addSID. In the following description, the variables i and j indicate column and row numbers, respectively, where the lower-left cell of the grid is (1, 1).

- byrow = TRUE and addSID = FALSE implies $PID = i + (j - 1) \times (m - 1)$
- byrow = FALSE and addSID = FALSE implies $PID = j + (i - 1) \times (n - 1)$
- byrow = TRUE and addSID = TRUE implies $PID = i, SID = j$
- byrow = FALSE and addSID = TRUE implies $PID = j, SID = i$

Value

PolySet with columns PID, SID (if addSID = TRUE), POS, X, and Y. The **PolySet** is a set of rectangular grid cells with vertices:

$(x_i, y_j), (x_{i+1}, y_j), (x_{i+1}, y_{j+1}), (x_i, y_{j+1})$.

See Also

[addPolys](#), [clipPolys](#), [combineEvents](#), [findCells](#), [findPolys](#), [PolySet](#), [thickenPolys](#).

Examples

```
#--- make a 10 x 10 grid
polyGrid <- makeGrid(x=0:10, y=0:10)
#--- plot the grid
plotPolys(polyGrid, density=0, projection=1)
```

makeProps

Make Polygon Properties

Description

Append a column for a polygon property (e.g., border or lty) to **PolyData** based on measurements in the **PolyData**'s Z column.

Usage

```
makeProps(pdata, breaks, propName="col", propVals=1:(length(breaks)-1))
```

Arguments

pdata	PolyData with a Z column.
breaks	either a vector of cut points or a scalar denoting the number of intervals that Z is to be cut into.
propName	name of the new column to append to pdata.
propVals	vector of values to associate with Z breaks.

Details

This function acts like the `cut` function to produce `PolyData` suitable for the `polyProps` plotting argument (see `addLabels`, `addLines`, `addPoints`, `addPolys`, `addStipples`, `plotLines`, `plotMap`, `plotPoints`, and `plotPolys`). The Z column of `pdata` is equivalent to the data vector `x` of the `cut` function.

Value

`PolyData` with the same columns as `pdata` plus an additional column `propName`.

See Also

`addLabels`, `addLines`, `addPoints`, `addPolys`, `addStipples`, `plotLines`, `plotMap`, `plotPoints`, `plotPolys`, `PolyData`, `PolySet`.

Examples

```
#--- create a PolyData object
pd <- data.frame(PID=1:10, Z=1:10)

#--- using 3 intervals, create a column named 'col' and populate it with
#--- the supplied values
makeProps(pdata = pd, breaks = 3, propName = "col",
          propVals = c(1:3))
```

makeTopography

Make Topography Data From Freely Available Online Data

Description

Make topography data suitable for the `contour` and `contourLines` functions using freely available global seafloor topography data.

Usage

```
makeTopography (dat, digits=2, func=NULL)
```

Arguments

<code>dat</code>	data frame with three optionally-named columns: X, Y, and Z. The columns must appear in that order.
<code>digits</code>	integer indicating the precision to be used by the function round on (X,Y) values.
<code>func</code>	function to summarize Z if (X,Y) points are duplicated. Defaults to <code>mean()</code> if no function is specified.

Details

Data obtained through the acquisition form at http://topex.ucsd.edu/cgi-bin/get_data.cgi is suitable for this function. `read.table` will import its ASCII files into R/S, creating the data argument for this function.

When creating data for regions with longitude values spanning -180° to 0° , consider subtracting 360 from the result's X coordinates (x).

When creating bathymetry data, consider negating the result's elevations (z) to give depths positive values.

Combinations of (X,Y) do not need to be complete ($z[x,y]=NA$) or unique ($z[x,y]=\text{func}(Z[x,y])$).

Value

List with elements x, y, and z. x and y are vectors, while z is a matrix with rownames x and colnames y. `contour` and `contourLines` expect data conforming to this list format.

See Also

`graphics::contour`, `grDevices::contourLines`, `convCP`.

Examples

```
#-- Example 1: Sample data frame and conversion.
file <- data.frame(X=c(1,1,2,2),Y=c(3,4,3,4),Z=c(5,6,7,8))
print(makeTopography(file));

#-- Example 2: Aleutian Islands bathymetry
require(PBSmapping);
isob <- c(100,500,1000,2500,5000);
icol <- rgb(0,0,seq(255,100,len=length(isob)),max=255);
afile <- paste(system.file(package="PBSmapping"),
"/Extra/aleutian.txt",sep="")
aleutian <- read.table(afile, header=FALSE,
  col.names=c("x","y","z"))
aleutian$x <- aleutian$x - 360
aleutian$z <- -aleutian$z
alBathy <- makeTopography(aleutian)
alCL <- contourLines(alBathy,levels=isob)
alCP <- convCP(alCL)
alPoly <- alCP$PolySet
attr(alPoly,"projection") <- "LL"
plotMap(alPoly,type="n");
addLines(alPoly,col=icol);
data(nepacLL); addPolys(nepacLL,col="gold");
legend(x="topleft",bty="n",col=icol,lwd=2,
  legend=as.character(isob));
```

nepacLL

Data: Shoreline of the NE Pacific Ocean (Normal Resolution)

Description

[PolySet](#) of polygons for the northeast Pacific Ocean shoreline.

Usage

```
data(nepacLL)
```

Format

Data frame consisting of 4 columns: PID = primary polygon ID, POS = position of each vertex within a given polygon, X = longitude coordinate, and Y = latitude coordinate. Attributes: projection = "LL".

Note

In R, the data must be loaded using the [data](#) function.

Source

Polygon data from the GSHHS (Global Self-consistent, Hierarchical, High-resolution Shoreline) database gshhs_h.b.

Download from <http://www.soest.hawaii.edu/wessel/gshhs/gshhs.html>

```
nepacLL <-importGSHHS("gshhs_h.b", xlim=c(-190,-110), ylim=c(34,72),  
                    level=1, n=15, xoff=-360)
```

References

Wessel, P. and Smith, W.H.F. (1996) A global, self-consistent, hierarchical, high-resolution shoreline database. *Journal of Geophysical Research* **101**, 8741–8743.

http://www.soest.hawaii.edu/pwessel/pwessel_pubs.html

See Also

Data: [nepacLLhigh](#), [worldLL](#), [worldLLhigh](#), [bcBathymetry](#)

[importGSHHS](#), [addPolys](#), [clipPolys](#), [plotPolys](#), [plotMap](#), [thickenPolys](#), [thinPolys](#)

`nepacLLhigh`*Data: Shoreline of the NE Pacific Ocean (High Resolution)*

Description

[PolySet](#) of polygons for the northeast Pacific Ocean shoreline.

Usage

```
data(nepacLLhigh)
```

Format

Data frame consisting of 4 columns: PID = primary polygon ID, POS = position of each vertex within a given polygon, X = longitude coordinate, and Y = latitude coordinate. Attributes: projection = "LL".

Note

In R, the data must be loaded using the [data](#) function.

Source

Polygon data from the GSHHS (Global Self-consistent, Hierarchical, High-resolution Shoreline) database `gshhs_f.b`.

Download from <http://www.soest.hawaii.edu/wessel/gshhs/gshhs.html>

```
nepacLLhigh <- importGSHHS("gshhs_f.b", xlim=c(-190,-110),
                          ylim=c(34,72), level=1, n=0, xoff=-360)
nepacLLhigh <- thinPolys(nepacLLhigh, tol=0.1, filter=3)
```

References

Wessel, P. and Smith, W.H.F. (1996) A global, self-consistent, hierarchical, high-resolution shoreline database. *Journal of Geophysical Research* **101**, 8741–8743.

http://www.soest.hawaii.edu/pwessel/pwessel_pubs.html

See Also

Data: [nepacLL](#), [worldLL](#), [worldLLhigh](#), [bcBathymetry](#)

[importGSHHS](#), [addPolys](#), [clipPolys](#), [plotPolys](#), [plotMap](#), [thickenPolys](#), [thinPolys](#)

PBSmapping

PBS Mapping: Draw Maps and Implement Other GIS Procedures

Description

This software has evolved from fisheries research conducted at the Pacific Biological Station (PBS) in Nanaimo, British Columbia, Canada. It extends the R language to include two-dimensional plotting features similar to those commonly available in a Geographic Information System (GIS). Embedded C code speeds algorithms from computational geometry, such as finding polygons that contain specified point events or converting between longitude-latitude and Universal Transverse Mercator (UTM) coordinates. It includes data for a global shoreline and other data sets in the public domain.

For a complete user's guide, see the file `PBSmapping-UG.pdf` in the R directory `.../library/PBSmapping/doc`.

PBSmapping includes 10 demos that appear as figures in the User's Guide. To see them, run the function `.PBSfigs()`.

More generally, a user can view all demos available from locally installed packages with the function `runDemos()` in our related (and recommended) package `PBSmodelling`.

PBSprint

Specify Whether to Print Summaries

Description

Specify whether PBS Mapping should print object summaries or not. If not, data objects are displayed as normal.

Usage

```
PBSprint
```

Details

If `PBSprint = TRUE`, the mapping software will print summaries rather than the data frames for `EventData`, `LocationSet`, `PolyData`, and `PolySet` objects. If `PBSprint = FALSE`, it will print the data frames.

This variable's default value is `FALSE`.

Value

`TRUE` or `FALSE`, depending on the user's preference.

See Also

[summary](#).

plotLines

*Plot a PolySet as PolyLines***Description**

Plot a [PolySet](#) as polylines.

Usage

```
plotLines (polys, xlim = NULL, ylim = NULL, projection = FALSE,
           plt = c(0.11, 0.98, 0.12, 0.88), polyProps = NULL,
           lty = NULL, col = NULL, bg = 0, axes = TRUE,
           tckLab = TRUE, tck = 0.014, tckMinor = 0.5 * tck, ...)
```

Arguments

polys	PolySet to plot (<i>required</i>).
xlim	range of X-coordinates.
ylim	range of Y-coordinates.
projection	desired projection when PolySet lacks a projection attribute; one of "LL", "UTM", or a numeric value. If Boolean, specifies whether to check polys for a projection attribute.
plt	four element numeric vector (x1, x2, y1, y2) giving the coordinates of the plot region measured as a fraction of the figure region. Set to NULL if mai in par is desired.
polyProps	PolyData specifying which polylines to plot and their properties. par parameters passed as direct arguments supersede these data.
lty	vector describing line types (cycled by PID).
col	vector describing colours (cycled by PID).
bg	background colour of the plot.
axes	Boolean value; if TRUE, plot axes.
tckLab	Boolean vector (length 1 or 2); if TRUE, label the major tick marks. If given a two-element vector, the first element describes the tick marks on the x-axis and the second element describes those on the y-axis.
tck	numeric vector (length 1 or 2) describing the length of tick marks as a fraction of the smallest dimension. If tckLab = TRUE, these tick marks will be automatically labelled. If given a two-element vector, the first element describes the tick marks on the x-axis and the second element describes those on the y-axis.
tckMinor	numeric vector (length 1 or 2) describing the length of tick marks as a fraction of the smallest dimension. These tick marks can not be automatically labelled. If given a two-element vector, the first element describes the tick marks on the x-axis and the second element describes those on the y-axis.
...	additional par parameters, or the arguments main, sub, xlab, or ylab for the title function.

Details

This function plots a [PolySet](#), where each unique (PID, SID) describes a polyline. It does not connect each polyline's last vertex to its first. Unlike [plotMap](#), the function ignores the aspect ratio. It clips polys to xlim and ylim before plotting.

The function creates a blank plot when polys equals NULL. In this case, the user must supply both xlim and ylim arguments. Alternatively, it accepts the argument type = "n" as part of ..., which is equivalent to specifying polys = NULL, but requires a [PolySet](#). In both cases, the function's behaviour changes slightly. To resemble the [plot](#) function, it plots the border, labels, and other parts according to [par](#) parameters such as col.

For additional help on the arguments lty and col, please see [par](#).

Value

[PolyData](#) consisting of the PolyProps used to create the plot.

Note

To satisfy the aspect ratio, this plotting routine resizes the plot region. Consequently, [par](#) parameters such as plt, mai, and mar will change. When the function terminates, these changes persist to allow for additions to the plot.

See Also

[addLines](#), [calcLength](#), [clipLines](#), [closePolys](#), [convLP](#), [fixBound](#), [fixPOS](#),
[locatePolys](#), [thinPolys](#), [thickenPolys](#).

Examples

```
#--- create a PolySet to plot
polys <- data.frame(PID=rep(1,4),POS=1:4,X=c(0,1,1,0),Y=c(0,0,1,1))
#--- plot the PolySet
plotLines(polys, xlim=c(-.5,1.5), ylim=c(-.5,1.5))
```

plotMap

Plot a PolySet as a Map

Description

Plot a [PolySet](#) as a map, using the correct aspect ratio.

Usage

```
plotMap (polys, xlim = NULL, ylim = NULL, projection = TRUE,
         plt = c(0.11, 0.98, 0.12, 0.88), polyProps = NULL,
         border = NULL, lty = NULL, col = NULL, colHoles = NULL,
         density = NA, angle = NULL, bg = 0, axes = TRUE,
         tckLab = TRUE, tck = 0.014, tckMinor = 0.5 * tck, ...)
```

Arguments

polys	PolySet to plot (<i>required</i>).
xlim	range of X-coordinates.
ylim	range of Y-coordinates.
projection	desired projection when PolySet lacks a projection attribute; one of "LL", "UTM", or a numeric value. If Boolean, specifies whether to check polys for a projection attribute.
plt	four element numeric vector (x1, x2, y1, y2) giving the coordinates of the plot region measured as a fraction of the figure region. Set to NULL if mai in par is desired.
polyProps	PolyData specifying which polygons to plot and their properties. par parameters passed as direct arguments supersede these data.
border	vector describing edge colours (cycled by PID).
lty	vector describing line types (cycled by PID).
col	vector describing fill colours (cycled by PID).
colHoles	vector describing hole colours (cycled by PID). The default, NULL, should be used in most cases as it renders holes transparent. colHoles is designed solely to eliminate retrace lines when images are converted to PDF format. If colHoles is specified, underlying information (i.e., previously plotted shapes) will be obliterated. If NA is specified, only outer polygons are drawn, consequently filling holes.
density	vector describing shading line densities (lines per inch, cycled by PID).
angle	vector describing shading line angles (degrees, cycled by PID).
bg	background colour of the plot.
axes	Boolean value; if TRUE, plot axes.
tckLab	Boolean vector (length 1 or 2); if TRUE, label the major tick marks. If given a two-element vector, the first element describes the tick marks on the x-axis and the second element describes those on the y-axis.
tck	numeric vector (length 1 or 2) describing the length of tick marks as a fraction of the smallest dimension. If tckLab = TRUE, these tick marks will be automatically labelled. If given a two-element vector, the first element describes the tick marks on the x-axis and the second element describes those on the y-axis.
tckMinor	numeric vector (length 1 or 2) describing the length of tick marks as a fraction of the smallest dimension. These tick marks can not be automatically labelled. If given a two-element vector, the first element describes the tick marks on the x-axis and the second element describes those on the y-axis.
...	additional par parameters, or the arguments main, sub, xlab, or ylab for the title function.

Details

This function plots a [PolySet](#), where each unique (PID, SID) describes a polygon. It connects each polygon's last vertex to its first. The function supports both borders (border, lty) and fills (col,

density, angle). When supplied with the appropriate arguments, it can draw only borders or only fills. Unlike [plotLines](#) and [plotPolys](#), it uses the aspect ratio supplied in the projection attribute of polys. If this attribute is missing, it attempts to use its projection argument. In the absence of both, it uses a default aspect ratio of 1:1. It clips polys to xlim and ylim before plotting.

The function creates a blank plot when polys equals NULL. In this case, the user must supply both xlim and ylim arguments. Alternatively, it accepts the argument type = "n" as part of ..., which is equivalent to specifying polys = NULL, but requires a [PolySet](#). In both cases, the function's behaviour changes slightly. To resemble the [plot](#) function, it plots the border, labels, and other parts according to [par](#) parameters such as col.

For additional help on the arguments border, lty, col, density, and angle, please see [polygon](#) and [par](#).

Value

[PolyData](#) consisting of the PolyProps used to create the plot.

Note

To satisfy the aspect ratio, this plotting routine resizes the plot region. Consequently, [par](#) parameters such as plt, mai, and mar will change. When the function terminates, these changes persist to allow for additions to the plot.

See Also

[addLabels](#), [addPolys](#), [addStipples](#), [clipPolys](#), [closePolys](#), [fixBound](#), [fixPOS](#), [locatePolys](#), [plotLines](#), [plotMap](#), [plotPoints](#), [thinPolys](#), [thickenPolys](#).

Examples

```
#--- create a PolySet to plot
polys <- data.frame(PID=rep(1,4),POS=1:4,X=c(0,1,1,0),Y=c(0,0,1,1))
#--- plot the PolySet
plotMap(polys,xlim=c(-.5,1.5),ylim=c(-.5,1.5),density=0,projection=1)
```

plotPoints

Plot EventData/PolyData as Points

Description

Plot [EventData/PolyData](#), where each unique EID or (PID, SID) describes a point.

Usage

```
plotPoints (data, xlim = NULL, ylim = NULL, projection = FALSE,
            plt = c(0.11, 0.98, 0.12, 0.88), polyProps = NULL,
            cex = NULL, col = NULL, pch = NULL, axes = TRUE,
            tckLab = TRUE, tck = 0.014, tckMinor = 0.5 * tck, ...)
```

Arguments

data	EventData or PolyData to plot (<i>required</i>).
xlim	range of X-coordinates.
ylim	range of Y-coordinates.
projection	desired projection when PolySet lacks a projection attribute; one of "LL", "UTM", or a numeric value. If Boolean, specifies whether to check polys for a projection attribute.
plt	four element numeric vector (x1, x2, y1, y2) giving the coordinates of the plot region measured as a fraction of the figure region. Set to NULL if mai in par is desired.
polyProps	PolyData specifying which points to plot and their properties. par parameters passed as direct arguments supersede these data.
cex	vector describing character expansion factors (cycled by EID or PID).
col	vector describing colours (cycled by EID or PID).
pch	vector describing plotting characters (cycled by EID or PID).
axes	Boolean value; if TRUE, plot axes.
tckLab	Boolean vector (length 1 or 2); if TRUE, label the major tick marks. If given a two-element vector, the first element describes the tick marks on the x-axis and the second element describes those on the y-axis.
tck	numeric vector (length 1 or 2) describing the length of tick marks as a fraction of the smallest dimension. If tckLab = TRUE, these tick marks will be automatically labelled. If given a two-element vector, the first element describes the tick marks on the x-axis and the second element describes those on the y-axis.
tckMinor	numeric vector (length 1 or 2) describing the length of tick marks as a fraction of the smallest dimension. These tick marks can not be automatically labelled. If given a two-element vector, the first element describes the tick marks on the x-axis and the second element describes those on the y-axis.
...	additional par parameters, or the arguments main, sub, xlab, or ylab for the title function.

Details

This function clips data to xlim and ylim before plotting. It only adds [PolyData](#) containing X and Y columns.

The function creates a blank plot when polys equals NULL. In this case, the user must supply both xlim and ylim arguments. Alternatively, it accepts the argument type = "n" as part of ..., which is equivalent to specifying polys = NULL, but requires a [PolySet](#). In both cases, the function's behaviour changes slightly. To resemble the [plot](#) function, it plots the border, labels, and other parts according to [par](#) parameters such as col.

For additional help on the arguments cex, col, and pch, please see [par](#).

Value

[PolyData](#) consisting of the PolyProps used to create the plot.

Note

To satisfy the aspect ratio, this plotting routine resizes the plot region. Consequently, `par` parameters such as `plt`, `mai`, and `mar` will change. When the function terminates, these changes persist to allow for additions to the plot.

See Also

[addPoints](#), [combineEvents](#), [convDP](#), [findPolys](#), [locateEvents](#).

Examples

```
#--- load the data (if using R)
if (!is.null(version$language) && (version$language == "R")) {
  data(nepacLL)
  data(surveyData)
}
#--- plot a map
plotMap(nepacLL, xlim=c(-136, -125), ylim=c(48, 57))
#--- add events
addPoints(surveyData, col=1:7)
```

plotPolys

Plot a PolySet as Polygons

Description

Plot a [PolySet](#) as polygons.

Usage

```
plotPolys (polys, xlim = NULL, ylim = NULL, projection = FALSE,
           plt = c(0.11, 0.98, 0.12, 0.88), polyProps = NULL,
           border = NULL, lty = NULL, col = NULL, colHoles = NULL,
           density = NA, angle = NULL, bg = 0, axes = TRUE,
           tckLab = TRUE, tck = 0.014, tckMinor = 0.5 * tck, ...)
```

Arguments

<code>polys</code>	PolySet to plot (<i>required</i>).
<code>xlim</code>	range of X-coordinates.
<code>ylim</code>	range of Y-coordinates.
<code>projection</code>	desired projection when PolySet lacks a projection attribute; one of "LL", "UTM", or a numeric value. If Boolean, specifies whether to check polys for a projection attribute.
<code>plt</code>	four element numeric vector (<code>x1</code> , <code>x2</code> , <code>y1</code> , <code>y2</code>) giving the coordinates of the plot region measured as a fraction of the figure region. Set to NULL if <code>mai</code> in <code>par</code> is desired.

<code>polyProps</code>	PolyData specifying which polygons to plot and their properties. <code>par</code> parameters passed as direct arguments supersede these data.
<code>border</code>	vector describing edge colours (cycled by PID).
<code>lty</code>	vector describing line types (cycled by PID).
<code>col</code>	vector describing fill colours (cycled by PID).
<code>colHoles</code>	vector describing hole colours (cycled by PID). The default, <code>NULL</code> , should be used in most cases as it renders holes transparent. <code>colHoles</code> is designed solely to eliminate retrace lines when images are converted to PDF format. If <code>colHoles</code> is specified, underlying information (i.e., previously plotted shapes) will be obliterated. If <code>NA</code> is specified, only outer polygons are drawn, consequently filling holes.
<code>density</code>	vector describing shading line densities (lines per inch, cycled by PID).
<code>angle</code>	vector describing shading line angles (degrees, cycled by PID).
<code>bg</code>	background colour of the plot.
<code>axes</code>	Boolean value; if <code>TRUE</code> , plot axes.
<code>tckLab</code>	Boolean vector (length 1 or 2); if <code>TRUE</code> , label the major tick marks. If given a two-element vector, the first element describes the tick marks on the x-axis and the second element describes those on the y-axis.
<code>tck</code>	numeric vector (length 1 or 2) describing the length of tick marks as a fraction of the smallest dimension. If <code>tckLab = TRUE</code> , these tick marks will be automatically labelled. If given a two-element vector, the first element describes the tick marks on the x-axis and the second element describes those on the y-axis.
<code>tckMinor</code>	numeric vector (length 1 or 2) describing the length of tick marks as a fraction of the smallest dimension. These tick marks can not be automatically labelled. If given a two-element vector, the first element describes the tick marks on the x-axis and the second element describes those on the y-axis.
<code>...</code>	additional <code>par</code> parameters, or the arguments <code>main</code> , <code>sub</code> , <code>xlab</code> , or <code>ylab</code> for the <code>title</code> function.

Details

This function plots a [PolySet](#), where each unique (PID, SID) describes a polygon. It connects each polygon's last vertex to its first. The function supports both borders (`border`, `lty`) and fills (`col`, `density`, `angle`). When supplied with the appropriate arguments, it can draw only borders or only fills. Unlike `plotMap`, it ignores the aspect ratio. It clips `polys` to `xlim` and `ylim` before plotting.

This function creates a blank plot when `polys` equals `NULL`. In this case, the user must supply both `xlim` and `ylim` arguments. Alternatively, it accepts the argument `type = "n"` as part of `...`, which is equivalent to specifying `polys = NULL`, but requires a [PolySet](#). In both cases, the function's behaviour changes slightly. To resemble the `plot` function, it plots the border, labels, and other parts according to `par` parameters such as `col`.

For additional help on the arguments `border`, `lty`, `col`, `density`, and `angle`, please see [polygon](#) and [par](#).

Value

[PolyData](#) consisting of the `PolyProps` used to create the plot.

Note

To satisfy the aspect ratio, this plotting routine resizes the plot region. Consequently, `par` parameters such as `plt`, `mai`, and `mar` will change. When the function terminates, these changes persist to allow for additions to the plot.

See Also

[addLabels](#), [addPolys](#), [addStipples](#), [clipPolys](#), [closePolys](#), [fixBound](#), [fixPOS](#), [locatePolys](#), [plotLines](#), [plotMap](#), [plotPoints](#), [thinPolys](#), [thickenPolys](#).

Examples

```
#--- create a PolySet to plot
polys <- data.frame(PID=rep(1,4),POS=1:4,X=c(0,1,1,0),Y=c(0,0,1,1))
#--- plot the PolySet
plotPolys(polys, xlim=c(-.5,1.5), ylim=c(-.5,1.5), density=0)
```

 PolyData

PolyData Objects

Description

PBS Mapping functions that expect PolyData will accept properly formatted data frames in their place (see 'Details').

`as.PolyData` attempts to coerce a data frame to an object with class PolyData.

`is.PolyData` returns TRUE if its argument is of class PolyData.

Usage

```
as.PolyData(x, projection = NULL, zone = NULL)
is.PolyData(x, fullValidation = TRUE)
```

Arguments

<code>x</code>	data frame to be coerced or tested.
<code>projection</code>	optional projection attribute to add to PolyData, possibly overwriting an existing attribute.
<code>zone</code>	optional zone attribute to add to PolyData, possibly overwriting an existing attribute.
<code>fullValidation</code>	Boolean value; if TRUE, fully test x.

Details

We define PolyData as a data frame with a first column named PID and (optionally) a second column named SID. Unlike a [PolySet](#), where each contour has many records corresponding to the vertices, a PolyData object must have only one record for each PID or each (PID, SID) combination. Conceptually, this object associates data with contours, where the data correspond to additional fields in the data frame. The R/S language conveniently allows data frames to contain fields of various atomic modes ("logical", "numeric", "complex", "character", and "null"). For example, PolyData with the fields (PID, PName) might assign character names to a set of primary polygons. Additionally, if fields X and Y exist (perhaps representing locations for placing labels), consider adding attributes zone and projection. Inserting the string "PolyData" as the class attribute's first element alters the behaviour of some functions, including `print` (if `PBSprint` is TRUE) and `summary`.

Our software particularly uses PolyData to set various plotting characteristics. Consistent with graphical parameters used by the R/S functions `lines` and `polygon`, column names can specify graphical properties:

- `lty` - line type in drawing the border and/or shading lines;
- `col` - line or fill colour;
- `border` - border colour;
- `density` - density of shading lines;
- `angle` - angle of shading lines.

When drawing polylines (as opposed to closed polygons), only `lty` and `col` have meaning.

Value

The `as.PolyData` method returns an object with classes "PolyData" and "data.frame", in that order.

See Also

[EventData](#), [LocationSet](#), [PolySet](#).

PolySet

PolySet Objects

Description

PBS Mapping functions that expect PolySet's will accept properly formatted data frames in their place (see 'Details').

`as.PolySet` attempts to coerce a data frame to an object with class PolySet.

`is.PolySet` returns TRUE if its argument is of class PolySet.

Usage

```
as.PolySet(x, projection = NULL, zone = NULL)
is.PolySet(x, fullValidation = TRUE)
```

Arguments

<code>x</code>	data frame to be coerced or tested.
<code>projection</code>	optional projection attribute to add to the PolySet, possibly overwriting an existing attribute.
<code>zone</code>	optional zone attribute to add to the PolySet, possibly overwriting an existing attribute.
<code>fullValidation</code>	Boolean value; if TRUE, fully test x.

Details

In our software, a PolySet data frame defines a collection of polygonal contours (i.e., line segments joined at vertices), based on four or five numerical fields:

- PID - the primary identification number for a contour;
- SID - optional, the secondary identification number for a contour;
- POS - the position number associated with a vertex;
- X - the horizontal coordinate at a vertex;
- Y - the vertical coordinate at a vertex.

The simplest PolySet lacks an SID column, and each PID corresponds to a different contour. By analogy with a child's "follow the dots" game, the POS field enumerates the vertices to be connected by straight lines. Coordinates (X, Y) specify the location of each vertex. Thus, in familiar mathematical notation, a contour consists of n points (x_i, y_i) with $i = 1, \dots, n$, where i corresponds to the POS index. A PolySet has two potential interpretations. The first associates a line segment with each successive pair of points from 1 to n , giving a *polyline* (in GIS terminology) composed of the sequential segments. The second includes a final line segment joining points n and 1, thus giving a *polygon*.

The secondary ID field allows us to define regions as composites of polygons. From this point of view, each primary ID identifies a collection of polygons distinguished by secondary IDs. For example, a single management area (PID) might consist of two fishing areas, each defined by a unique SID. A secondary polygon can also correspond to an inner boundary, like the hole in a doughnut. We adopt the convention that POS goes from 1 to n along an outer boundary, but from n to 1 along an inner boundary, regardless of rotational direction. This contrasts with other GIS software, such as ArcView (ESRI 1996), in which outer and inner boundaries correspond to clockwise and counter-clockwise directions, respectively.

The SID field in a PolySet with secondary IDs must have integer values that appear in ascending order for a given PID. Furthermore, inner boundaries must follow the outer boundary that encloses them. The POS field for each contour (PID, SID) must similarly appear as integers in strictly increasing or decreasing order, for outer and inner boundaries respectively. If the POS field erroneously contains floating-point numbers, `fixPOS` can renumber them as sequential integers, thus simplifying the insertion of a new point, such as point 3.5 between points 3 and 4.

A PolySet can have a `projection` attribute, which may be missing, that specifies a map projection. In the current version of PBS Mapping, projection can have character values "LL" or "UTM", referring to "Longitude-Latitude" and "Universal Transverse Mercator". We explain these projections more completely below. If projection is numeric, it specifies the aspect ratio r , the number of x units per y unit. Thus, r units of x on the graph occupy the same distance as one unit of y . Another optional attribute `zone` specifies the UTM zone (if `projection="UTM"`) or the preferred zone for conversion from Longitude-Latitude (if `projection="LL"`).

A data frame's class attribute by default contains the string "data.frame". Inserting the string "PolySet" as the class vector's first element alters the behaviour of some functions. For example, the `summary` function will print details specific to a PolySet. Also, when `PBSprint` is TRUE, the print function will display a PolySet's summary rather than the contents of the data frame.

Value

The `as.PolySet` method returns an object with classes "PolySet" and "data.frame", in that order.

References

Environmental Systems Research Institute (ESRI). (1996) *ArcView GIS: The Geographic Information System for Everyone*. ESRI Press, Redlands, California. 340 pp.

See Also

[EventData](#), [LocationSet](#), [PolyData](#).

print

Print PBS Mapping Objects

Description

This function displays information about a PBS Mapping object.

`summary.EventData`, `summary.LocationSet`, `summary.PolyData`, and `summary.PolySet` produce an object with class `summary.PBS`.

Usage

```
## S3 method for class 'EventData'
print(x, ...)
## S3 method for class 'LocationSet'
print(x, ...)
## S3 method for class 'PolyData'
print(x, ...)
## S3 method for class 'PolySet'
print(x, ...)
## S3 method for class 'summary.PBS'
print(x, ...)
```

Arguments

`x` a PBS Mapping object of appropriate class.
`...` additional arguments to `print`.

See Also

[EventData](#), [LocationSet](#), [PBSprint](#), [PolyData](#), [PolySet](#), [summary](#).

Examples

```
#--- load the data (if using R)
if (!is.null(version$language) && (version$language == "R"))
  data(nepacLL)
#--- change to summary printing style
PBSprint <- TRUE
#--- print the PolySet
print(nepacLL)
```

pythagoras

Data: Pythagoras' Theorem Diagram PolySet

Description

[PolySet](#) of shapes to prove Pythagoras' Theorem: $a^2 + b^2 = c^2$.

Usage

```
data(pythagoras)
```

Format

4 column data frame: PID = primary polygon ID, POS = position of each vertex within a given polyline, X = X-coordinate, and Y = Y-coordinate. Attributes: `projection = 1`.

Note

In R, the data must be loaded using the [data](#) function.

Source

An artificial construct to illustrate the proof of Pythagoras' Theorem using trigonometry.

See Also

[addPolys](#), [plotPolys](#), [plotMap](#), [PolySet](#).

summary

Summarize PBS Mapping Objects

Description

summary method for PBS Mapping classes.

Usage

```
## S3 method for class 'EventData'  
summary(object, ...)  
## S3 method for class 'LocationSet'  
summary(object, ...)  
## S3 method for class 'PolyData'  
summary(object, ...)  
## S3 method for class 'PolySet'  
summary(object, ...)
```

Arguments

object	a PBS Mapping object, such as EventData, a LocationSet, PolyData, or a PolySet.
...	further arguments passed to or from other methods.

Details

After creating a list of summary statistics, this function assigns the class "summary.PBS" to the output in order to accomplish formatted printing via `print.summary.PBS`.

Value

A list of summary statistics.

See Also

[EventData](#), [LocationSet](#), [PBSprint](#), [PolyData](#), [PolySet](#).

Examples

```
#--- load the data (if using R)  
if (!is.null(version$language) && (version$language == "R"))  
  data(surveyData)  
print(summary(surveyData))
```

surveyData

Data: Tow Information from Pacific Ocean Perch Survey

Description

[EventData](#) of Pacific ocean perch (POP) tow information (1966-89).

Usage

```
data(surveyData)
```

Format

Data frame consisting of 9 columns: PID = primary polygon ID, POS = position of each vertex within a given polygon, X = longitude coordinate, Y = latitude coordinate, trip = trip ID, tow = tow number in trip, catch = catch of POP (kg), effort = tow effort (minutes), depth = fishing depth (m), and year = year of survey trip. Attributes: projection = "LL", zone = 9.

Note

In R, the data must be loaded using the [data](#) function.

Source

The GFBio database, maintained at the Pacific Biological Station (Fisheries and Oceans Canada, Nanaimo, BC V9T 6N7), archives catches and related biological data from commercial groundfish fishing trips and research/assessment cruises off the west coast of British Columbia (BC).

The POP (*Sebastes alutus*) survey data were extracted from GFBio. The data extraction covers bottom trawl surveys that focus primarily on POP biomass estimation: 1966-89 for the central BC coast and 1970-85 for the west coast of Vancouver Island. Additionally, a 1989 cruise along the entire BC coast concentrated on the collection of biological samples. Schnute et al. (2001) provide a more comprehensive history of POP surveys including the subset of data presented here.

References

Schnute, J.T., Haigh, R., Krishka, B.A. and Starr, P. (2001) Pacific ocean perch assessment for the west coast of Canada in 2001. *Canadian Science Advisory Secretariat, Research Document 2001/138*, 90 pp.

See Also

[addPoints](#), [combineEvents](#), [EventData](#), [findPolys](#), [makeGrid](#), [plotPoints](#).

thickenPolys	<i>Thicken a PolySet of Polygons</i>
--------------	--------------------------------------

Description

Thicken a [PolySet](#), where each unique (PID, SID) describes a polygon.

Usage

```
thickenPolys (polys, tol = 1, filter = 3, keepOrig = TRUE,
             close = TRUE)
```

Arguments

polys	PolySet to thicken.
tol	tolerance (in kilometers when proj is "LL" and "UTM"; otherwise, same units as polys).
filter	minimum number of vertices per result polygon.
keepOrig	Boolean value; if TRUE, keep the original points in the PolySet .
close	Boolean value; if TRUE, create intermediate vertices between each polygon's last and first vertex, if necessary.

Details

This function thickens each polygon within polys according to the input arguments.

If `keepOrig = TRUE`, all of the original vertices appear in the result. It calculates the distance between two sequential original vertices, and if that distance exceeds `tol`, it adds a sufficient number of vertices spaced evenly between the two original vertices so that the distance between vertices no longer exceeds `tol`. If `close = TRUE`, it adds intermediate vertices between the last and first vertices when necessary.

If `keepOrig = FALSE`, only the first vertex of each polygon is guaranteed to appear in the results. From this first vertex, the algorithm walks the polygon summing the distance between vertices. When this cumulative distance exceeds `tol`, it adds a vertex on the line segment under inspection. After doing so, it resets the distance sum, and walks the polygon from this new vertex. If `close = TRUE`, it will walk the line segment from the last vertex to the first.

Value

[PolySet](#) containing the thickened data. The function recalculates the POS values for each polygon.

See Also

[thinPolys](#).

Examples

```
#--- load the data (if using R)
if (!is.null(version$language) && (version$language == "R"))
  data(nepacLL)
#--- plot Vancouver Island
plotMap(nepacLL[nepacLL$PID == 33, ])
#--- calculate a thickened version using a 30 kilometers tolerance,
#--- without keeping the original points
p <- thickenPolys(nepacLL[nepacLL$PID == 33, ], tol = 30, keepOrig = FALSE)
#--- convert the PolySet to EventData by dropping the PID column and
#--- renaming POS to EID
p <- p[-1];
names(p)[1] <- "EID";
#--- convert the now invalid PolySet into a data frame, and then into
#--- EventData
p <- as.EventData(as.data.frame(p), projection="LL");
#--- plot the results
addPoints(p, col=2, pch=19)
```

thinPolys

*Thin a PolySet of Polygons***Description**

Thin a [PolySet](#), where each unique (PID, SID) describes a polygon.

Usage

```
thinPolys (polys, tol = 1, filter = 3)
```

Arguments

polys	PolySet to thin.
tol	tolerance (in kilometers when proj is "LL" and "UTM"; otherwise, same units as polys).
filter	minimum number of vertices per result polygon.

Details

This function executes the Douglas-Peucker line simplification algorithm on each polygon within polys.

Value

[PolySet](#) containing the thinned data. The function recalculates the POS values for each polygon.

See Also

[thickenPolys](#).

Examples

```
#--- load the data (if using R)
if (!is.null(version$language) && (version$language == "R"))
  data(nepacLL)
#--- plot a thinned version of Vancouver Island (3 km tolerance)
plotMap(thinPolys(nepacLL[nepacLL$PID == 33, ], tol = 3))
#--- add the original Vancouver Island in a different line type to
#--- emphasize the difference
addPolys(nepacLL[nepacLL$PID == 33, ], border=2, lty=8, density=0)
```

towData

Data: Tow Information from Longspine Thornyhead Survey

Description

[PolyData](#) of tow information for a longspine thornyhead survey (2001).

Usage

```
data(towData)
```

Format

Data frame consisting of 8 columns: PID = primary polygon ID, POS = position of each vertex within a given polygon, X = longitude coordinate, Y = latitude coordinate, depth = fishing depth (m), effort = tow effort (minutes), distance = tow track distance (km), catch = catch of longspine thornyhead (kg), and year = year of survey. Attributes: projection = "LL", zone = 9.

Note

In R, the data must be loaded using the [data](#) function.

Source

The GFBio database, maintained at the Pacific Biological Station (Fisheries and Oceans Canada, Nanaimo, BC V9T 6N7), archives catches and related biological data from commercial groundfish fishing trips and research/assessment cruises off the west coast of British Columbia (BC). The longspine thornyhead (*Sebastolobus altivelis*) survey data were extracted from GFBio. Information on the first 45 tows from the 2001 survey (Starr et al. 2002) are included here. Effort is time (minutes) from winch lock-up to winch release.

References

Starr, P.J., Krishka, B.A. and Choromanski, E.M. (2002) Trawl survey for thornyhead biomass estimation off the west coast of Vancouver Island, September 15 - October 2, 2001. *Canadian Technical Report of Fisheries and Aquatic Sciences* **2421**, 60 pp.

See Also

[makeProps](#), [PolyData](#), [towTracks](#).

towTracks

Data: Tow Track Polylines from Longspine Thornyhead Survey

Description

[PolySet](#) of geo-referenced polyline tow track data from a longspine thornyhead survey (2001).

Usage

```
data(towTracks)
```

Format

Data frame consisting of 4 columns: PID = primary polygon ID, POS = position of each vertex within a given polyline, X = longitude coordinate, and Y = latitude coordinate. Attributes: projection = "LL", zone = 9.

Note

In R, the data must be loaded using the [data](#) function.

Source

The longspine thornyhead (*Sebastolobus altivelis*) tow track spatial coordinates are available at the Pacific Biological Station (Fisheries and Oceans Canada, Nanaimo, BC V9T 6N7). The geo-referenced coordinates of the first 45 tows from the 2001 survey (Starr et al. 2002) are included here. Coordinates are recorded once per minute between winch lock-up and winch release.

References

Starr, P.J., Krishka, B.A. and Choromanski, E.M. (2002) Trawl survey for thornyhead biomass estimation off the west coast of Vancouver Island, September 15 - October 2, 2001. *Canadian Technical Report of Fisheries and Aquatic Sciences* **2421**, 60 pp.

See Also

[addLines](#), [calcLength](#), [cliplines](#), [plotLines](#), [PolySet](#), [towData](#).

`worldLL`*Data: Shorelines of the World (Normal Resolution)*

Description

[PolySet](#) of polygons for the global shorelines.

Usage

```
data(worldLL)
```

Format

Data frame consisting of 4 columns: PID = primary polygon ID, POS = position of each vertex within a given polygon, X = longitude coordinate, and Y = latitude coordinate. Attributes: projection = "LL".

Note

In R, the data must be loaded using the [data](#) function.

Source

Polygon data from the GSHHS (Global Self-consistent, Hierarchical, High-resolution Shoreline) database `gshhs_1.b`.

Download from <http://www.soest.hawaii.edu/wessel/gshhs/gshhs.html>

```
worldLL <-importGSHHS("gshhs_1.b", xlim=c(-20,360), ylim=c(-90,90),
                    level=1, n=15, xoff=0)
worldLL <- .fixGSHHSWorld(worldLL)
```

References

Wessel, P. and Smith, W.H.F. (1996) A global, self-consistent, hierarchical, high-resolution shore-line database. *Journal of Geophysical Research* **101**, 8741–8743.

http://www.soest.hawaii.edu/pwessel/pwessel_pubs.html

See Also

Data: [worldLLhigh](#), [nepacLL](#), [nepacLLhigh](#)

[importGSHHS](#), [addPolys](#), [clipPolys](#), [plotPolys](#), [plotMap](#), [thickenPolys](#), [thinPolys](#)

`worldLLhigh`*Data: Shorelines of the World (High Resolution)*

Description

`PolySet` of polygons for the global shorelines.

Usage

```
data(worldLLhigh)
```

Format

Data frame consisting of 4 columns: PID = primary polygon ID, POS = position of each vertex within a given polygon, X = longitude coordinate, and Y = latitude coordinate. Attributes: projection = "LL".

Note

In R, the data must be loaded using the `data` function.

Source

Polygon data from the GSHHS (Global Self-consistent, Hierarchical, High-resolution Shoreline) database `gshhs_i.b`.

Download from <http://www.soest.hawaii.edu/wessel/gshhs/gshhs.html>

```
worldLLhigh <-importGSHHS("gshhs_i.b", xlim=c(-20,360),
                        ylim=c(-90,90), level=1, n=15, xoff=0)
worldLLhigh <- .fixGSHHSWorld(worldLLhigh)
```

References

Wessel, P. and Smith, W.H.F. (1996) A global, self-consistent, hierarchical, high-resolution shoreline database. *Journal of Geophysical Research* **101**, 8741–8743.

http://www.soest.hawaii.edu/pwessel/pwessel_pubs.html

See Also

Data: `worldLL`, `nepacLL`, `nepacLLhigh`

`importGSHHS`, `addPolys`, `clipPolys`, `plotPolys`, `plotMap`, `thickenPolys`, `thinPolys`

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