

# Package ‘loa’

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**Type** Package

**Title** Lattice Options and Add-Ins

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**Description** Various plots and functions that make use of the lattice/trellis plotting framework. The plots, which include loaPlot(), GoogleMap() and trianglePlot(), use panelPal(), a function that extends 'lattice' and 'hexbin' package methods to automate plot subscript and panel-to-panel and panel-to-key synchronization/management.

**Depends** R (>= 3.0.0), lattice

**Imports** methods, MASS, grid, png, RgoogleMaps, RColorBrewer, mgcv, plyr

**License** GPL (>= 2)

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## Description

The loa package contains various plots, options and add-ins for use with the [lattice](#) package.

## Details

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[lattice](#) provides an elegant and highly powerful implementation of the Trellis plotting structure described by Cleveland and colleagues. In particular the combination of `panel...` functions, which can be layered within plots to generate novel visualisations, and simple-to-use conditioning make it a hugely effective tool when working with data.

The loa package contains a number of plot functions developed to make use of this framework. These are summarized in section 1 of this manual, and include:

- 1.1. [loaPlot](#) for various XYZ plots.

- 1.2. [GoogleMap](#) and associated geoplottng functions.
- 1.3. [trianglePlot](#) and associated functions.
- 1.4. [stackPlot](#) and associated functions.
- 1.5. [loaBarPlot](#) and associated functions.

Other panel . . . functions and example data are summarized in sections 2 and 3, respectively:

- 2.1. Specialist panels, e.g. [panel.kernelDensity](#),
- 2.2. Specialist panels for polar plotting, e.g. [panel.polarPlot](#).
- 3.1. Example data, e.g. [lat.lon.meuse](#).

While such 'stand alone' plot functions are of obvious value, the code framework is of possibly wider interest because it provides a template for the rapid third-party development of novel visualization functions and a highly flexible 'test bed' for the comparison of different data handling strategies.

Therefore, the functions in this package have been written in a relatively disaggregated fashion so code can be easily rearranged or modified by others to quickly develop alternative plots within the [lattice](#) framework. Firstly, plot functions in section 1 have where possible been supplied as main plot functions and plot component functions that handle data, axes, panels, etc. Secondly, the workhorse functions, those common functions used through-out the package to simplify many routine operations have been grouped together and summarized in section 4:

- 4.1. [panelPal](#)
- 4.2. plot structure handlers: [formulaHandler](#), etc.
- 4.3. Plot lim(s) and scale(s) handlers: [limsHandler](#), [localScalesHandler](#), etc.
- 4.4. Plot conditioning handlers: [condsPanelHandler](#), etc.
- 4.5. Common plot argument handlers: [cexHandler](#), [colHandler](#), [zHandler](#), etc.
- 4.6. Key handlers: [keyHandler](#), etc.
- 4.7. Other panel functions: [getArgs](#), etc.
- 4.8. List handlers: [listHandler](#), etc.

And, finally, functions used for working with data post-plotting, are summarized in section 5:

- 5.1. Interactive functions for working with plot outputs: [getXY](#), etc.

This package is very much intended to be an evolutionary exercise. I use it on a routine basis to develop plots for use elsewhere and compare data visualization methods. However, that working pattern can generate some very 'developer-centric' code. So, I would be very pleased to hear from others - what they did and did not like about the package; what they would have liked to have been different; and, perhaps most interesting for me what they are using it to do.

## Author(s)

Karl Ropkins <k.ropkins@its.leeds.ac.uk>

## References

Functions in `loa` make extensive use of code developed by others. In particular, I gratefully acknowledge the huge contributions of:

`lattice`: Sarkar, Deepayan (2008) *Lattice: Multivariate Data Visualization with R*. Springer, New York. ISBN 978-0-387-75968-5

`Trellis Plotting`: Becker, R. A., Cleveland, W. S., Shyu, M. J. (1996). The Visual Design and Control of Trellis Display, *Journal of Computational and Graphical Statistics*, 5(2), 123-155. Cleveland, W.S. (1993) *Visualizing Data*, Hobart Press, Summit, New Jersey.

## See Also

[loaPlot](#), [GoogleMap](#), [trianglePlot](#)

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1.1.loaPlot

*loaPlot, XYZ plots for lattice*

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## Description

`loaPlot` is a standard XYZ plotting function, where X and Y are the axes of a conventional XY plot and Z is an element (or elements if supplied in the form  $Z1 + Z2 + Z3\dots$ ) visualized at associated XY coordinates. By default, `loaPlot` links Z to plot point size and color to generate a bubbleplot style output, or using modified plot calls other plot types.

## Usage

```
loaPlot(x, data = NULL, panel = panel.loaPlot,
        ..., local.scales = FALSE, reset.xylimits = TRUE,
        load.lists = NULL, by.group = NULL, by.zcase = NULL,
        preprocess = TRUE)
```

#standard panels

```
panel.loaPlot(..., loa.settings = FALSE)
panel.loaPlot2(..., loa.settings = FALSE)
```

#grids

```
panel.loaGrid(grid.x = NULL, grid.y = NULL,
              xlim = NULL, ylim = NULL, ...,
              grid = NULL, panel.scales = NULL)
```

**Arguments**

<code>x</code>	A formula with the general structure $z \sim x * y \mid \text{cond}$ applied like in the <code>lattice</code> function <code>levelplot</code> or a matrix. For a formula, <code>x</code> and <code>y</code> are the horizontal and vertical axes, <code>z</code> is any additional information to be used in point, symbol, surface or glyph generation, and <code>cond</code> is any additional conditioning to be applied. <code>x</code> and <code>y</code> are required elements; <code>z</code> and <code>cond</code> are typically optional. (Note: this element of the plot is handled by <code>formulaHandler</code> ).
<code>data</code>	If supplied, the assumed source of elements of <code>x</code> , typically a <code>data.frame</code> .
<code>panel</code>	<code>panel</code> is the function to be used when generating the content of the individual panels within the <code>lattice</code> plot. By default, this is the <code>loa</code> panel function <code>panel.loaPlot</code> .
<code>...</code>	Additional arguments are passed on to related functions. For <code>loaPlot</code> these are <code>colHandler</code> , <code>cexHandler</code> and the function set by <code>panel</code> . This mechanism provides access to most common plot parameters, e.g. <code>col</code> , <code>pch</code> , and <code>cex</code> for plot symbol color, type and size, respectively. By default, both data point color and size are z-scaled for <code>loaPlot</code> . If <code>z</code> is supplied, and <code>cex</code> and <code>col</code> are not set by the user in the plot command, these plot properties are managed by <code>cexHandler</code> and <code>colHandler</code> , respectively. <code>cexHandler</code> and <code>colHandler</code> arguments can be also be passed directly as part of the <code>loaPlot</code> command to fine-tune these, e.g. <code>cex.range</code> to change the <code>cex</code> range that <code>z</code> values are scaled to and <code>col.region</code> to change the color range that is applied to <code>z</code> when coloring points. See associated Help documents for further information.
<code>local.scales</code>	For <code>loaPlot</code> only, logical. If <code>TRUE</code> , this removes the standard <code>lattice</code> axis from the plot. It is intended to be used with <code>panel</code> functions which generate their own axes or have no axes.
<code>reset.xylimits</code>	For <code>loaPlot</code> only, logical or character vector. If a logical, if the panel outputs are preprocessed (using <code>panelPal</code> ), should the <code>x</code> and <code>y</code> limits be reset? If a character vector, one or more terms controlling post-processing plot range management: <code>refit.xylimits</code> , equivalent to <code>reset.xylimits = TRUE</code> ; and <code>max.xylimits</code> , to reset both <code>x</code> and <code>y</code> ranges to maximum. (Note: If <code>xlim</code> or <code>ylim</code> are supplied in the plot call, these will typically override all <code>reset.xylimits</code> settings.)
<code>load.lists</code>	For <code>loaPlot</code> only, character vector. In-development alternative to list based arguments. This option identifies plot call arguments that <code>loaPlot</code> should manage using <code>listLoad</code> . See associated help documentation for further details.
<code>by.group</code> , <code>by.zcase</code>	For <code>loaPlot</code> only. Arguments for routine by group and by <code>zcase</code> handling of plot inputs. Important: These are current under review.
<code>preprocess</code>	For <code>loaPlot</code> only, logical, passed to <code>panelPal</code> . If <code>TRUE</code> , and used with a correctly configured <code>panel</code> function, this processes the plot input before generating the plot. This means color scales in the different plot panels and the key are automatically aligned and the associated trellis object output contains the <code>panel</code> function outputs rather than the inputs. See <code>panelPal</code> Help documents for further information.

`loa.settings` For `panel...` functions only, logical, passed to `panelPal` to manage plot reworking. See associated Help documents for further information.

`grid.x`, `grid.y`, `xlim`, `ylim`, `grid`, `panel.scales`  
 For `panel.loaGrid` only, grid settings, typically recovered by `loaPlot`.

### Details

`loaPlot` provides lattice-style conditioning/handling for a range of commonly used XYZ plotting options. It is perhaps easiest pictured as a 'mid point' alternative somewhere between the standard `lattice` plot functions `xyplot` and `levelplot`.

The default form of the plot uses an extension of the subscripting methods described by Deepayan Sarkar in Chapter 5 of *Lattice* (see sections on scatterplots and extensions). The default output is a bubble plot (see example 1 below).

### Value

`loaPlot` returns a trellis object, much like a conventional `lattice` plot function.

### Note

`panel.loaPlot2` is an alternative versions of `panel.loaPlot` that is currently under revision. Please use with care.

`loaPlot` arguments `by.group` and `by.zcase` are currently in revision. Please use with care.

### Author(s)

Karl Ropkins

### References

These functions make extensive use of code developed by others.

`lattice`: Sarkar, Deepayan (2008) *Lattice: Multivariate Data Visualization with R*. Springer, New York. ISBN 978-0-387-75968-5

`RColorBrewer`: Erich Neuwirth <erich.neuwirth@univie.ac.at> (2011). `RColorBrewer: ColorBrewer palettes`. R package version 1.0-5. <http://CRAN.R-project.org/package=RColorBrewer>

### See Also

In `loa`: `panelPal`

In other packages, see

`lattice`: `xyplot`; and `levelplot`.

**Examples**

```
## Example 1
## Basic usage

loaPlot(Ozone~Solar.R*Temp|Wind>8,
        data=airquality,
        col.regions="Blues")

# Notes:
# Formula structure z ~ x * y |cond like levelplot.
# Data (and groups) assignment like in standard lattice plots.
# By default z is linked to col and cex.
# Unless overridden by user inputs or group or zcase setting.
# Plot passed via ...Handler functions to provide shortcut plot
# reworking, e.g. here colHandler handles color scales
# using col.region to generate a color range.
# (Here, arguments like "Blues" and c("green", "red") are
# allowed and handled using functions in the RColorBrewer
# package.)

# Formula structures:
# ~ x * y           like xyplot y ~ x
# ~ x * y | cond    like xyplot y ~ x | cond
# z ~ x * y         like xyplot y ~ x, col=f(z), cex=f(z)
# z ~ x * y | cond  like xyplot y ~ x | cond, col=f(z), cex=f(z)
# z ~ x * y, groups = g like xyplot y ~ x, groups=g, cex=f(z)
# z1 + z2 ~ x * y   (zcases)
# etc

## Example 2
## Basic modifications

loaPlot(Ozone~Solar.R*Temp, groups=airquality$Wind>8,
        data=airquality)

# When groups are applied, by default group id is linked to col.
# The follow example illustrates three options:

loaPlot(Ozone~Solar.R*Temp, groups=airquality$Wind>8,
        data=airquality,
        group.args=c("pch"), pch=c(1,4),
        col="blue")

# notes:
# Here, group.args is used to change the default group arguments.
# (So, pch rather than col is used to identify groups.)
# pch is then assigned by group rather than by (x,y) case or z case.
# (See panelPal Help further further details of assignments in loa.)
# col supplied by the user supercedes the default z linkage.
# (So, here cex remains z scales but col is fixed as blue.)
```

```

## Example 3
## Key handling

loadPlot(Ozone~Solar.R*Temp, data=airquality,
         col.regions=c("green", "red"))

# Key settings are by the key argument (as in lattice)
# or key... arguments via keyHandler and listLoad, so e.g.:

loadPlot(Ozone~Solar.R*Temp, data=airquality,
         col.regions=c("green", "red"),
         key.fun = draw loaColorKey)

# Notes:
# By default the loadPlot uses draw loaPlotZKey to generate
# its color key unless an alternative is supplied via key.fun.
# (Here, the draw.colorKey wrapper draw loaColorKey is used to
# generate a color bar similar to that in levelplot.)

## Example 4
## panels

loadPlot(Ozone~Solar.R*Temp|Wind>8, data=airquality,
         col.regions="Reds")

# The combined use of loadPlot, panelPal and appropriately configured
# panel functions provides automatic handling of a range of plot
# elements, e.g.:

loadPlot(Ozone~Solar.R*Temp|Wind>8, data=airquality,
         col.regions="Reds", panel=panel.binPlot)

# Notes:
# Here, the choice of default key is set by the panel... function;
# the panel by default bins data by location and for each bin cell
# calculates the mean Ozone concentration just like a standard
# lattice panel would, but it also tracks these values (calculated
# within the panels) and scales panel-to-panel and panel-to-key
# so users do not have to do that retrospectively; and, finally,
# it retains in-panel calculations so users can recover them.
# (See associated helps for further details: ?panelPal about methods;
# and ?panel.binPlot about the panel function.)

```



**Description**

Plotting georeferenced data on maps using lattice and RgoogleMaps

**Usage**

```

GoogleMap(x, data = NULL, panel = panel.loaPlot,
          map = NULL, map.panel = panel.GoogleMapsRaster,
          recolor.map = FALSE, ..., lon.lat = FALSE)

GoogleMap.old(x, data = NULL, map = NULL,
              map.panel = panel.GoogleMapsRaster,
              panel = panel.xyplot,
              recolor.map = FALSE, ...)

googleMap(...)

quickMap(lat, lon, show.data = FALSE, ...)

#map handlers
makeMapArg(ylim, xlim, aspect = NULL,
           recolor.map = FALSE, ...)
getMapArg(object = trellis.last.object())

#map panel handlers
panel.GoogleMapsRaster(map)
panel.GoogleMaps(map)

#axis handlers
xscale.components.GoogleMaps(lim, ..., map = map)
yscale.components.GoogleMaps(lim, ..., map = map)
axis.components.GoogleMaps(map, xlim = NULL, ylim = NULL, ...)

```

**Arguments**

- |                  |  |
|------------------|--|
| x                | For GoogleMap and GoogleMap.old only. A formula setting the plot structure, by default $z \sim \text{latitude} * \text{longitude} \mid \text{cond}$ . The axis elements latitude and longitude are required, while z and conditioning cond are optional.   |
| data             | For GoogleMap and GoogleMap.old only. If supplied, the assumed source of the elements of formula x, typically a data.frame.  |
| panel, map.panel | For GoogleMap and GoogleMap.old only. The panels to use when generating the plot data and map layers, respectively. panel is by default the standard loa scatter plot panel <code>panel.loaPlot</code> . map.panel can be the default <code>panel.GoogleMapsRaster</code> or the alternative <code>panel.GoogleMaps</code> . |
| map              | For GoogleMap and related functions only. If supplied, a modified RgoogleMaps output, to be used as the plot background. If NULL (default), this is generated us-  |

	ing the <code>RgoogleMaps</code> function <code>GetMap</code> , the supplied latitude, longitude ranges, and any additional <code>RgoogleMaps</code> arguments supplied within the call. The map is supplied via <code>makeMapArg</code> which modifies the <code>RgoogleMaps</code> output before returning it to simplify local handling and (lattice) plotting.
<code>recolor.map</code>	For <code>GoogleMap</code> and <code>RgoogleMapsWrapper</code> only. If supplied, a vector of elements that R can treat as colors, used as a color scale to recolor map. This uses standard <code>RColorBrewer</code> functions, so can handle arguments like <code>recolor.map = c("white", "grey")</code> for greyscale, etc. Disabled by the default <code>FALSE</code> or <code>NULL</code> .
<code>lon.lat</code>	For <code>GoogleMap</code> only, logical. If <code>TRUE</code> applies $z \sim \text{lon} * \text{lat}   \text{cond?}$ This operation is handled using the <code>formula.type</code> argument in <code>formulaHandler</code>
<code>lat,lon</code>	For <code>quickMap</code> only. Numeric vectors of latitude and longitude values.
<code>ylim, xlim, lim</code>	The latitude and longitude plot ranges. <code>ylim</code> and <code>xlim</code> are only required by <code>makeMapArg</code> , which uses these to set the requested map size. For the axis handlers ( <code>yscale...</code> and <code>xscale...</code> ) the local alternative <code>lim</code> is used for both <code>ylim</code> and <code>xlim</code> in generic code. In <code>GoogleMap</code> and <code>quickMap</code> , if supplied, <code>xlim</code> and <code>ylim</code> are passed to <code>lattice</code> function <code>xyplot</code> via <code>LatLon2XY.centered</code> to handle local scaling.
<code>aspect</code>	The aspect ratio of the plot. If not supplied (recommended), this is determined based on map size, but can be forced by user.
<code>show.data</code>	For <code>quickMap</code> only, a Logical. Should the <code>lat</code> , <code>lon</code> values supplied be plotted on the map ( <code>show.data = TRUE</code> ) or just be used to define the range/size of the map being generated? Default <code>show.data = FALSE</code> .
<code>object</code>	For <code>getMapArg</code> only, a <code>lattice</code> plot to recover an <code>RgoogleMaps</code> map from. (If not supplied, this is assumed to last <code>lattice</code> plot.)
<code>...</code>	Additional arguments are passed on to related functions. For, <code>quickMap</code> these are <code>makeMapArg</code> and the <code>lattice</code> function <code>xyplot</code> . For <code>GoogleMap</code> these are <code>makeMapArg</code> , <code>cexHandler</code> , <code>colHandler</code> and <code>xyplot</code> . <code>makeMapArg</code> uses the <code>RgoogleMaps</code> function <code>GetMap</code> . So, most <code>GetMap</code> arguments can be directly accessed from either <code>GoogleMap</code> or <code>quickMap</code> via this route, e.g. <code>maptype = "satellite"</code> . The returned object is then modified to simplify its handling by the associated panel and axis functions. By default both data point colour and size are z-scaled for <code>GoogleMap</code> . If <code>z</code> is supplied, and <code>cex</code> and <code>col</code> are not set by the user in the plot command, these plot properties are managed by <code>cexHandler</code> and <code>colHandler</code> , respectively. <code>cexHandler</code> and <code>colHandler</code> arguments can be passed direct as part of a <code>GoogleMap</code> command to fine-tune these, e.g. <code>cex.range</code> to change the <code>cex</code> range that <code>z</code> values are scaled to and <code>col.region</code> to change the color range that is applied to <code>z</code> . See associated Help documents for further information. Similarly, argument passing to <code>xyplot</code> in both <code>GoogleMap</code> and <code>quickMap</code> provides access to most common plot parameters, e.g. <code>col</code> , <code>pch</code> , and <code>cex</code> for plot symbol color, type and size, respectively. <code>getMapArg</code> recovers the map from a <code>lattice</code> plot generated with <code>GoogleMap</code> . Unless the plot object is supplied in the <code>getMapArg</code> call, this is assumed to be the last <code>lattice</code> (trellis) output.

**Details**

NOTE: GoogleMap and related panel and axis handling functions are currently in development functions and may be subject to changes.

GoogleMap provides lattice-style conditioning/handling for RgoogleMaps outputs. This uses [loaPlot](#) and the latest version of [panelPal](#) to manage default panel and key settings.

GoogleMap.old is the previous version of the GoogleMap which uses the previous version of [panelPal](#)

googleMap is a GoogleMap wrapper, included because this alternative form of the plot name was used in earlier versions of the package.

quickMap is crude map plotter intended to demonstrate the use of the other 'handler' functions when building dedicated mapping functions.

makeMapArg accepts latitude and longitude ranges and RgoogleMaps function [GetMap](#) arguments, and produces an output suitable for use with the `panel.GoogleMapsRaster` and `panel.GoogleMaps` panel functions or in subsequent GoogleMap calls if, e.g., the users wishes to reuse an existing map.

`panel.GoogleMapsRaster` and `panel.GoogleMaps` are lattice panel functions that generate map layers for a lattice plot using `makeMapArg` outputs.

`yscale.components.GoogleMaps` and `xscale.components.GoogleMaps` are y- and x-axis handlers for use with the above panels.

`axis.components.GoogleMaps` is a wrapper that combines `yscale.components.GoogleMaps` and `xscale.components.GoogleMaps` and allows both axis to be set from the `lattice` function argument `axis` rather than each individually, via `yscale.components` and `xscale.components`.

**Value**

GoogleMap and quickMap return trellis objects, much like conventional lattice plot functions.

`makeMapArg` returns a modified form of the RgoogleMaps function [GetMap](#) output suitable for use as the map argument with the above functions. Note: the automatic assignment of the RgoogleMaps function argument size

`getMapArg` recovers the map from an existing GoogleMap output.

`panel.GoogleMapsRaster` and `panel.GoogleMaps` generate panel outputs suitable for use in standard lattice panel functions.

`yscale.components.GoogleMaps`, `xscale.components.GoogleMaps` generate suitable latitude, longitude scales for use with map layers. `axis.components.GoogleMaps` is a wrapper for their routine use.

**Note**

Google Maps outputs are 2D projections of curve sections of the Earth's surface. Therefore, the assignment of points within panels and the annotation of latitudes and longitudes along axis needs to be locally handled to account for this.

GoogleMap and quickMaps use RgoogleMaps functions [LatLon2XY](#), [LatLon2XY.centered](#) and [XY2LatLon](#) to locally scale both axis and data.

Important: Users wanting to add data to these plots, e.g. using `update` or `layers` in `latticeExtra`, should first rescale the data. Likewise, users wanting to add maps to other plots will need to rescale plotted data to use these maps. See Example 1 below.

Important: The Google API returns a map panel larger than the data (latitude, longitude) range requested. However, it does this using a limited number of panel sizes. This means you may get back a map that is large than necessary. As `xlim` and `ylim` are passed to the API when they are called resetting these can produce similar effects (so you may not get exactly the map range you ask for! If you want to manually optimise the map ranges, the best option is currently to start with:

```
GoogleMap(..., size=c(640,640))
```

...and then reduce either or both of these values until you generate an appropriate map size.

### Author(s)

Karl Ropkins

### References

This function makes extensive use of code developed by others.

`lattice`: Sarkar, Deepayan (2008) *Lattice: Multivariate Data Visualization with R*. Springer, New York. ISBN 978-0-387-75968-5

`RColorBrewer`: Erich Neuwirth <erich.neuwirth@univie.ac.at> (2011). `RColorBrewer`: Color-Brewer palettes. R package version 1.0-5. <http://CRAN.R-project.org/package=RColorBrewer>

`RgoogleMaps`: Markus Loecher and Sense Networks (2011). `RgoogleMaps`: Overlays on Google map tiles in R. R package version 1.1.9.6. <http://CRAN.R-project.org/package=RgoogleMaps>

### See Also

In other packages, see

`RgoogleMaps`: [GetMap](#); [LatLon2XY](#); [LatLon2XY.centered](#); and, [XY2LatLon](#).

`lattice`: [xyplot](#); [panel.xyplot](#); and [panel.levelplot](#).

### Examples

```
## Example 1
## quickMap code
## as example of third-party use of functions

quickMap <- function(lat, lon, show.data = FALSE, ...){

  #get map
  map <- makeMapArg(lat, lon, ...)

  #scale axis for map projection
  map.axis.comps <- axis.components.GoogleMaps(map)
  map.axis <- function(components, ...)
    axis.default(components = map.axis.comps, ...)
```

```

#scale data for map projection
#see ?Rgooglemaps::LatLon2XY
temp <- LatLon2XY.centered(map, lat, lon)
lat <- temp$newY
lon <- temp$newX

#plot data on map
xyplot(lat~lon,
        xlim = map$xlim, ylim = map$ylim,
        aspect = map$aspect,
        axis = map.axis,
        panel = function(...){
          panel.GoogleMapsRaster(map)
          if(show.data)
            panel.xyplot(...)
        }, ...)
}

## Example 2
## Off-line GoogleMap examples

# Use a subsample of lat.lon.meuse
temp <- lat.lon.meuse[sample(1:155, 15),]

GoogleMap(zinc~latitude*longitude, col.regions=c("grey", "darkred"),
          data=temp, map=roadmap.meuse)

GoogleMap(zinc~latitude*longitude, col.regions=c("grey", "darkred"),
          panel=panel.binPlot,
          data=temp, map=roadmap.meuse)

GoogleMap(cadmium*50+copper*10+lead*2+zinc~latitude*longitude,
          col.regions=c("grey", "darkred"),
          key.z.main="Concentrations", panel.zcases = TRUE,
          data=temp, map=roadmap.meuse)

GoogleMap(cadmium*50+copper*10+lead*2+zinc~latitude*longitude,
          col.regions=c("grey", "darkred"), panel=panel.zcasePiePlot,
          data=temp, map=roadmap.meuse)

# Note 1:
# Here, the map argument is supplied so example works off-line.
# If not supplied and R is on-line, GoogleMap will get map
# from the Google API. Repeat any of above without map argument
# when on-line. For example:
## Not run:
  GoogleMap(zinc~latitude*longitude, col.regions=c("grey", "darkred"),
            data=lat.lon.meuse)
## End(Not run)
# (The map will appear slightly different because non-default
# size and mptype settings were used to make roadmap.meuse. See
# ?roadmap.meuse for details.)

```

```
# Note 2:
# To make a map for use with panel.GoogleMaps or panel.GoogleMapsRaster
# without plotting use makeMapArg(). To recover a map from a previously
# plotted loa GoogleMap use getMapArg().
```

---

### 1.3.trianglePlot *Triangle plots*

---

#### Description

Triangle plot functions for Lattice.

#### Usage

```
trianglePlot(x, data = NULL, ..., ref.cols = TRUE)

#standard panels

panel.trianglePlot(x = NULL, y = NULL, a0 = NULL, b0 = NULL,
  c0 = NULL, ..., loa.settings = FALSE, plot = TRUE,
  process = TRUE)
panel.trianglePlotFrame(..., grid = NULL, axes = NULL)
panel.trianglePlotGrid(alim = NULL, blim = NULL, clim = NULL,
  ..., grid = TRUE, panel.scales = NULL)
panel.trianglePlotAxes(alim = NULL, blim = NULL, clim = NULL,
  ..., axes = TRUE, ticks=TRUE, annotation=TRUE,
  panel.scales = NULL)

#other panels
panel.triangleByGroupPolygon(x = NULL, y = NULL, a0 = NULL,
  b0 = NULL, c0 = NULL, ..., loa.settings = FALSE,
  plot = TRUE, process = TRUE)
panel.triangleKernelDensity(x = NULL, y = NULL, a0 = NULL,
  b0 = NULL, c0 = NULL, ..., loa.settings = FALSE,
  plot = TRUE, process = TRUE)
panel.triangleSurfaceSmooth(x = NULL, y = NULL, z = NULL,
  a0 = NULL, b0 = NULL, c0 = NULL, ..., loa.settings = FALSE,
  plot = TRUE, process = TRUE)

#data handlers

triABC2XY(a, b = NULL, c = NULL, ..., force.abc = TRUE,
  if.na = "remove.row", if.neg = "remove.row",
  verbose = FALSE)
```

```

triXY2ABC(x, y = NULL, ..., force.xy = TRUE,
          verbose = FALSE)
triLimsReset(ans)
triABCsquareGrid(a, b = NULL, c = NULL, ..., n=100)

```

### Arguments

x	For trianglePlot only, a formula in structure $z \sim a0 + b0 + c0 \mid \text{cond}$ . The elements a0, b0 and c0, the inputs for the three axis on the triangle plot, are required, while z and conditioning (cond) are optional. (For other functions, x may be used as the pair to y. See y below.)
data	For trianglePlot only, if supplied, the assumed source of the elements of formula x, typically a data.frame.
...	Additional arguments.
ref.cols	Either a logical to turn off/on grid color-coding or a vector of colors to be applied to a0, b0 and c0 axes and grids. These are applied to the grid lines and axes tick and annotation components. Some users, particularly those less familiar with triangle plots, can find such color referencing helpful when analyzing such plots. By default, the colorings are quite subtle, so users can see the effect if they look for it but it does not take over the plot when it is not focused on. Finer control can be achieved using axes, ticks, grid, etc. (See below).
y, a, a0, b, b0, c, c0, z	(and x in relevant functions). a/a0, b/b0 and c/c0 are the three scales of the triangle plot, and x and y are the equivalent 2-D projections. The arguments are typically options in panel functions (panel... functions), conversion functions (triABC2XY and triXY2ABC) and the scaling function triLimsReset. z is the z-case from the plot formula.
loa.settings, plot, process	<a href="#">loaPlot</a> arguments used to manage <a href="#">panelPal</a> activity.
grid, axes, ticks, annotation	User-resets for the axes, grid, tick and annotation elements of the plots. These can be NULL or FALSE to turn off, TRUE to show, a vector (in which case they are assumed to be color assignments) or a list of standard plot parameters, e.g. col, lty, lwd, etc. for color, line type and line thickness, etc. Plot parameter assignments are applied to all axes unless specific axes are identified. For example, trianglePlot calls including grid.col = 2 make all axes red, while calls including grid.a0.col = 2 only recolor the first (a0) axis.
alim, blim, clim	Delimiters for a, b and c scales, equivalent to xlim and ylim in conventional plots, but less flexible. See Details below for more information.
panel.scales	A local argument, typically a list, that controls the appearance of the a0/b0/c0 axes. This is roughly equivalent to the scales argument used by conventional lattice plots to handle x and y axis, but intended for non-standard scales, such as the triangle axes used here. It can be set directly or used in combination with the local scale(s) handler function <a href="#">localScalesHandler</a> to override/hijack standard scales operations. (See note below).

<code>force.abc, force.xy</code>	Logicals. If a list or data.frame is supplied to <code>triABC2XY</code> or <code>triXY2ABC</code> as a source or <code>a/b/c</code> or <code>x/y</code> respectively should appropriately named elements be used regardless of order? See Note below.
<code>if.na</code>	Character. Handling method to be used if NAs are present. The default option <code>'remove.row'</code> replaces all entries in the same row with NAs. (Note: this is different from <code>na.omit</code> which would remove the whole row. Here, the row is retained as NAs to maintain indices for conditioning.) Other options currently include: <code>'make.zero'</code> to replace the NA with 0; and <code>'keep.as.is'</code> to leave unchanged.
<code>if.neg</code>	Character. Like <code>if.na</code> but for negative values: <code>'remove.row'</code> to replace all entries in the same row with NAs; <code>'make.zero'</code> to replace all negative values with 0; <code>'rescale.col'</code> rescales any column (i.e., a, b or c) that contains a negative from zero by subtracting the minimum.
<code>verbose</code>	Logical, default FALSE. Should a full output be returned? The alternative FALSE generates a minimal report.
<code>ans</code>	For <code>triLimsReset</code> only, a trellis output, e.g. a lattice plot, to be scaled and plotted based on the assumption that it is a <code>trianglePlot</code> .
<code>n</code>	For <code>triABCsquareGrid</code> only, number of points to divide each axes by when generating the data grid.

## Details

`trianglePlot` generates a triangle plot using the lattice framework.

`panel.trianglePlot...` functions handle the appearance of triangle plot outputs.

`panel.trianglePlot`, which is assigned as the default panel manages both the data layer of the plot and the plot frame (axes, grid, annotation, etc).

`panel.trianglePlotAxes` and `panel.trianglePlotGrid` generate axes and grid components of the plot, and `panel.trianglePlotFrame` is a wrapper for these. The data layer, which by default is `panel.loaPlot`, can be accessed separately using the `data.panel` argument.

`triangleKernelDensity` generates a kernel density surface for supplied `a0`, `b0` and `c0` cases.

`triangleSurfaceSmooth` generates a smoothed surface for supplied `a0`, `b0`, `c0` and `z` cases.

`triABC2XY` converts supplied (a, b, c) coordinates to an (x, y) scale suitable for use with triangle plot functions.

`triXY2ABC` converts supplied (x,y) coordinates from triangle plots to the associated proportional (a, b, c) scale.

There are various options for range limiting with `triABC2XY`, `triXY2ABC`, and therefore triangle plots. Firstly, limits can be set individually with `alim`, `blim` and `clim`, much like with `xlim` and `ylim` for conventional plots. However, they can also be set at once using `lims`, as in e.g. `lims = c(0, 1)` to set all axes to full ranges, or on the basis of minimum and maximum cut-offs using `abc.mins` and `abc.maxs`, respectively.

`trianglePlot` uses `localScalesHandler` to override normal lattice handling of scales. This allows parameters for axes other than 'x' and 'y' to be passed via the `scales` argument for axis generation within the plot panel itself. The function does this by recovering the information for



each of the local axes (here  $a_0$ ,  $b_0$  and  $c_0$ ) from scales, and passing this on to the plot as the argument `panel.scales` which can then be evaluated by an appropriate `panel...` function like `panel.trianglePlotAxes`. At the same time it also resets scales to stop the standard axes being generated. The intention here is two-fold. Firstly, to provide plot users with an axes control mechanism like the standard scales control of  $x$  and  $y$  that they already know. And, secondly, to provide developers with a simple framework for the quick addition of non-standard axes or scales. See [localScalesHandler](#) and `panel.localScale` for further details.

`trianglePlot` uses `getPlotArgs` to manage lattice defaults and plot developer and user resets for the different plot components (axes, ticks, grid, annotation). As with `localScalesHandler`, the intention here is to provide more routine access to higher level plot control.

### Value

`trianglePlot` returns trellis objects, much like conventional lattice plot functions.

`panel.trianglePlot...` functions are intended for use within a `trianglePlot` function call.

`triABC2XY` returns a list containing the named components  $x$  and  $y$ , which are the 2-D  $(x,y)$  transformations of supplied  $(a,b,c)$  `trianglePlot` elements.

`triXY2ABC` returns a list containing the named components  $a$ ,  $b$  and  $c$ , which are the  $(a,b,c)$  triangle plot coordinates associated with supplied 2-D  $(x,y)$  that `trianglePlot` would generate.

`resetTriLims` returns a supplied trellis object, rescaled based on the assumption that it is a triangle plot.

### Note

General:

With triangle plots, the  $(a_0, b_0, c_0)$  scales are proportional. So regardless of the absolute sizes of a coordinate set  $(a,b,c)$ , values are plotted and handled as proportions, i.e.  $a/(a+b+c)$ ,  $b/(a+b+c)$  and  $c/(a+b+c)$ , respectively. This means that absolute values of  $a$ ,  $b$  and  $c$  are lost when points are plotted on these axes. So, the function `triXY2ABC` returns the relative proportions of  $a$ ,  $b$  and  $c$ , not the absolute amounts, when translating a 2-D  $(x,y)$  coordinates into the associated  $(a, b, c)$  coordinates.

Development:

This is an in-development plot, and 'best handling' strategies have not been decided for several elements. So, future versions of these functions may differ significantly from the current version.

In particular:

Current axes assignments, e.g.  $(a, b, c)$  versus  $(a_0, b_0, c_0)$ , etc., have not been finalised. So, these may change in future versions of the function.

Currently, `trianglePlot` scale adjustment options have been limited. Options under evaluation include: (1) by `alim`, `blim`, `clim` setting, equivalent to `xlim` and `ylim`, (2) by `lims` to set all axes ranges the same, and (3) by `maxs` to setting all axes range maximums and `mins` to set all axes range minimums, etc.

These options are currently only available via the data converters.

One of the issues here is that the axes ranges are all inter-linked. The range of one axis is a function of the other two axes ranges. So, setting these can generate contradictions. For example, `lims=c(0, 0.1)` should in theory set all ranges to  $(0, 0.1)$ . But, the triangle  $a = b = c = c(0, 0.1)$

cannot exist. Therefore, the plot would attempt to recover the extended range that includes all the requested ranges ( $a = c(0, 0.1)$ ,  $b = c(0, 0.1)$  and  $c = c(0, 0.1)$ ), which in this case is the full range:  $a = b = c = c(0, 1)$ . Suggestions on this topic are very welcome.

`trianglePlot`:

As part of the `loa` version 0.2.19 update, `trianglePlot` was rewritten to run with the most recent version of `panelPal` function. This means all plot functions in `loa` now use the most recent version of `panelPal`.

This update should provide improved plot handling similar to recent versions of `loaPlot` and `GoogleMap` functions which both already (from versions 0.2.0 onwards) use the latest version of `panelPal`.

`panel.trianglePlotAxes`:

Code currently in revision. Please handle with care.

`triABC2XY`, `triABCsquareGrid`:

Code currently in revision. Please handle with care.

### Author(s)

Karl Ropkins

### References

These function makes extensive use of code developed by others.

Currently, several triangle plotting methods are under evaluation for use within this package. These are:

The tri-plot method of Graham and Mudgley:

Graham, David J. and Mudgley, Nicholas, G. Graphical representation of particle shape using triangular diagrams: An Excel spreadsheet method. *Earth Surface Processes and Landforms*, 25, 1473-1477, 2000.

The `triangle.param` method of Chessel (as coded in R package 'ade4')

Dray, S. and Dufour, A.B.(2007). The `ade4` package: implementing the duality diagram for ecologists. *Journal of Statistical, Software*. 22(4): 1-20.

Chessel, D. and Dufour, A.B. and Thioulouse, J. (2004). The `ade4` package - I - One-table methods. *R News*. 4: 5-10.

Dray, S. and Dufour, A.B. and Chessel, D. (2007). The `ade4` package-II: Two-table and K-table methods. *R News*. 7(2): 47-52.

And the trilinear plot of Allen as reported by Zhu:

Zhu, M. (2008). How to draw a trilinear Plot. *Statistical Computing & Graphics*, 19(1), June, 2008.

In this version of the package tri-plot methods are used for the `triABC2XY` and `triXY2ABC` transforms and a modification `triangle.param` methods is used to calculate suitable values for `alim`, `blim` and `clim`.

As elsewhere, the use of `lattice` is also gratefully acknowledged:

`lattice`: Sarkar, Deepayan (2008). *Lattice: Multivariate Data Visualization with R*. Springer, New York. ISBN 978-0-387-75968-5

**See Also**

In `loa`: For in-panel axis/scale generation, see [loaPlot](#), [panelPal](#), [localScalesHandler](#) and [panel.localScale](#).

In other packages: [xyplot](#) in [lattice](#).

**Examples**

```
## Example 1
## Basic triangle plot usage

trianglePlot(cadmium~copper+lead+zinc|lime,
             data=lat.lon.meuse)

# Notes:
# Formula structure  $z \sim a_0 + b_0 + c_0 | \text{cond}$ , where  $a_0$ ,  $b_0$  and
#  $c_0$  are the three axes of the triangle plot
# Data (and groups) assignment like in standard lattice plots.
# By default  $z$  is linked to col and cex.
# Unless overridden by user inputs or group or zcase setting.
# Plot handling is similar to loaPlot
# (So, see ?loaPlot and ?panelPal for further details.)

# Formula variations:
# basic triangle plot without  $z$  values assignment
# trianglePlot(~a0+b0+c0, ...)
# ... with  $z$  values set
# trianglePlot(z~a0+b0+c0, ...)
# ... with grouping
# trianglePlot(z~a0+b0+c0, groups=grps, ...)

## Example 2
## Basic frame (axes, grid, tick, annotation) management

trianglePlot(~1+1+1, type="n",
            grid.alpha = 0.2,
            ticks.alpha = 0.2)      ## grid and tick alpha reset

# notes:
# Here, grid and ticks arguments are used to remove or modify these
# elements of the plot frame individually.
# Setting can be management in list form like in normal lattice or
# in a loa shorthand where e.g. the argument grid.a0.lty = 1 is equivalent
# to grid = list(a0 = list(lty = 1))
# (So, quicker if you are only changing a small number of elements.)
```

## Description

Stack plot functions for Lattice.

## Usage

```
stackPlot(x, data = NULL, ...)

#standard panels

panel.stackPlot(..., process=TRUE, plot=TRUE,
                loa.settings = FALSE)

#data handlers
##currently not exported
```

## Arguments

x	For <code>stackPlot</code> only, a formula of general structure $y_1 + y_2 \sim x \mid \text{cond}$ , etc. The elements $y_1, y_2$ , etc are stacked on the y-axis, and plotted against x. Both are required.
data	For <code>stackPlot</code> only, if supplied, the assumed source of the elements of formula x, typically a <code>data.frame</code> .
...	Additional arguments.
loa.settings, plot, process	<a href="#">loaPlot</a> arguments used to manage <a href="#">panelPal</a> activity.

## Details

`stackPlot` generates a stack plot using the lattice framework.

`panel.stackPlot` handles the appearance of triangle plot outputs.

## Value

`stackPlot` returns trellis objects, much like conventional lattice plot functions.

`panel.stackPlot` is intended for use within a `trianglePlot` function call.

**Note**

Development:

This is an in-development plot, and 'best handling' strategies have not been decided for several elements. So, future versions of these functions may differ significantly from the current version.

In particular:

`stackPlot`:

The `stackPlot` argument `x` may include conditioning in the form `y ~ x | cond`. However, exact handling is has not been defined, so may subject to change.

To generate the stacks, `stackPlot` resets `y` values by applying `y - min(y)` to each layer and then stacks them. It also generates a second element `y0` of asociated baselines. This is then used in the form `x = c(x, rev(x)), y = c(y, rev(y0))` with `panel.polygon` to generate the stack layers.

`panel.stackPlot`:

Code currently in revision. Please handle with care.

**Author(s)**

Karl Ropkins

**References**

These function makes extensive use of code developed by others.

As elsewhere, the use of `lattice` is also gratefully acknowledged:

`lattice`: Sarkar, Deepayan (2008). *Lattice: Multivariate Data Visualization with R*. Springer, New York. ISBN 978-0-387-75968-5

**See Also**

In `loa`: [loaPlot](#) and [panelPal](#).

In other packages: [xyplot](#) and [panel.polygon](#) in `lattice`.

**Examples**

```
## Example 1
## Basic stack plot usage

## Not run:
stackPlot(lead~dist.m, data=lat.lon.meuse)
stackPlot(cadmium+copper+lead+zinc~dist.m, data=lat.lon.meuse)
## End(Not run)

stackPlot(cadmium*40+copper*5+lead+zinc~dist.m, data=lat.lon.meuse)
```

1.5.loaBarPlot

*loaBarPlot***Description**

Bar plot variation using for Student Project.

**Usage**

```
loaBarPlot(x, y=NULL, groups=NULL, cond=NULL,
           data=NULL, ..., drop.nas=TRUE, stat=NULL)
```

**Arguments**

<code>x</code>	Either the x case for the bar plot or a plot formula. If the x case, typically a vector of factor or grouping terms, used to assign x positions in bar plot. If a plot formula, a plot description in the format <code>y~xlcond</code> , where x is a factor or grouping term and y and cond are optional.
<code>y</code>	(Optional) The y case for the bar plot, typically a vector of numeric terms, used with <code>stat</code> when calculating summary information for bar plots.
<code>groups, cond</code>	(Optional) The group case for the bar plot, typically a vector of factor or grouping terms.
<code>data</code>	(Optional) if supplied, the assumed source of the plot elements, x, y, groups and cond, typically a <code>data.frame</code> .
<code>...</code>	Additional arguments, passed on to <code>lattice</code> function.
<code>drop.nas</code>	Option to drop NAs before plotting results, default TRUE.
<code>stat</code>	If supplied, the function used to summarise y data after grouping by x, groups and cond. By default, this counts x cases if y is not supplied, or calculates sum of y values if these are supplied.

**Details**

`loaBarPlot` summarises supplied plot data and generates a bar plot using the `lattice` framework.

**Value**

`loaBarPlot` returns trellis objects, much like conventional `lattice` plot functions.

**Note**

Development:

This is an in-development plot, and 'best handling' strategies have not been decided for several elements. So, future versions of these functions may differ significantly from the current version.

In particular:

`loaBarPlot`:

This is for student project, may not be staying.  
Code currently in revision. Please handle with care.

**Author(s)**

Karl Ropkins

**References**

These function makes extensive use of code developed by others.  
As elsewhere, the use of `lattice` is also gratefully acknowledged:  
`lattice`: Sarkar, Deepayan (2008). *Lattice: Multivariate Data Visualization with R*. Springer, New York. ISBN 978-0-387-75968-5

**See Also**

In `loa`: [listUpdate](#) and [colHandler](#).  
In other packages: [barchart](#) in `lattice`.

**Examples**

```
## Example 1
## Basic bar plot usage

  loaBarPlot(Species, Sepal.Width, data=iris, stat=mean)

#or equivalent using formula
## Not run:
  loaBarPlot(Sepal.Width~Species, data=iris, stat=mean)
## End(Not run)
```

---

2.1.specialist.panels *Special panel functions 01*

---

**Description**

Specialist panel functions for use with `lattice` and `loa` plots.

**Usage**

```
panel.loaLevelPlot(x = NULL, y = NULL, z = NULL, ...,
  loa.settings = FALSE)
```

```
panel.surfaceSmooth(x = NULL, y = NULL, z = NULL,
  breaks = 200, x.breaks = breaks, y.breaks = breaks,
  smooth.fun = NULL, too.far=0, ...,
  plot = TRUE, process = TRUE, loa.settings = FALSE)
```

```
panel.kernelDensity(x, y, z = NULL, ..., n = 20,
  local.wt = TRUE, kernel.fun = NULL, too.far = 0,
  panel.range = TRUE, process = TRUE, plot = TRUE,
  loa.settings = FALSE)
```

```
panel.binPlot(x = NULL, y = NULL, z = NULL,
  breaks=20, x.breaks = breaks, y.breaks = breaks,
  x1=NULL, x2=NULL, y1=NULL, y2=NULL,
  statistic = mean, pad.grid = FALSE, ...,
  plot = TRUE, process = TRUE, loa.settings = FALSE)
```

### Arguments

x, y, z	lattice function arguments passed down to individual panels.
...	Additional arguments, typically passed on. See below.
loa.settings, process, plot	For panel... functions that intended to be handled using <a href="#">panelPal</a> . loa.settings, a logical indicating if the safe mode setting should be returned. process and plot, logicals, indicating if the process and plot sections of the panel function should be run. See below and <a href="#">panelPal</a> help documents for further details.
breaks, x.breaks, y.breaks	(For panel.surfaceSmooth and panel.binPlot) How many break points to introduce when smoothing a surface or binning data. breaks can be used to set the same number of breaks on both axes, while x.breaks and y.breaks can be used to set these separately.
smooth.fun	(For panel.surfaceSmooth) A function that can fit a surface estimate to (x, y, z) data. See notes below for further details.
too.far	(For panel.surfaceSmooth and panel.kernelDensity) The distance from original data at which to stop predicting surface values. See notes below for further details.
n	(For panel.kernelDensity) the number of x and y cases to estimate when estimating density.
local.wt	(For panel.kernelDensity) A logical (default TRUE) indicating if kernel density estimates should be weighed relative to other groups, panels, etc.
kernel.fun	(For panel.kernelDensity) A function that can estimate kernel densities.
panel.range	(For panel.kernelDensity) A logical (default FALSE) indicating if the kernel density estimation data range should be forced to the full panel range. See Below.
x1, x2, y1, y2	(For panel.binPlot) Vectors giving the bin cell dimensions used when binning x and y elements. Typically ignored and calculated within the plot call.



<code>statistic</code>	(For <code>panel.binPlot</code> ) the function to use when calculating z values for each set of binned. By default, this is <code>mean</code> . So, if a z element is supplied in the plot call, the data is binned according to x and y values, and the mean of z values within each bin reported/plotted. If z is not supplied, <code>statistic</code> is reset to <code>length</code> to generate a frequency plot and a warning generated.
<code>pad.grid</code>	For <code>panel.binPlot</code> , Logical, should empty bins be reported? If <code>TRUE</code> , they are reported as NAs; if <code>FALSE</code> , they are not reported.

## Details

`panel.loaLevelPlot` is intended for plot data structured for use with the `lattice` function `levelplot`, e.g. regularised (x,y,z) or a matrix:

```
loaPlot(..., panel = panel.loaLevelPlot)
levelplot(...) #in lattice
```

Other specialist `panel...` functions can be used with the `lattice` function `xyplot`:

```
xyplot(..., panel = panel.kernelDensity)
xyplot(..., n = 50, panel = panel.kernelDensity)
xyplot(..., panel = function(...) panel.kernelDensity(..., n = 50))
#etc
```

However, they are intended for use with loa plots that incorporate `panelPal`. This combination provides a mechanism for the routine preprocessing of panel data, the association of specialist keys, and the routine alignment of panel and legend settings in cases where values are reworked within the panel function call:

```
loaPlot(..., panel = panel.kernelDensity)
#etc
```

`panel.surfaceSmooth` and other similar `panel...` functions generate smoothed surfaces using supplied (x,y,z) data and pass this to `panel.loaLevelPlot` to plot.

By default, `panel.surfaceSmooth` uses stats function `loess` to generate a surface. Alternative smooths can be set using the `smooth.fun` argument, and the surface range can to controlled using the `too.far` argument.

`panel.kernelDensity` generates kernel density estimations based on the supplied x and y data ranges. Because it is density plot, it counts the number of z values. So, z values are ignored. It is intended to be used in the form:

```
loaPlot(~x*y, ..., panel = panel.kernelDensity)
```

So, if any z information is supplied, users are warned that it has been ignored, e.g:

```
loaPlot(z~x*y, ..., panel = panel.kernelDensity)
#warning generated
```

`panel.binPlot` bins supplied z data according to x and y values and associated break points (set by `break` arguments), and then calculates the required statistic for each of these. By default, this is `mean`, but alternative functions can be set using the `statistic` argument. It is intended to be used in form:

```
loaPlot(z~x*y, ..., panel = panel.binPlot)
```

If no z values are supplied, as in:

```
loaPlot(~x*y, ..., panel = panel.binPlot)
```

... `panel.binPlot` resets statistic to `length` (again with a warning) and gives a count of the number of elements in each bin.

### Value

As with other `panel...` functions in this package, output are suitable for use as the `panel` argument in `loa` (and sometimes `lattice`) plot calls.

### Note

All these `panel...` functions treat `col` and `col.regions`, etc, as discrete arguments. Typically, `col` links to lines (contour lines for surfaces, bin borders for binned data) and `col.regions` links any generates surface region.

`panel.surfaceSmooth` passes additional arguments on to the `smooth.fun` to estimate surface smooths and the `lattice` function `panel.levelplot` to generate the associated plot. If no `kernel.fun` is supplied in the `panel` call, the `stats` function `loess` is used to estimate surface smooth. The `too.far` argument is based on same in `vis.gam` function in the `mgcv` package.

`panel.kernelDensity` passes additional arguments on to the `kernel.fun` to estimate kernel density and the `lattice` function `panel.contourplot` to generate the associated plot. If no `kernel.fun` is supplied in the `panel` call, the `MASS` function `kde2d` is used to estimate kernel density.

`panel.binPlot` passes limited arguments on to `lrect`.

`panel.kernelDensity` and `panel.binPlot` are currently under review.

### Author(s)

Karl Ropkins

### References

These function makes extensive use of code developed by others.

`lattice`: Sarkar, Deepayan (2008) *Lattice: Multivariate Data Visualization with R*. Springer, New York. ISBN 978-0-387-75968-5

(for `panel.kernelDensity`) `MASS` package: Venables, W. N. and Ripley, B. D. (2002) *Modern Applied Statistics with S*. Fourth edition. Springer.

(for `panel.surfaceSmooth`) `mgcv` package and `too.far` argument: Wood, S.N. (2004) Stable and efficient multiple smoothing parameter estimation for generalized additive models. *Journal of the American Statistical Association*. 99:673-686. Also <http://www.maths.bath.ac.uk/~sw283/>

### See Also

In `loa`: `panelPal`

In `lattice`: `xyplot`, `levelplot`, `panel.contourplot`, `lrect`

**Examples**

```
## Example 1
## for data already set up for levelplot

loaPlot(volcano, panel=panel.loaLevelPlot)

## Example 2
## Surface smooth

loaPlot(copper~longitude*latitude, data= lat.lon.meuse,
        panel=panel.surfaceSmooth, grid=TRUE,
        too.far=0.1, col.regions=3:2)
```

---

2.2.specialist.panels *Special panel functions 02*


---

**Description**

In development specialist panel functions for polar plotting

**Usage**

```
panel.polarPlot(x = NULL, y = NULL, r = NULL, theta = NULL, ...,
               data.panel = panel.loaPlot, loa.settings = FALSE,
               plot = TRUE, process = TRUE)

#grid, axes and axes labelling

panel.polarFrame(..., grid = TRUE, axes = TRUE, labels = TRUE,
                panel.scales = NULL, grid.panel = panel.polarGrid,
                axes.panel = panel.polarAxes, labels.panel = panel.polarLabels)

panel.polarAxes(axes.theta = NULL, axes.r = NULL, thetalim = NULL,
               rlim = NULL, ..., axes = NULL, panel.scales = NULL)

panel.polarGrid(grid.theta = NULL, grid.r = NULL,
               thetalim = NULL, rlim = NULL, ..., grid = NULL,
               panel.scales = NULL)

panel.polarLabels(labels.theta = NULL, labels.r = NULL,
                 thetalim = NULL, rlim = NULL, ..., labels = NULL,
                 panel.scales = NULL)
```

**Arguments**

<code>x, y</code>	The x and y coordinates of plot points.
<code>r, theta</code>	The equivalent polar coordinates of the plot points. If these are not supplied, x and y are assumed to be polar coordinates and these are calculated by the function.
<code>...</code>	Additional arguments, typically passed on. For <code>panel.polarPlot</code> these are passed to the <code>data.panel</code> . See below.
<code>data.panel</code>	The panel to use to handle data once polar coordinates have been checked for or generated. For <code>panel.polarPlot</code> , by default this is <code>panel.loaPlot</code> .
<code>loa.settings, plot, process</code>	<code>loa</code> panel management arguments, handled by <code>panelPal</code> . See associated help documentation for further details.
<code>grid, axes, labels</code>	plot management options for the grid, axis and axis label elements of the plot. These can be logicals (TRUE to include the element or FALSE to remove it) or lists of plot parameters.
<code>panel.scales</code>	<code>loa</code> plot management argument used when generating grids, axes and labels within the plot panel.
<code>grid.panel, axes.panel, labels.panel</code>	Used by the <code>panel...Frame</code> functions to identify the <code>panel...</code> functions to use when generating the grid, axes and axis labelling elements of the plot.
<code>axes.theta, axes.r, thetalim, rlim</code>	For <code>panel.polarAxes</code> axes settings. <code>axes.theta</code> and <code>axes.r</code> are the theta and r coordinates of the axis reference points, tick marks, etc. <code>thetalim</code> and <code>rlim</code> are the plot/axes ranges (like <code>xlim</code> and <code>ylim</code> in standard lattice plots).
<code>grid.theta, grid.r</code>	Like <code>axes.theta</code> and <code>axes.r</code> but for grid.
<code>labels.theta, labels.r</code>	Like <code>axes.theta</code> and <code>axes.r</code> but for labels.

**Details**

The `panel.polar...` series of the functions are intended for use with `loaPlot`.

`panel.polarPlot` generates a 'bubble plot' style output on polar coordinates. It generates axes and annotation within each plot panel using the other panel functions.

`panel.polarGrids`, `panel.polarAxes` and `panel.polarLabels` generate plot grid, axes and axes labelling elements of the plot. `panel.polarFrame` provides a wrapper for these plot elements.

Users can fine-tune axes, grids and labels by supplying additional arguments in plot calls, or replace these elements with purpose written functions to more completely modify plot appearance.

**Value**

The `panel.polar...` functions are intended to be used as the `panel` argument in `loa` plot calls. So, e.g.:

```
a <- 1:360
loaPlot(a~a*a, panel=panel.polarPlot)
```

They can also be used with relatively simple lattice plots. However, some features of loa plots managed by [panelPal](#), e.g. default plot appearance management, automatic grouping and panel and key alignment will not be available.

### Note

`panel.polarPlot` is in-development. Function arguments may change.

### Author(s)

Karl Ropkins

### References

This function makes extensive use of code developed by others.

`lattice`: Sarkar, Deepayan (2008) *Lattice: Multivariate Data Visualization with R*. Springer, New York. ISBN 978-0-387-75968-5

### See Also

In loa: [loaPlot](#); and [panelPal](#).

In other packages: [xyplot](#) in [lattice](#).

2.3.specialist.panels *Special panel functions 03*

### Description

In development specialist panel functions for generating zcase glyph structures.

### Usage

```
panel.zcasePiePlot(..., loa.settings = FALSE)
```

```
panel.zcasePieSegmentPlot(..., zcase.rescale=TRUE,
  loa.settings = FALSE)
```

### Arguments

...	Additional arguments, typically setting the color and properties of the plotted glyphs. See below.
<code>zcase.rescale</code>	Should the glyph element be rescaled? See below.
<code>loa.settings</code>	loa options, to be handled by <a href="#">panelPal</a> .

**Details**

All these `panel...` functions generate glyphs using `z` inputs and plot these at the associated `(x, y)` location. So, for example a called which used one of the panels and the plot formula:

$$z1 + z2 + z3 + z4 \sim x * y$$

... would plot a series of glyphs, each containing four elements that would be scaled according to `z1`, `z2`, `z3` and `z4`, and each at the associated `(x, y)` location. This means there will be one discrete glyph for each row of data supplied to the plot call.

`panel.zcasePiePlot` generates a series of `x/y` referenced pie graphs. By default, pie dimensions are assigned as: Pie size (radius) proportional to sum of `z` cases and scaled using `cexHandler` (`z1 + z2 + z3 + z4` for the above formula); Number of Pie segments equal to number of `z` cases (so, 4 for the above formula); Pie segment width (`phi`) proportional to the individual `zcase` (so, `z1 / (z1 + z2 + z3 + z4) * 360` for first pie segment for the above formula).

`panel.zcasePieSegmentPlot` is a variation on the conventional pie plot where segment radius rather than segment width is varying by `zcase`.

**Value**

These `panel...` functions are intended to be used as the `panel` argument in `loa` plot calls. So, e.g.:

```
a <- 1:10
b <- 10:1
loaPlot(a+b~a*a, panel=panel.zcasePiePlot)
loaPlot(a+b~a*a, panel=panel.zcasePieSegmentPlot)
```

**Note**

Functions in development. Arguments may change, e.g.:

`panel.zcasePieSegmentPlot` includes the argument `zcase.rescale`. This normalises data within each `zcase` before generating the pie segments. This might not stay.

**Author(s)**

Karl Ropkins

**References**

These function makes extensive use of code developed by others.

`lattice`: Sarkar, Deepayan (2008) *Lattice: Multivariate Data Visualization with R*. Springer, New York. ISBN 978-0-387-75968-5

**See Also**

In `loa`: [loaPlot](#), [panelPal](#)

In other packages: [xyplot](#) in `lattice`.

## Examples

```
## Example 1
## plotting georeferenced pie plots

# Using a subsample of lat.lon.meuse
temp <- lat.lon.meuse[sample(1:155, 15),]

## Not run:
# plot Cu/Pb/Zn pie plots at sampling locations
  loaPlot(copper+lead+zinc~longitude*latitude,
          panel=panel.zcasePiePlot, data=temp)
# then rescale smaller pie segments on the fly
## End(Not run)

loaPlot(copper*10+lead*4+zinc~longitude*latitude,
        panel=panel.zcasePiePlot, data=temp)
```

---

## 2.4.specialist.panels *Special panel functions 04*

---

### Description

In development specialist panel functions

### Usage

```
panel.compareZcases(x=x, y=y, z=NULL, ...,
                    loa.settings = FALSE)
```

### Arguments

<code>x, y, z</code>	Standard plot data series, typically vectors.
<code>...</code>	Additional arguments, typically passed on.
<code>loa.settings</code>	loa options, to be handled by panelPal.

### Details

The `panel.compareZcases` generates a simple plot which compares z and y elements.

**Value**

These `panel...` functions are intended to be used as the `panel` argument in `loa` plot calls. So, e.g.:

```
x <- 1:10
y <- 1:10
z <- y + rnorm(10)
loaPlot(z~x*y, panel=panel.compareZcases, col.regions="Reds")
```

**Note**

These are ad hoc `panel...` functions. Not sure of their life expectancy...

**Author(s)**

Karl Ropkins

**References**

This function makes extensive use of code developed by others.

`lattice`: Sarkar, Deepayan (2008) *Lattice: Multivariate Data Visualization with R*. Springer, New York. ISBN 978-0-387-75968-5

**See Also**

In `loa`: [loaPlot](#), [panelPal](#).

In other packages: [xyplot](#) in `lattice`.

---

3.1.example.data      *example data for use with loa*

---

**Description**

Example data intended for use with examples in `loa`.

**Usage**

```
lat.lon.meuse
```

```
roadmap.meuse
```



**Format**

lat.lon.meuse is a modified form of the meuse data set taken from the sp package. Here, coordinate (x,y) elements have been transformed to latitudes and longitudes and the object class has been converted from SpatialPointsDataFrame to data.frame.

roadmap.meuse is a previously downloaded map intended for use with off-line plot examples using lat.lon.meuse.

**Details**

lat.lon.meuse was generated using method based on mzn object production in <https://github.com/etes/Geoprocessing/blob/master/heatmap.R>.

```
library(sp); library(gstat); library(rgdal)
data(meuse)
coordinates(meuse) =~ x + y
proj4string(meuse) = CRS("+init=epsg:28992")
meuse1 = spTransform(meuse, CRS("+init=epsg:4326"))
meuse2=as.data.frame(meuse1)
mzn=meuse2[,c(14,13,4)]
names(mzn)<-c("Latitude", "Longitude", "zinc")
```

roadmap.meuse was generated using:

```
GoogleMap(zinc~latitude*longitude, data=lat.lon.meuse, size=c(450,500), mptype="roadmap")
roadmap.meuse <- getMapArg()
```

**References**

For meuse:

M G J Rikken and R P G Van Rijn, 1993. Soil pollution with heavy metals - an inquiry into spatial variation, cost of mapping and the risk evaluation of copper, cadmium, lead and zinc in the floodplains of the Meuse west of Stein, the Netherlands. Doctoraalveldwerkverslag, Dept. of Physical Geography, Utrecht University

P.A. Burrough, R.A. McDonnell, 1998. Principles of Geographical Information Systems. Oxford University Press.

Stichting voor Bodemkartering (Stiboka), 1970. Bodemkaart van Nederland : Blad 59 Peer, Blad 60 West en 60 Oost Sittard: schaal 1 : 50 000. Wageningen, Stiboka.

For sp:

Roger S. Bivand, Edzer J. Pebesma, Virgilio Gomez-Rubio, 2008. Applied spatial data analysis with R. Springer, NY. <http://www.asdar-book.org/>

Pebesma, E.J., R.S. Bivand, 2005. Classes and methods for spatial data in R. R News 5 (2), <http://cran.r-project.org/doc/Rnews/>.

**Examples**

```

## data structure of lat.lon.meuse

head(lat.lon.meuse)

## Use a subsample of lat.lon.meuse

temp <- lat.lon.meuse[sample(1:155, 15),]

## various loaPlot examples
## using lat.lon.meuse

loaPlot(~longitude*latitude, data=temp)

loaPlot(cadmium~longitude*latitude, data=temp)

loaPlot(cadmium~longitude*latitude, col.regions=c("green", "red"),
        data=temp)

loaPlot(cadmium*50+copper*10+lead*2+zinc~longitude*latitude, panel.zcases = TRUE,
        col.regions=c("green", "red"),
        key.z.main="Concentrations", data=temp)

## (off line) GoogleMap example
## using lat.lon.meuse and roadmap.meuse

GoogleMap(zinc~latitude*longitude, data=temp,
          map=roadmap.meuse, col.regions=c("grey", "darkred"))

# Note 1:
# With loaPlot and GoogleMap, note latitude, longitude axes
# assignments:
# loaPlot plots z ~ x * y | cond.
# GoogleMap plots z ~ lat * lon | cond (z ~ y * x | cond)

# Note 2:
# Here, the map argument is supplied so example works off-line.
# If not supplied and R is on-line, GoogleMap will (try to) get map
# from the Google API. Look at:
## Not run:
  GoogleMap(zinc~latitude*longitude, data=lat.lon.meuse,
            col.regions=c("grey", "darkred"))
## End(Not run)
# (The map will appear slightly different, because non-default
# size and mptype settings were used to make roadmap.meuse. See above.)

```

---

4.1.panel.pal	<i>panelPal</i>
---------------	-----------------

---

**Description**

lattice plot management using the loa function `panelPal`

**Usage**

```
panelPal(ans, panel = NULL, preprocess = FALSE,
         reset.xylims = FALSE, legend = NULL,
         by.group = NULL, by.zcase = NULL, ...)
```

```
panelPal.old(x, y, subscripts, at, col.regions, ...,
            panel = panel.xyplot, ignore = NULL,
            group.fun = NULL)
```

```
loaHandler(panel = NULL, ...)
```

**Arguments**

<code>ans</code>	For <code>panelPal</code> only, a standard trellis object, such as that generated by <code>lattice</code> function <code>xyplot</code> .
<code>panel</code>	A panel function, e.g. <code>panel.xyplot</code> . If supplied in <code>panelPal</code> call, typically the one used to generate <code>ans</code> . If supplied in <code>panelPal.old</code> , the panel that is intended to be used when generating a plot.
<code>preprocess</code> , <code>reset.xylims</code> , <code>legend</code> , <code>by.group</code> , <code>by.zcase</code>	For <code>panelPal</code> only, loa plot management arguments. <code>preprocess</code> : Logical, should the supplied panel function be preprocessed? <code>reset.xylims</code> : Logical, should the plot dimensions be reset if changed, e.g. by preprocessing? <code>legend</code> : the legend as with standard lattice plots, <code>by.group</code> : a vector of plot argument names to be linked to any group conditioning, <code>by.zcase</code> : a vector of plot argument names to be linked to any z case conditioning See Details below.
<code>...</code>	Additional arguments, typically passed on.
<code>x</code> , <code>y</code> , <code>subscripts</code> , <code>at</code> , <code>col.regions</code>	For <code>panelPal.old</code> only, panel arguments passed down to individual panels.
<code>ignore</code>	Any additional arguments that <code>panelPal.old</code> should ignore and pass on to panel unchecked/unmodified.
<code>group.fun</code>	Fine control of the standard lattice plot argument group. It can be a vector or list containing the same number of elements as there are groups. These can be functions (or the names of functions as characters) setting individual functions for group or sets of parameters to be evaluated using the panel function. For

example, the current NULL default generates a list of `col` and `pch` settings that produce a conventional grouped scatter plot output when the `group` argument is applied to the `panel` default `panel.xyplot`. See Details below.

## Details

`panelPal` provides a range of generic plot management features.

Firstly, it allows plot as well as panel defaults to be managed by the `panel...` function. This allows the panel developer to control plot-level components of the plot output, e.g. which key to use with the plot and what default settings to apply to it. See example 1 below.

Secondly, it uses a generalised extension of the subscripting methods described by Deepayan Sarkar in Chapter 5 of *Lattice* (see sections on scatterplots and extensions) to automatically handle plot argument subscripting, demonstrated in example 2 below.

Thirdly, it applies an extension of the method used by the hexbin lattice panel to pass hex cell counts (calculated in panels) to the plot key and standardise the assignment of associated parameters within all panels to provide more general panel-to-panel and panel-to-scale. The method is briefly discussed in Chapter 14 of Sarkar.

This method has also been extended by isolating processing and plot components of the `panel...` function operation allowing results of any calculations made in-panel to be retained rather than lost when plot is generated.

Fourthly, `group...` and `zcase...` arguments can be used to manage plot group and `zcase` based plot outputs.

Some `panelPal` are implemented if specially structured (or `loa`-friendly) `panel...` functions are supplied. These are illustrated in the final example below.

`loaHandler` is a workhorse that assesses information in '`loa`' friendly `panel...` functions. As well as checking this, `loaHandler` also checks the supplied panel for any default plot settings. This allows users to manage the appearance of a plot from the panel or automatically associated color keys.

## Value

Both `panelPal` and `panelPal.old` are intended to be used with `trellis` plot outputs.

`panelPal` should be employed retrospectively. So, for example:

```
p1 <- xyplot(...)
panelPanel(p1, ...)
```

The previous version, currently retained as `panelPal.old`, was employed developed as a `panel...` function wrapper and intended to be employed within the plot call. So, for example:

```
xyplot(..., panel = function(...) panelPal(..., panel=panel))
```

Because it is run within the plot call, and therefore within each panel called, it does not provide features that require panel-to-plot, panel-to-key or panel-to-panel communication.

`loaHandler` returns either a logical (`FALSE` if not `loa` 'friendly'; `TRUE` if `loa` 'friendly') or a list of default arguments to be used when plotting.

**Note**

The `by.group` and `by.zcase` arguments of `panelPal` and the `group.fun` argument of `panelPal.old` are all currently under review. Please do not use these.

**Author(s)**

Karl Ropkins

**References**

These function makes extensive use of code developed by others.

`lattice`:

Sarkar, Deepayan (2008) *Lattice: Multivariate Data Visualization with R*. Springer, New York. ISBN 978-0-387-75968-5

`hexbin`:

Dan Carr, ported by Nicholas Lewin-Koh and Martin Maechler (2013). `hexbin`: Hexagonal Binning Routines. R package version 1.26.2. <http://CRAN.R-project.org/package=hexbin>

`panelPal.old` and `panelPal` both apply an extension of the subscribing methods described by Deepayan Sarkar in Chapter 5 of *Lattice* (see sections on scatterplots and extensions) to automatically handle plot argument subscribing.

`panelPal` applies an extension of the method used by hex bin lattice panel to communicate hex cell counts (calculated in panels) panel-to-panel and panel-to-scale. The method is briefly discussed in Chapter 14 of Sarkar.

**See Also**

[lattice](#), [xyplot](#),

**Examples**

```
## the combination of panelPal and specially
## structured panel... functions provides
## several additional plot features:

## example 1
## plot management from the panel... functions.

# loaHandler can used to see if a panel is loa-friendly

loaHandler(panel.xyplot) #FALSE
loaHandler(panel.loaPlot) #panel defaults

# note that these include a list called
# default.settings. These are settings that are
# automatically added to the plot call.

# Here this assigns a specialist key to that
# panel. However, the same mechanism can also
```

```
# be used to turn off plot elements like the
# standard lattice axes, when using in panel
# alternatives

# first some silly data

a <- rnorm(1000)
b <- rnorm(1000)

# now compare:

# default plot
# note bubble plot style key

loaPlot(a~a*b)

# bin plot
# with classic color key

loaPlot(a~a*b, panel = panel.binPlot)

## example 2
## automatic subscripting with loa

# Other arguments are not automatically
# aligned with the main plots.

# For example, consider the data:

a <- 1:10
ref <- rep(1:2, each=5)

# and associated lattice xyplot output:

xyplot(a~a|ref, col=ref, pch=19)

# Here, the 'col' argument does not
# automatically track plot conditioning.

# With lattice plots you need to assign
# arguments you want to track in this
# manner using subscripts, as discussed
# in Lattice Chapter 5.

# Now compare a similar loaPlot:

loaPlot(~a*a|ref, col=ref, pch=19)

# Here, panelPal automatically handles
# such subscripting. It extends this
# assumption to all supplied arguments.
```

```
# For example, try
## Not run:
  loaPlot(~a*a|ref, col=ref, pch=ref)
  loaPlot(~a*a|ref, col=ref, pch=1:10)
## End(Not run)
```

---

## 4.2.plot.structure.handlers

*Handler functions for plot structure arguments.*

---

### Description

Function(s) for handling (front end) plot arguments like `x` and `strip` that manage the plot structure.

### Usage

```
formulaHandler(x, data = NULL, groups = NULL, ...,
  expand.plot.args = TRUE, formula.type = "z~x*y|cond", panel.zcases = FALSE,
  coord.conversion = NULL, lattice.like = NULL, check.xy.dimensions = TRUE,
  check.coord.dimensions = TRUE, get.zcase.dimensions = TRUE,
  output = "extra.args")

matrixHandler(x, data = NULL, row.values=NULL, column.values=NULL,
  ...)

stripHandler(..., striplab=NULL)

getZcaseDimensions(...)
```

### Arguments

<code>x</code>	(For <code>formulaHandler</code> ) A formula or matrix ( <code>matrixHandler</code> ) intended to be used to generate a lattice plot. See Below.
<code>data</code>	If supplied, the assumed source of the elements of formula <code>x</code> , typically a <code>data.frame</code> .
<code>groups</code>	If supplied, the grouping argument to be used with <code>x</code> and <code>data</code> .
<code>...</code>	Additional arguments are passed on to related functions.
<code>expand.plot.args</code>	For <code>formulaHandler</code> only, logical. Should any short elements of the plot structure be expanded?
<code>formula.type</code>	For <code>formulaHandler</code> only, character vector or function. The plot structure to be used when generating the plot, e.g. <code>z ~ x * y   cond</code> for <a href="#">loaPlot</a>
<code>panel.zcases</code>	For <code>formulaHandler</code> only, logical. Should <code>zcase</code> arguments, e.g. <code>z1</code> and <code>z2</code> in <code>z1 + z2 ~ x * y   cond</code> , be treated as panel conditioning terms rather than grouping terms?

<code>coord.conversion</code>	For <code>formulaHandler</code> only, function. If supplied, the conversion to use to convert coordinate information supplied using other coordinate systems to $(x, y)$ .
<code>lattice.like</code>	For <code>formulaHandler</code> only, list. For preprocessing, a list of plot terms that can be passed directly to <code>lattice/loa</code> plots.
<code>check.xy.dimensions</code> , <code>check.coord.dimensions</code>	For <code>formulaHandler</code> only, logicals. Should the formula structure be tested before attempting to generate the plot? See Note below.
<code>get.zcase.dimensions</code>	For <code>formulaHandler</code> only, logical. Should the dimensions of any multiple zcases be calculated? See Note below.
<code>output</code>	For <code>formulaHandler</code> only, character vector. The format to return function output in.
<code>row.values</code> , <code>column.values</code>	For <code>matrixHandler</code> only, row and column values to be assigned to supplied matrix <code>x</code> .
<code>striplab</code>	For <code>stripHandler</code> only, character vector. If supplied, the label to add to the panel strip when conditioning is applied. By default, it applies the standard <code>lattice</code> convention, i.e., show for numerics.

## Details

`formulaHandler` manages the formula component or `x` element of `loa` plot functions.

For example, for `loaPlot` it assumes the general formula structure  $z \sim x * y \mid \text{cond}$ , and applies it in a similar fashion to the `lattice` function `levelplot`.

Within the formula part of the plot call `x` and `y` are the horizontal and vertical axes, `z` is any additional information to be used in point, symbol, surface or glyph generation, and `cond` any additional conditioning to be applied. (The coordinates, `x` and `y`, are required elements; `z` and `cond` are typically optional.)

`matrixHandler` converts a matrix supplied as `x` element of a `loa` plot to a formula and associated data. If `row.values` and `column.values` are supplied, these are used as  $(x, y)$  values for the supplied matrix.

`stripHandler` manages the strip component of `loa` plot functions.

`getZcaseDimensions` tracks the dimensions of multiple `z`

## Value

`formulaHandler` returns a list, containing the plot elements defined in the supplied formula.

`matrixHandler` returns a list containing all supplied arguments, subject to the following modifications: matrix `x` converted to formula ( $z \sim x * y$ ); data, replaced with supplied matrix content; `xlim` and `ylim`, added is not supplied.

`stripHandler` returns a list containing all supplied arguments, subject to the following modifications: `strip`, Updated or generated if not supplied; `striplab`, added to `strip` via the `strip` argument `var.name`, if this is undeclared in call.



`getZcaseDimensions` returns a list containing all the supplied arguments, plus two additions arguments (if supplied in the call): `zcase.zlim` and `z.rowsum.lim`. `zcase.zlim` is a list of `lim` values, one for each `zcase`. `z.rowsum.lim` is the range of 'by-row' sums of `zcases`. These are calculated using any `zcase` information supplied in the call.

### Note

These function are in development and may be subject to changes.

The current version of `formulaHandler` includes code from the `stats` function `get_all_vars`. It operates in a similar fashion to the previous version but checks `zcase` dimensions.

The previous version of `formulaHandler` was a wrapper for the `lattice` function `latticeParseFormula`. This version of `formulaHandler` was updated to simplify multiple `z` argument handling.

The latest version of `formulaHandler` includes more flexible `formula.type` handling. For example, it can now handle formulas that have more than two coordinates. As a result the `check.xy.dimensions` argument was replaced with a `check.coord.dimensions` argument. The previous argument will however remain in the function formals and function as before until all related code has been updated.

The latest version of `formulaHandler` uses `getZcaseDimensions` to calculate the dimensions of `z` if it is multi-part, e.g.  $z_1 + z_2 + \text{etc} \sim x * y$  rather than  $z \sim x * y$ .

The current version of `matrixHandler` is based on code from `levelplot.matrix` in `lattice`. If used with `x` and `data` arguments it will overwrite `data` with the matrix content.

### Author(s)

Karl Ropkins

### References

This function makes extensive use of code developed by others.

`lattice`: Sarkar, Deepayan (2008) *Lattice: Multivariate Data Visualization with R*. Springer, New York. ISBN 978-0-387-75968-5

### See Also

In `loa`: [loaPlot](#); [panelPal](#)

In other packages: [levelplot](#) in `lattice`.

---

4.3.lims.and.scales.handlers

*Plot lims and scales handlers*

---

### Description

In development functions for lims and scales handling with `lattice` plots.

**Usage**

```

limsHandler(x=NULL, y=NULL, z=NULL, ..., lim.borders = 0.05)

localScalesHandler(scales = NULL, ..., allowed.scales =c("x", "y"),
  disallowed.scales = NULL, remove.box = FALSE)

panel.localScale(x.loc, y.loc, lim, ..., panel.scale = NULL,
  label.before = TRUE, x.offset = NULL, y.offset = NULL,
  axis = TRUE, ticks = TRUE, annotation = TRUE)

yscale.component.log10(...)

xscale.component.log10(...)

```

**Arguments**

<code>x, y, z</code>	<code>x, y</code> and/or <code>z</code> data series.
<code>lim.borders</code>	numeric vector, giving the relative border to extend <code>...lim</code> ranges by when generating axes or scales. The <code>lim.borders</code> are relative proportions. So, the default setting of 0.05 adds an extra +/- 5 are supplied the first three are treated as <code>codex</code> , <code>y</code> and <code>z</code> <code>lim.borders</code> , respectively. If less than three values are supplied, the three values are generated by wrapping. So, the default setting of 0.05 is equivalent to <code>c(0.05, 0.05, 0.05)</code> .
<code>scales, panel.scale</code>	A list of elements like the <code>scales</code> argument used with <code>lattice</code> functions. Current default elements <code>draw</code> (= TRUE), <code>arrows</code> (= FALSE), <code>tick.number</code> (= 5), <code>abbreviate</code> (= FALSE), <code>minlength</code> (= 4), and <code>tck</code> (= 1).
<code>...</code>	Additional arguments.
<code>allowed.scales</code>	A character vector containing the names of the axes to be generated for as part of a local axis.
<code>disallowed.scales</code>	A character vector containing the names of any axes that are not required. Note: If found, these are removed from <code>scales</code> before evaluation.
<code>remove.box</code>	A logical, default FALSE. Should the box <code>lattice</code> typically places around standard plots be removed? This can be useful if you are using a <code>panel...</code> function to generate axes within the plot.
<code>x.loc, y.loc, lim</code>	two member vectors setting the <code>x</code> and <code>y</code> locations where the scale is to be drawn ( <code>x.loc</code> and <code>y.loc</code> ), and the limits of the range to be annotated on the scale ( <code>lim</code> ). Note: These are currently handled 'as is', i.e. for both locations and limit, the first element is the start point and the second is the end point, and any other elements are ignored.

`label.before`, `x.offset`, `y.offset`

Scale annotation overrides. `label.before` is a logical, which controls the position of annotation, tick marks and/or arrows, etc relative to the scale line (i.e., above/left before or below/right after). By default `panel.localScale` generates tick marks and labels at right angles to the scale line/axis. `x.offset` and `y.offset` force the offsets when adding tick marks and annotation.

`axis`, `ticks`, `annotation`

If supplied, fine controls for the appearance of the axis line, axis tick marks and axis annotation on the generated scale. These can be vectors, in which they are assumed to be color assignments, or lists of common plot parameters, such as `col`, `lty`, `lwd`, etc.

## Details

`limsHandler` generates `xlim`, `ylim` and/or `zlim` ranges for axes or color scales for use in a range of plots.

`localScalesHandler` provides a relatively crude mechanism for the removal of conventional lattice plot axes and the generation of alternative axes using a `panel...` function like `panel.localScale`.

## Value

`limsHandler` returns a list containing `...lim` ranges for any of the elements `codex`, `y` and/or `z` supplied to it.

`localScalesHandler` returns a list containing the elements: `scales`, `panel.scales` and possibly `par.settings`. `scales` turns off the standard axes annotation. `panel.scales` is a list of named elements, one per named axis, describing the axis layout. If `remove.box = TRUE`, the additional argument `par.settings` is also supplied.

All arguments should be passed on to the associated plot.

`panel.scales` or axis-specific elements in `panel.scales` can then be evaluated by an associated `panel...` function run from within the lattice plot call. This would typically take the form:

```
panel.my.axis(panel.scale = panel.scale$axis, ...)
```

`panel.localScale` is a local axis/scale plotter. It can be used in combination with `localScalesHandler`, and should be called once for each axis that is required, or it can be used 'stand alone' `panel` to add a local scale to a lattice plot.

`yscale.component.log10` and `xscale.component.log10` are simple axis transforms for use with `log` to the base 10 transformed plot axes.

## Note

`panel.localScale` is currently in revision. Scale arrows are currently not available.

## Author(s)

Karl Ropkins

**References**

These function makes extensive use of code developed by others.

lattice: Sarkar, Deepayan (2008) Lattice: Multivariate Data Visualization with R. Springer, New York. ISBN 978-0-387-75968-5

**See Also**

In other packages: [xyplot](#) in [lattice](#).

**Examples**

```
## See trianglePlot Example 2 for example application
```

---

4.4.cond.handlers      *Plot Conditioning*

---

**Description**

Plot conditioning handling

**Usage**

```
condsPanelHandler(..., conds = NULL, panel = NULL,
  by.cond = NULL, process = TRUE, plot = TRUE)

groupsPanelHandler(..., groups = NULL, panel = NULL,
  by.group = NULL, process = TRUE, plot = TRUE)

zcasesPanelHandler(..., zcases = NULL, panel = NULL,
  by.zcase = NULL, process = TRUE, plot = TRUE)

groupsAndZcasesPanelHandler(panel=NULL, ...,
  plot = TRUE, process = TRUE)

groupsHandler(z = NULL, groups = NULL, ..., group.ids = NULL,
  handler = "zzz")

zcasesHandler(z = NULL, zcases = NULL, ..., zcases.ids = NULL,
  handler = "zzz")

groupsAndZcasesHandler(..., loa.settings = NULL)

stepwiseZcasesGlyphHandler(zcases = NULL, ..., zcase.ids = NULL,
  panel.elements = NULL, loaGlyph = NULL)
```

**Arguments**

...	Additional arguments. See Notes below.
conds, panel, by.cond	For all supplied additional arguments, conds is a vector of conditioning indices. This is typically a logical, numeric, factor or character vector which can be used to assign other elements undeclared call arguments to specific subsets. panel identifies the panel... function, and should also be supplied so loa can manage processing and plot activities correctly. by.cond identifies routine plot operations associated with the requested conditioning. This can be a list of plot arguments or panel... functions that should be associated with the requested conditioning. See process and plot below and associated Notes.
plot, process, loa.settings	Passed to and handled by <a href="#">panelPal</a> . For panels that can be preprocessed, plot and process turn off or on processing and the plotting steps of the panel code. See <a href="#">panelPal</a> Help documentation from further details.
groups, by.group	As conds and by.cond but for grouping.
zcases, by.zcase	As conds and by.cond but for zcase condition.
z, handler	The z data series and any associated plot arguments that need special handling.
group.ids, zcases.ids, zcase.ids	If given, vectors of the unique cases in groups and zcases, respectively.
panel.elements	If given, the names of all plot arguments that have been vectorized by panelPal.
loaGlyph	(For <a href="#">stepwiseZcasesGlyphHandler</a> only), the loa glyph to draw. See <a href="#">loa.glyphs</a> for further details.

**Details**

NOTE: These functions are currently in development and may be subject to changes.

`condsPanelHandler` is a general purpose function that can be used to routinely manage plot conditioning within a `panel...` function call. It takes all undeclared arguments are supplied to it, and subsets them by unique case in the supplied `conds` argument. Then it modifies each of these based on the associated elements of `by.cond` and processes and/or plots the results depending on `process` and `plot` settings.

`groupsPanelHandler` is similar but is intended for use with the plot call argument `groups`.

`zcasesPanelHandler` is similar but is intended for use with arguments conditioned within the `z` term of the plot formula. So, for example, for unique handling of `z1` and `z2` cases in the plot `loaPlot(z1+z2~x*y)`.

`groupsAndZcasesPanelHandler` is a wrapper for `groups` and `zcase` that allows users to simultaneously and uniquely handle both types of conditioning.

`stepwiseZcasesGlyphHandler` is a `...Handler` function for generating glyph plots based on multiple `z` inputs.

**Value**

All `..PanelHandlers` functions are intended for use with `panelPal`. Using different combinations of these allows plot developers a high degree of flexibility.

**Note**

This function is in development and may be subject to changes.

**Author(s)**

Karl Ropkins

**References**

This function makes extensive use of code developed by others.

`lattice`: Sarkar, Deepayan (2008) *Lattice: Multivariate Data Visualization with R*. Springer, New York. ISBN 978-0-387-75968-5

**See Also**

[panelPal](#)

For information on related functions in other packages, see

[lattice: xyplot](#); [panel.xyplot](#); and [panel.levelplot](#).

---

4.5.plot.argument.handlers

*Common plot argument handlers*

---

**Description**

Functions for use the routine handling of some common plot arguments.

**Usage**

```
cexHandler(z = NULL, cex = NULL,  
           cex.range = NULL, expand.outputs = TRUE,  
           ref = NULL, ..., xlim = NULL)
```

```
colHandler(z = NULL, col = NULL,  
           region = NULL, colorkey = FALSE, legend = NULL,  
           pretty = FALSE, at = NULL, cuts = 20,  
           col.regions = NULL, alpha.regions = NULL,  
           expand.outputs = TRUE, ref = NULL,  
           ..., xlim = NULL, output="col")
```

```
colRegionsHandler(...)
```

```
pchHandler(z = NULL, pch = NULL, pch.order = NULL,
           expand.outputs = TRUE, ref = NULL, ...,
           xlim = NULL)
```

```
scalesHandler(...)
```

```
zHandler(z = NULL, expand.outputs = TRUE,
         ref = NULL, ...)
```

## Arguments

- z** If supplied, a vector of values intended to be used as a scale when assigning a property. For `cexHandler`, the `cex` of, e.g., points on a scatter plot. Here, size scales are managed using a reference range `cex.range`, but superseded by `cex` settings, if also supplied. For `colHandler`, the color of, e.g., points on a scatter plot. Here, color scales are managed using a `colorkey` method similar to that used by the `lattice` function `levelplot`, see below (arguments `region`, `colorkey`, `pretty`, `at`, `cuts`, `col.regions` and `alpha.regions`). If `z` is `NULL` or not supplied, all colors are set by `col` if supplied or as the default `lattice` symbol color if both `z` and `col` are not supplied. For `pchHandler`, the `pch` of, e.g., points on a scatter plot. Here, plot symbols are managed using a reference vector `pch.order`, but superseded by `pch` settings, if also supplied. For `zHandler`, any vector that should be expanded by wrapping to a given length, e.g. the length of the `x` (or `y`) data series to plotting.
- cex, col, pch** For associated handlers, the parameter value(s) to be managed (i.e., `cex` for `cexHandler`, etc. Note: In all cases if these are not `NULL` these supersede any supplied `z` or `...Handler` modification.
- cex.range** If supplied, the range for `z` to be rescaled to when using this to generate a `cex` scale. NOTE: `cex.range = FALSE` disables this `cex` scaling and uses `z` values directly; `cex.range = TRUE` applied default scaling, equivalent to `cex.range = c(0.75, 3)`.
- region, colorkey, legend, pretty, at, cuts, col.regions, alpha.regions** The colorscale settings to be used when generating a `colorkey`. The most useful of these are probably `col.regions` which can be used to reset the color scale, `alpha.regions` which sets the `col.region` alpha transparency (0 for invisible to 1 for solid) and `colorkey` which can be a logical (forcing the `colorkey` on or off) or a list of components that can be used to fine-tune the appearance of the `colorkey`. Note: The generation of colorscales is handled by `RColorBrewer`.
- pch.order** A vector of symbol ids (typically the numbers 1 to 24) to be used when plotting points if, e.g. using a scatter plot. By default, all points are plotted using the first of these `pch` ids unless any conditioning (e.g. grouping or `zcase` handling)

	is declared and linked to <code>pch</code> , in which symbols are assigned in series from <code>pch.order</code> .
<code>expand.outputs</code> , <code>ref</code>	<code>expand.outputs</code> is a Logical (default TRUE): should outputs be expanded to the same length as <code>ref</code> ? This can be useful if, e.g., coloring points on a scatter plot that may be conditioned and therefore may require subscript handling, in which case <code>ref</code> could be the x or y data series, or any other vector of the same length. Note: if <code>ref</code> is not supplied <code>expand.outputs</code> is ignored.
<code>zlim</code>	The range over which the scale is to be applied if not <code>range(z)</code> .
<code>output</code>	For <code>colHandler</code> . The function output. Either the <code>col</code> vector alone ( <code>output='col'</code> ) or the full list of color parameters.
<code>...</code>	Additional arguments, often ignored.

### Details

The `...Handler` functions are argument handlers intended to routinely handle some common activities associated with plotting data.

`cexHandler` manages symbol sizes. It generates a (hopefully) sensible `cex` scale for handling plot symbol size based on a supplied input (`z`).

`colHandler` manages colors. It works like the `colorkey` in `levelplot` in `lattice`, to generate a colorscale based on a supplied input (`z`).

`colRegionsHandler` is a wrapper for `colHandler` that can be used to with the `col.regions` argument.

`scalesHandler` is a crude method to avoid `scales` argument list structures.

`zHandler` expands (by wrapping) or foreshortens vectors.

### Value

`cexHandler` returns a vector, which can be used as the `cex` argument in many common plotting functions (e.g. `plot`, `xyplot`).

`colHandler` depending on `output` setting returns either the `col` vector or a list containing elements (`z`, `col`, `legend`, `at`, `col.regions` and `alpha.regions`), which can be used to create a `col` series scaled by `z` and an associated `colorkey` like that generated by `levelplot` for other `lattice` functions (e.g. `xyplot`).

`colRegionsHandler` returns a vector of color values suitable for use with the `col.regions` argument.

`scalesHandler` returns the supplied arguments modified as follows: all `scales...` arguments are converted into a single `list(...)`; all `scales.x...` and `scales.y...` argument are converted into `list(x=list(...))` and `list(y=list(...))`, respectively. so e.g. `scales.x.rot=45` generates `scales=list(x=list(rot=45))`.

`pchHandler` returns a vector of `pch` values of an appropriate length, depending on `expand.outputs` and `ref` settings.



**Note**

cexHandler recently revised. Default cex range now smaller, in line with feedback.  
scalesHandler might not be staying.

**Author(s)**

Karl Ropkins

**References**

These function makes extensive use of code developed by others.

lattice: Sarkar, Deepayan (2008) Lattice: Multivariate Data Visualization with R. Springer, New York. ISBN 978-0-387-75968-5

RColorBrewer: Erich Neuwirth <erich.neuwirth@univie.ac.at> (2011). RColorBrewer: Color-Brewer palettes. R package version 1.0-5. <http://CRAN.R-project.org/package=RColorBrewer>

**See Also**

In other packages: See [xyplot](#) in [lattice](#).

**Examples**

```
#some trivial data
a <- 1:10

## Example 1
## Simple plot with cex handling

myplot1 <- function(x, y, z = NULL, cex = NULL,
                   cex.range = NULL, ...){

  #set cex
  cex <- cexHandler(z, cex, cex.range)

  #plot
  xyplot(y~x, cex = cex,...)
}

myplot1(a, a, a)

# compare
## Not run:
myplot1(a, a)           #like plot(x, y)
myplot1(a, a, a*100)   #as myplot1(a, a, a)
                        #because cex scaled by range

myplot1(a, b, c,
        cex.range = c(1,5)) #cex range reset
myplot1(a, b, c,
```

```

        cex.range = c(10,50),
        cex = 1)          #cex supersedes all else if supplied
## End(Not run)

## Example2
## plot function using lists/listUpdates

myplot2 <- function(x, y, z = NULL, ...){

  #my default plot
  default.args <- list(x = y~x, z = z,
                      pch = 20, cex = 4)

  #update with whatever user supplied
  plot.args <- listUpdate(default.args, list(...))

  #col Management
  plot.args$col <- do.call(colHandler, plot.args)
  do.call(xyplot, plot.args)
}

#with colorkey based on z case
myplot2(a, a, a)

# compare
## Not run:
myplot2(a, b, c,
        col.regions = "Blues") #col.regions recoloring
myplot2(a, b, c,
        col = "red")          ##but (again) col supersedes if supplied
## End(Not run)

# Note:
# See also example in ?listUpdate

```

### Description

Workhorse functions for routine use of keys in plots.

### Usage

```
keyHandler(key = NULL, ..., output = "key")
```

```
#keys

draw.loaPlotZKey(key = NULL, draw = FALSE, vp = NULL, ...)
draw.loaColorKey(key = NULL, draw = FALSE, vp = NULL, ...)
draw.loaColorRegionsKey(key = NULL, draw = FALSE, vp = NULL, ...)
draw.zcasePlotKey(key = NULL, draw = FALSE, vp = NULL, ...)
draw.ycasePlotKey(key = NULL, draw = FALSE, vp = NULL, ...)
draw.groupPlotKey(key = NULL, draw = FALSE, vp = NULL, ...)
draw.key.log10(key = NULL, draw = FALSE, vp = NULL, ...)
```

### Arguments

key	The key to be used.
...	Any additional arguments to be used to modify the the key before plotting.
output	The format to return the function output in. This is 'key' for all routine (in plot) use.
draw, vp	lattice and grid arguments using when plotting GROB objects. Generally, these can be ignored.

### Details

keyHandler is a general function that routine generates defaults arguments for add a key to a plot. draw...key functions are all specialist plots keys. They are typically modifications of or variations on similar functions in lattice, e.g. draw.key and draw.colorkey.

draw.loaPlotZKey is the default 'bubble plot' key used with [loaPlot](#).

draw.loaColorKey and draw.loaColorRegionsKey are variations on the draw.colorkey function in [lattice](#).

draw.zcasePlotKey, draw.ycasePlotKey and draw.groupPlotKey are simple legends based on zcase, ycase and group annotation.

draw.key.log10 is a simple legend for use with log to the base 10 transformed z scale.

### Value

keyHandler return a list of plot arguments to be used to generate a key .

When run within plot calls, the draw...key functions associated color keys. If they are used with loa plots and suitable panel... functions, color scales are automatically aligned.

### Note

In Development: Function structures may change in future package updates.

**Author(s)**

Karl Ropkins

**References**

These functions make extensive use of code developed by others.

`lattice`: Sarkar, Deepayan (2008) *Lattice: Multivariate Data Visualization with R*. Springer, New York. ISBN 978-0-387-75968-5

**See Also**

In other packages: See `xyplot` in `lattice`.

---

4.7.other.panel.functions

*Other panel functions argument handlers*

---

**Description**

In development panel functions for use with `lattice`

**Usage**

```
parHandler(scheme = NULL, ...)
```

```
#related
```

```
getArgs(source = TRUE, local.resets = TRUE,  
        user.resets = TRUE, is.scales.lines = FALSE,  
        elements = NULL, ..., defaults = list(),  
        defaults.only = FALSE)
```

```
getPlotArgs(defaults.as = "axis.line", source = TRUE,  
            local.resets = TRUE, user.resets = TRUE,  
            elements = NULL, ..., is.scales.lines = NULL,  
            defaults.only = TRUE)
```

```
isGood4LOA(arg)
```

**Arguments**

- `scheme` The color scheme to apply. This can be a list of parameters to apply or a character vector for a pre-defined scheme. Current pre-defined schemes include 'greyscale' (for black and white figures).
- `source`, `local.resets`, `user.resets` When recovering plot arguments with `getArgs` or `getPlotArgs`, places to search for relevant parameters. If supplied these would typically be vectors or lists. If vectors, they are assumed to be `col` setting. If lists, they are assumed to be lists of named parameters for inclusion. There are two cases that need to be handled specially: (1) some sources, `local.resets` and/or `user.resets` may contain both axis-specific and general information, e.g. For a scales list, parameters to be applied just to the x axis in `scales$x` and parameters to be applied to all scales in `scales`. In such cases these need to be checked in order (see `elements` below.) (2) Some sources, e.g. axis scales, contain both text and line parameters, with e.g. line settings declared as `col.line`, etc., rather than `col`, etc., (which are intended for use with text.) When supplied these need to be handled correctly (see `is.scales.lines` below). `local.resets` and `user.resets` are intended as overrides for the code developer and user, respectively. These can be logicals as well as vectors or lists. If logicals they turn on/off the associated plot components (using `isGood4LOA`). The check/update order is `source`, then `source$element`, then `local.reset`, then `local.reset$element`, then `user.reset`, then `user.reset$element`. This means that the developer always has last say regarding the default appearance of a plot component and the user always has the very last say from the command line if the `local.reset` is included as a formal argument in that plot.
- `is.scales.lines` When recovering arguments with `getArgs` or `getPlotArgs`, should `source` be treated as a lattice scales list? If so, and `source` is checked for line parameters, line-specific terms such as `col.line`, etc., will be recovered as `col`, etc., while general terms (meant for text in scales lists) will be ignored. (Note: `getPlotArgs` guesses this based on `defaults.as` if not supplied.)
- `elements` When recovering arguments with `getArgs` or `getPlotArgs`, this identifies the elements in `source`, `local.resets` and `user.resets` that may contain case-specific information. As with lattice handling of scales axis-specific information in `source$element(s)` is assumed to take priority over general information in `source`. (Note: if `elements` are not declared only general/top level information in `source`, `local.resets` and `user.resets` is considered at present.)
- ... Other arguments, often ignored.
- `defaults`, `defaults.only`, `defaults.as` When recovering arguments with `getArgs`, `defaults` is an optional 'fall-back' in case nothing is recovered from `source`, `local.resets` and `user.resets`. `defaults.only` is a logical: if TRUE only parameters named in `defaults` are searched for, otherwise all parameters are recovered. With `getPlotArgs`, `defaults.as` selects an appropriate default. This should be a trellis parameter name, e.g. 'axis.line', 'axis.text', etc. The function uses this to identify appropriate

plot parameters to search for/select, e.g. `pch`, `col`, `cex`, etc for `'plot.symbol'`, and to identify default values for each of these (if `defaults.only = TRUE`).

`arg` For `isGood4LOA` a plot argument that can used to turn a plot panel or panel component on or off.

### Details

`getArgs` returns a list of parameters/values based on `lattice`, developer and user settings. If multiple elements are identified as containing case-specific information, the list will contain one list of plot parameters for each named element.

`getPlotArgs` is a variation of `getArgs` intended for use with `panel...` and `l...` type `lattice` functions. It returns a list of plot parameters for different plot components, e.g. symbols, lines, or text.

`isGood4LOA` is a simple workhorse that checks if a supplied `arg` should be used by `loa`. (See value and note below.)

`parHandler` manages the default appearance of plots.

### Value

`getArgs` and `getPlotArgs` return lists of located parameters/values. For example, the call

```
getPlotArgs(default.as = "axis.line")
```

returns a list containing the `lattice` defaults for an axis line (`alpha`, `col`, `lty` and `lwd`) These can then be used in combination with appropriate `x` and `y` values in `l.lines`, or `panel.lines` calls. The arguments `local.resets` and `user.resets` can be added into the call to provide developer and user overrides. (See note below.)

`isGood4LOA` returns a logical (TRUE or FALSE), depending on the type of a supplied argument. This returns FALSE for NULL, for all FALSE logicals, and any `arg` that has previously been tagged as 'not wanted'.

`parHandler` returns a list a list suitable for use as `par.settings` with most `lattice` plots.

### Note

`getPlotArgs` is intended as a 'workhorse' for plot developers, to recover `lattice` settings, impose their own preferences on these, and in turn to provide users with similar options to quickly override developer settings.

`isGood4LOA` only exists because I, perhaps wrongly, equate `arg = NULL` with `arg = FALSE` when that argument is a component of a plot defined in the plot formal. For example, in `trianglePlot` I want `grids = NULL` to turn off the plot grids much like `grids = FALSE`, but got fed up always writing the same everywhere. Does not mean it is right, particularly useful or even clever...

The `getPlotArgs/isGood4LOA` combination is a first attempt at providing plot developers with a simple tool to integrate plot argument management by `lattice`, the plot developer and the plot user. It is intended to be applied in the form shown in the Examples below.

Axis, tick, grid and annotation handling in `trianglePlot` is intended to illustrate this type of application.

**Author(s)**

Karl Ropkins

**References**

These function makes extensive use of code developed by others.

lattice: Sarkar, Deepayan (2008) Lattice: Multivariate Data Visualization with R. Springer, New York. ISBN 978-0-387-75968-5

**See Also**

In other packages: See [xyplot](#) in [lattice](#).

**Examples**

```
#getPlotArgs/isGood4LOA notes

#in formals
#my.plot <- function(..., user.reset = TRUE, ...)

#in main code body
#local.resets <- [what developer wants]
#plot.arg <- getPlotArgs("[type]", source, local.reset, user.reset)

#in panel call
#(for panel off/on control)
#if(isGood4LOA(plot.arg)) panel...(..., plot.arg,...)

#in panel... function
#for panel component off/on control)
#if(isGood4LOA(plot.arg1)) panel...(..., plot.arg1,...)
#if(isGood4LOA(plot.arg2)) l...(..., plot.arg2,...)
#etc.
```

**Description**

Workhorse functions for routine list handling in loa and elsewhere.

**Usage**

```
listHandler(a, use = NULL, ignore = NULL,
            drop.dots=TRUE)
```

```
listUpdate(a, b, use = NULL, ignore = NULL,
            use.a = use, use.b = use,
            ignore.a = ignore, ignore.b = ignore,
            drop.dots = TRUE)
```

```
listExpand(a, ref = NULL, use = NULL,
            ignore = NULL, drop.dots = TRUE)
```

```
listLoad(..., load = NULL)
```

**Arguments**

a	A required list. The list to be modified.
b	For listUpdate only, a required second list, the contents of which are used to update a with.
use, use.a, use.b	Vectors, all defaults NULL. If supplied, a vector of the names of list entries to be used. Other entries are then discarded. use is applied to all supplied lists, while use.a, use.b, etc. can be used to subset a and b lists individually.
ignore, ignore.a, ignore.b	Vectors, default NULL. As with use, etc, but for entries to be ignored/not passed on for modification.
ref	For listExpand only, a vector, default NULL. A reference data series, the length of which is used as the expansion length to be applied when wrapping of list entries.
drop.dots	Logical, default TRUE. If TRUE, this removes "... " entries from list names before updating.
...	For listLoad only, any additional arguments.
load	For listLoad only, a character vector, default NULL. The names of any lists to be automatically generated from the additional arguments supplied as part of the command call.

**Details**

listHandler is a general function used by other list... functions for routine list preprocessing.

listUpdate is a list handler intended for use when managing user updates for default options (see examples).

listExpand is a list handler that expands vectors to a given reference length, intended for use for data wrapping.

listLoad is a list generator. See Note below.



**Value**

By default, all `list...` functions return results as lists.

`listHandler`, `listUpdate` and `listExpand` functions all return a modified (or updated) version of supplied list `a`.

`listLoad` (in-development) returns modified (or updated) version of additional arguments as a list. See Note below.

**Note**

`listLoad` is an in-development workhorse function that generates lists based on the supplied `load` argument.

It assumes each element of `load` is the name of an expected list and searches the associated additional arguments for arguments to populate it with using the rule `'[load].[arg]` is an element of list `[load]`'. So, for example, for a call including the arguments `load = 'key'` and `key.fun = draw.colorkey`, it would strip out both arguments and return `key = list(fun=draw.colorkey)`. Used in functions, it allowed list-in-list args that can be commonplace when modifying, for example, key elements of conventional `lattice` plots to be simplified.

**Author(s)**

Karl Ropkins

**References**

These functions make extensive use of code developed by others.

`lattice`: Sarkar, Deepayan (2008) *Lattice: Multivariate Data Visualization with R*. Springer, New York. ISBN 978-0-387-75968-5

**See Also**

[lattice](#), [xyplot](#),

**Examples**

```
## Example 1
## general

# two lists
list1 <- list(a = 1:10, b = FALSE)
list2 <- list(b = TRUE, c = "new")

# updating a with b
# keeps unchanged list1 entry, a
# updates changed list1 entry, b
# adds new (list2) entry, c
listUpdate(list1, list2)
```

```

## Example2
## use in plot functions
## to simplify formals

## Not run:
# some data
a <- 1:10
b <- rnorm(10,5,2)

#a bad plot function

badplot <- function(x, ...){

  #setting defaults in xyplot call itself
  xyplot(x = x, pch = 20, col = "red",
        panel = function(...){
          panel.grid(-1, -1)
          panel.xyplot(...)
          panel.abline(0,1)
        }, ...)
}

badplot(a~b)          #OK

# compare with
badplot(a~b, xlim = c(1,20)) #OK
badplot(a~b, col = "blue")  #not OK

# because col hardcoded into badplot function
# It is duplicated in call and '...'
# so user cannot update col

#a standard correction

stdplot <- function(x, pch = 20, col = "red", ...){

  #setting defaults in xyplot call itself
  xyplot(x = x, pch = 20, col = "red",
        panel = function(x=x, pch=pch, col=col, ...){
          panel.grid(-1, -1)
          panel.xyplot(x=x, pch=pch, col=col, ...)
          panel.abline(0,1)
        }, ...)
}

stdplot(a~b)          #OK
stdplot(a~b, col = "blue",
        xlim=c(1:20)) #also OK

# An alternative correction using lists and
# listUpdate that removes the need for formal

```

```

# definition of all modified plot arguments

myplot <- function(x, ...){

  #defaults I set for myplot form of xyplot
  mylist <- list(x = x, pch = 20, col = "red",
                panel = function(...){
                  panel.grid(-1, -1)
                  panel.xyplot(...)
                  panel.abline(0,1)
                })
  #plot
  do.call(xyplot, listUpdate(mylist, list(...)))
}

myplot(a~b)          #OK
myplot(a~b, col = "blue",
        xlim = c(1,20))  #also OK

## End(Not run)

```

---

4.9.loa.shapes

*loa shapes*


---

### Description

Simple shapes.

### Usage

```
loaPolygon(x, y, z=NULL, rot=NULL, ...,
           polygon = NULL, loa.scale = NULL)
```

```
loaCircle(..., polygon = NULL, radius = 1)
```

```
loaPieSegment(..., polygon = NULL, start = 0,
              angle=360, radius = 1, center=TRUE)
```

### Arguments

x, y	The x and y points at which to plot the requested shape.
z	If supplied a z term, most often used to set the size of the polygon.
rot	The angle to rotate the polygon by before drawing it.
...	Any additional arguments, usually passed on.

<code>polygon</code>	A list with elements <code>x</code> and <code>y</code> giving the polygon/shape to be plotted.
<code>loa.scale</code>	A list of parameters that can be used to fine-tune the polygon plotting.
<code>radius</code>	The radius to used when drawing either circles or pie segments.
<code>start, angle</code>	When drawing pie segments, <code>angle</code> the angle of of the segment and <code>start</code> point.
<code>center</code>	Should the segement begin and end at the center?

### Details

`loaPolygon` is a general function for drawing polygons. It is intended as an alternative to `lpolygon`, and other standard `loa...` shapes are typically wrappers for this function.

`loaCircle` draws a circle with an origin at  $(x, y)$ .

`loaPieSegment` draws a pie segment (or slice of cake) shape. It is typically used as building block for pie plots and other similar glyph structures.

### Value

All these functions generate simple shapes and are intended to be run within `panel...` functions as building blocks for more complex glyph type structures.

### Author(s)

Karl Ropkins

### References

These functions make extensive use of code developed by others.

`lattice`: Sarkar, Deepayan (2008) *Lattice: Multivariate Data Visualization* with R. Springer, New York. ISBN 978-0-387-75968-5

### See Also

In other packages: See [lrect](#), and similar, in [lattice](#)

---

5.1.plot.interactives *Interactive plot functions*

---

### Description

Recovering information from existing `lattice` plots.

**Usage**

```

getXY(n = -1, ..., unit = "native", scale.correction = NULL)

getLatLon(..., map = NULL, object = trellis.last.object(),
           scale.correction = function(x) {
             temp <- XY2LatLon(map, x$x, x$y)
             as.list(as.data.frame(temp))
           })

screenLatticePlot(object = trellis.last.object(), ...)

```

**Arguments**

n	If positive, the maximum number of points to locate. If negative (default), unlimited.
unit	The unit to use when reporting located points, by default "native".
scale.correction	The correction to apply if the plot has locally scaled axes. See Note below.
map, object	For getLatLon only. The plot layer as generated makeMapArg and the plot object generated by GoogleMap. The map is strictly required as a reference when converting plot points to associated latitude, longitude values, and can be supplied directly as map, or recovered from the plot, which can be supplied as object. If neither are supplied (as in default use), the function attempts to recover map from the last lattice plot via trellis.last.object.
...	Additional arguments, passed on to related functions. These may be subject to revision, but are currently: <a href="#">trellis.focus</a> for panel selection (if working with multi-panel plots) and <a href="#">lpoints</a> to set point properties (if marking selected points). For getLatLon, additional arguments are also passed to <a href="#">XY2LatLon</a> for x, y to latitude, longitude rescaling.

**Details**

getXY is an interactive function which returns the locations of points on a plot selected using the mouse (left click to select points; right click and stop to end point collection; escape to abort without returning any values).

It is a wrapper for the [grid](#) function [grid.locator](#) that behaves more like [locator](#), the equivalent function intended for use with [plot](#) outputs.

By default getXY selections are not automatically marked. Adding common plot parameters to the function call overrides this behaviour, e.g. to add red symbols and lines.

```
ans <- getXY(col = "red", pch = 4, type = "b")
```

getXY also provides a mechanism to handle data plotted on locally scaled axes. See Note below.

getLatLon is wrapper for getXY for use with GoogleMap outputs and other similarly georeferenced plots. See Note below.

screenLatticePlot is a crude plot screening function. It is currently in development.

**Value**

`getXY` returns the x and y coordinates of the selected points on a plot as a list containing two components, x and y.

`getLatLon` returns the latitude and longitude values of the selected points on a map as a list containing two components, lat and lon.

**Note**

`getXY` recovers the (x, y) coordinates of points selected on a previously generated plot.

Some plots, use local scaling. For example, when plotting latitude, longitude data on a map a scale correction may be used to account for the curvature of the Earth. Similarly, if different data series are plotted on primary and secondary axes in a single plot, some or all data may be normalised. In such cases scaling may be local, i.e. what you actually plot may not be exactly what the annotation says it is.

Using `getXY` on such plots would recover the actual (x, y) coordinates of the points selected.

However, corrections can be applied using `scale.correction`, if it is supplied, to convert these to the same scale as the axes annotation. The correction should be a function that can be applied directly to a standard `getXY` output (a list of x and y values) and rescale x and y to give their 'corrected' values.

`getLatLon` provides an example of the mechanism, and is for use with georeferenced plots that have been locally scaled using `RgoogleMaps` functions like `LatLon2XY`. `getLatLon` uses `XY2LatLon` to rescale x and y values and then `as...` functions to convert the outputs of this step to a list format like that generated by `locator`, `grid.locator` or `getXY`.

**Author(s)**

Karl Ropkins

**References**

This function makes extensive use of code developed by others.

`lattice`: Sarkar, Deepayan (2008) *Lattice: Multivariate Data Visualization with R*. Springer, New York. ISBN 978-0-387-75968-5

`RgoogleMaps`: Markus Loecher and Sense Networks (2011). *RgoogleMaps: Overlays on Google map tiles in R*. R package version 1.1.9.6. <http://CRAN.R-project.org/package=RgoogleMaps>

**See Also**

In other packages: See `grid.locator`; `trellis.focus` and `lpoints` in `lattice`. See `LatLon2XY` in `RgoogleMap`.

---

**5.2.plot.add.functions***General plot add... functions*

---

**Description**

General functions for adding extra layers to existing plots.

**Usage**

```
add_loaPanel(lattice.plot=trellis.last.object(),
             preprocess = NULL, panel = NULL,
             postprocess = NULL, ...)
```

**Arguments**

<code>lattice.plot</code>	The plot to add an extra layer or layers to. This defaults to last lattice plot produced if not set in call.
<code>preprocess</code>	If set, a function to apply to <code>lattice.plot</code> before adding new panel.
<code>panel</code>	If set, a new <code>panel...</code> function to add to <code>lattice.plot</code> after the current <code>lattice.plot</code> panel.
<code>postprocess</code>	If set, a function to apply to <code>lattice.plot</code> after adding new panel.
<code>...</code>	Additional arguments, passed on to related functions.

**Details**

`add_loaPanel` is a frame work function for adding additional layers to an exiting lattice plot.

**Value**

`add_loaPanel` returns the supply (or last) lattice plot with an additional plot layer added.

**Note**

`add_loaPanel` is in-development aand may change significantly in future pacakge versions

**Author(s)**

Karl Ropkins

## References

This function makes extensive use of code developed by others.

lattice: Sarkar, Deepayan (2008) Lattice: Multivariate Data Visualization with R. Springer, New York. ISBN 978-0-387-75968-5

Grid (part of base R): R Core Team (2017). R: A language and environment for statistical computing. R Foundation for Statistical Computing, Vienna, Austria. URL <https://www.R-project.org/>.

5.3.plot.addXYFit.functions

*Plot addXYFit... functions*

## Description

Functions for adding XY fit model layers to existing plots.

## Usage

```
#main functions

addXYLMFit(lattice.plot = trellis.last.object(),
           preprocess = addXYFit_prep,
           panel=panel.loaXYFit, ...)

addXYLOESSFit(lattice.plot = trellis.last.object(),
              preprocess = addXYFit_prep, model.method = loaXYFit_loess,
              panel = panel.loaXYFit, ...)

#others

addXYFit_prep(lattice.plot = trellis.last.object(),
              model.method = loaXYFit_lm, ...)

loaXYFit_lm(x, y, output.len = 25, formula.signif = 2,
            r2.digits = 3, group.id = NULL, ...)

loaXYFit_loess(x, y, output.len = 25, r2.digits = 3,
              group.id = NULL, ...)

panel.loaXYFit(...)
```

## Arguments

`lattice.plot` The plot to add an extra layer or layers to. This defaults to last lattice plot produced if not set in call.



<code>preprocess</code>	If set, a function to apply to <code>lattice.plot</code> before adding new panel.
<code>panel</code>	If set, a new <code>panel...</code> function to add to <code>lattice.plot</code> after the current <code>lattice.plot</code> panel.
<code>...</code>	Additional arguments, passed on to daughter functions.
<code>model.method</code>	Function to use on XY data to generate fit reports.
<code>x, y</code>	XY data from plot, typically filtered by both <code>panel</code> and <code>group</code> .
<code>output.len</code>	Number of XY pairs to use when making fit line standard error bands.
<code>formula.signif, r2.digits</code>	When number of significant numbers and digits to use when reporting the fit formula and <code>r.squared</code> , respectively.
<code>group.id</code>	When XY data is grouped, the group identity.

### Details

`addXYLMFit` and `addXYLMFit` add linear and loess model fit surfaces to exiting lattice plot.

Both functions are intended to be used in form:

[make plot, e.g. `loaPlot(...)` then] `addXYLMFit(...)`

`loaXYFit_lm` and `loaXYFit_lm` generate fit and report information for linear and loess fit surfaces.

Other functions contain code structure common to all `addXYFit` functions.

### Value

`addXYLMFit` returns the supply (or last) lattice plot with an added linear fit surface layer, generated using `lm()`.

`addXYLMFit` returns the supply (or last) lattice plot with an added linear fit surface layer, generated using `loess()`.

The elements of the XYFit layers are `fit` the fit line, `se` the standard error bands, and `report` the fit report panels. These track properties, e.g. `col`, etc, of the plot they add to unless these terms are reset in the `addXYFit` call. They can also be removed by setting to `FALSE`, e.g. `fit=FALSE`, or updated individually using e.g. `fit.col=2` to change just the fit line color to red.

The report position within the plot panel can be set using `report.position` which uses a relative (`x=0-1, y=0-1`) scale to position the report.

### Note

These functions are in-development and may change significantly in future package versions.

### Author(s)

Karl Ropkins

## References

This function makes extensive use of code developed by others.

lattice: Sarkar, Deepayan (2008) Lattice: Multivariate Data Visualization with R. Springer, New York. ISBN 978-0-387-75968-5

lm() and loess() in stats (part of base R): R Core Team (2017). R: A language and environment for statistical computing. R Foundation for Statistical Computing, Vienna, Austria. URL <https://www.R-project.org/>.

## Examples

```
# add lm fit a loaPlot
addXYLMFit(loaPlot(cadmium~copper*zinc, data=lat.lon.meuse))

## Not run:

loaPlot(cadmium~copper*zinc, data=lat.lon.meuse)
addXYLOESSFit(col="red")

p <- loaPlot(cadmium~copper*zinc, data=lat.lon.meuse)
addXYLMFit(p, se.col="grey", report.position=c(0.95, 0.22))

## End(Not run)
```

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