

# Package ‘HilbertVisGUI’

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**Title** HilbertVisGUI

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**Depends** R (>= 2.6.0), HilbertVis (>= 1.1.6)

**SystemRequirements** gtkmm-2.4

**Suggests** lattice, IRanges

**Description** An interactive tool to visualize long vectors of integer data by means of Hilbert curves

**License** GPL (>= 3)

**URL** <http://www.ebi.ac.uk/~anders/hilbert>

**biocViews** Visualization

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dotsapply                      *List apply for '...' arguments*

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## Description

A kludge to get around a certain problem in using lapply with '...' ellipsis function arguments.

**Usage**

```
dotsapply( fun, ... )
```

**Arguments**

fun	A function that takes one argument
...	Arguments to which the function should be applied

**Details**

"dotsapply(fun,...)" gives the same result as `lapply( list(...), fun)`. However, the construction with "list", when used for variables enumerated explicitly, will result in their duplication.

Assume, for example, that you have three very large vectors a, b, and c, whose lengths you wish to know. If you write "`lapply( list(a,b,c),length )`", R will duplicate all three vectors in memory when constructing the list, which results in unnecessary use of memory. The alternative "`dotsapply( length, a, b, c )`" avoids this. Of course, you could also write "`list( length(a), length(b), length(c) )`", which neither causes duplication.

This last possibility is, however, not an option, if, instead of "a,b,c", you have ellipsed function arguments, i.e. literally "...". In this special case, dotsapply comes in handy, and as this case arose in [hilbertDisplay](#), I implemented dotsapply as a kludge and export it from the package just in case somebody has use for it.

**Value**

A list of function values

**Author(s)**

Simon Anders, EMBL-EBI, <sanders@fs.tum.de>

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hilbertDisplay

*Hilbert curve visualization*

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**Description**

Display one or several long integer vectors in an interactive fashion by means of the Hilbert curve.

See the vignette for a full explanation.

**Usage**

```

hilbertDisplay(
  ...,
  palettePos = colorRampPalette( c( "white", "red" ) )( 300 ),
  paletteNeg = colorRampPalette( c( "white", "blue" ) )( length(palettePos) ),
  maxPaletteValue = NULL,
  naColor = "gray",
  plotFunc = simpleLinPlot,
  names=NULL,
  sameScale=TRUE,
  pow2=FALSE,
  portrait=TRUE,
  fullLengths = NULL )

```

**Arguments**

...	The data vectors to be visualized. This may be integer or real vectors, either as ordinary R vectors or as R1e vectors (as defined in the IRanges packages). Care is taken within <code>hilbertDisplay</code> that these vectors do not get duplicated, so that you can pass very large vectors.
<code>palettePos</code>	The color palette used to visualize positive valued bins in the data. This must be either be a character vector of valid color names. If no palette is supplied, a ramp from white to red is used.
<code>paletteNeg</code>	The color palette for bins with negative values. Both palettes must have the same number of colours.
<code>maxPaletteValue</code>	The absolute bin value that should correspond to the end of the two palettes. (The beginning of the palettes always represents 0.) Within the graphical user interface, buttons are provided to adjust this interactively. If no value is provided, the median of the absolute values of maxima and minima of the data vectors is used.
<code>naColor</code>	The color to be used for bins which contain NAs or correspond to data outside the limits of the data vector. Pass a color name or a triple of RGB values. By default, "gray" is used.
<code>plotFunc</code>	An R function that is called if you use the "Linear plot" function offer by <code>HilbertCurveDisplay</code> 's GUI. If you enable this function and then click on a pixel in the display, the function supplied as <code>plotFunc</code> is called. If you do not supply this parameter, the function <code>simpleLinPlot</code> (part of this package) is used. If you supply your own function, it must accept two parameters: <code>data</code> and <code>info</code> . <code>data</code> will be the currently displayed data vector. Be careful that your function does not duplicate it (check with <code>tracemem</code> , if in doubt) in order to avoid performance problems when dealing with large data. The second argument, <code>info</code> , is a list, supplying the following fields, all of which, except for the last one, are single integers: <code>info\$binLo</code> , <code>info\$bin</code> , and <code>info\$binHi</code> are the lower, middle, and upper coordinate (i.e., vector index) of the bin represented by the pixel onto which the user has clicked. <code>info\$dispLo</code> and <code>info\$dispHi</code> are the lowest and highest index of the part of the vector currently displayed. <code>info\$seqIdx</code>

is the index of the currently displayed vector (i.e., its position in the `'...'` argument) and `info$seqName` is its name. All indices are one-based. Your function should plot a region of interest around `data[info$bin]`, or do some other useful operation. Any return value is ignored. For a very simple example, see the body of [simpleLinPlot](#).

<code>names</code>	The names of the sequences. A character vector. If not supplied, the expressions used in the <code>'...'</code> argument, as reported by <a href="#">substitute</a> are used.
<code>sameScale</code>	Setting this argument to TRUE pads all but the largest vector with NAs such that all vectors have the same length. (The padding is done "virtually", i.e. no duplication in memory occurs.) The purpose of this is to make sure that the bin size (i.e. the number of values depicted by one pixel) stays constant, when using the DisplayHilbertViewer GUI's "Prev" and "Next" buttons, which switch the display through the supplied data vectors.
<code>pow2</code>	Setting this argument to TRUE pads all vectors virtually with NAs such that their length becomes a power of 2. The purpose of this becomes apparent if you zoom in so much that several pixels correspond to the same data vector element. Then, without this option, the values take on strange fractal forms, while they are square in case of a power-of-2 length.
<code>portrait</code>	Setting this option to FALSE changes the GUI layout such that the controls appear to the right of the curve display ("landscape layout") as opposed to the usual case of them appearing below ("portrait layout"). This is useful for small screens as the GUI window may be two tall to fit on the screen in portrait mode.
<code>fullLengths</code>	This option allows you to manually control the padding of vectors with NAs if you do not like the result of the <code>sameScale</code> or <code>pow2</code> option. Supply an integer vector with as many values as there are vectors in the <code>'...'</code> argument, specifying the length including padding for each data vector. Passing numbers smaller than the length of the data vector results in only the beginning of the vector being displayed.

### Value

Returns an invisible NULL.

### Author(s)

Simon Anders, EMBL-EBI, <sanders@fs.tum.de>

### See Also

[simpleLinPlot](#)

### Examples

```
random <- c( as.integer( runif(100000)*30 ) )
ramp <- c( as.integer( 0:19999/100 ) )
try( hilbertDisplay( random, ramp ) )
```

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`hilbertDisplayThreeChannel`*Show up to three data vectors in the Hilbert curve display GUI.*

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**Description**

This is a variant of `hilbertDisplay` that takes up to three long data vector but does not display them in different panels (allowing to switch with the “Previous” and “Next” buttons but instead put them in one picture, using the three colour channels to overlay them. This is look for correlations in spatial distribution of the different data vectors.

**Usage**

```
hilbertDisplayThreeChannel(  
  dataRed,  
  dataGreen = 0,  
  dataBlue = 0,  
  naColor = col2rgb("gray"),  
  fullLength = max(length(dataRed), length(dataGreen), length(dataBlue)),  
  portrait = FALSE)
```

**Arguments**

<code>dataRed</code>	The data to be displayed in the red colour channel, to be provided as a numeric vector with values between 0 (black) and 1 (bright red). You may either pass an ordinary R vector or an R1e vector as defined in the IRanges package.
<code>dataGreen</code>	As <code>dataRed</code> but for the green channel.
<code>dataBlue</code>	As <code>dataRed</code> but for the blue channel.
<code>naColor</code>	The color to use to indicate NA.
<code>fullLength</code>	The vector length that should correspond to the full display. By default, the length of the longest data vector. It may make sense to round up to the next power of two.
<code>portrait</code>	Whether to arrange the buttons to the right of the display. Useful on small screens.

**Value**

Invisible NULL.

**Note**

This function is a bit less mature than the function `hilbertDisplay`. Especially, the GUI functions “Linear plot”, “Previous”, and “Next” are displayed but defunct.

**Author(s)**

Simon Anders

**See Also**[hilbertDisplay](#)

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`simpleLinPlot`*Default plot callback function for hilbertDisplay*

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**Description**

Plots a small part of the supplied vector.

**Usage**

```
simpleLinPlot( data, info )
```

**Arguments**

<code>data</code>	The data vector, a vector of integers.
<code>info</code>	A list with fields describing the context, as described in the help page for <a href="#">hilbertDisplay</a> .

**Details**

This is a very simple function that is called by default by [hilbertDisplay](#) when the user chooses the "Linear plot" option and clicks on a pixel in the display. It displays a piece of 2000 values of the data vector, centered around `info$bin`. You will often want to replace this function by one tailored to your needs.

**Value**

Null.

**Author(s)**

Simon Anders, EMBL-EBI, <sanders@fs.tum.de>

**See Also**[hilbertDisplay](#)

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