

Package ‘Rtoolbox’

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Title Toolbox with R scripts

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Depends reshape2

Description Toolbox with R scripts for mostly bioinformatic purposes

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OverviewPlot	<i>Create heatmaps of segmentation values from a DNACopy object.</i>
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Description

Creates color-based plots of the segmentation values from a provided DNACopy object.

Usage

```
OverviewPlot(DNACopy.object, samples, range.CNA = c(-2, 2), color.palette = colorRampPalette(c("b
```

Arguments

<code>DNAcopy.object</code>	DNAcopy object with the segmentation values that need to be plotted.
<code>samples</code>	character vector with the names of the samples that need to be plotted. Names need to be as given in the DNAcopy object. This is an optional argument; all samples will be plotted by default.
<code>range.CNA</code>	the range of the segmentation values that will be plotted. Segmentation values outside of <code>range.CNA</code> will be capped to the minimum and maximum <code>range.CNA</code> values. This is an optional argument; the default range is <code>c(-2, 2)</code> .
<code>color.palette</code>	the color palette that will be used for heatmaps. This is an optional argument; the default color palette is <code>colorRampPalette(c("blue", "white", "white", "red"))(49)</code> .

Details

`OverviewPlot` will print heatmaps of the segmentation values in a DNAcopy object. The `samples` argument can be used to limit the amount of plotted samples. The `range.CNA` vector determines where the segmentation values are capped and the range of segmentation values that is plotted. Finally, the `color.palette` argument can be used to create heatmaps with custom colors. All arguments except `DNA.copy.object` are optional.

Author(s)

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Examples

```
## Generate heatmaps with \code{\link{OverviewPlot}}.

## Not run: OverviewPlot(segment.CNA.object) ## Plot using default settings.
## Not run: OverviewPlot(segment.CNA.object, samples = unique(segment.CNA.object$output$ID)[1:3]) ## Plot on
## Not run: OverviewPlot(segment.CNA.object, range.CNA = c(-1, 1)) ## Plot with values from -1 to 1 (outside o
## Not run: OverviewPlot(segment.CNA.object, color.palette = colorRampPalette(c("blue", "white", "red"))(49))
```

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