Package ‘affylmGUI’

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affyGUI-package

The affyGUI Package: Graphical User Interface for limma analysis of Affymetrix microarrays

Description

The affyGUI package provides a Graphical User Interface (GUI) for analysis of Affymetrix microarray data using the limma package (Linear Modes for MicroArray data).

Details

The GUI is launched by typing `affyGUI()` at the R prompt. For detailed help, including example sessions, see the affyGUI home page at http://bioinf.wehi.edu.au/affyGUI.

The limma package offers R users a command-line interface to state-of-the-art techniques for linear modelling of microarray data and for identifying differentially expressed genes (Ritchie et al, 2015). The affyGUI package, while not as powerful as limma to the expert user, offers a simple point-and-click interface to many of the commonly-used limma and affy functions.

To use the affyGUI package, you need to have R 1.8.1 or later, Tcl/Tk 8.3 or later (ActiveTcl for Windows/Linux or Fink Tcl/Tk for MacOSX (X11)) and the limma, affyGUI and tkrplot R packages. At least one of the affy, gcma or affyPLM Bioconductor packages are also required for normalization and probe-set summarization. The R2HTML (CRAN) and xtable (CRAN) packages are recommended. affyGUI has been tested succesfully on Windows, Red Hat Linux, and on Mac OSX with X11.


Author(s)


References


affylmGUI

See Also

The GUI is launched by `affylmGUI()`.
More documentation is available from http://bioinf.wehi.edu.au/affylmGUI.
`showChangeLog()` shows the most recent lines from the affylmGUI package changeLog.
See `limma-package` for help about the limma package.

Examples

```r
library(limma)
changeLog(package="affylmGUI")
```

---

**affylmGUI**

Graphical User Interface for the limma microarray package

**Description**

Graphical User Interface for limma package analysis of Affymetrix microarrays.

**Usage**

```r
affylmGUI(BigfontsForaffylmGUIpresentation=FALSE)
```

**Arguments**

`BigfontsForaffylmGUIpresentation`

logical, if `TRUE` then larger fonts are used. However, some font sizes are not
controlled by affylmGUI and so must be adjusted in the operating system, e.g.,
in the Control Panel in Windows under Display, Appearance.

**Details**

The `affylmGUI` function launches a Graphical User Interface for the affy, gcrma, affyPLM and
limma packages. The GUI uses Tk widgets (via the R TclTk interface by Peter Dalgaard) in order
to provide a simple interface to limma functions for linear modelling of Affymetrix microarrays and
identification of differentially expressed genes.

Almost all users will type `affylmGUI()`. The use of `affylmGUI(Bigfonts=TRUE)` is only intended
for the preparation of a talk about the affylmGUI package.

**Author(s)**

Created by James Wettenhall between 2003 and September 2006. Maintained by Keith Satterley

**See Also**

`affylmGUI-package`, `internal`
Internal Functions

Description

Internal functions defined by the affylmGUI package.

Usage

AboutaffylmGUI()
affyHelp()
affylmGUIhelp()
affyPlotMA()
affyPlotMAcontrast()
ChooseCDF()
ChooseContrastParameterization()
chooseDir()
ChooseEbayesStatistic()
ComputeContrasts()
ComputeLinearModelFit()
CopyGraph(img)
DeleteContrastParameterization()
DensityPlot()
DensityPlotAll()
evalRcode()
ExportHTMLreport()
ExportNormalizedExpressionValues()
ExportTopTable()
fixSeps(string)
generalPlotFunction(code="",WindowTitle="")
GetComponentsToExportInHTMLreport(contrastParameterizationIndex=NULL)
GetContrast(contrastParameterizationIndex)
GetContrastNamesForHeatDiagram(numContrasts=2,ContrastNames=c("Contrast 1","Contrast 2"))
GetContrastParameterizationName()
GetContrasts(NumContrasts=0)
GetDEcutoff()
GetGeneLabelsOptions()
GetJpegOrPngParams(graphFileType)
GetJpegOrPngX11Params(graphFileType)
GetlimmaDataSetName()
GetLogPLMDataChoice()
GetMultipleContrasts(contrastParameterizationIndex)
GetNormalizationMethod()
getPackageVersion(pkgName)
GetParameterizationName()
GetPlotLabels(plottitle="",xlabel="",ylabel="")
GetPlotSize()
internal

GetPlotTitle(plottitle="")
GetPValueCutoff(p.value=0.01)
GetResidualTypeChoice()
GetRNATypesFrom.ContrastsFromDropDowns.String(string)
GetSetNames(numSets=2,set1="",set2="",set3="")
GetSlideNum(all=FALSE)
GetSlideNums()
GetWhichProbes(includeBoth=FALSE)
GetWtAreaParams()
GetWtFlagParams()
HeatDiagramDialog(parameterName)
HeatDiagramPlot()
HTMLPlotUsingFunction(Caption = "", File = "plot.html", GraphRelativeDirectory = ".", GraphAbsoluteDirectory = NULL, ... = "png", GraphBorder = 1, Align = "center", plotFunction = NULL,Width=600,Height=600,PointSize=12,BG="white",res=72,...)
ImageArrayPlot()
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IntensityHistogramAll()
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NormalizeNow()
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NUSEPlotAll()
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onExit()
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OpenCDFandTargetsfiles()
OpenCDFFile()
OpenLimmaFile()
OpenTargetsFile()
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QQTplot()
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Require(pkg)
Resize(img,plotFunction)
RLEPlotAll()
RNADegradationPlotAll()
SaveAsLimmaFile()
SaveGraphAsJpeg(initialfile, plotFunction)
SaveGraphAsPDF(initialfile, plotFunction)
SaveGraphAsPNG(initialfile, plotFunction)
SaveGraphAsPostscript(initialfile, plotFunction)
SaveLimmaFile()
SetupPlotKeyBindings(tt,img)
SetupPlotMenus(tt,initialfile,plotFunction,img)
SetWD()
showChangeLog()
showCitations()
showTopTable(..., export=FALSE)
SimplifyContrastsExpression(string)
strstr(haystack, needle)
tclArrayVar()
TclRequire(tclPkg)
Try(expr)
TryReadImgProcFile(expr)
UpDownOrBoth()
vennDiagramaafyImGUI(object, include = "both", names, cex = 1.5, mar = rep(1, 4), ...)
VennDiagramPlot()
ViewContrastsMatrixAsPairs(contrastsMatrix, contrastsMatrixList, contrastParameterizationIndex = NULL)
ViewContrastsMatrixInTable(contrastsMatrixList, contrastParameterizationIndex = NULL)
ViewExistingContrastParameterization()
ViewRNATargets()

Arguments

... HTMLplotUsingFunction:arg14:and:showTopTable:arg1:and:vennDiagramaafyImGUI:arg6
Align HTMLplotUsingFunction:arg8
all GetSlideNum:arg1
BG HTMLplotUsingFunction:arg12
Caption HTMLplotUsingFunction:arg1
code generalPlotFunction: arg1: a code
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contrastParameterizationIndex

GetComponentsToExportInHTMLreport:arg1:and:contrastParameterizationIndex:arg1:and:GetContrast:

export showTopTable:arg2
expr Try:arg1:and:TryReadImgProcFile:arg1
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GraphBorder HTMLplotUsingFunction:arg7
GraphFileName HTMLplotUsingFunction:arg5
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GraphRelativeDirectory HTMLplotUsingFunction:arg3

GraphSaveAs HTMLplotUsingFunction:arg6
haystack nstrstr:arg1:and:strstr:arg1
Height HTMLplotUsingFunction:arg11
Details

These functions are called by `affyLinGUI()` via the GUI and are not intended to be called directly by users.
Author(s)


See Also

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