"C:\Program Files\R\R-3.0.1\bin\x64\R.exe" -f "C:\Users\visita-mas\AppData\Local\Apps\2.0\3BP1WRJV.VTB\3CDR9RGT.D1P\skyl..tion\_e4141a2a22107248\_0013.0000\_8870d1d2982721de\Tools\QuaSAR-1\_33\QuaSAR-Skyline.R" --slave --no-save --args "C:\Users\visita-mas\AppData\Local\Apps\2.0\3BP1WRJV.VTB\3CDR9RGT.D1P\skyl..tion\_e4141a2a22107248\_0013.0000\_8870d1d2982721de\Tools\QuaSAR-1\_33\QuaSAR.R" "C:\Users\visita-mas\AppData\Local\Apps\2.0\3BP1WRJV.VTB\3CDR9RGT.D1P\skyl..tion\_e4141a2a22107248\_0013.0000\_8870d1d2982721de\Tools\QuaSAR-1\_33\common.R" "C:\Users\visita-mas\AppData\Local\Temp\QuaSAR\_QuaSAR\_Input.csv" NULL teste "light Area" 1 "heavy Area" fmol/ul 1 1 -1 1 0 0 0 150 150 1 0.2 0 NULL teste 0

[1] $Id: QuaSAR-Skyline.R 149 2014-10-23 13:56:13Z manidr $

[1] $Id: QuaSAR.R 149 2014-10-23 13:56:13Z manidr $

Attaching package: 'boot'

The following object is masked from 'package:lattice':

melanoma

The following object is masked from 'package:gtools':

inv.logit, logit

KernSmooth 2.23 loaded

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Attaching package: 'gplots'

The following object is masked from 'package:stats':

lowess

[1] Processing arguments ...

[1] >>> Processing part 1 of 1

[1] Initializing ...

[1] Running AuDIT ...

Error in `colnames<-`(`\*tmp\*`, value = c("peptide", "sample", "replicate", :

attempt to set 'colnames' on an object with less than two dimensions

Calls: tryCatch ... tapply -> lapply -> FUN -> FUN -> run.AuDIT -> colnames<-

Execution halted

Finished!