Package: affyContam (via r-universe)

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Title structured corruption of affymetrix cel file data

Version 1.71.0

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Description structured corruption of cel file data to demonstrate QA effectiveness

Depends R (>= 2.7.0), tools, methods, utils, Biobase, affy, affydata

Suggests hgu95av2cdf

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LazyLoad yes

biocViews Infrastructure

Config/pak/sysreqs zlib1g-dev

Repository https://bioc.r-universe.dev

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setRectRegion *set a rectangular or circular region in an affybatch to a specified set of values*

Description

set a rectangular or circular region in an affybatch to a specified set of values

Usage

```
setRectRegion(x, chip=1, xinds=251:350, yinds=251:350, vals=10, valgen=NULL)
setCircRegion(x, chip=1, center=c(350,350), rad=100, vals=10, valgen=NULL)
getRectRegion(x, chip=1, xinds=251:350, yinds=251:350)
getCircRegion(x, chip=1, center=c(350,350), rad=100)
```

Arguments

x	AffyBatch instance
chip	sample index
xinds	x coordinates to be contaminated
yinds	y coordinates to be contaminated
vals	values to be assigned to rectangle elements
center	geometric center of circle to be altered
rad	radius of circle to be altered, in xy units of the chip addressing system used by xy2indices in the cdf package
valgen	function of parameter n that generates n values to be inserted in the altered region

Value

set* functions return AffyBatch instance with intensities modified as requested

get* functions return numeric vectors of intensities as requested.

Author(s)

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Examples

```
library(affydata)
data(Dilution)
opar = par(no.readonly=TRUE)
par(mfrow=c(2,2))
hist(Dilution, main="original")
image(Dilution[,1], main="original")
#
# we will contaminate in two ways: thin line at fixed low intensity, and
```

```
# circular blob at moderate random intensity
#
ab = setRectRegion(Dilution, 1, xinds=25:30, yinds=1:620,
  vals=10)
ab = setCircRegion(ab, 1, valgen=function(n){
   rnorm(n, 350,50)})
hist(ab, main="chip 1 contaminated by normal")
image(ab[,1], main="chip 1 contaminated")
ex = getCircRegion(Dilution, 1)
length(ex)
ab = setCircRegion(Dilution, 1, vals=pmin(2*ex,65535))
image(ab[,1], main="chip 1 contaminated by doubling")
par(opar)
```

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