

Package ‘mvoutData’

October 12, 2020

Title affy and illumina raw data for assessing outlier detector performance

Description affy and illumina raw data for assessing outlier detector performance

Version 1.24.0

Author VJ Carey <stvjc@channing.harvard.edu>

Depends R (>= 2.10.0), methods, Biobase (>= 2.5.5), affy (>= 1.23.4), lumi

Maintainer VJ Carey <stvjc@channing.harvard.edu>

License Artistic-2.0

biocViews ExperimentData, MicroarrayData

git_url <https://git.bioconductor.org/packages/mvoutData>

git_branch RELEASE_3_11

git_last_commit c432f5b

git_last_commit_date 2020-04-27

Date/Publication 2020-10-12

R topics documented:

s12c 1

Index 3

s12c *raw expression data for evaluation of outlier detection algorithms*

Description

raw expression data for evaluation of outlier detection algorithms

Usage

```
data(s12c); data(ILM1)
```

Format

s12c is an AffyBatch created from SpikeIn U133a, with first two samples contaminated. ILM1 is the result of lumi::lumiR applied to the CSV file supplied by Illumina Inc. for the MAQC study (19 arrays).

Examples

```
data(s12c)
s12c
```

Index

*** datasets**

s12c, 1

ILM1 (s12c), 1

s12c, 1