

Package ‘bronchialIL13’

December 10, 2024

Version 1.45.0

Title time course experiment involving il13

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Depends R(>= 2.10.0), affy (>= 1.23.4)

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Description derived from CNMC (pepr.cnmcresearch.org)
http://pepr.cnmcresearch.org/browse.do?action=list_prj_exp&projectId=95
Human Bronchial Cell line A549

License GPL-2

biocViews ExperimentData, MicroarrayData

URL <http://www.biostat.harvard.edu/~carey>

git_url <https://git.bioconductor.org/packages/bronchialIL13>

git_branch devel

git_last_commit 4e828b9

git_last_commit_date 2024-10-29

Repository Bioconductor 3.21

Date/Publication 2024-12-10

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HAHrma

data on Affymetrix U133a chips regarding time course response of human bronchial cell line A549 to IL13

Description

data on Affymetrix U133a chips regarding time course response of human bronchial cell line A549 to IL13

Usage

```
data(HAHrma)
data(HAH)
```

Format

The format is a Biobase exprSet structure. phenoData variables are id, trt and time (hours). HAH is derived from a ReadAffy of 15 CEL files, and HAHrma is derived from rma(HAH), with manual construction of the phenoData based on the filenames. The CEL files are in inst/cel/dataoq.zip.

Source

http://pepr.cnmcresearch.org/browse.do?action=list_prj_exp&projectId=95

Examples

```
data(HAHrma)
table(HAHrma$time, HAHrma$trt)
```

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* **data**

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