Package 'bronchialIL13'

February 26, 2026

Version 1.48.0

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 RELEASE_3_22

git_last_commit 818c46b

git_last_commit_date 2025-10-29

Repository Bioconductor 3.22

Date/Publication 2026-02-26

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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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