

bronchialIL13

April 2, 2010

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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*Topic **data**

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