

bronchialIL13

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HARma

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(HARma)
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 HARma 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(HARma)
table(HARma$time, HARma$trt)
```

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