

Package ‘serumStimulation’

October 18, 2022

Version 1.32.0

Date 2011-08-09

Title serumStimulation is a data package which is used by examples in package pcaGoPromoter

Author Morten Hansen <mhansen@sund.ku.dk>

Maintainer Morten Hansen, <mhansen@sund.ku.dk>

Description Contains 13 micro array data results from a serum stimulation experiment

biocViews ExperimentData, MicroarrayData

LazyLoad yes

License GPL (>= 2)

Depends R (>= 2.10)

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

git_branch RELEASE_3_15

git_last_commit 726e86a

git_last_commit_date 2022-04-26

Date/Publication 2022-10-18

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serumStimulation	<i>Data set from serum stimulation DNA micro array</i>
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Description

The serumStimulation data set is from an DNA micro array analysis of 13 samples from a serum stimulation experiment. There is 5 controls, 5 serum stimulated with inhibitor and 3 serum stimulation without inhibitor.

The original .CEL files have been read with ReadAffy and normalized with rma.

The data is the output of `exprs(rma(ReadAffy()))`

Usage

```
serumStimulation
```

Format

See `exprs` for description of output.

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