

Package ‘serumStimulation’

June 20, 2024

Version 1.40.0

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_19

git_last_commit 118dd67

git_last_commit_date 2024-04-30

Repository Bioconductor 3.19

Date/Publication 2024-06-20

Contents

serumStimulation	2
Index	3

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

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.

Index

* **datasets**

serumStimulation, [2](#)

serumStimulation, [2](#)