

Package ‘serumStimulation’

September 11, 2025

Version 1.45.0

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 devel

git_last_commit 54425cc

git_last_commit_date 2025-04-15

Repository Bioconductor 3.22

Date/Publication 2025-09-11

Contents

serumStimulation	1
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, [1](#)

serumStimulation, [1](#)