

Package ‘serumStimulation’

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

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

biocViews ExperimentData, MicroarrayData

LazyLoad yes

License GPL (>= 2)

Depends R (>= 2.10)

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R topics documented:

serumStimulation 1

Index 3

serumStimulation *Data set from serum stimulation DNA micro array*

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

*Topic **datasets**

 serumStimulation, [1](#)

 serumStimulation, [1](#)