

# Package ‘serumStimulation’

April 4, 2014

**Version** 0.99.5

**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

**LazyLoad** yes

**License** GPL (>= 2)

**Depends** R (>= 2.10)

## R topics documented:

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

\*Topic **datasets**

serumStimulation, 1

serumStimulation, 1