

Package ‘SNAGEEdata’

December 18, 2025

Version 1.46.0
Date 2012-01-26
Title SNAGEE data
Author David Venet <davenet@ulb.ac.be>
Maintainer David Venet <davenet@ulb.ac.be>
Depends R (>= 2.6.0)
Suggests ALL, hgu95av2.db, SNAGEE
Description SNAGEE data - gene list and correlation matrix
License Artistic-2.0
biocViews MicroarrayData
URL <http://fleming.ulb.ac.be/SNAGEE>
git_url <https://git.bioconductor.org/packages/SNAGEEdata>
git_branch RELEASE_3_22
git_last_commit 36029d4
git_last_commit_date 2025-10-29
Repository Bioconductor 3.22
Date/Publication 2025-12-18

Contents

SNAGEEdata-package	1
getCC	2
Index	3

SNAGEEdata-package	<i>SNAGEE - data</i>
--------------------	----------------------

Description

Signal-to-Noise applied to Gene Expression Experiments - database of gene correlations.

Details

Package: SNAGEEdata
 Version: 0.99.0
 Date: 2012-01-26
 Depends: R (>= 2.6.0)
 Suggests: SNAGEE
 License: Artistic-2.0
 URL: <http://fleming.ulb.ac.be/SNAGEE>

Index:

getCC Gene-gene correlations and list of genes

Author(s)

David Venet <davenet@ulb.ac.be>

Maintainer: David Venet <davenet@ulb.ac.be>

Examples

```
# the gene-gene correlations
cc = getCC();
```

getCC	<i>Gene-gene correlations</i>
-------	-------------------------------

Description

Get the gene-gene correlations and the list of genes.

Usage

```
getCC(mode="complete")
```

Arguments

mode Which correlations should be recovered. complete: calculated with all platforms; woAffy: calculated without the Affymetrix platforms.

Value

A list with two elements: g is the list of gene IDs, cc is the upper triangular part of the correlation matrix.

Examples

```
# Get the list of genes
geneList = getCC()$g;
```

Index

getCC, [2](#)

SNAGEEdata (SNAGEEdata-package), [1](#)

SNAGEEdata-package, [1](#)