

Package ‘SNAGEEdata’

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Version 1.36.0

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Title SNAGEE data

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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>

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

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getCC	Gene-gene correlations and list of genes
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Author(s)

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Examples

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

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

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