Package ‘breastCancerTRANSBIG’

February 1, 2017

Type Package

Title Gene expression dataset published by Desmedt et al. [2007] (TRANSBIG).

Version 1.12.0

Date 2011-02-10

Description Gene expression data from a breast cancer study published by Desmedt et al. in 2007, provided as an eSet.

biocViews ExperimentData, CancerData, BreastCancerData, MicroarrayData, GEO

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Depends R (>= 2.5.0)

Suggests survcomp, genefu, Biobase

LazyLoad yes

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URL http://compbio.dfci.harvard.edu/

NeedsCompilation no

R topics documented:

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transbig Gene expression, annotations and clinical data from TRANSBIG 2006

Description

This dataset contains the gene expression, annotations and clinical data as published from TRANSBIG in 2006.
Usage

data(transbig)

Format

ExpressionSet with 22283 features and 198 samples, containing:

- exprs(transbig): Matrix containing gene expressions as measured by Affymetrix hgu133a technology (single-channel, oligonucleotides).
- fData(transbig): AnnotatedDataFrame containing annotations of Affy microarray platform hgu133a.
- pData(transbig): AnnotatedDataFrame containing Clinical information of the breast cancer patients whose tumors were hybridized.
- experimentalData(transbig): MIAME object containing information about the dataset.
- annotation(transbig): Name of the affy chip.

Details

This dataset represent the study published by Desmedt et al. in 2006.

- Abstract: Purpose: Recently, a 76-gene prognostic signature able to predict distant metastases in lymph node-negative (N-) breast cancer patients was reported. The aims of this study conducted by TRANSBIG were to independently validate these results and to compare the outcome with clinical risk assessment. Experimental Design: Gene expression profiling of frozen samples from 198 N- systemically untreated patients was done at the Bordet Institute, blinded to clinical data and independent of Veridex. Genomic risk was defined by Veridex, blinded to clinical data. Survival analyses, done by an independent statistician, were done with the genomic risk and adjusted for the clinical risk, defined by Adjuvant! Online. Results: The actual 5- and 10-year time to distant metastasis were 98% (88-100%) and 94% (83-98%), respectively, for the good profile group and 76% (68-82%) and 73% (65-79%), respectively, for the poor profile group. The actual 5- and 10-year overall survival were 98% (88-100%) and 87% (73-94%), respectively, for the good profile group and 84

Source


References


Examples

```r
## load Biobase package
library(Biobase)
## load the dataset
data(transbig)
```
## show the first 5 rows and columns of the expression data

```r
exprs(transbig)[1:5,1:5]
```

## show the first 6 rows of the phenotype data

```r
head(pData(transbig))
```

## show first 20 feature names

```r
featureNames(transbig)[1:20]
```

## show the experiment data summary

```r
experimentData(transbig)
```

## show the used platform

```r
annotation(transbig)
```

## show the abstract for this dataset

```r
abstract(transbig)
```
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