

Package ‘lungExpression’

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Title ExpressionSets for Parmigiani et al., 2004 Clinical Cancer
Research paper

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Depends R (>= 2.4.0), Biobase (>= 2.5.5)

Description Data from three large lung cancer studies provided as ExpressionSets

LazyLoad yes

biocViews ExperimentData, CancerData, LungCancerData

License GPL (>= 2)

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harvard

A Harvard study on lung cancer gene expression

Description

A Harvard study on lung cancer gene expression. Data is represented as an ExpressionSet.

Usage

data(harvard)

Details

Annotation for the phenoData will be updated.

References

Bhattacharjee et al., Classification of human lung carcinomas by mRNA expression profiling reveals distinct adenocarcinoma subclasses, PNAS 2001, 98:13790-5.

Parmigiani et al., A cross-study comparison of gene expression studies for the molecular classification of lung cancer, Clinical Cancer Research, 10:2922-2927, 2004.

Examples

data(harvard)

michigan

A Michigan study on lung cancer gene expression

Description

A Michigan study on lung cancer gene expression. Data is represented as an ExpressionSet.

Usage

data(michigan)

Details

Annotation for the phenoData will be updated.

References

Beer et al., Gene expression profiles predict survival of patients with lung adenocarcinoma. Nature Medicine 8(8):816-824 (2002).

Parmigiani et al., A cross-study comparison of gene expression studies for the molecular classification of lung cancer, Clinical Cancer Research, 10:2922-2927, 2004.

Examples

```
data(michigan)
```

```
stanford
```

Public lung cancer data from the Stanford study

Description

Public lung cancer data from the Stanford study represented as an ExpressionSet

Usage

```
data(stanford)
```

Details

Annotation for the phenoData will be updated.

References

Garber et al., Diversity of Gene Expression in Adenocarcinoma of the Lung, PNAS, 2001, 98(24):13784-9.

Parmigiani et al., A cross-study comparison of gene expression studies for the molecular classification of lung cancer, Clinical Cancer Research, 10:2922-2927, 2004.

Examples

```
data(stanford)
```

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