

Package ‘charmData’

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Type Package

Title An example dataset for use with the charm package

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Description An example dataset for use with the charm package

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Depends R(>= 2.11.0), charm, pd.charm.hg18.example

biocViews ExperimentData

ZipData no

LazyLoad yes

R topics documented:

charmData-package 1

Index 3

charmData-package	<i>An example dataset for the charm package</i>
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Description

This is a small example DNA methylation microarray dataset for use with the charm package. It contains Nimblegen xys (raw data) files for 8 samples and the corresponding sample description file.

Author(s)

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Examples

```
dataDir <- system.file("data", package="charmData")  
setwd(dataDir)  
dir()
```

Index

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