

# Package ‘charmData’

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**Type** Package

**Title** An example dataset for use with the charm package

**Version** 1.1.1

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**Description** An example dataset for use with the charm package

**License** LGPL (>= 2)

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

**biocViews** ExperimentData

**ZipData** no

**LazyLoad** yes

## R topics documented:

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charmData-package      *An example dataset for the charm package*

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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()
```

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