

charmData

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

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.

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Examples

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

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