

# Package ‘MethylAidData’

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**Type** Package

**Title** MethylAid-summarized data on 2800 Illumina 450k array samples

**Version** 1.0.0

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**Author** Maarten van Iterson, Elmar. Tobi, Roderick Slieker, Wouter den Hollander, Rene Luijk and Bas Heijmans

**Maintainer** M. van Iterson <m.van\_iterson@lumc.nl>

**Description** A data package containing summarized Illumina 450k array data on 2800 samples. The data can be use as a background data set in the interactive application.

**URL** <http://shiny.bioexp.nl/MethylAid>

**License** GPL (>= 2)

**VignetteBuilder** knitr

**biocViews** ExperimentData

**Depends** MethylAid, R (>= 3.2)

**Suggests** BiocParallel, BiocStyle, knitr

**NeedsCompilation** no

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exampleDataLarge      *summarizedData object on 2800 450k Human Methylation samples*

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**Description**

Summarized 450K data on 2800 samples. The data can be used as a background data set in the interactive application.

**Usage**

```
exampleDataLarge
```

**Format**

```
summarizedData-object
```

**Value**

Pre-summarizedData object on 2800 450k Human Methylation samples.

**Examples**

```
data(exampleDataLarge)
data(exampleData)
## Not run: visualize(exampleData, background=exampleDataLarge)
```

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