

# minfiData

March 23, 2012

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MsetEx

*An example dataset for Illumina's Human Methylation 450k dataset, after preprocessing.*

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## Description

This contains the raw data for 6 samples from Illumina's Human Methylation 450k dataset. The data has been preprocessed.

## Usage

```
data (MsetEx)
```

## Format

An object of class "MethylSet"

## Details

Currently, the pheno data for these 6 samples are masked. Scripts for creating the object is found in the `scripts` directory of the package and `extdata` contains the IDAT files. The data has been preprocessed using `preprocessRaw`.

## See Also

[MethylSet-class](#) for the class definition, [preprocessRaw](#) for the preprocessing function, [RGsetEx](#) for the companion raw data.

## Examples

```
data (MsetEx)  
pData (MsetEx)
```

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`RGsetEx`*An example dataset for Illumina's Human Methylation 450k dataset.*

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**Description**

This contains the raw data for 6 samples from Illumina's Human Methylation 450k dataset.

**Usage**

```
data (RGsetEx)
```

**Format**

An object of class "RGChannelSet "

**Details**

Currently, the pheno data for these 6 samples are masked. Scripts for creating the object is found in the `scripts` directory of the package and `extdata` contains the IDAT files.

**See Also**

[RGChannelSet-class](#) for the class definition, [MsetEx](#) for the comparison preprocessed data.

**Examples**

```
data (RGsetEx)
pData (RGsetEx)
```

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