

Package ‘minfiDataEPIC’

June 4, 2026

Version 1.39.0

Title Example data for the Illumina Methylation EPIC array

Description

Data from 3 technical replicates of the cell line GM12878 from the EPIC methylation array.

Author Jean-Philippe Fortin, Kasper Daniel Hansen

Maintainer Kasper Daniel Hansen <kasperdanielhansen@gmail.com>

License Artistic-2.0

Depends R (>= 3.3), minfi (>= 1.21.2),
IlluminaHumanMethylationEPICmanifest,
IlluminaHumanMethylationEPICanno.ilm10b2.hg19

LazyData yes

biocViews Homo_sapiens_Data, MethylationArrayData, MicroarrayData

NeedsCompilation no

git_url <https://git.bioconductor.org/packages/minfiDataEPIC>

git_branch devel

git_last_commit 35f2e75

git_last_commit_date 2026-04-28

Repository Bioconductor 3.24

Date/Publication 2026-06-04

Contents

MsetEPIC	2
RGsetEPIC	2
Index	4

MsetEPIC	<i>An example dataset for Illumina's Human Methylation EPIC dataset, after preprocessing.</i>
----------	---

Description

This contains the raw data for 3 technical replicates of the cell line GM12878 from the Illumina's Human Methylation EPIC platform. The data has been preprocessed with `preprocessRaw`.

Usage

```
data(MsetEPIC)
```

Format

An object of class "MethylSet"

Details

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

Examples

```
data(MsetEPIC)
pData(MsetEPIC)
```

RGsetEPIC	<i>An example dataset for the Illumina's Human Methylation EPIC platform.</i>
-----------	---

Description

This contains the raw data for 3 technical replicates of the cell line GM12878 from the Illumina's Human Methylation EPIC platform.

Usage

```
data(RGsetEPIC)
```

Format

An object of class "RGChannelSet"

Details

Scripts for creating the object is found in the `scripts` directory of the package and `extdata` contains the IDAT files.

See Also

[RGChannelSet](#) for the class definition, [MsetEPIC](#) for the companion preprocessed data.

Examples

```
data(RGsetEPIC)  
pData(RGsetEPIC)
```

Index

* datasets

MsetEPIC, [2](#)

RGsetEPIC, [2](#)

MethylSet, [2](#)

MsetEPIC, [2](#), [3](#)

preprocessRaw, [2](#)

RGChannelSet, [3](#)

RGsetEPIC, [2](#), [2](#)