

ILLUMINAHUMANMETHYLATIONEPICv2manifest

November 27, 2024

ILLUMINAHUMANMETHYLATIONEPICv2manifest

Manifest object for the ILLUMINAHUMANMETHYLATIONEPIC v2.0 array

Description

This object contains the array design for Illumina's Human Methylation EPIC v2.0 microarray.

Usage

```
data(ILLUMINAHUMANMETHYLATIONEPICv2manifest)
```

Format

An object of class `ILLUMINAMETHYLATIONMANIFEST`

Details

This package is based on the file <https://support.illumina.com/content/dam/illumina-support/documents/downloads/productfiles/methylationepic/MethylationEPIC%20v2%20Files.zip> from https://support.illumina.com/array/array_kits/infinium-methylationepic-beadchip-kit/downloads.html.

A script for creating the data object in this package is contained in `system.files("scripts", "manifest.R", package = "ILLUMINAHUMANMETHYLATIONEPICv2manifest")`, which is slightly adjusted from the original script in the "ILLUMINAHUMANMETHYLATIONEPICmanifest" package.

References

None.

See Also

The description of the class in the `minfi` package, [ILLUMINAMETHYLATIONMANIFEST-class](#).

Examples

```
ILLUMINAHUMANMETHYLATIONEPICv2manifest
```

Index

* datasets

 IlluminaHumanMethylationEPICv2manifest,

[1](#)

IlluminaHumanMethylationEPICv2manifest,

[1](#)