# IlluminaHumanMethylationEPICv2manifest

September 30, 2024

# 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

 $\label{eq:linear} Illumina {\tt Human} {\tt Methylation} {\tt EPICv2manifest}$ 

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# \* datasets

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