

Package ‘IlluminaHumanMethylation450kmanifest’

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Version 0.2.1

Title Annotation for Illumina’s 450k methylation arrays

Description Manifest for Illumina’s 450k array data

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Depends R (>= 2.13.0), minfi

LazyData yes

biocViews IlluminaChip, Homo_sapiens, DNAMethylation

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IlluminaHumanMethylation450kmanifest
Manifest object for IlluminaHumanMethylation450kmanifest

Description

This object contains the array design for Illumina’s Human Methylation 450k microarray.

Usage

```
data(IlluminaHumanMethylation450kmanifest)
```

Format

An object of class “IlluminaMethylationManifest”

References

Bibikova, M. et al. (2011) Genomics. High density DNA methylation array with single CpG site resolution. <http://dx.doi.org/10.1016/j.ygeno.2011.07.007>

See Also

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

Examples

```
data(IlluminaHumanMethylation450kmanifest)
```

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