

# ILLUMINAHUMANMETHYLATION450KMANIFEST

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ILLUMINAHUMANMETHYLATION450KMANIFEST

*Manifest object for the ILLUMINAHUMANMETHYLATION450K array*

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## Description

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

## Usage

```
data(ILLUMINAHUMANMETHYLATION450KMANIFEST)
```

## Format

An object of class `ILLUMINAMETHYLATIONMANIFEST`

## Details

This package is based on the file `HUMANMETHYLATION450_15017482_v.1.2.csv`.

## 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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## \* datasets

    IlluminaHumanMethylation450kmanifest,

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