

# illuminaHumanMethylation450kmanifest

April 2, 2013

---

illuminaHumanMethylation450kmanifest

*Manifest object for the illuminaHumanMethylation450k array*

---

## 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)
```

# Index

## \*Topic **datasets**

    IlluminaHumanMethylation450kmanifest,  
    [1](#)

IlluminaHumanMethylation450kmanifest, [1](#)