

# IlluminaHumanMethylationMSAmanifest

January 21, 2026

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## IlluminaHumanMethylationMSAmanifest

*Manifest object for the IlluminaHumanMethylationMSA array*

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### Description

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

### Usage

```
data(IlluminaHumanMethylationMSAmanifest)
```

### 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/infinummethylationscreening/MSA-48v1-0\\_20102838\\_A1.csv](https://support.illumina.com/content/dam/illumina-support/documents/downloads/productfiles/infinummethylationscreening/MSA-48v1-0_20102838_A1.csv)

The script used to generate this package can be found in `system.files("scripts", "manifest.R", package = "IlluminaHumanMethylationMSAmanifest")`

### See Also

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

### Examples

```
data(IlluminaHumanMethylationMSAmanifest)
```

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