

# illuminaHumanMethylationMSAmanifest

December 11, 2024

---

illuminaHumanMethylationMSAmanifest

*Manifest object for the illuminaHumanMethylationMSA array*

---

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

# Index

## \* datasets

    IlluminaHumanMethylationMSAmanifest,  
    [1](#)

IlluminaHumanMethylationMSAmanifest, [1](#)