

# ILLUMINAHUMANMETHYLATIONMSAMANIFEST

June 3, 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/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)
```

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