

# hgu133afrmovecs

August 27, 2025

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|--------------------|------------------------------------------------------|
| hgu133abarcodevecs | <i>Vectors used by barcode function for hgu133a.</i> |
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## Description

Parameters of the background distribution for use with the barcode function. These correspond to version 3.0 of the Gene Expression Barcode website.

## Usage

```
data(hgu133abarcodevecs)
```

## Format

A list with 3 elements.

|         |                                |
|---------|--------------------------------|
| mu      | background means               |
| tau     | background standard deviations |
| entropy | observed gene entropy          |

## Examples

```
data(hgu133abarcodevecs)  
str(hgu133abarcodevecs)
```

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|                 |                                          |
|-----------------|------------------------------------------|
| hgu133afrmovecs | <i>Vectors used by fRMA for hgu133a.</i> |
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## Description

Frozen parameter vectors for use with the frma and GNUMSE functions.

## Usage

```
data(hgu133afrmovecs)  
data(hgu133ahsentrezgfrmovecs)
```

**Format**

A list with 6 or 7 elements.

|                 |                                                       |
|-----------------|-------------------------------------------------------|
| normVec         | normalization vector                                  |
| probeVec        | probe effect vector                                   |
| probeVarWithin  | within batch probe variance                           |
| probeVarBetween | between batch probe variance                          |
| probesetSD      | within probeset standard deviation                    |
| medianSE        | median standard error for gene expression estimates   |
| version         | the Entrez Gene alternative CDF version if applicable |

**Examples**

```
data(hgu133afrmvects)
str(hgu133afrmvects)
```

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