

# hgu133afrmavecs

October 2, 2015

---

hgu133abarcovecs      *Vectors used by barcode function for hgu133a.*

---

## 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(hgu133abarcovecs)
```

## Format

A list with 3 elements.

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

## Examples

```
data(hgu133abarcovecs)  
str(hgu133abarcovecs)
```

---

hgu133afrmavecs      *Vectors used by fRMA for hgu133a.*

---

## Description

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

**Usage**

```
data(hgu133afirmavecs)
data(hgu133ahsentrezgfirmavecs)
```

**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(hgu133afirmavecs)
str(hgu133afirmavecs)
```

# Index

## \*Topic **datasets**

hgu133abarcovecs, [1](#)

hgu133afrmavecs, [1](#)

hgu133abarcovecs, [1](#)

hgu133afrmavecs, [1](#)

hgu133ahsentrezgfrmovecs  
(hgu133afrmavecs), [1](#)