

# hgu133a2frmavecs

September 27, 2024

---

hgu133a2barcodevecs     *Vectors used by barcode function for hgu133a2.*

---

## 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(hgu133a2barcodevecs)
```

## Format

A list with 3 elements.

mu	background means
tau	background standard deviations
entropy	observed gene entropy

## Examples

```
data(hgu133a2barcodevecs)  
str(hgu133a2barcodevecs)
```

---

hgu133a2frmavecs     *Vectors used by fRMA for hgu133a2.*

---

## Description

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

## Usage

```
data(hgu133a2frmavecs)  
data(hgu133a2hsentrezgfrmavecs)
```

**Format**

A list with 6 or 7 elements.

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

**Examples**

```
data(hgu133a2frmavecs)  
str(hgu133a2frmavecs)
```

# Index

## \* datasets

hgu133a2barcodevecs, [1](#)

hgu133a2frmavecs, [1](#)

hgu133a2barcodevecs, [1](#)

hgu133a2frmavecs, [1](#)

hgu133a2hsentrezgfrmavecs  
(hgu133a2frmavecs), [1](#)