

hgu133plus2frmavecs

February 25, 2026

hgu133plus2barcodevecs

Vectors used by barcode function for hgu133plus2.

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(hgu133plus2barcodevecs)
```

Format

A list with 3 elements.

mu	background means
tau	background standard deviations
entropy	observed gene entropy

Examples

```
data(hgu133plus2barcodevecs)  
str(hgu133plus2barcodevecs)
```

hgu133plus2frmavecs *Vectors used by fRMA for hgu133plus2.*

Description

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

Usage

```
data(hgu133plus2frmavecs)
data(hgu133plus2hsentrezgfrmavecs)
```

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(hgu133plus2frmavecs)
str(hgu133plus2frmavecs)
```

Index

* datasets

hgu133plus2barcodevecs, [1](#)

hgu133plus2frmavecs, [2](#)

hgu133plus2barcodevecs, [1](#)

hgu133plus2frmavecs, [2](#)

hgu133plus2hsentrezgfrmavecs
(hgu133plus2frmavecs), [2](#)