

# Package ‘hgu133plus2barcodevecs’

July 2, 2024

**Title** hgu133plus2 data for barcode

**Version** 1.42.0

**Author** Matthew N. McCall <mmccall@jhsp.h.edu>, Rafael A. Irizarry <rafa@jhu.edu>

**Description** Data used by the barcode package for microarrays of type hgu133plus2.

**Maintainer** Matthew N. McCall <mmccall@jhsp.h.edu>

**Depends** R (>= 2.10.0)

**License** GPL (>= 2)

**biocViews** Homo\_sapiens\_Data, MicroarrayData

**git\_url** <https://git.bioconductor.org/packages/hgu133plus2barcodevecs>

**git\_branch** RELEASE\_3\_19

**git\_last\_commit** 5db696c

**git\_last\_commit\_date** 2024-04-30

**Repository** Bioconductor 3.19

**Date/Publication** 2024-07-02

## Contents

hgu133plus2barcodevecs . . . . .	1
<b>Index</b>	<b>3</b>

---

hgu133plus2barcodevecs  
*hgu133plus2 data for barcode*

---

## Description

Data used by the barcode package for microarrays of type hgu133plus2.

**Usage**

```
data(bcparams-GPL570)
```

**Format**

A matrix of parameters estimates used in the barcode function for Affymetrix hgu133plus2 microarray data.

# Index

## \* datasets

hgu133plus2barcodevecs, [1](#)

bcparams (hgu133plus2barcodevecs), [1](#)

hgu133plus2barcodevecs, [1](#)