

# Package ‘hgu133abarcovecs’

October 16, 2018

**Title** hgu133a data for barcode

**Version** 1.18.0

**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 hgu133a.

**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/hgu133abarcovecs>

**git\_branch** RELEASE\_3\_7

**git\_last\_commit** c8b482f

**git\_last\_commit\_date** 2018-04-30

**Date/Publication** 2018-10-16

## R topics documented:

hgu133abarcovecs . . . . . 1

**Index** 2

---

hgu133abarcovecs	<i>hgu133a data for barcode</i>
------------------	---------------------------------

---

## Description

Data used by the barcode package for microarrays of type hgu133a.

## Usage

```
data(bcparams-GPL96)
```

## Format

A matrix of parameters estimates used in the barcode function for Affymetrix hgu133a microarray data.

# Index

## \*Topic **datasets**

hgu133abarcovecs, [1](#)

bcparams (hgu133abarcovecs), [1](#)

hgu133abarcovecs, [1](#)