

Package ‘hapmap100khind’

September 17, 2024

Title Sample data - Hapmap 100K HIND Affymetrix
Version 1.46.0
Author Hapmap Consortium
Suggests oligo
Maintainer Benilton Carvalho <Benilton.Carvalho@cancer.org.uk>
Description Sample dataset obtained from <http://www.hapmap.org>
biocViews ExperimentData, HapMap, CancerData
License GPL
git_url <https://git.bioconductor.org/packages/hapmap100khind>
git_branch RELEASE_3_19
git_last_commit f7436a5
git_last_commit_date 2024-04-30
Repository Bioconductor 3.19
Date/Publication 2024-09-17

Contents

hapmap100khind-package	1
Index	3

hapmap100khind-package
Sample HapMap 100K Hind

Description

Sample dataset on the 100K Hind Affymetrix platform. Data obtained from <http://www.hapmap.org>. The package is meant to be used only for demonstration of BioConductor tools.

Details

Package: hapmap100khind
Type: Package
Version: 1.0
Date: March/2007
License: GPL

Details on the data are available at <http://www.hapmap.org> .

The maintainer does not warrant the accuracy of the data.

Author(s)

Data generated by the HapMap Consortium.

Package maintained by Benilton Carvalho <Benilton.Carvalho@cancer.org.uk>.

References

The International HapMap Consortium. The International HapMap Project. Nature 426, 789-796 (2003).

Examples

```
## library(oligo)
## library(hapmap100khind)
## the.path <- system.file("celFiles", package="hapmap100khind")
## cels <- list.celfiles(path=the.path, full.names=TRUE)
## temporaryDir <- tempdir()
## rawData <- read.celfiles(fullfilenames, tmpdir=temporaryDir)
```

Index

* **package**

hapmap100khind-package, [1](#)

hapmap100khind

(hapmap100khind-package), [1](#)

hapmap100khind-package, [1](#)