Package 'hapmapsnp5'

November 12, 2026

Title Sample data - Hapmap SNP 5.0 Affymetrix

Version 1.55.0

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, SNPData

License GPL

git_url https://git.bioconductor.org/packages/hapmapsnp5

git_branch devel

git_last_commit 949e1ae

git_last_commit_date 2026-04-28

Repository Bioconductor 3.24

Date/Publication 2026-11-12

Contents

hapmapsnp5-package 1

Index **3**

`hapmapsnp5-package` *Sample HapMap SNP 5.0*

Description

Sample dataset on the SNP 5.0 Affymetrix platform. Data obtained from http://www.hapmap.org . The package is meant to be used only for demonstration of BioConductor tools.

Details

Package:	hapmapsnp5
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(hapmapsnp5)
## the.path <- system.file("celFiles", package="hapmapsnp5")
## cels <- list.celfiles(path=the.path, full.names=TRUE)
## temporaryDir <- tempdir()
## rawData <- read.celfiles(fullFilenames, tmpdir=temporaryDir)
```

Index

∗ **package**
hapmapsnp5-package, 1

hapmapsnp5 *(*hapmapsnp5-package*)*, 1
hapmapsnp5-package, 1