Package 'hapmap500ksty'

April 24, 2025

Title Sample data - Hapmap 500K STY Affymetrix

Version 1.50.0

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/hapmap500ksty

git_branch RELEASE_3_21

git_last_commit d3b6505

git_last_commit_date 2025-04-15

Repository Bioconductor 3.21

Date/Publication 2025-04-24

Contents

hapmap500ksty-package 1

Index **3**

`hapmap500ksty-package` *Sample HapMap 500K Sty*

Description

Sample dataset on the 500K Sty Affymetrix platform. Data obtained from http://www.hapmap.org . The package is meant to be used only for demonstration of BioConductor tools.

Details

Package:	hapmap500ksty
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(hapmap500ksty)
## the.path <- system.file("celFiles", package="hapmap500ksty")
## cels <- list.celfiles(path=the.path, full.names=TRUE)
## temporaryDir <- tempdir()
## rawData <- read.celfiles(fullFilenames, tmpdir=temporaryDir)
```

Index

∗ **package**
hapmap500ksty-package, 1

hapmap500ksty *(*hapmap500ksty-package*)*, 1
hapmap500ksty-package, 1