

Package ‘hapmap500ksty’

April 13, 2022

Title Sample data - Hapmap 500K STY Affymetrix
Version 1.36.0
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_14
git_last_commit 423345b
git_last_commit_date 2021-10-26
Date/Publication 2022-04-13

R topics documented:

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](#)