

Package ‘hapmap100kxba’

April 11, 2019

Title Sample data - Hapmap 100K XBA Affymetrix

Version 1.24.0

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/hapmap100kxba>

git_branch RELEASE_3_8

git_last_commit 33d3b9b

git_last_commit_date 2018-10-30

Date/Publication 2019-04-11

R topics documented:

hapmap100kxba-package 1

Index 3

hapmap100kxba-package *Sample HapMap 100K Xba*

Description

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

Details

Package: hapmap100kxba
Type: Package
Version: 1.0
Date: 2007-03-03
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(hapmap100kxba)
## the.path <- system.file("celFiles", package="hapmap100kxba")
## cels <- list.celfiles(path=the.path, full.names=TRUE)
## temporaryDir <- tempdir()
## rawData <- read.celfiles(fullfilenames, tmpdir=temporaryDir)
```

Index

*Topic **package**

hapmap100kxba-package, [1](#)

hapmap100kxba (hapmap100kxba-package), [1](#)

hapmap100kxba-package, [1](#)