

Package ‘hapmap100kxba’

October 18, 2022

Title Sample data - Hapmap 100K XBA Affymetrix
Version 1.38.0
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_15
git_last_commit 97f0621
git_last_commit_date 2022-04-26
Date/Publication 2022-10-18

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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)
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