Package 'hapmap100kxba'

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Title Sample data - Hapmap 100K XBA Affymetrix

Version 1.54.0

Author Hapmap Consortium

Suggests oligo

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Description Sample dataset obtained from http://www.hapmap.org

biocViews ExperimentData, HapMap, CancerData

License GPL

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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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