hapmap100khind

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hapmap100khind-package

Sample HapMap 100K Hind

Description

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

Details

Package:	hapmap100khind
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 <bcarvalh@jhsph.edu>.

References

The International HapMap Consortium. The International HapMap Project. Nature 426, 789-796 (2003).

Examples

```
## library(oligo)
## library(hapmap100khind)
## the.path <- system.file("celFiles", package="hapmap100khind")
## cels <- list.celfiles(path=the.path, full.names=TRUE)
```

```
## temporaryDir <- tempdir()
## rawData <- read.celfiles(fullFilenames, tmpdir=temporaryDir)
```

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