

Package ‘hapmap100khind’

February 3, 2026

Title Sample data - Hapmap 100K HIND Affymetrix

Version 1.53.0

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/hapmap100khind>

git_branch devel

git_last_commit 45e32ee

git_last_commit_date 2025-10-29

Repository Bioconductor 3.23

Date/Publication 2026-02-03

Contents

hapmap100khind-package	1
Index	3

hapmap100khind-package
<i>Sample HapMap 100K Hind</i>

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 <Benilton.Carvalho@cancer.org.uk>.

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)
```

Index

* **package**

hapmap100khind-package, [1](#)

hapmap100khind

(hapmap100khind-package), [1](#)

hapmap100khind-package, [1](#)