

# Package ‘hapmap100khind’

October 18, 2017

**Title** Sample data - Hapmap 100K HIND Affymetrix

**Version** 1.18.0

**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

**NeedsCompilation** no

## R topics documented:

hapmap100khind-package . . . . . 1

**Index** . . . . . 3

---

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 <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

\*Topic **package**

hapmap100khind-package, [1](#)

hapmap100khind

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

hapmap100khind-package, [1](#)