

hapmapsnp6

August 10, 2009

hapmapsnp6-package *Sample HapMap SNP 6.0*

Description

Sample dataset on the SNP 6.0 Affymetrix platform. Data obtained from <http://www.hapmap.org> .
The package is meant to be used only for demonstration of BioConductor tools.

Details

Package: hapmapsnp6
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@jhsp.edu>.

References

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

Examples

```
## library(oligo)
## library(hapmapsnp6)
## the.path <- system.file("celFiles", package="hapmapsnp6")
## cels <- list.celfiles(path=the.path, full.names=TRUE)
## temporaryDir <- tempdir()
## rawData <- read.celfiles(fullfilenames, tmpdir=temporaryDir)
```

Index

*Topic **package**

hapmapsnp6-package, [1](#)

hapmapsnp6 (*hapmapsnp6-package*), [1](#)

hapmapsnp6-package, [1](#)