

# hapmapsnp5

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hapmapsnp5-package *Sample HapMap SNP 5.0*

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## Description

Sample dataset on the SNP 5.0 Affymetrix platform. Data obtained from <http://www.hapmap.org> .  
The package is meant to be used only for demonstration of BioConductor tools.

## Details

Package: hapmapsnp5  
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](mailto:bcarvalh@jhsp.edu)>.

## References

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

## Examples

```
## library(oligo)
## library(hapmapsnp5)
## the.path <- system.file("celFiles", package="hapmapsnp5")
## cels <- list.celfiles(path=the.path, full.names=TRUE)
## temporaryDir <- tempdir()
## rawData <- read.celfiles(fullfilenames, tmpdir=temporaryDir)
```

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