Package ‘hapmapsnp6’

February 20, 2024

Title  Sample data - Hapmap SNP 6.0 Affymetrix
Version 1.44.0
Author Hapmap Consortium
Depends R (>= 2.15)
Suggests oligo, oligoClasses
Maintainer Benilton Carvalho <Benilton.Carvalho@cancer.org.uk>
Description Sample dataset obtained from http://www.hapmap.org
biocViews  ExperimentData, HapMap, SNPData
License GPL
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R topics documented:

  hapmapsnp6-package .................................................. 2
crlmmResult ............................................................ 3

Index  ................................................................. 4
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

<table>
<thead>
<tr>
<th>Package:</th>
<th>hapmapsnp6</th>
</tr>
</thead>
<tbody>
<tr>
<td>Type:</td>
<td>Package</td>
</tr>
<tr>
<td>Version:</td>
<td>1.0</td>
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<td>March/2007</td>
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<td>License:</td>
<td>GPL</td>
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</table>

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


Examples

```r
# 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)
```
crlmmResult

Results for crlmm on 90 CEU HapMap samples

Description

Results for crlmm on 90 CEU HapMap samples. These results were produced straight from the CEL files available on hapmap.org and are meant to be used for demonstration.

Usage

data(crlmmResult)

Format

SnpSet object.

Source

http://www.hapmap.org

Examples

data(crlmmResult)
Index

* datasets
  * crlmmResult, 3
* package
  * hapmapsnp6-package, 2

crlmmResult, 3

hapmapsnp6 (hapmapsnp6-package), 2
hapmapsnp6-package, 2