# Package 'hapmap500knsp'

April 24, 2025

**Title** Sample data - Hapmap 500K NSP Affymetrix

**Version** 1.50.0

**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/hapmap500knsp

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`hapmap500knsp-package` *Sample HapMap 500K Nsp*

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### Description

Sample dataset on the 500K Nsp Affymetrix platform. Data obtained from http://www.hapmap.org . The package is meant to be used only for demonstration of BioConductor tools.

## Details

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|---|---|
| Package: | hapmap500knsp |
| 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(hapmap500knsp)
## the.path <- system.file("celFiles", package="hapmap500knsp")
## cels <- list.celfiles(path=the.path, full.names=TRUE)
## temporaryDir <- tempdir()
## rawData <- read.celfiles(fullFilenames, tmpdir=temporaryDir)
```

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