

# hapmap370k

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hapmap370k-package      *Example HapMap data from Illumina 370k BeadChips*

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## Description

Example HapMap dataset from the Illumina 370k platform. Arrays were processed at CIDR using samples obtained from the International HapMap Consortium This package is only intended for use in the demonstration of BioConductor tools.

## Details

Package: hapmap370k  
Type: Package  
Version: 1.0  
Date: May/2009  
License: GPL

Details on the data are available from <http://www.hapmap.org>.

The maintainer does not warrant the accuracy of the data.

## Author(s)

Package maintained by Matthew Ritchie <mritchie@wehi.edu.au>

## References

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

## Examples

```
## library(cr1mm)
```

```
## library(hapmap370k)
## the.path <- system.file("idatFiles", package="hapmap370k")
## RG = readIdatFiles(path=the.path)
```

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