# Package 'SNPhoodData'

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Title Additional and more complex example data for the SNPhood package	
<b>Version</b> 1.34.0	
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<b>Description</b> This companion package for SNPhood provides some example datasets of a larger size than allowed for the SNPhood package. They include full and real-world examples for performing analyses with the SNPhood package.	
VignetteBuilder knitr	
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<b>Depends</b> R (>= $3.2$ )	
biocViews ExperimentData	
License LGPL (>= 3)	
LazyData true	
BugReports christian.arnold@embl.de	
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## Description

This companion package for SNPhood provides some example datasets of a larger size than allowed for the SNPhood package. They include full and real-world examples for performing analyses with SNPhood.

#### **Details**

```
For a list of files that the package provides, type
```

```
list.files(pattern = "*", system.file("extdata", package = "SNPhoodData"),full.names
= TRUE)
```

See also the package vignette (browseVignettes("SNPhoodData")).

#### Data

For an even more detailed overview of the data that the SNPhoodData package provides, see the vignette. Briefly, the example dataset consists of the following files:

- 14,000 previously identified H3K27ac QTLs for individuals from the YRI population [1,2] (file cisQ.H3K27AC.chr21.txt)
- H3K27ac ChIP-Seq data in BAM format for two individuals (GM10847, GM12890) obtained from Kasowski et al [2] from the CEU population (two replicates each, files SNYDER\_HG19\_\*). The reads have previously been mapped to the personalized phased genomes of these individuals [2].
- Corresponding genotypes for the SNPs were obtained from the \*1000 Genomes Project\* [3] (file genotypes.vcf.gz)

For the references, see the corresponding package vignette by typing browseVignettes("SNPhoodData").

## Usage

You may use the example files to run a full SNPhood analysis. For a full example, see the workflow vignette in the SNPhood package (browseVignettes("SNPhood"))

#### **Contact Information**

We value all the feedback that we receive and will try to reply in a timely manner. Please report any bug that you encounter as well as any feature request that you may have to <SNPhood@gmail.com>.

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