

Package: genetics.binaRies (via r-universe)

September 14, 2024

Title Binaries for Tools Used in Genetics in an R Package

Version 0.1.1

Description A convenient way to make binaries such as plink, bcftools, and others available to other packages. This package forms part of the MRC IEU OpenGWAS system. The system includes other R packages, including; gwasglue, ieugwasr, gwasvcf, and TwoSampleMR.

License MIT + file LICENSE

URL <https://github.com/MRCIEU/genetics.binaRies>

BugReports <https://github.com/MRCIEU/genetics.binaRies/issues>

Imports utils

Suggests testthat (>= 3.0.0)

Encoding UTF-8

Roxygen list(markdown = TRUE)

RoxygenNote 7.2.3

Config/testthat/edition 3

Repository <https://mrcieu.r-universe.dev>

RemoteUrl <https://github.com/MRCIEU/genetics.binaRies>

RemoteRef HEAD

RemoteSha b0324f180476d80c43bba2ab026b72c5be426a92

Contents

get_bcftools_binary	2
get_gcta_binary	2
get_plink_binary	3

Index	4
--------------	----------

get_bcftools_binary *Find binary for bcftools*

Description

Find binary for bcftools

Usage

```
get_bcftools_binary()
```

Value

Path to bcftools

Examples

```
if (Sys.info()[["sysname"]] != "Windows") get_bcftools_binary()
```

get_gcta_binary *Find binary for gcta*

Description

Find binary for gcta

Usage

```
get_gcta_binary()
```

Value

path to gcta

Examples

```
get_gcta_binary()
```

<code>get_plink_binary</code>	<i>Find binary for plink</i>
-------------------------------	------------------------------

Description

Find binary for plink

Usage

```
get_plink_binary()
```

Value

Path to plink

Examples

```
get_plink_binary()
```

Index

`get_bcftools_binary`, [2](#)
`get_gcta_binary`, [2](#)
`get_plink_binary`, [3](#)