

Package: genetics.binaRies (via r-universe)

October 19, 2026

Title Binaries for Tools Used in Genetics in an R Package

Version 0.1.2

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.

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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 7.3.2

Config/testthat/edition 3

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

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

RemoteRef HEAD

RemoteSha 2fcd3ee3088b729c7eb34cf2aac9dc2e04fe4412

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get_bcftools_binary	<i>Find binary for bcftools</i>
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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	<i>Find binary for gcta</i>
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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>
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Description

Find binary for plink

Usage

```
get_plink_binary()
```

Value

Path to plink

Examples

```
get_plink_binary()
```

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