

Package: EpipwR.data (via r-universe)

October 17, 2026

Type Package

Title EpipwR.data: Reference data for EpipwR

Version 1.7.0

Description This package provides reference data for EpipwR. EpipwR is a fast and efficient power analysis for continuous and binary phenotypes of epigenomic-wide association studies. This package is only meant to be used in conjunction with EpipwR.

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Encoding UTF-8

LazyData false

URL <https://github.com/jbarth216/EpipwR.data>

BugReports <https://github.com/jbarth216/EpipwR.data>

Imports ExperimentHub

Suggests knitr, rmarkdown, sessioninfo

biocViews ExperimentHub, MethylationArrayData, MicroarrayData, TissueMicroarrayData, Tissue, ExperimentData

VignetteBuilder knitr

Config/pak/sysreqs libicu-dev libpng-dev libssl-dev zlib1g-dev

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

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RemoteUrl <https://github.com/bioc/EpipwR.data>

RemoteRef HEAD

RemoteSha cba09c2ff4e0e71f6bbe2f7195c7de76692e1a66

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`EpiwR.data`*EpiwR.data: Reference data for EpiwR packages*

Description

This package provides reference data required for the EpiwR package. Additional information about the source of each reference data set can be found in the vignette: `vignette("EpiwR.data")`

Value

`EpiwR.data` gives the user access to experiment data needed for the EpiwR R package

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