

Package ‘EpipwR.data’

May 26, 2026

Type Package

Title EpipwR.data: Reference data for EpipwR

Version 1.6.0

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

git_url <https://git.bioconductor.org/packages/EpipwR.data>

git_branch RELEASE_3_23

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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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