Package 'EpipwR.data'

November 10, 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.

License Artistic-2.0

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

git_last_commit feb6ed4

git_last_commit_date 2026-04-28

Repository Bioconductor 3.23

Date/Publication 2026-11-10

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

Description

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

Value

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

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