

Package ‘EpipwR.data’

January 22, 2026

Type Package

Title EpipwR.data: Reference data for EpipwR

Version 1.4.0

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_22

git_last_commit fbead4a

git_last_commit_date 2025-10-29

Repository Bioconductor 3.22

Date/Publication 2026-01-22

Author Jackson Barth [aut, cre] (ORCID:
<<https://orcid.org/0009-0009-6307-9928>>),
Austin Reynolds [aut]

Maintainer Jackson Barth <Jackson_Barth@Baylor.edu>

Contents

EpipwR.data	2
Index	3

`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

Index

EpipwR.data, [2](#)