

Package ‘EpipwR.data’

May 14, 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.

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 devel

git_last_commit cba09c2

git_last_commit_date 2026-04-28

Repository Bioconductor 3.24

Date/Publication 2026-05-14

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

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

Index

EpipwR.data, [2](#)