

Package ‘pwrEWAS.data’

April 13, 2022

Title pwrEWAS.data: Reference data accompanying pwrEWAS

Version 1.8.0

Description This package provides reference data required for pwrEWAS.
pwrEWAS is a user-friendly tool to estimate power in EWAS as a function of sample and effect size for two-group comparisons of DNAm (e.g., case vs control, exposed vs non-exposed, etc.).

Author Stefan Graw

Maintainer Stefan Graw <shgraw@uams.edu>

License Artistic-2.0

LazyData FALSE

Imports ExperimentHub

Suggests knitr, RUnit, BiocGenerics

biocViews ExperimentHub, MethylationArrayData, MicroarrayData, TissueMicroarrayData, Tissue

Encoding UTF-8

RoxygenNote 6.1.1

VignetteBuilder knitr

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

git_branch RELEASE_3_14

git_last_commit 560f9f5

git_last_commit_date 2021-10-26

Date/Publication 2022-04-13

R topics documented:

pwrEWAS.data	2
Index	3

pwrEWAS.data

pwrEWAS.data: Reference data accompanying pwrEWAS

Description

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

Index

- * **DNAm**

 - pwrEWAS.data, [2](#)

- * **microarray**

 - pwrEWAS.data, [2](#)

- * **tissue**

 - pwrEWAS.data, [2](#)

pwrEWAS.data, [2](#)