

Package ‘pwrEWAS.data’

October 26, 2021

Title pwrEWAS.data: Reference data accompanying pwrEWAS

Version 1.7.0

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 master

git_last_commit 5c3664e

git_last_commit_date 2021-05-19

Date/Publication 2021-10-26

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](#)