

# Package ‘pwrEWAS.data’

October 16, 2021

**Title** pwrEWAS.data: Reference data accompanying pwrEWAS

**Version** 1.6.0

**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.).

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**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\_13

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*pwrEWAS.data*

*pwrEWAS.data: Reference data accompanying pwrEWAS*

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**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")`

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