

Package ‘epimutacionsData’

June 5, 2025

Title Data for epimutacions package

Version 1.12.0

Description This package includes the data necessary to run functions and examples in epimutacions package.
Collection of DNA methylation data.
The package contains 2 datasets:
(1) Control (GEO: GSE104812), (GEO: GSE97362) case samples; and
(2) reference panel (GEO: GSE127824).
It also contains candidate regions to
be epimutations in 450k methylation arrays.

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biocViews ExperimentHub, Homo_sapiens_Data, MethylationArrayData, MicroarrayData

Suggests rmarkdown, BiocStyle, knitr, ExperimentHub, minfi

VignetteBuilder knitr

Encoding UTF-8

Depends R (>= 4.2.0)

URL <https://github.com/LeireAbarrategui/epimutacionsData>

BugReports <https://github.com/LeireAbarrategui/epimutacionsData/issues>

NeedsCompilation no

RoxygenNote 7.1.2

git_url <https://git.bioconductor.org/packages/epimutacionsData>

git_branch RELEASE_3_21

git_last_commit c7a24c4

git_last_commit_date 2025-04-15

Repository Bioconductor 3.21

Date/Publication 2025-06-05

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