

Package ‘epimutacionsData’

July 2, 2024

Title Data for epimutacions package

Version 1.8.0

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_19

git_last_commit c57824f

git_last_commit_date 2024-04-30

Repository Bioconductor 3.19

Date/Publication 2024-07-02

Author Leire Abarrategui [aut, cre],
Juan R. Gonzalez [aut],
Carlos Ruiz-Arenas [aut],
Carles Hernandez-Ferrer [aut]

Maintainer Leire Abarrategui <abarrategui.leire@gmail.com>

Contents