Package 'epimutacionsData'

November 10, 2026

Title Data for epimutacions package

Version 1.16.0

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

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