Package ‘smokingMouse’

January 25, 2024

Title Provides access to smokingMouse project data
Version 1.0.0
Date 2023-07-21

Description This is an ExperimentHub package that provides access to the data at the gene, exon, transcript and junction level used in the analyses of the smokingMouse project. See https://github.com/LieberInstitute/smokingMouse_Indirects. This datasets contain the expression counts of genes, transcripts, exons and exon-exon junctions across 208 mice samples from pup and adult brains and adult blood. They also contain relevant information of these samples and features, such as conditions, QC metrics and if they were used after filtering steps and also if the features were differently expressed in the different experiments.

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BugReports https://support.bioconductor.org/t/smokingMouse

Suggests ExperimentHub, AnnotationHubData, BiocStyle, covr, ExperimentHubData, knitr, RefManageR, rmarkdown, sessioninfo, testthat (>= 3.0.0)

biocViews ExperimentHub, ExpressionData, Mus_musculus_Data, RNASeqData

Encoding UTF-8

Roxygen list(markdown = TRUE)

RoxygenNote 7.2.3

Config/testthat/edition 2

URL https://github.com/LieberInstitute/smokingMouse

VignetteBuilder knitr
git_url https://git.bioconductor.org/packages/smokingMouse
git_branch RELEASE_3_18
git_last_commit cb8ad94
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R topics documented:

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smokingMouse-package  Provides access to data of the smokingMouse project

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Author(s)

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