Package ‘tximportData’

March 19, 2024

Title       tximportData
Version     1.30.0
Author      Michael Love
Maintainer  Michael Love <michaelisaiahlove@gmail.com>
Description This package provides the output of running various transcript abundance quantifiers on a set of 6 RNA-seq samples from the GEUVADIS project. The quantifiers were Cufflinks, RSEM, kallisto, Salmon and Sailfish. alevin example output is also included. For details on version numbers, sample information, and details on calls, see the package vignette.
biocViews   ExperimentData, SequencingData, RNASeqData
License     GPL (>= 2)
Suggests    knitr, markdown
VignetteBuilder knitr
NeedsCompilation no
git_url     https://git.bioconductor.org/packages/tximportData
git_branch  RELEASE_3_18
git_last_commit  57f359c
git_last_commit_date  2023-10-24
Repository   Bioconductor 3.18
Date/Publication  2024-03-19

\R topics documented: