Package ‘tximportData’

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Title  tximportData
Version  1.30.0
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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
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R topics documented: