

Package: tximportData (via r-universe)

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Title tximportData

Version 1.41.1

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Description This package provides the output of running various transcript abundance quantifiers on a set of 2 RNA-seq samples from the GEUVADIS project. The quantifiers were Cufflinks, RSEM, kallisto, Salmon and Sailfish. alevin example output is also included. Inferential replicate data from earlier versions is archived at <[doi:10.5281/zenodo.22982575](https://doi.org/10.5281/zenodo.22982575)>. 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

Repository <https://bioc.r-universe.dev>

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RemoteRef HEAD

RemoteSha 59766ff72a3c545424a085049458f15a0df4fd2c