

# 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:**