

# Package ‘tximportData’

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**Title** tximportData

**Version** 1.23.0

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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. 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** master

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**R topics documented:**