

# Package ‘tximportData’

October 16, 2018

**Title** tximportData

**Version** 1.8.0

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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

**VignetteBuilder** knitr

**NeedsCompilation** no

**git\_url** <https://git.bioconductor.org/packages/tximportData>

**git\_branch** RELEASE\_3\_7

**git\_last\_commit** 88949cb

**git\_last\_commit\_date** 2018-04-30

**Date/Publication** 2018-10-16

**R topics documented:**