Package 'tximportData'

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Title tximportData

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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

R topics documented: