

Package ‘RnaSeqGeneEdgeRQL’

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Title Gene-level RNA-seq differential expression and pathway analysis
using Rsubread and the edgeR quasi-likelihood pipeline

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Description A workflow package for RNA-Seq experiments

Depends R (>= 3.3.0), edgeR, gplots, org.Mm.eg.db, GO.db

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VignetteBuilder knitr

biocViews WorkflowStep

Suggests knitr, knitcitations

NeedsCompilation no

URL <http://f1000research.com/articles/5-1438/v2>

R topics documented: