

# Package ‘RNAseqRtools’

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**Type** Package

**Title** Epigenomics NGS Data Analysis with R.

**Version** 0.1.0

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**Maintainer** The package maintainer <andrea.lauria104@gmail.com>

**Description** A Collection of Tools for Next-  
Generation Sequencing Data Analysis, with a focus on Epigenomics research.

- RNA-seq
- scRNA-seq
- ChIP-seq
- BS-seq

**Encoding** UTF-8

**LazyData** true

**Imports** utilsRtools, ComplexHeatmap, edgeR, DESeq2, sva, cluster,  
vegan, gProfileR, clusterProfiler, org.Mm.eg.db, org.Hs.eg.db

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**R topics documented:**