CummeRbund
RRID:SCR_014568
Type: Tool

Proper Citation

CummeRbund (RRID:SCR_014568)

Resource Information

URL: http://compbio.mit.edu/cummeRbund/index.html

Description: A package of the R software which is used for simplifying and analyzing Cufflink RNA-Seq output. This program takes various output files from a cuffdiff run and creates a SQLite database of the results that will describe the appropriate relationships between the genes, transcripts, transcription start sites and CDS regions.

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Resource Type: Resource, software resource, data processing software, data analysis software, sequence analysis software, software application

Keywords: r software, cufflink, rna-seq, sqlite, gene, transcript, transcription start site, cds region, r, rnaseq, rna seq

Resource ID: SCR_014568

Parent Organization: Harvard University; Cambridge; United States, Massachusetts Institute of Technology; Massachusetts; USA

Availability: Open Source

Website Status: Last checked up

Mentions Count: 255

Ratings and Alerts
No rating or validation information has been found for CummeRbund.

No alerts have been found for CummeRbund.

Data and Source Information
Source: SciCrunch Registry

Usage and Citation Metrics
We found 255 mentions in open access literature.

**Listed below are recent publications.** The full list is available at scicrunch.


