FlipFlop

RRID:SCR_000625
Type: Tool

Proper Citation

FlipFlop (RRID:SCR_000625)

Resource Information

URL: http://bioconductor.org/packages/release/bioc/html/flipflop.html

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Description: Software that discovers which isoforms of a gene are expressed in a given sample together with their abundances, based on RNA-Seq read data.

Synonyms: flipflop - Fast lasso-based isoform prediction as a flow problem

Resource Type: software resource

Defining Citation: PMID:24813214

Keywords: standalone software, unix/linux, mac os x, windows, r, rna-seq, bio.tools

Funding:

Availability: GNU General Public License, v3

Resource Name: FlipFlop

Resource ID: SCR_000625

Alternate IDs: OMICS_04028, biotools:flipflop

Alternate URLs: https://bio.tools/flipflop

Record Creation Time: 20220129T080202+0000

Record Last Update: 20250420T013958+0000

Ratings and Alerts

No rating or validation information has been found for FlipFlop.

No alerts have been found for FlipFlop.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.