

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 9, 2026

FlipFlop

RRID:SCR_000625

Type: Tool

Proper Citation

FlipFlop (RRID:SCR_000625)

Resource Information

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

Proper Citation: FlipFlop (RRID:SCR_000625)

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

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: FlipFlop

Resource ID: SCR_000625

Alternate IDs: OMICS_04028, biotools:flipflop

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

License: GNU General Public License, v3

Record Creation Time: 20220129T080202+0000

Record Last Update: 20260905T132422+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.