

# Resource Summary Report

Generated by [ASWG](#) on Aug 2, 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:** 20260801T190122+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.