## **Resource Summary Report**

Generated by <u>NIF</u> on May 4, 2025

# <u>sailfish</u>

RRID:SCR\_024326 Type: Tool

**Proper Citation** 

sailfish (RRID:SCR\_024326)

#### **Resource Information**

URL: https://www.cs.cmu.edu/~ckingsf/software/sailfish/

Proper Citation: sailfish (RRID:SCR\_024326)

**Description:** Software tool that implements novel, alignment free algorithm for estimation of isoform abundances directly from set of reference sequences and RNA-seq reads.

Resource Type: software resource, software application

Defining Citation: PMID:24752080

Keywords: estimation of isoform abundances, reference sequences, RNA-seq reads,

Funding:

Availability: Free, Available for download, Freely available,

Resource Name: sailfish

Resource ID: SCR\_024326

Alternate IDs: OMICS\_03939

Old URLs: https://sources.debian.org/src/sailfish/

Record Creation Time: 20230830T050217+0000

Record Last Update: 20250503T061143+0000

**Ratings and Alerts** 

No rating or validation information has been found for sailfish.

No alerts have been found for sailfish.

## Data and Source Information

Source: <u>SciCrunch Registry</u>

### **Usage and Citation Metrics**

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at <u>NIF</u>.

Fashemi BE, et al. (2024) IFRD1 is required for maintenance of bladder epithelial homeostasis. iScience, 27(12), 111282.