

Resource Summary Report

Generated by [RRID](#) on Aug 2, 2026

[sailfish](#)

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,

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: 20260801T191119+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: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [RRID](#) .

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