Resource Summary Report

Generated by <u>dkNET</u> on Apr 28, 2025

<u>HSA</u>

RRID:SCR_001809 Type: Tool

Proper Citation

HSA (RRID:SCR_001809)

Resource Information

URL: https://github.com/vlcc/HSA

Proper Citation: HSA (RRID:SCR_001809)

Description: A splice alignment software tool of RNA-Seq reads mapping.

Abbreviations: HSA

Resource Type: software resource

Defining Citation: PMID:24564867

Keywords: rna-seq, next generation sequencing

Funding:

Resource Name: HSA

Resource ID: SCR_001809

Alternate IDs: OMICS_01944

Record Creation Time: 20220129T080209+0000

Record Last Update: 20250420T014040+0000

Ratings and Alerts

No rating or validation information has been found for HSA.

No alerts have been found for HSA.

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>dkNET</u>.

Bu J, et al. (2013) HSA: a heuristic splice alignment tool. BMC systems biology, 7 Suppl 2(Suppl 2), S10.