

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 19, 2026

HSA

RRID:SCR_001809

Type: Tool

Proper Citation

HSA (RRID:SCR_001809)

Resource Information

URL: <http://www.nesys.uio.no/Atlas3D/>

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

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: HSA

Resource ID: SCR_001809

Alternate IDs: OMICS_01944

Record Creation Time: 20220129T080209+0000

Record Last Update: 20260912T195531+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: [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 [PRECISE-TBI](#) .

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