

Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 9, 2026

HeurAA

RRID:SCR_013212

Type: Tool

Proper Citation

HeurAA (RRID:SCR_013212)

Resource Information

URL: <http://sourceforge.net/projects/heuraa/>

Proper Citation: HeurAA (RRID:SCR_013212)

Description: Software for accurate and fast detection of genetic variations with a novel heuristic amplicon aligner program for next generation sequencing.

Abbreviations: HeurAA

Synonyms: heurAA - NGS multiplexed amplicon aligner

Resource Type: software resource

Defining Citation: [PMID:23349847](#)

Keywords: unix/linux, bio.tools

Resource Name: HeurAA

Resource ID: SCR_013212

Alternate IDs: OMICS_00097, biotools:heuraa

Alternate URLs: <https://bio.tools/heuraa>

Record Creation Time: 20220129T080314+0000

Record Last Update: 20260808T190020+0000

Ratings and Alerts

No rating or validation information has been found for HeurAA.

No alerts have been found for HeurAA.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.