

# Resource Summary Report

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## 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.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.