

# Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 7, 2026

## VarMatch

RRID:SCR\_024375

Type: Tool

### Proper Citation

VarMatch (RRID:SCR\_024375)

### Resource Information

**URL:** <https://github.com/medvedevgroup/varmatch>

**Proper Citation:** VarMatch (RRID:SCR\_024375)

**Description:** Software tool for variant matching problem.Used for robust matching of small variant datasets using flexible scoring schemes

**Synonyms:** varmatch

**Resource Type:** data analysis software, data processing software, software application, software resource

**Defining Citation:** [PMID:28011786](#)

**Keywords:** variant matching, matching of small variant datasets, flexible scoring schemes,

**Availability:** Free, Available for download, Freely available,

**Resource Name:** VarMatch

**Resource ID:** SCR\_024375

**Alternate IDs:** OMICS\_14435

**Old URLs:** <https://sources.debian.org/src/varmatch/>

**License:** Apache License 2.0

**Record Creation Time:** 20230830T050217+0000

**Record Last Update:** 20260905T133030+0000

### Ratings and Alerts

No rating or validation information has been found for VarMatch.

No alerts have been found for VarMatch.

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## Data and Source Information

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

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## 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](#) .

Tello D, et al. (2019) NGSEP3: accurate variant calling across species and sequencing protocols. Bioinformatics (Oxford, England), 35(22), 4716.