

# Resource Summary Report

Generated by [RRID](#) on Sep 10, 2026

## SRMA

RRID:SCR\_000669

Type: Tool

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### Proper Citation

SRMA (RRID:SCR\_000669)

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### Resource Information

**URL:** <http://sourceforge.net/projects/srma/>

**Proper Citation:** SRMA (RRID:SCR\_000669)

**Description:** A post-alignment micro re-aligner for next-generation high throughput sequencing data.

**Abbreviations:** SRMA

**Synonyms:** Short Read Micro re-Aligner

**Resource Type:** software resource

**Defining Citation:** [PMID:20932289](#)

**Keywords:** matlab, sequence re-alignment, command-line, java, next generation sequencing, bio.tools

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

**Resource Name:** SRMA

**Resource ID:** SCR\_000669

**Alternate IDs:** biotools:srma, OMICS\_01079

**Alternate URLs:** <https://bio.tools/srma>

**License:** GNU General Public License, v2

**Record Creation Time:** 20220129T080202+0000

**Record Last Update:** 20260905T132423+0000

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### Ratings and Alerts

No rating or validation information has been found for SRMA.

No alerts have been found for SRMA.

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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.