

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

SRMA

RRID:SCR_000669

Type: Tool

Proper Citation

SRMA (RRID:SCR_000669)

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: 20260725T190450+0000

Ratings and Alerts

No rating or validation information has been found for SRMA.

No alerts have been found for SRMA.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.