Resource Summary Report

Generated by <u>dkNET</u> on May 20, 2025

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

Funding:

Availability: GNU General Public License, v2

Resource Name: SRMA

Resource ID: SCR_000669

Alternate IDs: biotools:srma, OMICS_01079

Alternate URLs: https://bio.tools/srma

Record Creation Time: 20220129T080202+0000

Record Last Update: 20250519T203059+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: <u>SciCrunch Registry</u>

Usage and Citation Metrics

We have not found any literature mentions for this resource.