Resource Summary Report

Generated by <u>NIF</u> on May 15, 2025

SBARS

RRID:SCR_002371 Type: Tool

Proper Citation

SBARS (RRID:SCR_002371)

Resource Information

URL: https://github.com/mpyatkov/sbars

Proper Citation: SBARS (RRID:SCR_002371)

Description: Bioinformatics tool for searching different types of long repeats in sequences comparable by size with chromosomes.

Synonyms: S.B.A.R.S, S.B.A.R.S., S.B.A.R.S - Spectral-Based Approach for Repeats Search, Spectral-Based Approach for Repeats Search

Resource Type: software resource

Defining Citation: PMID:24532721

Keywords: linux, windows, bio.tools

Funding:

Resource Name: SBARS

Resource ID: SCR_002371

Alternate IDs: OMICS_03432, biotools:sbars

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

Record Creation Time: 20220129T080213+0000

Record Last Update: 20250420T014102+0000

Ratings and Alerts

No rating or validation information has been found for SBARS.

No alerts have been found for SBARS.

Data and Source Information

Source: <u>SciCrunch Registry</u>

Usage and Citation Metrics

We have not found any literature mentions for this resource.