

Resource Summary Report

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

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

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: 20260725T190521+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: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.