

# Resource Summary Report

Generated by ASWG on Aug 3, 2026

## Bamformatics

RRID:SCR\_013041

Type: Tool

### Proper Citation

Bamformatics (RRID:SCR\_013041)

### Resource Information

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

**Proper Citation:** Bamformatics (RRID:SCR\_013041)

**Description:** Software that provides a coherent and consistent approach to analysis of high-throughput sequencing data.

**Abbreviations:** Bamformatics

**Synonyms:** Bamformatics - Toolkit and GUI for sequencing data analysis

**Resource Type:** software resource

**Keywords:** matlab

**Resource Name:** Bamformatics

**Resource ID:** SCR\_013041

**Alternate IDs:** OMICS\_00283

**Record Creation Time:** 20220129T080314+0000

**Record Last Update:** 20260801T190455+0000

### Ratings and Alerts

No rating or validation information has been found for Bamformatics.

No alerts have been found for Bamformatics.

### Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [ASWG](#) .

Kerzendorfer C, et al. (2015) A thesaurus of genetic variation for interrogation of repetitive genomic regions. Nucleic acids research, 43(10), e68.