

# Resource Summary Report

Generated by [NIF](#) on Sep 11, 2026

## BSA4Yeast

RRID:SCR\_017113

Type: Tool

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### Proper Citation

BSA4Yeast (RRID:SCR\_017113)

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### Resource Information

**URL:** <https://bsa4yeast.lcsb.uni.lu>

**Proper Citation:** BSA4Yeast (RRID:SCR\_017113)

**Description:** Web application for Quantitative Trait Loci mapping via bulk segregant analysis of yeast sequencing data. Application provides automated data processing, annotations, and web interface to explore identified QTLs.

**Synonyms:** Bulk Segregant Analysis for Yeast

**Resource Type:** analysis service resource, data access protocol, data analysis service, production service resource, service resource, software resource, web service

**Keywords:** quantitative, trait, loci, mapping, bulk, segregant, analysis, yeast, sequencing, data, processing, annotation, bio.tools

**Funding:** Fonds Nationale de la Recherche (FNR) Luxembourg

**Availability:** Free, Freely available, Registration required

**Resource Name:** BSA4Yeast

**Resource ID:** SCR\_017113

**Alternate IDs:** biotools:BSA4Yeast

**Alternate URLs:** <https://bio.tools/BSA4Yeast>

**Record Creation Time:** 20220129T080333+0000

**Record Last Update:** 20260905T132818+0000

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### Ratings and Alerts

No rating or validation information has been found for BSA4Yeast.

No alerts have been found for BSA4Yeast.

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## 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 [NIF](#) .

Zhang Z, et al. (2019) BSA4Yeast: Web-based quantitative trait locus linkage analysis and bulk segregant analysis of yeast sequencing data. GigaScience, 8(6).