## **Resource Summary Report**

Generated by <u>RRID</u> on Apr 28, 2025

# **BSA4Yeast**

RRID:SCR\_017113 Type: Tool

**Proper Citation** 

BSA4Yeast (RRID:SCR\_017113)

#### **Resource Information**

URL: https://bsa4yeast.lcsb.uni.lu

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**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:** production service resource, software resource, data analysis service, service resource, data access protocol, analysis service 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

#### **Ratings and Alerts**

No rating or validation information has been found for BSA4Yeast.

No alerts have been found for BSA4Yeast.

#### Data and Source Information

Source: <u>SciCrunch Registry</u>

### **Usage and Citation Metrics**

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at <u>RRID</u>.

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