## **Resource Summary Report**

Generated by NIF on Apr 27, 2025

# **BLINK**

RRID:SCR\_016288

Type: Tool

### **Proper Citation**

BLINK (RRID:SCR\_016288)

#### Resource Information

URL: http://zzlab.net/blink/index.html

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**Description:** Software for next level of genome wide association studies with both individuals and markers in millions. The method releases the requirement that causative genes are evenly distributed on genome and consequently boosts statistical power.

**Synonyms:** BLINK (Bayesian-information and Linkage-disequilibrium Iteratively Nested Keyway)

**Resource Type:** data processing software, data analysis software, algorithm resource, software resource, software application

**Keywords:** GWAS, SNP, dataset, r, genome, bayesian, linkage, nested, keyway, statistic, bio.tools

**Funding:** 

Availability: Free, Available for download, Tutorial available

Resource Name: BLINK

Resource ID: SCR\_016288

Alternate IDs: biotools:BLINK

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

License URLs: https://github.com/Menggg/BLINK

**Record Creation Time:** 20220129T080329+0000

**Record Last Update:** 20250426T060535+0000

## Ratings and Alerts

No rating or validation information has been found for BLINK.

No alerts have been found for BLINK.

### Data and Source Information

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at NIF.

Huang M, et al. (2019) BLINK: a package for the next level of genome-wide association studies with both individuals and markers in the millions. GigaScience, 8(2).