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 <u>NIF</u>.

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).