

Resource Summary Report

Generated by [RRID](#) on Apr 9, 2025

UNKNOWN

RRID:SCR_009423

Type: Tool

Proper Citation

UNKNOWN (RRID:SCR_009423)

Resource Information

URL: <https://github.com/gaow/genetic-analysis-software/blob/master/pages/UNKNOWN.md>

Proper Citation: UNKNOWN (RRID:SCR_009423)

Description: Software application (entry from Genetic Analysis Software)

Abbreviations: UNKNOWN

Synonyms: LINKAGE - general pedigrees

Resource Type: software resource, software application

Keywords: gene, genetic, genomic

Funding:

Resource Name: UNKNOWN

Resource ID: SCR_009423

Alternate IDs: nlx_154684

Record Creation Time: 20220129T080252+0000

Record Last Update: 20250404T060815+0000

Ratings and Alerts

No rating or validation information has been found for UNKNOWN.

No alerts have been found for UNKNOWN.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 10 mentions in open access literature.

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

Fiorillo L, et al. (2023) U-Sleep's resilience to AASM guidelines. NPJ digital medicine, 6(1), 33.

Stahl KA, et al. (2022) Insurance Status and Travel Distance to Single Treatment Facility Predictive of Mastectomy. The Journal of surgical research, 270, 22.

Orr RJS, et al. (2022) Paleozoic origins of cheilostome bryozoans and their parental care inferred by a new genome-skimmed phylogeny. Science advances, 8(13), eabm7452.

Momattin H, et al. (2021) Robotic Pharmacy Implementation and Outcomes in Saudi Arabia: A 21-Month Usability Study. JMIR human factors, 8(3), e28381.

Carter MW, et al. (2019) Increasing Rates of Opioid Misuse Among Older Adults Visiting Emergency Departments. Innovation in aging, 3(1), igz002.

Faria JP, et al. (2016) Computing and Applying Atomic Regulons to Understand Gene Expression and Regulation. Frontiers in microbiology, 7, 1819.

Dong L, et al. (2016) Population-weighted efficiency in transportation networks. Scientific reports, 6, 26377.

Tito RY, et al. (2012) Insights from characterizing extinct human gut microbiomes. PloS one, 7(12), e51146.

Lin SH, et al. (2009) Clustering by neurocognition for fine mapping of the schizophrenia susceptibility loci on chromosome 6p. Genes, brain, and behavior, 8(8), 785.

Clark C, et al. (2008) Identifying smokers with a medical extraction system. Journal of the American Medical Informatics Association : JAMIA, 15(1), 36.