Resource Summary Report

Generated by <u>dkNET</u> on Apr 30, 2025

GWAMA

RRID:SCR_006624 Type: Tool

Proper Citation

GWAMA (RRID:SCR_006624)

Resource Information

URL: http://www.geenivaramu.ee/en/tools/gwama

Proper Citation: GWAMA (RRID:SCR_006624)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on February 28,2023. Software tool for meta analysis of whole genome association data.

Abbreviations: GWAMA

Synonyms: Genome-Wide Association Meta Analysis

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

Defining Citation: PMID:20509871, DOI:10.1186/1471-2105-11-288

Keywords: meta, analysis, genome, association, bio.tools

Funding:

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: GWAMA

Resource ID: SCR_006624

Alternate IDs: biotools:gwama, OMICS_00235

Alternate URLs: https://bio.tools/gwama, https://sources.debian.org/src/gwama/

Old URLs: http://www.well.ox.ac.uk/GWAMA/

Record Creation Time: 20220129T080237+0000

Record Last Update: 20250429T055105+0000

Ratings and Alerts

No rating or validation information has been found for GWAMA.

No alerts have been found for GWAMA.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 172 mentions in open access literature.

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

Marques IF, et al. (2025) Associations of maternal night shift work during pregnancy with DNA methylation in offspring: a meta-analysis in the PACE consortium. Clinical epigenetics, 17(1), 12.

Zhang Y, et al. (2025) The causal relationship between steroid hormones and risk of stroke: evidence from a two-sample Mendelian randomization study. Molecular brain, 18(1), 6.

Munsch G, et al. (2024) Genomic Landscape of Thrombosis Recurrence Risk Across Venous Thromboembolism Subtypes. medRxiv : the preprint server for health sciences.

Hrytsenko Y, et al. (2024) Obstructive sleep apnea mediates genetic risk of Diabetes Mellitus: The Hispanic Community Health Study/Study of Latinos. medRxiv : the preprint server for health sciences.

Eissman JM, et al. (2024) Sex-specific genetic architecture of late-life memory performance. Alzheimer's & dementia : the journal of the Alzheimer's Association, 20(2), 1250.

Burrows K, et al. (2024) A framework for conducting time-varying genome-wide association studies: An application to body mass index across childhood in six multiethnic cohorts. medRxiv : the preprint server for health sciences.

Wu XP, et al. (2024) Association between migraine and venous thromboembolism: a Mendelian randomization and genetic correlation study. Frontiers in genetics, 15, 1272599.

Schurz H, et al. (2024) Multi-ancestry meta-analysis of host genetic susceptibility to tuberculosis identifies shared genetic architecture. eLife, 13.

Keener R, et al. (2024) Validation of human telomere length multi-ancestry meta-analysis association signals identifies POP5 and KBTBD6 as human telomere length regulation genes. Nature communications, 15(1), 4417.

Simmonds E, et al. (2024) Chromosome X-wide association study in case control studies of pathologically confirmed Alzheimer's disease in a European population. Translational psychiatry, 14(1), 358.

Quester J, et al. (2024) High SHBG and Low Bioavailable Testosterone are Strongly Causally Associated with Increased Forearm Fracture Risk in Women: An MR Study Leveraging Novel Female-Specific Data. Calcified tissue international, 115(5), 648.

Rämö JT, et al. (2024) Rare genetic variation in VE-PTP is associated with central serous chorioretinopathy, venous dysfunction and glaucoma. medRxiv : the preprint server for health sciences.

Pujol-Gualdo N, et al. (2024) Circulating anti-Müllerian hormone levels in pre-menopausal women: novel genetic insights from a genome-wide association meta-analysis. Human reproduction (Oxford, England), 39(7), 1564.

Landvreugd A, et al. (2024) Using Polygenic Scores for Circadian Rhythms to Predict Wellbeing, Depressive Symptoms, Chronotype, and Health. Journal of biological rhythms, 39(3), 270.

Burrows K, et al. (2024) A framework for conducting GWAS using repeated measures data with an application to childhood BMI. Nature communications, 15(1), 10067.

Svyatova G, et al. (2024) Genetic Predisposition to Prediabetes in the Kazakh Population. Current issues in molecular biology, 46(10), 10913.

Wang Z, et al. (2024) Analysis of C-reactive protein omics-measures associates methylation risk score with sleep health and related health outcomes. medRxiv : the preprint server for health sciences.

Uddin MM, et al. (2024) Long-term longitudinal analysis of 4,187 participants reveals insights into determinants of clonal hematopoiesis. Nature communications, 15(1), 7858.

Koel M, et al. (2023) GWAS meta-analyses clarify the genetics of cervical phenotypes and inform risk stratification for cervical cancer. Human molecular genetics, 32(12), 2103.

Lagou V, et al. (2023) GWAS of random glucose in 476,326 individuals provide insights into diabetes pathophysiology, complications and treatment stratification. Nature genetics, 55(9), 1448.