Resource Summary Report

Generated by dkNET on Apr 17, 2025

SAGE

RRID:SCR_009302

Type: Tool

Proper Citation

SAGE (RRID:SCR_009302)

Resource Information

URL: http://darwin.cwru.edu/sage/

Proper Citation: SAGE (RRID:SCR_009302)

Description: Software application that provides researchers with the tools necessary for various types of statistical genetic analysis of human family data. (entry from Genetic

Analysis Software)

Abbreviations: SAGE

Synonyms: Statistical Analysis for Genetic Epidemiology

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, c, version 4.0 will be in c++, unix, (dec unix/solaris), ms-

windows, (95/nt), linux

Funding:

Resource Name: SAGE

Resource ID: SCR_009302

Alternate IDs: nlx 154606

Record Creation Time: 20220129T080252+0000

Record Last Update: 20250416T063543+0000

Ratings and Alerts

No rating or validation information has been found for SAGE.

No alerts have been found for SAGE.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 970 mentions in open access literature.

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

Monib WK, et al. (2025) Microlearning beyond boundaries: A systematic review and a novel framework for improving learning outcomes. Heliyon, 11(2), e41413.

Liu H, et al. (2025) Individual and joint associations between sleep duration and physical activity with cognitive function: A longitudinal analysis among middle-aged and older adults in China. Alzheimer's & dementia: the journal of the Alzheimer's Association, 21(1), e14212.

Panieri G, et al. (2025) Sanctuary for vulnerable Arctic species at the Borealis Mud Volcano. Nature communications, 16(1), 504.

Namdar Areshtanab H, et al. (2024) Developing the questionnaire of general population knowledge, attitudes and practices towards the COVID-19 outbreak. Nursing open, 11(3), e2143.

Gorla A, et al. (2024) Epigenetic patient stratification via contrastive machine learning refines hallmark biomarkers in minoritized children with asthma. Research square.

Milte CM, et al. (2024) Cross-sectional associations between fruit and vegetable intake and successful ageing across six countries: findings from the WHO Study on global AGEing and adult health (SAGE). Public health nutrition, 27(1), e244.

Agrawal R, et al. (2024) Understanding the association of disability with multimorbidity, and healthcare utilization in India's older adult population: insights from cross-sectional evidence of SAGE-2. Frontiers in public health, 12, 1435315.

Oduro JK, et al. (2024) Psychological and physical factors related to social integration of older adults in Ghana. BMC geriatrics, 24(1), 363.

Luna Santamaría M, et al. (2024) Digital RNA sequencing using unique molecular identifiers enables ultrasensitive RNA mutation analysis. Communications biology, 7(1), 249.

Fasanghari M, et al. (2024) Effect of alternative and complementary medicine on male infertility: An umbrella review. Health science reports, 7(6), e2118.

Wen B, et al. (2024) Assessment of false discovery rate control in tandem mass spectrometry analysis using entrapment. bioRxiv: the preprint server for biology.

Muyas F, et al. (2024) De novo detection of somatic mutations in high-throughput single-cell profiling data sets. Nature biotechnology, 42(5), 758.

Lekatou AG, et al. (2024) Simulating porcelain firing effect on the structure, corrosion and mechanical properties of Co-Cr-Mo dental alloy fabricated by soft milling. Odontology, 112(2), 372.

Schubert PJ, et al. (2024) DeepFocus: fast focus and astigmatism correction for electron microscopy. Nature communications, 15(1), 948.

Hughes O, et al. (2024) Genome-wide study investigating effector genes and polygenic prediction for kidney function in persons with ancestry from Africa and the Americas. Cell genomics, 4(1), 100468.

Al-Awabdeh E, et al. (2024) Panic disorder during pregnancy: A scoping review. Heliyon, 10(7), e28999.

Sweeney A, et al. (2024) ChemEM: Flexible Docking of Small Molecules in Cryo-EM Structures. Journal of medicinal chemistry, 67(1), 199.

Rosenberg PS, et al. (2024) Cancer Incidence Trends in Successive Social Generations in the US. JAMA network open, 7(6), e2415731.

Javed MA, et al. (2024) Telomere-to-telomere Genome Assembly of the Clubroot Pathogen Plasmodiophora Brassicae. Genome biology and evolution, 16(6).

Davoudi-Monfared E, et al. (2024) Adverse events of cell therapy clinical trials in human chronic spinal cord injury, a systematic review and meta-analysis. Regenerative therapy, 27, 381.