Resource Summary Report

Generated by NIF on Aug 20, 2026

RVTESTS

RRID:SCR_007639
Type: Tool

Proper Citation

RVTESTS (RRID:SCR_007639)

Resource Information

URL: http://genome.sph.umich.edu/wiki/RvTests

Proper Citation: RVTESTS (RRID:SCR_007639)

Description: Software application (entry from Genetic Analysis Software)

Abbreviations: RVTESTS

Synonyms: Rare Variants TESTS

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, c++

Resource Name: RVTESTS

Resource ID: SCR_007639

Alternate IDs: nlx_154603

Record Creation Time: 20220129T080242+0000

Record Last Update: 20260818T043311+0000

Ratings and Alerts

No rating or validation information has been found for RVTESTS.

No alerts have been found for RVTESTS.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 90 mentions in open access literature.

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

Sun Y, et al. (2026) Distinctive genetic architecture of infantile epileptic spasms syndrome compared to self-limited infantile epilepsy by trios whole-exome sequencing. Epilepsia open, 11(1), 280.

Li R, et al. (2026) Whole-genome sequence-based association analysis of African American individuals with bipolar disorder and schizophrenia. HGG advances, 7(1), 100499.

Aliyev E, et al. (2026) The biomedical landscape of genomic structural variation in the qatari population. Nature communications, 17(1), 1019.

Li J, et al. (2026) Circulating metabolites, genetics and lifestyle factors in relation to future risk of type 2 diabetes. Nature medicine, 32(2), 660.

Rosenberger A, et al. (2026) Genes associated with genetic and rare lung diseases and the risk of lung cancer. BMC cancer, 26(1).

Martín-Bórnez M, et al. (2025) Does COMT Play a Role in Parkinson's Disease Susceptibility across Diverse Ancestral Populations? Movement disorders : official journal of the Movement Disorder Society, 40(12), 2819.

Prijatelj V, et al. (2025) Genetic and epidemiologic assessment of mandibular cortical indices and bone mineral density in peripubertal children: the Generation R study. Clinical oral investigations, 29(12), 585.

Martín-Bórnez M, et al. (2025) Does COMT Play a Role in Parkinson's Disease Susceptibility Across Diverse Ancestral Populations? medRxiv : the preprint server for health sciences.

Wijnbergen D, et al. (2025) Multi-omics analysis in inclusion body myositis identifies mir-16 responsible for HLA overexpression. Orphanet journal of rare diseases, 20(1), 27.

Farashi S, et al. (2025) HTRA1/lncRNA HTRA1-AS1 dominates in age-related macular degeneration reticular pseudodrusen genetic risk with no complement involvement. Nature communications, 16(1), 10854.

Sun W, et al. (2025) TMEM175, SCARB2 and CTSB associations with Parkinson's disease risk across populations. NPJ Parkinson's disease, 11(1), 348.

Godrich D, et al. (2025) Genome-wide association studies of TDP-43 proteinopathy and hippocampal sclerosis reveal shared genetic associations with APOE and TMEM106B. Alzheimer's & dementia : the journal of the Alzheimer's Association, 21(11), e70760.

Haydar S, et al. (2025) A Genome-Wide Association Study of Taste Liking in the Danish Population. The Journal of nutrition, 155(8), 2591.

Zhong Y, et al. (2025) Common and rare variant analyses implicate late-infancy cerebellar development and immune genes in ADHD. Journal of neurodevelopmental disorders, 17(1), 34.

Lee MK, et al. (2025) Genome scan reveals several loci associated with torus palatinus. Orthodontics & craniofacial research, 28(1), 159.

Khani M, et al. (2025) Biobank-scale genetic characterization of Alzheimer's disease and related dementias across diverse ancestries. Nature communications, 16(1), 7554.

Sarnowski C, et al. (2024) Ancestrally diverse genome-wide association analysis highlights ancestry-specific differences in genetic regulation of plasma protein levels. medRxiv : the preprint server for health sciences.

Bayram E, et al. (2024) Genetic analysis of the X chromosome in people with Lewy body dementia nominates new risk loci. NPJ Parkinson's disease, 10(1), 39.

Haukka JK, et al. (2024) Whole-exome and whole-genome sequencing of 1064 individuals with type 1 diabetes reveals novel genes for diabetic kidney disease. Diabetologia, 67(11), 2494.

Tangtanatakul P, et al. (2024) Association of genetic variation on X chromosome with systemic lupus erythematosus in both Thai and Chinese populations. Lupus science & medicine, 11(1).