

Resource Summary Report

Generated by RRID on Aug 5, 2026

HARDY

RRID:SCR_009107

Type: Tool

Proper Citation

HARDY (RRID:SCR_009107)

Resource Information

URL: <http://www.stat.washington.edu/thompson/Genepi/Hardy.shtml>

Proper Citation: HARDY (RRID:SCR_009107)

Description: Markov chain Monte Carlo program for association in two-dimensional contingency tables, and for testing Hardy-Weinberg equilibrium. (entry from Genetic Analysis Software)

Abbreviations: HARDY

Synonyms: PANGAEA

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, c, unix, (dec-unix/..)

Resource Name: HARDY

Resource ID: SCR_009107

Alternate IDs: nlx_154205

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260805T044218+0000

Ratings and Alerts

No rating or validation information has been found for HARDY.

No alerts have been found for HARDY.

Data and Source Information

Usage and Citation Metrics

We found 7329 mentions in open access literature.

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

Le SV, et al. (2026) Signature of Selection Analysis Reveals Candidate Genes Related to Climate Adaptation and Production Traits in Lao Native Goats. *Journal of animal breeding and genetics = Zeitschrift fur Tierzuchtung und Zuchtungsbiologie*, 143(1), 152.

Morales-González E, et al. (2025) Maintenance of genetic diversity in subdivided populations using genomic coancestry matrices. *Molecular ecology resources*, 25(5), e13781.

López A, et al. (2025) Evaluating restriction enzyme selection for reduced representation sequencing in conservation genomics. *Molecular ecology resources*, 25(5), e13865.

Thia JA, et al. (2025) 'Drifting' Buchnera genomes track the microevolutionary trajectories of their aphid hosts. *Insect molecular biology*, 34(1), 19.

Mlambo ZP, et al. (2025) Association of angiogenic factors (placental growth factor and soluble FMS-like tyrosine kinase-1) in preeclamptic women of African ancestry comorbid with HIV infection. *Archives of gynecology and obstetrics*, 311(2), 259.

Li H, et al. (2025) Predictors of growth hormone level on postoperative day one in patients with acromegaly. *Endocrine*, 88(1), 249.

Zavarykina T, et al. (2025) The association of rs25487 of the XRCC1 gene and rs13181 of the ERCC2 gene polymorphisms with the ovarian cancer risk. *Biomolecules & biomedicine*, 25(5), 1197.

He C, et al. (2025) BbGSD: Black-boned Sheep Genome SNP Database. *Database : the journal of biological databases and curation*, 2025.

Sadeqi MB, et al. (2025) The Significant Effects of Threshold Selection for Advancing Nitrogen Use Efficiency in Whole Genome of Bread Wheat. *Plant direct*, 9(1), e70036.

Djordjevic N, et al. (2025) ABCB1 Polymorphism Is Associated with Higher Carbamazepine Clearance in Children. *Pediatric reports*, 17(1).

Lawrence AJ, et al. (2025) Maintenance of Genetic Diversity Despite Population Fluctuations in the Lesser Prairie-Chicken (*Tympanuchus pallidicinctus*). *Ecology and evolution*, 15(1), e70879.

Yao X, et al. (2025) Genomic Landscape and Prediction of Udder Traits in Saanen Dairy Goats. *Animals : an open access journal from MDPI*, 15(2).

Tiuc? RA, et al. (2025) NOS3 Gene Polymorphisms (rs2070744 and rs1799983) and Differentiated Thyroid Cancer: Investigating Associations with Clinical Outcomes.

International journal of molecular sciences, 26(2).

Tang J, et al. (2025) A genealogy-based approach for revealing ancestry-specific structures in admixed populations. *bioRxiv : the preprint server for biology*.

Gálvez-Montosa F, et al. (2025) Polymorphisms within autophagy-related genes as susceptibility biomarkers for pancreatic cancer: A meta-analysis of three large European cohorts and functional characterization. *International journal of cancer*, 156(2), 339.

Goulet D, et al. (2025) Mediation of genetic susceptibility to obesity through eating behaviours in children. *Pediatric obesity*, 20(4), e13180.

Huang SY, et al. (2025) Genome-wide association study unravels mechanisms of brain glymphatic activity. *Nature communications*, 16(1), 626.

Edmunds KJ, et al. (2025) BDNF expression mediates verbal learning and memory in women in a cohort enriched with risk for Alzheimer's disease. *Alzheimer's & dementia (Amsterdam, Netherlands)*, 17(1), e70062.

Alghashmari IH, et al. (2025) Knockdown-resistant mutations in head lice (*Pediculus humanus capitis*) collected from schoolchildren in Riyadh, Saudi Arabia. *Scientific reports*, 15(1), 2412.

Visscher PM, et al. (2025) Heritable polygenic editing: the next frontier in genomic medicine? *Nature*, 637(8046), 637.