

Resource Summary Report

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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

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 5677 mentions in open access literature.

Listed below are recent publications. The full list is available at [FDI Lab - SciCrunch.org](#).

Ban M, et al. (2024) Expression profiling of cerebrospinal fluid identifies dysregulated antiviral mechanisms in multiple sclerosis. *Brain : a journal of neurology*, 147(2), 554.

Zhang X, et al. (2024) Associations of genetic variations in the M3 receptor with salt sensitivity, longitudinal changes in blood pressure and the incidence of hypertension in Chinese adults. *Journal of clinical hypertension (Greenwich, Conn.)*, 26(1), 36.

Versoza CJ, et al. (2024) Novel Insights into the Landscape of Crossover and Noncrossover Events in Rhesus Macaques (*Macaca mulatta*). *Genome biology and evolution*, 16(1).

Saraçaydın G, et al. (2024) Shared genetic etiology between ADHD, task-related behavioral measures and brain activation during response inhibition in a youth ADHD case-control study. *European archives of psychiatry and clinical neuroscience*, 274(1), 45.

Huang Y, et al. (2024) Risk factors associated with age at onset of Parkinson's disease in the UK Biobank. *NPJ Parkinson's disease*, 10(1), 3.

Zhernakova DV, et al. (2024) Host genetic regulation of human gut microbial structural variation. *Nature*, 625(7996), 813.

Dementieva NV, et al. (2024) Risk of Sperm Disorders and Impaired Fertility in Frozen-Thawed Bull Semen: A Genome-Wide Association Study. *Animals : an open access journal from MDPI*, 14(2).

Rivera-Cameras A, et al. (2024) Association of the rs1126616 and rs9138 Variants in the SPP1 Gene among Mexican Patients with Systemic Lupus Erythematosus and Lupus Nephritis. *International journal of molecular sciences*, 25(2).

Chen S, et al. (2024) Cross-Species Comparative DNA Methylation Reveals Novel Insights into Complex Trait Genetics among Cattle, Sheep, and Goats. *Molecular biology and evolution*, 41(2).

Ghoneim ME, et al. (2024) Impact of CD 28, CD86, CTLA-4 and PD-1 genes polymorphisms on acute renal allograft rejection and graft survival among Egyptian recipients. *Scientific reports*, 14(1), 2047.

Gill R, et al. (2024) Genomic heterozygosity is associated with a lower risk of osteoarthritis. *BMC genomics*, 25(1), 85.

Ruiz-Montoya L, et al. (2024) Mass-Rearing Conditions Do Not Always Reduce Genetic Diversity: The Case of the Mexican Fruit Fly, *Anastrepha ludens* (Diptera: Tephritidae). *Insects*, 15(1).

Abecasis D, et al. (2024) Multidisciplinary estimates of connectivity and population structure suggest the use of multiple units for the conservation and management of meagre, *Argyrosomus regius*. *Scientific reports*, 14(1), 873.

Emanuelsson F, et al. (2024) Hyperglycaemia, diabetes and risk of fragility fractures: observational and Mendelian randomisation studies. *Diabetologia*, 67(2), 301.

Larrue R, et al. (2024) Integrating rare genetic variants into DPYD pharmacogenetic testing may help preventing fluoropyrimidine-induced toxicity. *The pharmacogenomics journal*, 24(1), 1.

Ximenez JPB, et al. (2024) Intestinal P-gp activity is reduced in postmenopausal women under breast cancer therapy. *Clinical and translational science*, 17(1), e13713.

Guo J, et al. (2024) Inherited polygenic effects on common hematological traits influence clonal selection on JAK2V617F and the development of myeloproliferative neoplasms. *Nature genetics*, 56(2), 273.

Wang F, et al. (2024) Association of cigarette smoking with cerebrospinal fluid biomarkers of insulin sensitivity and neurodegeneration. *Brain and behavior*, 14(2), e3432.

Zhang W, et al. (2024) Correlations of SDF-1 α and XRCC1 gene polymorphisms with the risk of renal cancer development and bioinformatics studies of SDF-1 α and XRCC1 and the prognosis of renal cancer. *Scientific reports*, 14(1), 3367.

Arunachalam V, et al. (2024) Novel genetic markers for chronic kidney disease in a geographically isolated population of Indigenous Australians: Individual and multiple phenotype genome-wide association study. *Genome medicine*, 16(1), 29.