Resource Summary Report

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LDMatrix

RRID:SCR_017391
Type: Tool

Proper Citation

LDMatrix (RRID:SCR_017391)

Resource Information

URL: https://ldlink.nci.nih.gov/?tab=ldmatrix

Proper Citation: LDMatrix (RRID:SCR_017391)

Description: Software tool to create interactive heatmap matrix of pairwise linkage disequilibrium statistics.

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

Keywords: Interactive, heatmap, matrix, pairwise, linkage, disequilibrium, statistics

Availability: Free, Freely available

Resource Name: LDMatrix

Resource ID: SCR_017391

Record Creation Time: 20220129T080335+0000

Record Last Update: 20260903T235434+0000

Ratings and Alerts

No rating or validation information has been found for LDMatrix.

No alerts have been found for LDMatrix.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 38 mentions in open access literature.

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

Li H, et al. (2026) Population-specific ABO haplotypes reveal distinct venous thromboembolism risk in East Asians: insights from a large-scale genetic study. The Lancet regional health. Western Pacific, 66, 101781.

Lin X, et al. (2025) Lack of association between MALAT1 rs591291 and rs3200401 polymorphisms and venous malformation risk in the Chinese pediatric population. Biochemistry and biophysics reports, 43, 102209.

Del Bosque-Plata L, et al. (2025) LD block disorder-specific pleiotropic roles of novel CRHR1 in type 2 diabetes and depression disorder comorbidity. European archives of psychiatry and clinical neuroscience, 275(4), 1025.

Sangphukieo A, et al. (2024) Human Leukocyte Antigen Markers for Distinguishing Pustular Psoriasis and Adult-Onset Immunodeficiency with Pustular Reaction. Genes, 15(3).

Kilic A, et al. (2024) Vitamin D constrains inflammation by modulating the expression of key genes on Chr17q12-21.1. eLife, 12.

Santo EE, et al. (2023) FOXO3A-short is a novel regulator of non-oxidative glucose metabolism associated with human longevity. Aging cell, 22(3), e13763.

Horimoto ARVR, et al. (2023) Admixture mapping implicates 13q33.3 as ancestry-of-origin locus for Alzheimer disease in Hispanic and Latino populations. HGG advances, 4(3), 100207.

Abondio P, et al. (2023) Apolipoprotein E (APOE) Haplotypes in Healthy Subjects from Worldwide Macroareas: A Population Genetics Perspective for Cardiovascular Disease, Neurodegeneration, and Dementia. Current issues in molecular biology, 45(4), 2817.

Patel A, et al. (2023) MendelianRandomization v0.9.0: updates to an R package for performing Mendelian randomization analyses using summarized data. Wellcome open research, 8, 449.

Chu YJ, et al. (2023) HBV genotype-dependent association of HLA variants with the serodecline of HBsAg in chronic hepatitis B patients. Scientific reports, 13(1), 359.

Jaros RK, et al. (2023) Comorbidity genetic risk and pathways impact SARS-CoV-2 infection outcomes. Scientific reports, 13(1), 9879.

Zheng Y, et al. (2023) Lipocalin 2 - mutation screen and serum levels in patients with anorexia nervosa or obesity and in lean individuals. Frontiers in endocrinology, 14, 1137308.

Alcalá-Santiago Á, et al. (2022) Vitamin D Deficiency and COVID-19: A Biological Database Study on Pathways and Gene-Disease Associations. International journal of molecular sciences, 23(22).

Wang J, et al. (2022) Causality of telomere length associated with calcific aortic valvular stenosis: A Mendelian randomization study. Frontiers in medicine, 9, 1077686.

Ko CL, et al. (2022) Genome-wide association study reveals ethnicity-specific SNPs associated with ankylosing spondylitis in the Taiwanese population. Journal of translational medicine, 20(1), 589.

Wang R, et al. (2022) Genetic variation of interleukin-1 receptor type 1 is associated with severity of COVID-19 disease. The Journal of infection, 84(2), e19.

Su W, et al. (2022) Mendelian Randomization Study on Causal Association of Pyroglutamine with COVID-19. Journal of epidemiology and global health, 12(4), 541.

Cerván-Martín M, et al. (2022) Common Variation in the PIN1 Locus Increases the Genetic Risk to Suffer from Sertoli Cell-Only Syndrome. Journal of personalized medicine, 12(6).

Theofanopoulou C, et al. (2022) Oxytocin and vasotocin receptor variation and the evolution of human prosociality. Comprehensive psychoneuroendocrinology, 11, 100139.

Liu C, et al. (2022) Associations of ATP-Sensitive Potassium Channel's Gene Polymorphisms With Type 2 Diabetes and Related Cardiovascular Phenotypes. Frontiers in cardiovascular medicine, 9, 816847.