Resource Summary Report

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HAPSTAT

RRID:SCR_013382
Type: Tool

Proper Citation

HAPSTAT (RRID:SCR_013382)

Resource Information

URL: http://www.bios.unc.edu/~lin/hapstat/

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Description: Software interface for the statistical analysis of haplotype-disease association. HAPSTAT allows the user to estimate or test haplotype effects and haplotype-environment interactions by maximizing the (observed-data) likelihood that properly accounts for phase uncertainty and study design. The current version considers cross-sectional, case-control and cohort studies. (entry from Genetic Analysis Software)

Abbreviations: HAPSTAT

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, ms-windows, (xp/2000/nt/98)

Funding:

Resource Name: HAPSTAT

Resource ID: SCR_013382

Alternate IDs: nlx_154395

Record Creation Time: 20220129T080315+0000

Record Last Update: 20250416T063641+0000

Ratings and Alerts

No rating or validation information has been found for HAPSTAT.

No alerts have been found for HAPSTAT.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 36 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.

Jang JK, et al. (2024) Clinical implications of genetic polymorphisms in blepharospasm. Experimental and therapeutic medicine, 28(2), 332.

Shi Y, et al. (2023) Lack of association between the VEGFA gene polymorphisms and preterm birth in Korean women. Genomics & informatics, 21(3), e29.

Cho SH, et al. (2021) Genetic Polymorphisms in miR-604A>G, miR-938G>A, miR-1302-3C>T and the Risk of Idiopathic Recurrent Pregnancy Loss. International journal of molecular sciences, 22(11).

Kim IJ, et al. (2021) Association between HOTAIR lncRNA Polymorphisms and Coronary Artery Disease Susceptibility. Journal of personalized medicine, 11(5).

Ryu CS, et al. (2020) MiR-10a, 27a, 34b/c, and 300 Polymorphisms are Associated with Ischemic Stroke Susceptibility and Post-Stroke Mortality. Life (Basel, Switzerland), 10(12).

Park HS, et al. (2020) The Synergistic Effect of Plasminogen Activator Inhibitor-1 (PAI-1) Polymorphisms and Metabolic Syndrome on Coronary Artery Disease in the Korean Population. Journal of personalized medicine, 10(4).

Park HS, et al. (2020) A study of associations between CUBN, HNF1A, and LIPC gene polymorphisms and coronary artery disease. Scientific reports, 10(1), 16294.

Ryu CS, et al. (2020) MPG and NPRL3 Polymorphisms are Associated with Ischemic Stroke Susceptibility and Post-Stroke Mortality. Diagnostics (Basel, Switzerland), 10(11).

Ahn TK, et al. (2020) 3'-UTR Polymorphisms of Vitamin B-Related Genes Are Associated with Osteoporosis and Osteoporotic Vertebral Compression Fractures (OVCFs) in Postmenopausal Women. Genes, 11(6).

An HJ, et al. (2020) Association between Platelet-Specific Collagen Receptor Glycoprotein 6 Gene Variants, Selected Biomarkers, and Recurrent Pregnancy Loss in Korean Women. Genes, 11(8).

Cho HY, et al. (2019) Association of Complement Factor D and H Polymorphisms with Recurrent Pregnancy Loss. International journal of molecular sciences, 21(1).

Park HS, et al. (2019) Association Study between the Polymorphisms of Matrix Metalloproteinase (MMP) Genes and Idiopathic Recurrent Pregnancy Loss. Genes, 10(5).

Park HS, et al. (2019) The microRNApolymorphisms inmiR-150 and miR-1179 are associated with risk of idiopathic recurrent pregnancy loss. Reproductive biomedicine online, 39(2), 187.

Ryu CS, et al. (2019) The association of AGO1 (rs595961G>A, rs636832A>G) and AGO2 (rs11996715C>A, rs2292779C>G, rs4961280C>A) polymorphisms and risk of recurrent implantation failure. Bioscience reports, 39(11).

Ko EJ, et al. (2019) Analysis of the Association Between MicroRNA Biogenesis Gene Polymorphisms and Venous Thromboembolism in Koreans. International journal of molecular sciences, 20(15).

Salehi S, et al. (2018) Lack of Evidence of the Role of APOA5 3'UTR Polymorphisms in Iranian Children and Adolescents with Metabolic Syndrome. Diabetes & metabolism journal, 42(1), 74.

Ahn TK, et al. (2018) 3'-UTR Polymorphisms of MTHFR and TS Associated with Osteoporotic Vertebral Compression Fracture Susceptibility in Postmenopausal Women. International journal of molecular sciences, 19(3).

Park YS, et al. (2017) The Role of RNF213 4810G>A and 4950G>A Variants in Patients with Moyamoya Disease in Korea. International journal of molecular sciences, 18(11).

Rah H, et al. (2017) miR-27a and miR-449b polymorphisms associated with a risk of idiopathic recurrent pregnancy loss. PloS one, 12(5), e0177160.

Kim ES, et al. (2017) MTHFR 3'-untranslated region polymorphisms contribute to recurrent pregnancy loss risk and alterations in peripheral natural killer cell proportions. Clinical and experimental reproductive medicine, 44(3), 152.