Resource Summary Report

Generated by RRID on Aug 5, 2026

HPLUS

RRID:SCR_009239
Type: Tool

Proper Citation

HPLUS (RRID:SCR_009239)

Resource Information

URL: http://cougar.fhcrc.org/hplus/

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Description: An analysis tool for performing haplotype estimation on genetic markers such as SNPs and microsatellites. It is able to handle datasets that include case-control status as well as covariates and marker location variables (such as gene name, chromosome location, etc). (entry from Genetic Analysis Software)

Abbreviations: HPLUS

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, ms-windows, linux

Resource Name: HPLUS

Resource ID: SCR_009239

Alternate IDs: nlx_154400

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260805T044220+0000

Ratings and Alerts

No rating or validation information has been found for HPLUS.

No alerts have been found for HPLUS.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 12 mentions in open access literature.

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

Onyango CO, et al. (2023) Human NCR3 gene variants rs2736191 and rs11575837 alter longitudinal risk for development of pediatric malaria episodes and severe malarial anemia. BMC genomics, 24(1), 542.

Bonella F, et al. (2021) Potential clinical utility of MUC5B und TOLLIP single nucleotide polymorphisms (SNPs) in the management of patients with IPF. Orphanet journal of rare diseases, 16(1), 111.

Nourkami-Tutdibi N, et al. (2021) Genetic Association With Pseudomonas aeruginosa Acquisition in Cystic Fibrosis: Influence of Surfactant Protein D and Mannose-Binding Lectin. Frontiers in immunology, 12, 587313.

Gannoun MBA, et al. (2021) MMP-2 and MMP-9 Polymorphisms and Preeclampsia Risk in Tunisian Arabs: A Case-Control Study. Journal of clinical medicine, 10(12).

Kisia LE, et al. (2019) Genetic variation in interleukin-7 is associated with a reduced erythropoietic response in Kenyan children infected with Plasmodium falciparum. BMC medical genetics, 20(1), 140.

Munde EO, et al. (2017) Haplotype of non-synonymous mutations within IL-23R is associated with susceptibility to severe malaria anemia in a P. falciparum holoendemic transmission area of Kenya. BMC infectious diseases, 17(1), 291.

Zidi S, et al. (2015) Relationship of common vascular endothelial growth factor polymorphisms and haplotypes with the risk of cervical cancer in Tunisians. Cytokine, 74(1), 108.

Biros E, et al. (2015) Association between the Advanced Glycosylation End Product-Specific Receptor Gene and Cardiovascular Death in Older Men. PloS one, 10(7), e0134475.

Ben Ali Gannoun M, et al. (2015) Association of common eNOS/NOS3 polymorphisms with preeclampsia in Tunisian Arabs. Gene, 569(2), 303.

Ide S, et al. (2014) Haplotypes of P2RX7 gene polymorphisms are associated with both cold pain sensitivity and analgesic effect of fentanyl. Molecular pain, 10, 75.

Harms KC, et al. (2013) Association of TNF-?? polymorphism rs1800629 with multisomatoform disorder in a group of German patients and healthy controls: an explorative study. Cytokine, 61(2), 389.

Hildebrand F, et al. (2009) No association between CALCA polymorphisms and clinical outcome or serum procalcitonin levels in German polytrauma patients. Cytokine, 47(1), 30.