Resource Summary Report

Generated by NIF on Aug 20, 2026

FINETTI

RRID:SCR_009179
Type: Tool

Proper Citation

FINETTI (RRID:SCR_009179)

Resource Information

URL: http://ihg.gsf.de/cgi-bin/hw/hwa1.pl (testing part)

Proper Citation: FINETTI (RRID:SCR_009179)

Description: Software application that tests for deviation from Hardy-Weinberg equilibrium and tests for association in case controls studies; Plot genotype frequencies graphically using a de Finetti diagram. (entry from Genetic Analysis Software)

Abbreviations: FINETTI

Synonyms: de FINETTI generator

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, pascal, (web implementation: perl, c, php), web-based, ms-dos, ms-windows, (32), linux, (for stand-alone version)

Resource Name: FINETTI

Resource ID: SCR_009179

Alternate IDs: nlx_154316

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260818T043337+0000

Ratings and Alerts

No rating or validation information has been found for FINETTI.

No alerts have been found for FINETTI.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 299 mentions in open access literature.

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

Liu X, et al. (2025) Investigation of high-mobility group box 1 variants with lymph node status and colorectal cancer risk. World journal of gastrointestinal oncology, 17(4), 102584.

Cadena-Sandoval D, et al. (2024) Interactions between TNFAIP3, PTPN22, and TRAF1-C5 gene polymorphisms in patients with primary Sjögren's syndrome. Archives of rheumatology, 39(1), 60.

Palomba NP, et al. (2023) Common and Rare Variants in TMEM175 Gene Concur to the Pathogenesis of Parkinson's Disease in Italian Patients. Molecular neurobiology, 60(4), 2150.

Toledo-Lozano CG, et al. (2023) Individual and Combined Effect of MAO-A/MAO-B Gene Variants and Adverse Childhood Experiences on the Severity of Major Depressive Disorder. Behavioral sciences (Basel, Switzerland), 13(10).

García-Martín E, et al. (2023) Vitamin D receptor and binding protein genes variants in patients with migraine. Annals of clinical and translational neurology, 10(10), 1824.

Aljarba NH, et al. (2023) Association between interleukin-27 gene polymorphisms and Plasmodium falciparum Malaria. Innate immunity, 29(5), 83.

Strauss E, et al. (2023) Hypoxia-Inducible Pathway Polymorphisms and Their Role in the Complications of Prematurity. Genes, 14(5).

Boutros A, et al. (2023) The predictive and prognostic role of single nucleotide gene variants of PD-1 and PD-L1 in patients with advanced melanoma treated with PD-1 inhibitors. Immuno-oncology technology, 20, 100408.

Alzeer HS, et al. (2023) Genetic Variants of HOTAIR Associated with Colorectal Cancer: A Case-Control Study in the Saudi Population. Genes, 14(3).

Daghestani M, et al. (2023) Single Nucleotide Polymorphisms Associated with Rheumatoid Arthritis in Saudi Patients. Journal of clinical medicine, 12(15).

Liu K, et al. (2023) Association study of WNK1 genetic variants and essential hypertension risk in the Northern Han Chinese in Beijing. Frontiers in genetics, 14, 1234536.

García-Martín E, et al. (2023) Lack of Association between Common LAG3/CD4 Variants and Risk of Migraine. International journal of molecular sciences, 24(2).

Ampuero S, et al. (2022) IL-7/IL7R axis dysfunction in adults with severe community-acquired pneumonia (CAP): a cross-sectional study. Scientific reports, 12(1), 13145.

Zecca C, et al. (2022) Clinic and genetic predictors in response to erenumab. European journal of neurology, 29(4), 1209.

Raina JK, et al. (2022) "Association of MTHFR and MS/MTR gene polymorphisms with congenital heart defects in North Indian population (Jammu and Kashmir): a case-control study encompassing meta-analysis and trial sequential analysis". BMC pediatrics, 22(1), 223.

García-Martín E, et al. (2022) Association between LAG3/CD4 gene variants and risk of Parkinson's disease. European journal of clinical investigation, 52(11), e13847.

Meza G, et al. (2022) IFNL4 genotype influences the rate of HIV-1 seroconversion in men who have sex with men. Virulence, 13(1), 757.

Refisch A, et al. (2022) Analysis of CACNA1C and KCNH2 Risk Variants on Cardiac Autonomic Function in Patients with Schizophrenia. Genes, 13(11).

Liu C, et al. (2022) Association between miR-146a rs2910164, miR-196a2 rs11614913, and miR-499 rs3746444 polymorphisms and the risk of esophageal carcinoma: A case-control study. Cancer medicine, 11(21), 3949.

Gabriel-Medina P, et al. (2022) Influence of Type 2 Diabetes in the Association of PNPLA3 rs738409 and TM6SF2 rs58542926 Polymorphisms in NASH Advanced Liver Fibrosis. Biomedicines, 10(5).