Resource Summary Report

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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)

Funding:

Resource Name: FINETTI

Resource ID: SCR_009179

Alternate IDs: nlx_154316

Record Creation Time: 20220129T080251+0000

Record Last Update: 20250416T063539+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 283 mentions in open access literature.

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

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).

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

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.

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.

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.

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.

Aziz MA, et al. (2022) Association of ACE1 I/D rs1799752 and ACE2 rs2285666 polymorphisms with the infection and severity of COVID-19: A meta-analysis. Molecular genetics & genomic medicine, 10(11), e2063.

Hernández-Díaz Y, et al. (2021) Association between polymorphisms of FKBP5 gene and suicide attempt in a Mexican population: A case-control study. Brain research bulletin, 166, 37.

Pabalan N, et al. (2021) Ethnic and age-specific acute lung injury/acute respiratory distress syndrome risk associated with angiotensin-converting enzyme insertion/deletion polymorphisms, implications for COVID-19: A meta-analysis. Infection, genetics and evolution : journal of molecular epidemiology and evolutionary genetics in infectious diseases, 88, 104682.

Liu X, et al. (2021) Association of UCP1 and UCP2 variants with diabetic retinopathy susceptibility in type-2 diabetes mellitus patients: a meta-analysis. BMC ophthalmology, 21(1), 81.

Ying D, et al. (2021) Association Between Macrophage Migration Inhibitory Factor -173 G>C Gene Polymorphism and Childhood Idiopathic Nephrotic Syndrome: A Meta-Analysis. Frontiers in pediatrics, 9, 724258.

Janik MK, et al. (2021) MARC1 p.A165T variant is associated with decreased markers of liver injury and enhanced antioxidant capacity in autoimmune hepatitis. Scientific reports, 11(1), 24407.

Serrano-Rísquez C, et al. (2021) CD46 Genetic Variability and HIV-1 Infection Susceptibility. Cells, 10(11).

Pabalan N, et al. (2021) Associations of CB1 cannabinoid receptor (CNR1) gene polymorphisms with risk for alcohol dependence: Evidence from meta-analyses of genetic and genome-wide association studies. Medicine, 100(43), e27343.