Resource Summary Report

Generated by RRID on Aug 2, 2026

SNPedia

RRID:SCR_006125
Type: Tool

Proper Citation

SNPedia (RRID:SCR_006125)

Resource Information

URL: http://www.snpedia.com/index.php/SNPedia

Proper Citation: SNPedia (RRID:SCR_006125)

Description: Wiki investigating human genetics including information about the effects of variations in DNA, citing peer-reviewed scientific publications. It is used by Promethease to analyze and help explain your DNA. It is based on a wiki model in order to foster communication about genetic variation and to allow interested community members to help it evolve to become ever more relevant. As the cost of genotyping (and especially of fully determining your own genomic sequence) continues to drop, we""ll all want to know more - a lot more - about the meaning of these DNA variations and SNPedia will be here to help. SNPedia has been launched to help realize the potential of the Human Genome Project to connect to our daily lives and well-being. For more information see the Wikipedia page, http://en.wikipedia.org/wiki/SNPedia * Download URL: http://www.SNPedia.com/index.php/Bulk * Web Service URL: http://bots.SNPedia.com/api.php

Abbreviations: SNPedia

Resource Type: data or information resource, database

Defining Citation: PMID:22140107

Keywords: dna, genetics, gene, genome, genoset, genotype, medicine, medical condition, genetic variation, dna, genetic variation, genomics, single nucleotide polymorphism, medical association, phenotypic association, genealogical association, variation, genome annotation, phenotype, web service, bio.tools, FASEB list

Availability: Creative Commons Attribution-NonCommercial-ShareAlike License, v3

Resource Name: SNPedia

Resource ID: SCR_006125

Alternate IDs: biotools:snpedia, grid.465250.0, nlx_151604

Alternate URLs: https://ror.org/0253rdk33, https://bio.tools/snpedia

Record Creation Time: 20220129T080234+0000

Record Last Update: 20260801T190936+0000

Ratings and Alerts

No rating or validation information has been found for SNPedia.

No alerts have been found for SNPedia.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 62 mentions in open access literature.

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

Chen H, et al. (2026) rs2032582 polymorphism of ATP binding cassette subfamily B member 1 affect the susceptibility of inflammatory bowel disease: a systematical meta-analysis based on 14,876 subjects. International journal of colorectal disease, 41(1), 6.

Menon S, et al. (2025) Meta-analysis and in-silico functional characterization of the SNCA variant rs356220 in Parkinson's disease. Scientific reports, 15(1), 23358.

Van Schalkwyk M, et al. (2025) Pharmacokinetics of first-line tuberculosis drugs rifampin, isoniazid, ethambutol, and pyrazinamide during pregnancy and postpartum with and without efavirenz-based antiretroviral treatment: IMPAACT P1026s study. Antimicrobial agents and chemotherapy, 69(9), e0005225.

Pardo-Seco J, et al. (2025) Whole-Genome Sequencing in Galicia Reveals Male-Biased Pre-Islamic North African Ancestry, Subtle Population Structure, and Microgeographic Patterns of Disease Risk. FASEB journal : official publication of the Federation of American Societies for Experimental Biology, 39(24), e71253.

Fontani F, et al. (2025) Archaeogenetics reconstructs demography and extreme parental consanguinity in a Bronze Age community from Southern Italy. Communications biology, 8(1), 1766.

Nordfors U, et al. (2025) Archaeogenetics reveals fine-scale genetic continuity and patterns of kinship and health in medieval Finland. iScience, 28(8), 113086.

Alem M, et al. (2025) Immuno-informatics voyage through molecular mimicry of Heat Shock Proteins: Potential IBD immunopathogenesis. PloS one, 20(10), e0333618.

Perini JA, et al. (2024) Single-Nucleotide Polymorphisms Associated with Mercury Levels and Neurological Symptoms: An Overview. Toxics, 12(3).

Coad?? CA, et al. (2024) Association of Glycoprotein IIIa PlA1/A2 Polymorphism with Risk of Stroke: Updated Meta-Analysis. Current issues in molecular biology, 46(6), 5364.

Pilli E, et al. (2024) Ancient DNA challenges prevailing interpretations of the Pompeii plaster casts. Current biology : CB, 34(22), 5307.

Shade LMP, et al. (2024) GWAS of multiple neuropathology endophenotypes identifies new risk loci and provides insights into the genetic risk of dementia. Nature genetics, 56(11), 2407.

Mulero-Hern??ndez J, et al. (2024) Integration of chromosome locations and functional aspects of enhancers and topologically associating domains in knowledge graphs enables versatile queries about gene regulation. Nucleic acids research, 52(15), e69.

Amin USM, et al. (2023) Type 2 diabetes linked FTO gene variant rs8050136 is significantly associated with gravidity in gestational diabetes in a sample of Bangladeshi women: Meta-analysis and case-control study. PloS one, 18(11), e0288318.

Chaudhary H, et al. (2023) Association of FTO gene variant rs9939609 with polycystic ovary syndrome from Gujarat, India. BMC medical genomics, 16(1), 216.

Shanazarov N, et al. (2023) Association of Gene Polymorphisms with Breast Cancer Risk in the Kazakh Population. Asian Pacific journal of cancer prevention : APJCP, 24(12), 4195.

Vidic Z, et al. (2023) Association of OPRM1, MIR23B, and MIR107 genetic variability with acute pain, chronic pain and adverse effects after postoperative tramadol and paracetamol treatment in breast cancer. Radiology and oncology, 57(1), 111.

Pilo J, et al. (2023) 8-Oxoguanine DNA Glycosylase 1 Upregulation as a Risk Factor for Obesity and Colorectal Cancer. International journal of molecular sciences, 24(6).

Szakats S, et al. (2023) Sex-biased gene and microRNA expression in the developing mouse brain is associated with neurodevelopmental functions and neurological phenotypes. Biology of sex differences, 14(1), 57.

Kruszewski M, et al. (2022) Association of Myostatin Gene Polymorphisms with Strength and Muscle Mass in Athletes: A Systematic Review and Meta-Analysis of the MSTN rs1805086 Mutation. Genes, 13(11).

Atzeni R, et al. (2022) VariantAlert: A web-based tool to notify updates in genetic variant annotations. Human mutation, 43(12), 1808.