Resource Summary Report

Generated by SPARC SAWG on Sep 17, 2026

SNPALYZE

RRID:SCR_009401
Type: Tool

Proper Citation

SNPALYZE (RRID:SCR_009401)

Resource Information

URL: https://www.dynacom.co.jp/english/product/snpalyze_e/

Proper Citation: SNPALYZE (RRID:SCR_009401)

Description: Software application (entry from Genetic Analysis Software)

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, ms-windows, (98/me/nt4.0/2000/xp)

Resource Name: SNPALYZE

Resource ID: SCR_009401

Alternate IDs: nlx_154640

Record Creation Time: 20220129T080252+0000

Record Last Update: 20260912T200246+0000

Ratings and Alerts

No rating or validation information has been found for SNPALYZE.

No alerts have been found for SNPALYZE.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 78 mentions in open access literature.

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

Bayaraa O, et al. (2026) The synergistic interaction between ACE and TMPRSS2 polymorphisms increases the risk of severe COVID-19. PloS one, 21(2), e0343590.

Omi T, et al. (2026) Identification of SORCS1 as a candidate gene associated with canine behavioral traits: Insights from guide dog training outcomes. PloS one, 21(2), e0342346.

Atefrad S, et al. (2025) The association between NLGN4 gene variants and the incidence of autism spectrum disorders in Guilan, Iran. IBRO neuroscience reports, 18, 306.

Altankhuyag A, et al. (2024) The interactions between ARMS2, CFH, VEGF-A and environmental factors on the risk of age-related macular degeneration. Molecular vision, 30, 320.

Yamaguchi N, et al. (2024) Autophagy-Related Gene ATG7 Polymorphism Could Potentially Serve as a Biomarker of the Progression of Atrophic Gastritis. Journal of clinical medicine, 13(2).

Chen R, et al. (2024) Association of IL13 polymorphisms with susceptibility to myocardial infarction: A case-control study in Chinese population. PloS one, 19(8), e0308081.

Amano S, et al. (2023) Prediction and association analyses of skin phenotypes in Japanese females using genetic, environmental, and physical features. Skin research and technology : official journal of International Society for Bioengineering and the Skin (ISBS) [and] International Society for Digital Imaging of Skin (ISDIS) [and] International Society for Skin Imaging (ISSI), 29(1), e13231.

Wang D, et al. (2022) Chemerin levels and its genetic variants are associated with Gestational diabetes mellitus: A hospital-based study in a Chinese cohort. Gene, 807, 145888.

Soflaei SS, et al. (2022) Association of Paraoxonase-1 Genotype and Phenotype with Angiogram Positive Coronary Artery Disease. Arquivos brasileiros de cardiologia, 119(4), 593.

Park JW, et al. (2022) Gene Dose-Dependent and Additive Effects of ABCG2 rs2231142 and SLC2A9 rs3733591 Genetic Polymorphisms on Serum Uric Acid Levels. Metabolites, 12(12).

Jeon YJ, et al. (2021) 3'-UTR Polymorphisms in Thymidylate Synthase with Colorectal Cancer Prevalence and Prognosis. Journal of personalized medicine, 11(6).

Kondyarpu A, et al. (2021) Association of ISL1 polymorphisms and eosinophilic levels among otitis media patients. Journal of clinical laboratory analysis, 35(3), e23702.

Seo JE, et al. (2021) Association Between CLOCK Gene Variants and Restless Legs Syndrome in Koreans. Psychiatry investigation, 18(11), 1125.

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.

Fukunaga K, et al. (2021) Functional Characterization of the Effects of N-acetyltransferase 2 Alleles on N-acetylation of Eight Drugs and Worldwide Distribution of Substrate-Specific Diversity. Frontiers in genetics, 12, 652704.

Ohwaki A, et al. (2020) Placental Genetic Variants in the Upstream Region of the FLT1 Gene in Pre-eclampsia. Journal of reproduction & infertility, 21(4), 240.

Oh J, et al. (2020) Association between Five Common Plasminogen Activator Inhibitor-1 (PAI-1) Gene Polymorphisms and Colorectal Cancer Susceptibility. International journal of molecular sciences, 21(12).

Minn AKK, et al. (2020) Association study of long non-coding RNA HOTAIR rs920778 polymorphism with the risk of cancer in an elderly Japanese population. Gene, 729, 144263.

Kalantari S, et al. (2019) Single and multi-locus association study of TCF7L2 gene variants with susceptibility to type 2 diabetes mellitus in an Iranian population. Gene, 696, 88.

Matsusue A, et al. (2019) VNTR polymorphism in the monoamine oxidase A promoter region and cerebrospinal fluid catecholamine concentrations in forensic autopsy cases. Neuroscience letters, 701, 71.