

Resource Summary Report

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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 resource, software application

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: 20260801T191100+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 76 mentions in open access literature.

Listed below are recent publications. The full list is available at [ASWG](#) .

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.

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).

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.

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.

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).

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.

Kondyarpur 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.

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

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.

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.

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).

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.

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.

Kobayashi H, et al. (2019) Adiponectin Receptor gene Polymorphisms are Associated with Kidney Function in Elderly Japanese Populations. *Journal of atherosclerosis and thrombosis*, 26(4), 328.

An HJ, et al. (2019) 3'-UTR Polymorphisms in the Vascular Endothelial Growth Factor Gene (VEGF) Contribute to Susceptibility to Recurrent Pregnancy Loss (RPL). *International journal of molecular sciences*, 20(13).