

# Resource Summary Report

Generated by [RRID](#) on Aug 5, 2026

## TAGSNP

RRID:SCR\_008623

Type: Tool

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### Proper Citation

TAGSNP (RRID:SCR\_008623)

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### Resource Information

**URL:** <http://www-rcf.usc.edu/~stram/tagSNPs.html>

**Proper Citation:** TAGSNP (RRID:SCR\_008623)

**Description:** Software application (entry from Genetic Analysis Software)

**Abbreviations:** TAGSNP

**Resource Type:** software application, software resource

**Keywords:** gene, genetic, genomic, fortran90

**Resource Name:** TAGSNP

**Resource ID:** SCR\_008623

**Alternate IDs:** nlx\_154671

**Record Creation Time:** 20220129T080248+0000

**Record Last Update:** 20260805T044211+0000

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### Ratings and Alerts

No rating or validation information has been found for TAGSNP.

No alerts have been found for TAGSNP.

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### Data and Source Information

**Source:** [SciCrunch Registry](#)

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### Usage and Citation Metrics

We found 69 mentions in open access literature.

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

Choi SH, et al. (2025) AKT1 Functional Polymorphism (rs2494750) Influences Clinical Outcomes in Patients With Advanced Lung Cancer Treated With EGFR Inhibitors. Journal of Korean medical science, 40(31), e184.

Adjemout M, et al. (2024) From Genome-wide Association Studies to Functional Variants: ARL14 Cis-regulatory Variants Are Associated With Severe Malaria. The Journal of infectious diseases, 230(3), e743.

Shilenok I, et al. (2024) C11orf58 (Hero20) Gene Polymorphism: Contribution to Ischemic Stroke Risk and Interactions with Other Heat-Resistant Obscure Chaperones. Biomedicines, 12(11).

Cheng S, et al. (2024) Harnessing landrace diversity empowers wheat breeding. Nature, 632(8026), 823.

Hebbar P, et al. (2023) Linkage analysis using whole exome sequencing data implicates SLC17A1, SLC17A3, TATDN2 and TMEM131L in type 1 diabetes in Kuwaiti families. Scientific reports, 13(1), 14978.

Lee S, et al. (2023) Genetic variants of NEUROD1 target genes are associated with clinical outcomes of small-cell lung cancer patients. Thoracic cancer, 14(13), 1145.

Lee JH, et al. (2023) Genetic variants in key necroptosis regulators predict prognosis of non-small cell lung cancer after surgical resection. Thoracic cancer, 14(26), 2678.

Lee JH, et al. (2022) Epigenetic readers and lung cancer: the rs2427964C>T variant of the bromodomain and extraterminal domain gene BRD3 is associated with poorer survival outcome in NSCLC. Molecular oncology, 16(3), 750.

Choi SH, et al. (2022) Genetic variants in LKB1/AMPK/mTOR pathway are associated with clinical outcomes of chemotherapy in non-small cell lung cancer. Thoracic cancer, 13(23), 3322.

Warburton ML, et al. (2022) Comparative Analysis of Multiple GWAS Results Identifies Metabolic Pathways Associated with Resistance to A. flavus Infection and Aflatoxin Accumulation in Maize. Toxins, 14(11).

Lee JH, et al. (2021) Impact of immune checkpoint gene CD155 Ala67Thr and CD226 Gly307Ser polymorphisms on small cell lung cancer clinical outcome. Scientific reports, 11(1), 1794.

Wittenburg D, et al. (2021) Grouping of genomic markers in populations with family structure. BMC bioinformatics, 22(1), 79.

Parra-Galindo MA, et al. (2021) Pathway-based analysis of anthocyanin diversity in diploid potato. PloS one, 16(4), e0250861.

Benna C, et al. (2021) Gender-specific associations between polymorphisms of the circadian gene RORA and cutaneous melanoma susceptibility. *Journal of translational medicine*, 19(1), 57.

Zhou W, et al. (2021) A case-control study on correlation between the single nucleotide polymorphism of CLEC4E and the susceptibility to tuberculosis among Han people in Western China. *BMC infectious diseases*, 21(1), 788.

Kang MK, et al. (2021) Prognostic significance of genetic variants in GLUT1 in stage III non-small cell lung cancer treated with radiotherapy. *Thoracic cancer*, 12(6), 874.

Mohr S, et al. (2021) The alternative serotonin transporter promoter P2 impacts gene function in females with irritable bowel syndrome. *Journal of cellular and molecular medicine*, 25(16), 8047.

Volz NB, et al. (2020) Polymorphisms within Immune Regulatory Pathways Predict Cetuximab Efficacy and Survival in Metastatic Colorectal Cancer Patients. *Cancers*, 12(10).

Guo K, et al. (2020) Bone morphogenetic protein 9, and its genetic variants contribute to susceptibility of idiopathic pulmonary arterial hypertension. *Aging*, 12(3), 2123.

Li Z, et al. (2020) Association of Genetic and Environmental Factors with Non-Alcoholic Fatty Liver Disease in a Chinese Han Population. *International journal of environmental research and public health*, 17(14).