

Resource Summary Report

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TAGGER

RRID:SCR_009419

Type: Tool

Proper Citation

TAGGER (RRID:SCR_009419)

Resource Information

URL: <http://archive.broadinstitute.org/mpg/tagger/>

Proper Citation: TAGGER (RRID:SCR_009419)

Description: Software application (entry from Genetic Analysis Software)

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, web-based

Resource Name: TAGGER

Resource ID: SCR_009419

Alternate IDs: nlx_154669

Record Creation Time: 20220129T080252+0000

Record Last Update: 20260905T133252+0000

Ratings and Alerts

No rating or validation information has been found for TAGGER.

No alerts have been found for TAGGER.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 93 mentions in open access literature.

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

Bonacina G, et al. (2025) Protocol for low-level laser therapy (LLLT) to control post-endodontic treatment pain in patients with apical periodontitis: a randomized controlled trial. *BMC oral health*, 25(1), 1182.

Araida J, et al. (2024) rs12411980 single-nucleotide polymorphism related to PRTFDC1 expression is significantly associated with phantom tooth pain. *Molecular pain*, 20, 17448069241272215.

Camilo NG, et al. (2023) Influence of Chitosan 0.2% in Various Final Cleaning Methods on the Bond Strength of Fiberglass Post to Intrarradicular Dentin. *Polymers*, 15(22).

Morii M, et al. (2023) The rs216009 single-nucleotide polymorphism of the CACNA1C gene is associated with phantom tooth pain. *Molecular pain*, 19, 17448069231193383.

Al Ali L, et al. (2023) Fetuin-A and its genetic association with cardiometabolic disease. *Scientific reports*, 13(1), 21469.

Grissa D, et al. (2022) Diseases 2.0: a weekly updated database of disease-gene associations from text mining and data integration. *Database : the journal of biological databases and curation*, 2022.

Isayeva U, et al. (2022) Exploring the association between brain-derived neurotrophic factor levels and longitudinal psychopathological and cognitive changes in Sardinian psychotic patients. *European psychiatry : the journal of the Association of European Psychiatrists*, 65(1), e71.

Zhang Y, et al. (2022) Gene-environment interactions between CREB1 and childhood maltreatment on aggression among male Chinese adolescents. *Scientific reports*, 12(1), 1326.

Hawkins NT, et al. (2022) Systematic tissue annotations of genomics samples by modeling unstructured metadata. *Nature communications*, 13(1), 6736.

Djordjevic A, et al. (2022) Tag Variants of LGALS-3 Containing Haplotype Block in Advanced Carotid Atherosclerosis. *Journal of stroke and cerebrovascular diseases : the official journal of National Stroke Association*, 31(1), 106212.

Xu ZM, et al. (2022) Using population-specific add-on polymorphisms to improve genotype imputation in underrepresented populations. *PLoS computational biology*, 18(1), e1009628.

Soeda M, et al. (2022) Single-nucleotide polymorphisms of the SLC17A9 and P2RY12 genes are significantly associated with phantom tooth pain. *Molecular pain*, 18, 17448069221089592.

Schroor MM, et al. (2021) Associations between SNPs in Intestinal Cholesterol Absorption and Endogenous Cholesterol Synthesis Genes with Cholesterol Metabolism. *Biomedicines*, 9(10).

Wang Y, et al. (2021) LRRC3B Polymorphisms Contributed to Breast Cancer Susceptibility in Chinese Han Population. *Frontiers in oncology*, 11, 657168.

Wang Z, et al. (2021) Research Note: Fine mapping of sequence variants associated with body weight of Lueyang black-boned chicken in the CCKAR gene. *Poultry science*, 100(11), 101448.

He J, et al. (2021) The Effects of WISP1 Polymorphisms on the Prognosis of Lung Cancer Patients with Platinum-Based Chemotherapy. *Pharmacogenomics and personalized medicine*, 14, 1193.

Xu S, et al. (2021) Association of FOXO3a gene polymorphisms and ankylosing spondylitis susceptibility in Eastern Chinese Han population. *Gene*, 800, 145832.

Hartman T, et al. (2020) Customization scenarios for de-identification of clinical notes. *BMC medical informatics and decision making*, 20(1), 14.

Mahmuda NA, et al. (2020) One Single Nucleotide Polymorphism of the TRPM2 Channel Gene Identified as a Risk Factor in Bipolar Disorder Associates with Autism Spectrum Disorder in a Japanese Population. *Diseases (Basel, Switzerland)*, 8(1).

Castro-Santos P, et al. (2020) Association analysis in a Latin American population revealed ethnic differences in rheumatoid arthritis-associated SNPs in Caucasian and Asian populations. *Scientific reports*, 10(1), 7879.