Resource Summary Report

Generated by NIF on Apr 19, 2025

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

Funding:

Resource Name: TAGGER

Resource ID: SCR_009419

Alternate IDs: nlx_154669

Record Creation Time: 20220129T080252+0000

Record Last Update: 20250419T055209+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 90 mentions in open access literature.

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

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).

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

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

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.

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.

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.

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.

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

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.

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.

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

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.

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

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.

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).

Pan L, et al. (2020) CMIP SNPs and their haplotypes are associated with dyslipidaemia and clinicopathologic features of IgA nephropathy. Bioscience reports, 40(10).