

Resource Summary Report

Generated by SPARC SAWG on Aug 6, 2026

GASSOC

RRID:SCR_013136

Type: Tool

Proper Citation

GASSOC (RRID:SCR_013136)

Resource Information

URL: http://mayoresearch.mayo.edu/mayo/research/schaid_lab/software.cfm

Proper Citation: GASSOC (RRID:SCR_013136)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on May 24,2023. Software application for statistical methods for disease and genetic marker associations using cases and their parents. These methods include an extension of the transmission/disequilibrium test (TDT) for multiple marker alleles, as well as additional general tests sensitive to associations that depend on dominant or recessive genetic mechanisms. (entry from Genetic Analysis Software)

Synonyms: Genetic ASSOCiation analysis software for cases and parent

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, c, unix, sunos, solaris

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: GASSOC

Resource ID: SCR_013136

Alternate IDs: nlx_154077

Record Creation Time: 20220129T080314+0000

Record Last Update: 20260805T044308+0000

Ratings and Alerts

No rating or validation information has been found for GASSOC.

No alerts have been found for GASSOC.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 13 mentions in open access literature.

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

Wakil SM, et al. (2014) A new susceptibility locus for myocardial infarction, hypertension, type 2 diabetes mellitus, and dyslipidemia on chromosome 12q24. Disease markers, 2014, 291419.

Zhang Z, et al. (2014) Functional genetic variants of TNFSF15 and their association with gastric adenocarcinoma: a case-control study. PloS one, 9(9), e108321.

Al-Najai M, et al. (2013) Association of the angiotensinogen gene polymorphism with atherosclerosis and its risk traits in the Saudi population. BMC cardiovascular disorders, 13, 17.

Brighina L, et al. (2013) Analysis of vesicular monoamine transporter 2 polymorphisms in Parkinson's disease. Neurobiology of aging, 34(6), 1712.e9.

Alshahid M, et al. (2013) New susceptibility locus for obesity and dyslipidaemia on chromosome 3q22.3. Human genomics, 7(1), 15.

Hodži? A, et al. (2013) Genetic variation in circadian rhythm genes CLOCK and ARNTL as risk factor for male infertility. PloS one, 8(3), e59220.

Muiya N, et al. (2013) The 3'-UTR of the adiponectin Q gene harbours susceptibility loci for atherosclerosis and its metabolic risk traits. BMC medical genetics, 14, 127.

Razzaghi H, et al. (2012) Population-Based Resequencing of LIPG and ZNF202 Genes in Subjects with Extreme HDL Levels. Frontiers in genetics, 3, 89.

Beebe-Dimmer JL, et al. (2012) Genetic variation in glutathione S-transferase omega-1, arsenic methyltransferase and methylene-tetrahydrofolate reductase, arsenic exposure and bladder cancer: a case-control study. Environmental health : a global access science source, 11, 43.

Zhang X, et al. (2012) Association of single nucleotide polymorphisms in TCF2 with type 2 diabetes susceptibility in a Han Chinese population. PloS one, 7(12), e52938.

Guo X, et al. (2011) Mucin variable number tandem repeat polymorphisms and severity of cystic fibrosis lung disease: significant association with MUC5AC. PloS one, 6(10), e25452.

Del Greco M F, et al. (2011) Genome-wide association analysis and fine mapping of NT-proBNP level provide novel insight into the role of the MTHFR-CLCN6-NPPA-NPPB gene cluster. Human molecular genetics, 20(8), 1660.

Wang SS, et al. (2010) Common genetic variants and risk for HPV persistence and progression to cervical cancer. PloS one, 5(1), e8667.