

Resource Summary Report

Generated by T1D on Aug 5, 2026

metagen

RRID:SCR_003443

Type: Tool

Proper Citation

metagen (RRID:SCR_003443)

Resource Information

URL: <http://www.compgen.org/tools/metagen>

Proper Citation: metagen (RRID:SCR_003443)

Description: Software program providing a method for meta-analysis of case-control genetic association studies using random-effects logistic regression.

Abbreviations: metagen

Resource Type: software resource

Defining Citation: [PMID:17605724](#)

Keywords: bio.tools

Availability: Free, Available for download, Freely available

Resource Name: metagen

Resource ID: SCR_003443

Alternate IDs: OMICS_00238, biotools:metagen

Alternate URLs: <https://www.rdocumentation.org/packages/meta/versions/4.9-6/topics/metagen>

Record Creation Time: 20220129T080219+0000

Record Last Update: 20260801T190221+0000

Ratings and Alerts

No rating or validation information has been found for metagen.

No alerts have been found for metagen.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 18 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#).

Muratspahi? E, et al. (2025) De novo design of miniprotein agonists and antagonists targeting G protein-coupled receptors. bioRxiv : the preprint server for biology.

Liu L, et al. (2024) Genome-wide studies define new genetic mechanisms of IgA vasculitis. medRxiv : the preprint server for health sciences.

Veloso FCS, et al. (2024) Neonatal death prediction scores: a systematic review and meta-analysis. BMJ paediatrics open, 8(1).

Crucianelli S, et al. (2024) Effects of sulphur thermal water inhalations in long-COVID syndrome: Spa-centred, double-blinded, randomised case-control pilot study. Clinical medicine (London, England), 24(6), 100251.

Aida M, et al. (2024) Heyndrickxia coagulans strain SANK70258 suppresses symptoms of upper respiratory tract infection via immune modulation: a randomized, double-blind, placebo-controlled, parallel-group, comparative study. Frontiers in immunology, 15, 1389920.

Kawamata T, et al. (2023) Natto consumption suppresses atherosclerotic plaque progression in LDL receptor-deficient mice transplanted with iRFP-expressing hematopoietic cells. Scientific reports, 13(1), 22469.

Zharkova EK, et al. (2023) Bacterial Communities of Lamiacea L. Medicinal Plants: Structural Features and Rhizosphere Effect. Microorganisms, 11(1).

de Lavallaz JDF, et al. (2023) Risk factors for the development of premature ventricular complex-induced cardiomyopathy: a systematic review and meta-analysis. Journal of interventional cardiac electrophysiology : an international journal of arrhythmias and pacing, 66(5), 1145.

Petrovi? T, et al. (2022) IgG N-glycome changes during the course of severe COVID-19: An observational study. EBioMedicine, 81, 104101.

O'Brien W, et al. (2022) The Assessment of Functional Movement in Children and Adolescents: A Systematic Review and Meta-Analysis. Sports medicine (Auckland, N.Z.), 52(1), 37.

Kaushal R, et al. (2021) Dicer-like proteins influence Arabidopsis root microbiota independent of RNA-directed DNA methylation. Microbiome, 9(1), 57.

Runacres A, et al. (2021) Impact of the COVID-19 Pandemic on Sedentary Time and Behaviour in Children and Adults: A Systematic Review and Meta-Analysis. *International journal of environmental research and public health*, 18(21).

Strausz S, et al. (2021) Sleep apnoea is a risk factor for severe COVID-19. *BMJ open respiratory research*, 8(1).

Zhou Z, et al. (2020) Genome diversification in globally distributed novel marine Proteobacteria is linked to environmental adaptation. *The ISME journal*, 14(8), 2060.

Wang X, et al. (2019) The associations between screen time-based sedentary behavior and depression: a systematic review and meta-analysis. *BMC public health*, 19(1), 1524.

Ter Veer E, et al. (2016) The efficacy and safety of S-1-based regimens in the first-line treatment of advanced gastric cancer: a systematic review and meta-analysis. *Gastric cancer : official journal of the International Gastric Cancer Association and the Japanese Gastric Cancer Association*, 19(3), 696.

Li H, et al. (2015) Association of matrix metalloproteinase family gene polymorphisms with lung cancer risk: logistic regression and generalized odds of published data. *Scientific reports*, 5, 10056.

He W, et al. (2012) MDM2 SNP309 polymorphism is associated with lung cancer risk in women: A meta-analysis using METAGEN. *Experimental and therapeutic medicine*, 4(4), 569.