Resource Summary Report

Generated by <u>RRID</u> on Apr 19, 2025

<u>metagen</u>

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

Funding:

Resource Name: metagen

Resource ID: SCR_003443

Alternate IDs: OMICS_00238, biotools:metagen

Alternate URLs: https://bio.tools/metagen

Record Creation Time: 20220129T080219+0000

Record Last Update: 20250410T065007+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 17 mentions in open access literature.

Listed below are recent publications. The full list is available at <u>RRID</u>.

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

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.

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

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.