Resource Summary Report

Generated by NIF on Apr 20, 2025

GEOquery

RRID:SCR_000146

Type: Tool

Proper Citation

GEOquery (RRID:SCR_000146)

Resource Information

URL: http://www.bioconductor.org/packages/release/bioc/html/GEOquery.html

Proper Citation: GEOquery (RRID:SCR_000146)

Description: Software that establishes a bridge between GEO and BioConductor.

Abbreviations: GEOquery

Synonyms: GEOquery - Get data from NCBI Gene Expression Omnibus (GEO)

Resource Type: software resource

Defining Citation: PMID:17496320

Keywords: microarray, bio.tools

Funding:

Availability: GNU General Public License, v2

Resource Name: GEOquery

Resource ID: SCR_000146

Alternate IDs: biotools:geoquery, OMICS_01972

Alternate URLs: https://bio.tools/geoquery, https://sources.debian.org/src/r-bioc-geoquery/

Record Creation Time: 20220129T080159+0000

Record Last Update: 20250420T013931+0000

Ratings and Alerts

No rating or validation information has been found for GEOquery.

No alerts have been found for GEOquery.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 11 mentions in open access literature.

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

Farin HF, et al. (2023) Colorectal Cancer Organoid-Stroma Biobank Allows Subtype-Specific Assessment of Individualized Therapy Responses. Cancer discovery, 13(10), 2192.

Habgood-Coote D, et al. (2023) Diagnosis of childhood febrile illness using a multi-class blood RNA molecular signature. Med (New York, N.Y.), 4(9), 635.

Pieri V, et al. (2023) Aberrant L-Fucose Accumulation and Increased Core Fucosylation Are Metabolic Liabilities in Mesenchymal Glioblastoma. Cancer research, 83(2), 195.

Rohan P, et al. (2023) High Expression of THY1 in Intestinal Gastric Cancer as a Key Factor in Tumor Biology: A Poor Prognosis-Independent Marker Related to the Epithelial-Mesenchymal Transition Profile. Genes, 15(1).

Lu C, et al. (2022) Identification of CXCL10 and CXCL11 as the candidate genes involving the development of colitis-associated colorectal cancer. Frontiers in genetics, 13, 945414.

Birt IA, et al. (2021) Genetic Liability for Internalizing Versus Externalizing Behavior Manifests in the Developing and Adult Hippocampus: Insight From a Meta-analysis of Transcriptional Profiling Studies in a Selectively Bred Rat Model. Biological psychiatry, 89(4), 339.

Schmidt M, et al. (2020) Deconvolution of cellular subsets in human tissue based on targeted DNA methylation analysis at individual CpG sites. BMC biology, 18(1), 178.

Roy A, et al. (2019) PI3K-Yap activity drives cortical gyrification and hydrocephalus in mice. eLife, 8.

Mishra S, et al. (2017) Efficacy of ?-lactam/?-lactamase inhibitor combination is linked to WhiB4-mediated changes in redox physiology of Mycobacterium tuberculosis. eLife, 6.

Duan Y, et al. (2016) Prediction of key genes and miRNAs responsible for loss of muscle

force in patients during an acute exacerbation of chronic obstructive pulmonary disease. International journal of molecular medicine, 38(5), 1450.

Wang L, et al. (2016) Analyzing gene expression profiles in dilated cardiomyopathy via bioinformatics methods. Brazilian journal of medical and biological research = Revista brasileira de pesquisas medicas e biologicas, 49(10), e4897.