Resource Summary Report

Generated by NIF on Apr 29, 2025

OrderedList

RRID:SCR_001834
Type: Tool

Proper Citation

OrderedList (RRID:SCR_001834)

Resource Information

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

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Description: An R / bioconductor package for detecting similarity in ordered gene lists. Thereby, either simple lists can be compared or gene expression data can be used to deduce the lists. Significance of similarities is evaluated by shuffling lists or by resampling in microarray data, respectively.

Synonyms: OrderedList - Similarities of Ordered Gene Lists

Resource Type: software resource

Defining Citation: PMID:16844712

Keywords: standalone software, mac os x, unix/linux, windows, r, differential expression, microarray, multiple comparison

Funding:

Availability: GNU General Public License, v2 or greater

Resource Name: OrderedList

Resource ID: SCR_001834

Alternate IDs: OMICS_03525

Alternate URLs: http://compdiag.molgen.mpg.de/software/OrderedList.shtml

Record Creation Time: 20220129T080209+0000

Record Last Update: 20250420T014041+0000

Ratings and Alerts

No rating or validation information has been found for OrderedList.

No alerts have been found for OrderedList.

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.

Glastad KM, et al. (2023) Long ant life span is maintained by a unique heat shock factor. Genes & development, 37(9-10), 398.

Gospocic J, et al. (2021) Kr-h1 maintains distinct caste-specific neurotranscriptomes in response to socially regulated hormones. Cell, 184(23), 5807.

Auray G, et al. (2020) High-Resolution Profiling of Innate Immune Responses by Porcine Dendritic Cell Subsets in vitro and in vivo. Frontiers in immunology, 11, 1429.

Rosiles T, et al. (2020) Registered Report: Transcriptional Analysis of Savings Memory Suggests Forgetting is Due to Retrieval Failure. eNeuro, 7(6).

Goulart MR, et al. (2019) Phenotypic and transcriptomic characterization of canine myeloid-derived suppressor cells. Scientific reports, 9(1), 3574.

Talker SC, et al. (2018) Precise Delineation and Transcriptional Characterization of Bovine Blood Dendritic-Cell and Monocyte Subsets. Frontiers in immunology, 9, 2505.

Tarkkonen K, et al. (2017) Comparative analysis of osteoblast gene expression profiles and Runx2 genomic occupancy of mouse and human osteoblasts in vitro. Gene, 626, 119.

Rao M, et al. (2015) Enteric glia express proteolipid protein 1 and are a transcriptionally unique population of glia in the mammalian nervous system. Glia, 63(11), 2040.

Yang XH, et al. (2015) Identification of epigenetic modifications that contribute to pathogenesis in therapy-related AML: Effective integration of genome-wide histone

modification with transcriptional profiles. BMC medical genomics, 8 Suppl 2(Suppl 2), S6.

Voyle N, et al. (2015) Blood Protein Markers of Neocortical Amyloid-? Burden: A Candidate Study Using SOMAscan Technology. Journal of Alzheimer's disease : JAD, 46(4), 947.

Kim M, et al. (2015) Increased expression of interferon signaling genes in the bone marrow microenvironment of myelodysplastic syndromes. PloS one, 10(3), e0120602.