

Resource Summary Report

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GeneTerm Linker

RRID:SCR_006385

Type: Tool

Proper Citation

GeneTerm Linker (RRID:SCR_006385)

Resource Information

URL: <http://gtlinker.cnb.csic.es/>

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Description: Web application that filters and links enriched output data identifying sets of associated genes and terms, producing metagroups of coherent biological significance. The method uses fuzzy reciprocal linkage between genes and terms to unravel their functional convergence and associations. It can also be accessed through its web service.

Abbreviations: GTLinker

Synonyms: GeneTerm Linker - post enrichment functional association by non-redundant reciprocal linkage

Resource Type: analysis service resource, data analysis service, production service resource, service resource

Defining Citation: [PMID:21949701](#)

Keywords: gene, functional annotation, function, functional metagroup, p-value, annotation, web service

Availability: Acknowledgement requested

Resource Name: GeneTerm Linker

Resource ID: SCR_006385

Alternate IDs: OMICS_02227

Record Creation Time: 20220129T080235+0000

Record Last Update: 20260821T042656+0000

Ratings and Alerts

No rating or validation information has been found for GeneTerm Linker.

No alerts have been found for GeneTerm Linker.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Sánchez-Luis E, et al. (2020) Deciphering Master Gene Regulators and Associated Networks of Human Mesenchymal Stromal Cells. *Biomolecules*, 10(4).

González-Velasco O, et al. (2020) Transcriptomic landscape, gene signatures and regulatory profile of aging in the human brain. *Biochimica et biophysica acta. Gene regulatory mechanisms*, 1863(6), 194491.