

# Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

## 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:** 20260905T133135+0000

### Ratings and Alerts

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

No alerts have been found for GeneTerm Linker.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

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

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.

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