

Resource Summary Report

Generated by T1D on Aug 5, 2026

REViGO

RRID:SCR_005825

Type: Tool

Proper Citation

REViGO (RRID:SCR_005825)

Resource Information

URL: <http://revigo.irb.hr/>

Proper Citation: REViGO (RRID:SCR_005825)

Description: Web server that summarizes lists of Gene Ontology terms by removing redundant terms and visualizing the remaining ones in scatterplots, interactive graphs, treemaps, or tag clouds. Platform: Online tool, THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

Abbreviations: REViGO

Synonyms: Reduce + Visualize Gene Ontology, REViGO - Reduce Visualize Gene Ontology, REViGO - Reduce & Visualize Gene Ontology, REViGO - Reduce and Visualize Gene Ontology, REViGO - Reduce + Visualize Gene Ontology

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

Defining Citation: [PMID:21789182](#), [PMID:20585573](#)

Keywords: gene ontology, visualization, statistical analysis, ontology or annotation visualization

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: REViGO

Resource ID: SCR_005825

Alternate IDs: OMICS_02283, nlx_149332

Record Creation Time: 20220129T080232+0000

Record Last Update: 20260805T044127+0000

Ratings and Alerts

No rating or validation information has been found for REViGO.

No alerts have been found for REViGO.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2299 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#).

Arcego DM, et al. (2026) Sex-specific interaction effects of Syntaxin 1A coexpression network and childhood trauma on adult depressive symptoms. *EBioMedicine*, 123, 106062.

Voigt O, et al. (2025) Genetic parallels in biomineralization of the calcareous sponge *Sycon ciliatum* and stony corals. *eLife*, 14.

Azam S, et al. (2025) Advancing the Indian cattle pangenome: characterizing non-reference sequences in *Bos indicus*. *Journal of animal science and biotechnology*, 16(1), 21.

Oriowo TO, et al. (2025) A chromosome-level, haplotype-resolved genome assembly and annotation for the Eurasian minnow (*Leuciscidae: Phoxinus phoxinus*) provide evidence of haplotype diversity. *GigaScience*, 14.

Aoki N, et al. (2025) Influence of melanin and macrophage activation on hearing loss in *SLC26A4* deficient mice. *Neurobiology of disease*, 212, 106962.

Stomper J, et al. (2025) Sex differences in DNMT3A-mutant clonal hematopoiesis and the effects of estrogen. *Cell reports*, 44(4), 115494.

Yim KM, et al. (2025) Cell-type-specific dysregulation of gene expression due to *Chd8* haploinsufficiency during mouse cortical development. *Cell genomics*, 100986.

Spealman P, et al. (2025) Multilevel Gene Expression Changes in Lineages Containing Adaptive Copy Number Variants. *Molecular biology and evolution*, 42(2).

Pierson Smela M, et al. (2025) Rapid human oogonia-like cell specification via transcription factor-directed differentiation. *EMBO reports*, 26(4), 1114.

Aggarwal B, et al. (2025) CellSP: Module discovery and visualization for subcellular spatial transcriptomics data. *bioRxiv : the preprint server for biology*.

Nazeen S, et al. (2025) NERINE reveals rare variant associations in gene networks across multiple phenotypes and implicates an SNCA-PRL-LRRK2 subnetwork in Parkinson's

disease. bioRxiv : the preprint server for biology.

Zhang N, et al. (2025) Deciphering the molecular logic of WOX5 function in the root stem cell organizer. *The EMBO journal*, 44(1), 281.

Feng X, et al. (2025) Cortical arealization of interneurons defines shared and distinct molecular programs in developing human and macaque brains. *Nature communications*, 16(1), 672.

Carmelet-Rescan D, et al. (2025) Metabolic differentiation of brushtail possum populations resistant and susceptible to plant toxins revealed via differential gene expression. *Journal of comparative physiology. B, Biochemical, systemic, and environmental physiology*, 195(1), 103.

Okafor A, et al. (2025) Transcriptome analysis reveals a de novo DNA element that may interact with chromatin-associated proteins in *Plasmodium berghei* during erythrocytic development. *Scientific reports*, 15(1), 18621.

Matan-Lithwick S, et al. (2025) A Transcriptomic Signature of Depressive Symptoms in Late Life. *Biological psychiatry global open science*, 5(3), 100448.

Baty SM, et al. (2025) Strong Signatures of Selection on Candidate Genes Underlying Core Speciation Mechanisms in Desert Tortoises. *Molecular ecology resources*, 25(6), e14098.

Wang JT, et al. (2025) Loss of Sarm1 Mitigates Axonal Degeneration and Promotes Neuronal Repair After Ischemic Stroke. *bioRxiv : the preprint server for biology*.

Park S, et al. (2025) Skin ECM Provides a Bio-Derived Platform for Supporting Dermal Renewal and Matrix Synthesis. *Journal of microbiology and biotechnology*, 35, e2505015.

Marulanda-Gomez AM, et al. (2025) Combined cellular and proteomics approach suggests differential processing of a native and a foreign vibrio in the sponge *Halicondria panicea*. *mBio*, 16(8), e0147425.