AmiGO

RRID:SCR_002143
Type: Tool

Proper Citation

AmiGO (RRID:SCR_002143)

Resource Information

**URL:** [http://amigo.geneontology.org/](http://amigo.geneontology.org/)

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**Description:** Web tool to search, sort, analyze, visualize and download data of interest. Along with providing details of the ontologies, gene products and annotations, features a BLAST search, Term Enrichment and GO Slimmer tools, the GO Online SQL Environment and a user help guide. Used at the Gene Ontology (GO) website to access the data provided by the GO Consortium. Developed and maintained by the GO Consortium.

**Abbreviations:** AmiGO

**Synonyms:** AmiGO 2, AmiGene Ontology, AmiGO, Gene Ontology Database, GO Database, The Gene Ontology Consortium, Gene Ontology Consortium, GO Consortium

**Resource Type:** database, analysis service resource, service resource, data or information resource, production service resource, data analysis service

**Defining Citation:** PMID:19033274

**Keywords:** search, sort, analyze, visualize, data, ontology, gene, annotation, FASEB list

**Funding Agency:** NHGRI

**Availability:** Free, Available for download, Freely available

**Resource Name:** AmiGO

**Resource ID:** SCR_002143
Alternate IDs: nif-0000-20935, OMICS_02266

Alternate URLs: http://sourceforge.net/projects/geneontology/

Ratings and Alerts

No rating or validation information has been found for AmiGO.

No alerts have been found for AmiGO.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 1064 mentions in open access literature.

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

Šimon M, et al. (2023) Genome-wide screening for genetic variants in polyadenylation signal (PAS) sites in mouse selection lines for fatness and leanness. Mammalian genome: official journal of the International Mammalian Genome Society, 34(1), 12.


Iyer MS, et al. (2023) A Systems Biology Approach To Disentangle the Direct and Indirect Effects of Global Transcription Factors on Gene Expression in Escherichia coli. Microbiology


