QuickGO

RRID:SCR_004608
Type: Tool

Proper Citation

QuickGO (RRID:SCR_004608)

Resource Information

URL: http://www.ebi.ac.uk/QuickGO/

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Description: A web-based browser for Gene Ontology terms and annotations, which is provided by the UniProtKB-GOA group at the EBI. It is able to offer a range of facilities including bulk downloads of GO annotation data which can be extensively filtered by a range of different parameters and GO slim set generation. The software for QuickGO is freely available under the Apache 2 license. QuickGO can supply GO term information and GO annotation data via REST web services.

Abbreviations: QuickGO

Synonyms: Quick GO

Resource Type: software resource, database, data access protocol, web service, ontology, data or information resource, controlled vocabulary

Defining Citation: PMID:19744993, PMID:20157483

Keywords: gene, ontology, annotation, browser, visualization, search engine, slimmer-type tool, ontology or annotation browser, ontology or annotation search engine, ontology or annotation visualization, database or data warehouse, windows, mac os x, linux, unix, gold standard, bio.tools

Funding Agency: BBSRC

Availability: Apache License, v2, Free for academic use
Resource Name: QuickGO
Resource ID: SCR_004608
Alternate IDs: nlx_60318, OMICS_02276, biotools:quickgo
Alternate URLs: https://bio.tools/quickgo

Ratings and Alerts

No rating or validation information has been found for QuickGO.

No alerts have been found for QuickGO.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 398 mentions in open access literature.

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


Crozet F, et al. (2023) Filopodia-like protrusions of adjacent somatic cells shape the developmental potential of oocytes. Life science alliance, 6(6).


Muller PR, et al. (2023) Proteomic Analysis of Female Synovial Fluid to Identify Novel Biomarkers for Osteoarthritis. Life (Basel, Switzerland), 13(3).

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.

Guan Q, et al. (2023) In Silico Analysis and Immune Response of YaeT Protein Against...
Riemerella anatipestifer in Ducks. Applied biochemistry and biotechnology, 1.


Dishnica K, et al. (2023) Novel insights into the somatic proteome of Strongyloides stercoralis infective third-stage larvae. Parasites & vectors, 16(1), 45.


Börner J, et al. (2023) Ribonuclease E strongly impacts bacterial adaptation to different growth conditions. RNA biology, 20(1), 120.


Euclide PT, et al. (2022) Further evidence from common garden rearing experiments of heritable traits separating lean and siscowet lake charr (Salvelinus namaycush) ecotypes. Molecular ecology, 31(12), 3432.
