

Resource Summary Report

Generated by SPARC SAWG on Sep 12, 2026

GOfetcher

RRID:SCR_005681

Type: Tool

Proper Citation

GOfetcher (RRID:SCR_005681)

Resource Information

URL: <http://mcbc.usm.edu/gofetcher/>

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Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented on June 29, 2012. We developed a web application, GOfetcher, with a very comprehensive search facility for the GO project and a variety of output formats for the results. GOfetcher has three different levels for searching the GO: Quick Search, Advanced Search, and Upload Files for searching. The application includes a unique search option which generates gene information given a nucleotide or protein accession number which can then be used in generating gene ontology information. The output data in GOfetcher can be saved into several different formats; including spreadsheet, comma-separated values, and the Extensible Markup Language (XML) format. Platform: Online tool

Abbreviations: GOfetcher

Synonyms: GOfetcher: a database with complex searching facility for gene ontology

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

Defining Citation: [PMID:18728045](#)

Keywords: gene, nucleotide, protein, ontology, ontology or annotation browser

Funding: NSF EPS-0556308;
U.S. Army ;
Environmental Quality Program contract #W912HZ-05-P-0145

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: GOfetcher

Resource ID: SCR_005681

Alternate IDs: nlx_149124

Old URLs: <http://mcbc.usm.edu/gofetcher/>

Record Creation Time: 20220129T080231+0000

Record Last Update: 20260912T195629+0000

Ratings and Alerts

No rating or validation information has been found for GOfetcher.

No alerts have been found for GOfetcher.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.