

# Resource Summary Report

Generated by [NIF](#) on Apr 19, 2025

## [go-db-perl](#)

RRID:SCR\_005721

Type: Tool

### Proper Citation

go-db-perl (RRID:SCR\_005721)

### Resource Information

**URL:** <http://search.cpan.org/~cmungall/go-db-perl/>

**Proper Citation:** go-db-perl (RRID:SCR\_005721)

**Description:** Software resource that extends the functionality of go-perl (on which it depends) with GO Database access functionality. go-db-perl comes bundled with various scripts and a shell command line interface that can be used as standalone tools. Installation is more involved than for go-perl; you will need a MySQL database plus the requisite DBI and DBD Perl modules. Full installation instructions are included in the download. go-db-perl is in use both to drive AmiGO and internally within Ensembl. Platform: Windows compatible, Mac OS X compatible, Linux compatible, Unix compatible

**Abbreviations:** go-db-perl

**Resource Type:** software resource

**Keywords:** software library, database, gene ontology, database or data warehouse

**Funding:**

**Availability:** Free for academic use

**Resource Name:** go-db-perl

**Resource ID:** SCR\_005721

**Alternate IDs:** nlx\_149180

**Record Creation Time:** 20220129T080232+0000

**Record Last Update:** 20250420T014300+0000

---

## Ratings and Alerts

No rating or validation information has been found for go-db-perl.

No alerts have been found for go-db-perl.

---

## Data and Source Information

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

---

## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Carbon S, et al. (2009) AmiGO: online access to ontology and annotation data. *Bioinformatics* (Oxford, England), 25(2), 288.