

Resource Summary Report

Generated by [kravitz2](#) on Aug 1, 2026

GOArray

RRID:SCR_005785

Type: Tool

Proper Citation

GOArray (RRID:SCR_005785)

Resource Information

URL: <http://g2im.u-clermont1.fr/serimour/goarrays.html>

Proper Citation: GOArray (RRID:SCR_005785)

Description: GOArray is a Perl program which inputs a lists of genes annotated as of interest (GOI) or not, and determines if any associated GO terms have an overrepresentation of GOI. A permutation test is optionally used to assess confidence in the results. Output includes multiple visualizations and supplementary information and, for future reference, a summary of the statistical methods used. Platform: Windows compatible, Mac OS X compatible, Linux compatible, Unix compatible

Abbreviations: GOArray

Resource Type: data processing software, data analysis software, software application, software resource

Keywords: perl, gene, visualization, gene ontology, statistical analysis

Availability: Free for academic use

Resource Name: GOArray

Resource ID: SCR_005785

Alternate IDs: nlx_149259

Old URLs: <http://goarray.med.yale.edu/GOArray/>

Record Creation Time: 20220129T080232+0000

Record Last Update: 20260801T042617+0000

Ratings and Alerts

No rating or validation information has been found for GOArray.

No alerts have been found for GOArray.

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 [kravitz2](#) .

Comtet-Marre S, et al. (2018) FibroChip, a Functional DNA Microarray to Monitor Cellulolytic and Hemicellulolytic Activities of Rumen Microbiota. *Frontiers in microbiology*, 9, 215.