

# Resource Summary Report

Generated by [dkNET](#) on Aug 30, 2026

## GOArray

RRID:SCR\_005785

Type: Tool

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### Proper Citation

GOArray (RRID:SCR\_005785)

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### Resource Information

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

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**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 analysis software, data processing 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:** 20260829T182229+0000

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### Ratings and Alerts

No rating or validation information has been found for GOArray.

No alerts have been found for GOArray.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

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.