

# Resource Summary Report

Generated by T1D on Aug 7, 2026

## GeneCommittee

RRID:SCR\_004168

Type: Tool

### Proper Citation

GeneCommittee (RRID:SCR\_004168)

### Resource Information

**URL:** <http://sing.ei.uvigo.es/GC/>

**Proper Citation:** GeneCommittee (RRID:SCR\_004168)

**Description:** Tool for extensively testing the discriminatory power of biologically relevant gene sets in microarray data classification. While the user can work with different gene set collections and several microarray data files to configure specific classification experiments, the tool is able to run several tests in parallel. It is able to render valuable information for diagnostic analyses and clinical management decisions based on systematically evaluating custom hypothesis over different data sets using complementary classifiers, a key aspect in clinical research.

**Abbreviations:** geneCommittee

**Resource Type:** software resource, service resource, production service resource, data analysis service, analysis service resource

**Defining Citation:** [PMID:24475928](https://pubmed.ncbi.nlm.nih.gov/24475928/)

**Keywords:** dna, microarray, rna-seq, gene expression, classification, gene set, gene, gene set enrichment, functional annotation

**Availability:** Acknowledgement requested

**Resource Name:** GeneCommittee

**Resource ID:** SCR\_004168

**Alternate IDs:** OMICS\_02291

**Alternate URLs:** <https://github.com/michada/GeneCommittee>

**Record Creation Time:** 20220129T080223+0000

**Record Last Update:** 20260807T042602+0000

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## Ratings and Alerts

No rating or validation information has been found for GeneCommittee.

No alerts have been found for GeneCommittee.

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## Data and Source Information

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

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.