

Resource Summary Report

Generated by T1D on Sep 5, 2026

THEA - Tools for High-throughput Experiments Analysis

RRID:SCR_005802

Type: Tool

Proper Citation

THEA - Tools for High-throughput Experiments Analysis (RRID:SCR_005802)

Resource Information

URL: <http://thea.unice.fr/index-en.html>

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Description: THIS RESOURCE IS NO LONGER IN SERVICE, on documented July 16, 2012. An integrated information processing system dedicated to the analysis of post-genomic data. It allows automatic annotation of data issued from classification systems with selected biological information (including the Gene Ontology). Users can either manually search and browse through these annotations, or automatically generate meaningful generalizations according to statistical criteria (data mining). Platform: Windows compatible, Mac OS X compatible, Linux compatible, Unix compatible

Abbreviations: THEA

Synonyms: Tools for High-throughput Experiments Analysis

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

Defining Citation: [PMID:15130932](#)

Keywords: high-throughput, analysis, ontology, microarray, genomic, annotation, gene ontology, data mining, statistical analysis

Funding: French Ministry of Higher Education and Research ;
Bioinformatic Program

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: THEA - Tools for High-throughput Experiments Analysis

Resource ID: SCR_005802

Alternate IDs: nlx_149290

Record Creation Time: 20220129T080232+0000

Record Last Update: 20260903T234822+0000

Ratings and Alerts

No rating or validation information has been found for THEA - Tools for High-throughput Experiments Analysis.

No alerts have been found for THEA - Tools for High-throughput Experiments Analysis.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.