CLENCH
RRID:SCR_005735
Type: Tool

Proper Citation

CLENCH (RRID:SCR_005735)

Resource Information

URL: http://www.stanford.edu/~nigam/cgi-bin/dokuwiki/doku.php?id=clench

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Description: Cluster Enrichment (CLENCH) allows A. thaliana researchers to perform automated retrieval of GO annotations from TAIR and calculate enrichment of GO terms in gene group with respect to a reference set. Before calculating enrichment, CLENCH allows mapping of the returned annotations to arbitrary coarse levels using GO slim term lists (which can be edited by the user) and a local installation of GO. Platform: Windows compatible, Linux compatible,

Abbreviations: CLENCH

Synonyms: Cluster Enrichment, Cluster Enrichment (CLENCH), CLENCH - Cluster Enrichment, CLENCH: A program for calculating cluster enrichment using the Gene Ontology

Resource Type: source code, software resource, software application, data processing software

Defining Citation: PMID:14764555

Keywords: gene, microarray, function, functional categorization, statistical analysis, slimmer-type tool, gene ontology, annotation

Availability: Free for academic use

Resource Name: CLENCH
Ratings and Alerts

No rating or validation information has been found for CLENCH.

No alerts have been found for CLENCH.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 3 mentions in open access literature.

Listed below are recent publications. The full list is available at RRID.

