SOURCE
RRID:SCR_005799
Type: Tool

Proper Citation

SOURCE (RRID:SCR_005799)

Resource Information

URL: http://smd.stanford.edu/cgi-bin/source/sourceSearch

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Description: SOURCE compiles information from several publicly accessible databases, including UniGene, dbEST, UniProt Knowledgebase, GeneMap99, RHdb, GeneCards and LocusLink. GO terms associated with LocusLink entries appear in SOURCE. The mission of SOURCE is to provide a unique scientific resource that pools publicly available data commonly sought after for any clone, GenBank accession number, or gene. SOURCE is specifically designed to facilitate the analysis of large sets of data that biologists can now produce using genome-scale experimental approaches. Platform: Online tool

Abbreviations: SOURCE

Resource Type: database, analysis service resource, service resource, data or information resource, production service resource, data analysis service

Defining Citation: PMID:12519986

Keywords: genomic, functional annotation, ontology, gene expression, gene, genome, statistical analysis, bio.tools, FASEB list

Funding Agency: NIGMS, NCI, NIGMS

Availability: Restricted

Resource Name: SOURCE

Resource ID: SCR_005799
Alternate IDs: nlx_149287, biotools:source

Alternate URLs:
https://login.stanford.edu/idp/profile/SAML2/Redirect/SSO?execution=e1s1,
https://bio.tools/source

Ratings and Alerts

No rating or validation information has been found for SOURCE.

No alerts have been found for SOURCE.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 62 mentions in open access literature.

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


Wu J, et al. (2017) Nomogram integrating gene expression signatures with