

Resource Summary Report

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CiteXplore literature searching

RRID:SCR_010676

Type: Tool

Proper Citation

CiteXplore literature searching (RRID:SCR_010676)

Resource Information

URL: <http://www.ebi.ac.uk/citexplore/>

Proper Citation: CiteXplore literature searching (RRID:SCR_010676)

Description: CiteXplore combines literature search with text mining tools for biology. Search results are cross referenced to EBI applications based on publication identifiers. Links to full text versions are provided where available. The underlying database is populated with data from Medline, Patents, C.B.A. and Citeseer. The site offers a quick search as well as an advanced search. Results of queries can be saved in formats compatible with commonly-used bibliographic management software and exported. We provide a Simple Object Access Protocol (SOAP) based service to retrieve data from the Citation database. Text mining is provided internally by Whatizit and externally by iHOP.

Synonyms: CiteXplore

Resource Type: web service, database, data access protocol, data or information resource, software resource

Keywords: gold standard

Resource Name: CiteXplore literature searching

Resource ID: SCR_010676

Alternate IDs: nlx_76807

Record Creation Time: 20220129T080300+0000

Record Last Update: 20260804T043433+0000

Ratings and Alerts

No rating or validation information has been found for CiteXplore literature searching.

No alerts have been found for CiteXplore literature searching.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 7 mentions in open access literature.

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

Palmblad M, et al. (2017) Spatiotemporal analysis of tropical disease research combining Europe PMC and affiliation mapping web services. Tropical medicine and health, 45, 33.

Sarajli?? A, et al. (2013) Network topology reveals key cardiovascular disease genes. PloS one, 8(8), e71537.

de Matos P, et al. (2013) The Enzyme Portal: a case study in applying user-centred design methods in bioinformatics. BMC bioinformatics, 14, 103.

Daugherty LC, et al. (2012) Gene family matters: expanding the HGNC resource. Human genomics, 6(1), 4.

Magrane M, et al. (2011) UniProt Knowledgebase: a hub of integrated protein data. Database : the journal of biological databases and curation, 2011, bar009.

Lu Z, et al. (2011) PubMed and beyond: a survey of web tools for searching biomedical literature. Database : the journal of biological databases and curation, 2011, baq036.

Page GP, et al. (2008) Bioinformatic tools for inferring functional information from plant microarray data: tools for the first steps. International journal of plant genomics, 2008, 147563.