## **Resource Summary Report**

Generated by NIF on Apr 21, 2025

# **EvidenceFinder**

RRID:SCR\_013764

Type: Tool

## **Proper Citation**

EvidenceFinder (RRID:SCR\_013764)

#### **Resource Information**

URL: http://labs.europepmc.org/evf

**Proper Citation:** EvidenceFinder (RRID:SCR\_013764)

**Description:** A web application to assist in the identification of articles and research related to literature search terms. The search covers full text articles in the Europe PMC repository. Relevant papers are suggested to users based on the scientific term searched and the selection of questions, generated by the application, relevant to term searched.

Resource Type: web application, software resource

**Defining Citation:** DOI:10.1093/nar/gku1061

**Keywords:** web application, software resource, literature search, bio.tools

Funding: Wellcome Trust 098231

Availability: Free, Public

Resource Name: EvidenceFinder

Resource ID: SCR\_013764

Alternate IDs: biotools:evidence finder

**Alternate URLs:** https://bio.tools/evidence\_finder

License URLs: http://www.ebi.ac.uk/about/terms-of-use

**Record Creation Time: 20220129T080317+0000** 

**Record Last Update:** 20250420T015238+0000

### **Ratings and Alerts**

No rating or validation information has been found for EvidenceFinder.

No alerts have been found for EvidenceFinder.

#### 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 NIF.

Kostareva A, et al. (2016) Genetic Spectrum of Idiopathic Restrictive Cardiomyopathy Uncovered by Next-Generation Sequencing. PloS one, 11(9), e0163362.

Rouillard AD, et al. (2016) The harmonizome: a collection of processed datasets gathered to serve and mine knowledge about genes and proteins. Database: the journal of biological databases and curation, 2016.

Ananiadou S, et al. (2015) Event-based text mining for biology and functional genomics. Briefings in functional genomics, 14(3), 213.