

# Resource Summary Report

Generated by [dkNET](#) on Aug 26, 2026

## 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:** software resource, web application

**Defining Citation:** [DOI:10.1093/nar/gku1061](https://doi.org/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](https://bio.tools/evidence_finder)

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

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

**Record Last Update:** 20260821T194007+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 [dkNET](#) .

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.