# **Resource Summary Report**

Generated by NIF on Apr 23, 2025

# **EBIMed**

RRID:SCR\_005314

Type: Tool

### **Proper Citation**

EBIMed (RRID:SCR\_005314)

#### Resource Information

URL: http://www.ebi.ac.uk/Rebholz-srv/ebimed/

**Proper Citation:** EBIMed (RRID:SCR\_005314)

**Description:** A web application that combines Information Retrieval and Extraction from Medline. EBIMed finds Medline abstracts in the same way PubMed does. Then it goes a step beyond and analyses them to offer a complete overview on associations between UniProt protein/gene names, GO annotations, Drugs and Species. The results are shown in a table that displays all the associations and links to the sentences that support them and to the original abstracts. By selecting relevant sentences and highlighting the biomedical terminology EBIMed enhances your ability to acquire knowledge, relate facts, discover implications and, overall, have a good overview economizing the effort in reading.

**Abbreviations:** EBIMed

Resource Type: service resource

**Keywords:** protein, gene, annotation, drug, specie, association, database

**Funding:** 

Resource Name: EBIMed

Resource ID: SCR\_005314

Alternate IDs: OMICS\_01180

**Record Creation Time:** 20220129T080229+0000

Record Last Update: 20250420T014248+0000

## **Ratings and Alerts**

No rating or validation information has been found for EBIMed.

No alerts have been found for EBIMed.

#### **Data and Source Information**

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 1 mentions in open access literature.

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

Titz B, et al. (2014) Proteomics for systems toxicology. Computational and structural biotechnology journal, 11(18), 73.