# **Resource Summary Report**

Generated by dkNET on May 21, 2025

# eSLDB - eukaryotic Subcellular Localization database

RRID:SCR\_000052

Type: Tool

### **Proper Citation**

eSLDB - eukaryotic Subcellular Localization database (RRID:SCR\_000052)

#### **Resource Information**

URL: http://gpcr.biocomp.unibo.it/esldb

Proper Citation: eSLDB - eukaryotic Subcellular Localization database

(RRID:SCR\_000052)

**Description:** THIS RESOURCE IS NO LONGER IN SERVICE. Documented on August 22,2022. database of protein subcellular localization annotation for eukaryotic organisms. It contains experimental annotations derived from primary protein databases, homology based annotations and computational predictions.

Abbreviations: eSLDB

Synonyms: eukaryotic Subcellular Localization database

Resource Type: data or information resource, database

**Defining Citation: PMID:17108361** 

**Keywords:** proteome, protein, homology

Funding: European Union VI Framework Programme

Availability: THIS RESOURCE IS NO LONGER IN SERVICE.

Resource Name: eSLDB - eukaryotic Subcellular Localization database

Resource ID: SCR 000052

**Alternate IDs:** nif-0000-02815

**Record Creation Time:** 20220129T080159+0000

**Record Last Update:** 20250521T060725+0000

## **Ratings and Alerts**

No rating or validation information has been found for eSLDB - eukaryotic Subcellular Localization database.

No alerts have been found for eSLDB - eukaryotic Subcellular Localization database.

#### 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 dkNET.

Fischbach A, et al. (2018) The C-terminal domain of p53 orchestrates the interplay between non-covalent and covalent poly(ADP-ribosyl)ation of p53 by PARP1. Nucleic acids research, 46(2), 804.