Resource Summary Report

Generated by NIF 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 NIF.

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.