

# Resource Summary Report

Generated by [kravitz2](#) on Jul 31, 2026

## Functional Coverage of the Proteome

RRID:SCR\_007654

Type: Tool

### Proper Citation

Functional Coverage of the Proteome (RRID:SCR\_007654)

### Resource Information

**URL:** <http://cgl.imim.es/fcp/>

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**Description:** FCP is a publicly accessible web tool dedicated to analyzing the current state and trends of available proteome structures along the classification schemes of enzymes and nuclear receptors. It offers both graphical and quantitative data on the degree of functional coverage in that portion of the proteome by existing structures and on the bias observed in the distribution of those structures among proteins. Users can choose to search the website based on structures or ligands, and can also sort by enzyme or receptor. Users can also view data based on structural and population (species) filters.

**Synonyms:** FCP

**Resource Type:** data or information resource, database

**Keywords:** enzyme, nuclear receptor, protein, proteome, proteome structure

**Resource Name:** Functional Coverage of the Proteome

**Resource ID:** SCR\_007654

**Alternate IDs:** nif-0000-02834

**Record Creation Time:** 20220129T080243+0000

**Record Last Update:** 20260729T044153+0000

### Ratings and Alerts

No rating or validation information has been found for Functional Coverage of the Proteome.

No alerts have been found for Functional Coverage of the Proteome.

## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [kravitz2](#) .

Vella M, et al. (2014) libNeuroML and PyLEMS: using Python to combine procedural and declarative modeling approaches in computational neuroscience. Frontiers in neuroinformatics, 8, 38.

McDonald KK, et al. (2012) Exome analysis of two limb-girdle muscular dystrophy families: mutations identified and challenges encountered. PloS one, 7(11), e48864.