

Resource Summary Report

Generated by [RRID](#) on Aug 15, 2026

ProteinCartography

RRID:SCR_027230

Type: Tool

Proper Citation

ProteinCartography (RRID:SCR_027230)

Resource Information

URL: <https://github.com/Arcadia-Science/ProteinCartography>

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Description: Software pipeline to build similarity maps of protein space. Identifies proteins related to query protein using sequence- and structure-based searches, compares all protein structures, and creates navigable map that can be used to look at protein relationships and make hypotheses about function.

Resource Type: software toolkit, source code, software resource

Defining Citation: [DOI:10.57844/arcadia-a5a6-1068](https://doi.org/10.57844/arcadia-a5a6-1068)

Keywords: Identifies proteins, build similarity maps, protein space, protein relationships,

Availability: Free, Available for download, Freely available

Resource Name: ProteinCartography

Resource ID: SCR_027230

License: MIT license

Record Creation Time: 20250723T062012+0000

Record Last Update: 20260815T183257+0000

Ratings and Alerts

No rating or validation information has been found for ProteinCartography.

No alerts have been found for ProteinCartography.

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 [RRID](#) .

Yang LY, et al. (2026) The LiPP Benchmark Set for Modeling Lipid-Protein Complexes: Comparison of Co-Folding and Docking Methods. Journal of chemical information and modeling, 66(12), 7231.