

Resource Summary Report

Generated by [NIF](#) on Aug 21, 2026

ProLinks Database of Functional Linkages

RRID:SCR_003185

Type: Tool

Proper Citation

ProLinks Database of Functional Linkages (RRID:SCR_003185)

Resource Information

URL: <https://omictools.com/prolinks-tool>

Proper Citation: ProLinks Database of Functional Linkages (RRID:SCR_003185)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented July 7, 2017. Collection of inference methods used to predict functional linkages between proteins. These methods include the Phylogenetic Profile method which uses the presence and absence of proteins across multiple genomes to detect functional linkages; the Gene Cluster method which uses genome proximity to predict functional linkage; Rosetta Stone which uses a gene fusion event in a second organism to infer functional relatedness; and the Gene Neighbor method which uses both gene proximity and phylogenetic distribution to infer linkage.

Synonyms: Prolinks

Resource Type: software toolkit, software resource

Defining Citation: [PMID:15128449](#)

Keywords: functional linkage, protein linkage, inference method

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: ProLinks Database of Functional Linkages

Resource ID: SCR_003185

Alternate IDs: nif-0000-00580

Old URLs: <http://prl.mbi.ucla.edu/prlbeta/prolinks.jsp>
<http://dip.mbi.ucla.edu/dipbeta/prolinks.jsp>

Record Creation Time: 20220129T080217+0000

Record Last Update: 20260819T044101+0000

Ratings and Alerts

No rating or validation information has been found for ProLinks Database of Functional Linkages.

No alerts have been found for ProLinks Database of Functional Linkages.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.