

Resource Summary Report

Generated by [kravitz2](#) on Sep 4, 2026

Berkeley Bioinformatics Open-Source Projects

RRID:SCR_006704

Type: Tool

Proper Citation

Berkeley Bioinformatics Open-Source Projects (RRID:SCR_006704)

Resource Information

URL: <http://www.berkeleybop.org/>

Proper Citation: Berkeley Bioinformatics Open-Source Projects (RRID:SCR_006704)

Description: The BBOP, located at the Lawrence Berkeley National Labs, is a diverse group of scientific researchers and software engineers dedicated to developing tools and applying computational technologies to solve biological problems. Members of the group contribute to a number of projects, including the Gene Ontology, OBO Foundry, the Phenotypic Quality Ontology, modENCODE, and the Generic Model Organism Database Project. Our group is focused on the development, use, and integration of ontologies into biological data analysis. Software written or maintained by BBOP is accessible through the site.

Abbreviations: BBOP

Synonyms: Berkeley BOP

Resource Type: data or information resource, portal, software resource, topical portal

Keywords: ontology, biology, computation, data analysis

Funding: NHGRI ;
NIGMS

Resource Name: Berkeley Bioinformatics Open-Source Projects

Resource ID: SCR_006704

Alternate IDs: nlx_149171

Record Creation Time: 20220129T080237+0000

Record Last Update: 20260903T234850+0000

Ratings and Alerts

No rating or validation information has been found for Berkeley Bioinformatics Open-Source Projects.

No alerts have been found for Berkeley Bioinformatics Open-Source Projects.

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 [kravitz2](#) .

Van Auken K, et al. (2012) Text mining in the biocuration workflow: applications for literature curation at WormBase, dictyBase and TAIR. Database : the journal of biological databases and curation, 2012, bas040.