

Resource Summary Report

Generated by [dkNET](#) on Sep 3, 2026

DBAli. A Database of Structure Alignments.

RRID:SCR_013418

Type: Tool

Proper Citation

DBAli. A Database of Structure Alignments. (RRID:SCR_013418)

Resource Information

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

Proper Citation: DBAli. A Database of Structure Alignments. (RRID:SCR_013418)

Description: DBAli is a database that includes a comprehensive all-against-all comparison of protein structures in the PDB database. It is not currently being updated; however, updates should resume in the near future. All pairwise structural comparisons in DBAli have been obtained using the MAMMOTH program developed in the group of Prof. Angel R. Ortiz. All multiple structure alignments in DBAli have been obtained using the SALIGN command in MODELLER developed in the group of Prof. Andrej Sali.

Synonyms: DBAli

Resource Type: data or information resource, database

Keywords: protein, protein structure comparison

Resource Name: DBAli. A Database of Structure Alignments.

Resource ID: SCR_013418

Alternate IDs: nif-0000-02725

Record Creation Time: 20220129T080316+0000

Record Last Update: 20260904T000344+0000

Ratings and Alerts

No rating or validation information has been found for DBAli. A Database of Structure Alignments..

No alerts have been found for DBAli. A Database of Structure Alignments..

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.