

Resource Summary Report

Generated by T1D on Aug 3, 2026

CPDB - the Circular Permutation Database

RRID:SCR_002261

Type: Tool

Proper Citation

CPDB - the Circular Permutation Database (RRID:SCR_002261)

Resource Information

URL: <http://sarst.life.nthu.edu.tw/cpdb/>

Proper Citation: CPDB - the Circular Permutation Database (RRID:SCR_002261)

Description: A database of circular permutation (CP) in proteins that provides resources for studying circular permutation (CP) and circular permutation relationships among protein structures. This site also offers viable CP site predictions in order to facilitate the application of CP in academic researches and biotechnological developments.

Abbreviations: CPDB

Synonyms: Circular Permutation Database

Resource Type: database, data or information resource

Defining Citation: [PMID:18842637](#)

Keywords: circular permutation, circular permutation site prediction, cp cluster, cp site prediction, permutant, protein structure

Funding: National Science Council Taiwan 96-3112-B-007-006;
National Science Council Taiwan 97-2752-B-007-003-PAE

Resource Name: CPDB - the Circular Permutation Database

Resource ID: SCR_002261

Alternate IDs: nif-0000-02693

Record Creation Time: 20220129T080212+0000

Record Last Update: 20260803T040329+0000

Ratings and Alerts

No rating or validation information has been found for CPDB - the Circular Permutation Database.

No alerts have been found for CPDB - the Circular Permutation Database.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.