## **Resource Summary Report**

Generated by dkNET on Apr 29, 2025

# **CATCHprofiles**

RRID:SCR\_001182

Type: Tool

### **Proper Citation**

CATCHprofiles (RRID:SCR\_001182)

#### **Resource Information**

URL: http://catch.cmbi.ru.nl/

Proper Citation: CATCHprofiles (RRID:SCR\_001182)

**Description:** Software tool for exploring patterns in Chromatin Immuno Precipitation (ChIP) profiling data. The CATCH algorithm performs a hierarchical clustering of the profile patterns with an exhaustive alignment at each step. The algorithm has a user-friendly graphical interface that makes it easy to browse results.

**Abbreviations:** CATCHprofiles

**Synonyms:** CATCH - Unsupervised clustering of ChIP profiles

**Resource Type:** software resource

**Defining Citation: PMID:22238575** 

Keywords: cluster, chip, alignment, chip profile

**Funding:** 

Availability: Acknowledgement requested, GNU General Public License

**Resource Name:** CATCHprofiles

Resource ID: SCR\_001182

Alternate IDs: OMICS\_02171

**Record Creation Time:** 20220129T080206+0000

Record Last Update: 20250420T014022+0000

## **Ratings and Alerts**

No rating or validation information has been found for CATCHprofiles.

No alerts have been found for CATCHprofiles.

### **Data and Source Information**

Source: SciCrunch Registry

# **Usage and Citation Metrics**

We have not found any literature mentions for this resource.