

# Resource Summary Report

Generated by [RRID](#) on Sep 10, 2026

## 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

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**Resource Name:** CATCHprofiles

**Resource ID:** SCR\_001182

**Alternate IDs:** OMICS\_02171

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

**Record Last Update:** 20260905T132430+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.