

Resource Summary Report

Generated by SPARC SAWG on Sep 18, 2026

TargetCaller

RRID:SCR_000414

Type: Tool

Proper Citation

TargetCaller (RRID:SCR_000414)

Resource Information

URL: <http://cellnet.cecad.uni-koeln.de/15763.html>

Proper Citation: TargetCaller (RRID:SCR_000414)

Description: R software package for transcription factor (TF) target gene prediction based on ChIP-seq data. Version 0.4 contains code to compute FDR-corrected q-values via permutations of the peak-to-gene assignments (ClosestGene only).

Abbreviations: TargetCaller

Resource Type: software resource

Defining Citation: [PMID:24278002](https://pubmed.ncbi.nlm.nih.gov/24278002/)

Keywords: chip-seq, transcription factor, target gene

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: TargetCaller

Resource ID: SCR_000414

Alternate IDs: OMICS_01807

Record Creation Time: 20220129T080201+0000

Record Last Update: 20260912T195509+0000

Ratings and Alerts

No rating or validation information has been found for TargetCaller.

No alerts have been found for TargetCaller.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.