# Resource Summary Report

Generated by Kravitz on Sep 16, 2026

## ChIPModule

RRID:SCR_010877
Type: Tool

### Proper Citation

ChIPModule (RRID:SCR_010877)

### Resource Information

**URL:** http://www.cs.ucf.edu/~xiaoman/ChIPModule/ChIPModule.html

**Proper Citation:** ChIPModule (RRID:SCR_010877)

**Description:** A software tool for systematic discovery of transcription factors and their cofactors from ChIP-seq data.

**Abbreviations:** ChIPModule

**Synonyms:** ChIPModule: Systematic discovery of transcription factors and their cofactors from ChIP-seq data

**Resource Type:** software resource

**Resource Name:** ChIPModule

**Resource ID:** SCR_010877

**Alternate IDs:** OMICS_00477

**Record Creation Time:** 20220129T080301+0000

**Record Last Update:** 20260912T195725+0000

### Ratings and Alerts

No rating or validation information has been found for ChIPModule.

No alerts have been found for ChIPModule.

### Data and Source Information

**Source:** SciCrunch Registry

## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at Kravitz .

Pon JR, et al. (2015) MEF2B mutations in non-Hodgkin lymphoma dysregulate cell migration by decreasing MEF2B target gene activation. Nature communications, 6, 7953.