

# Resource Summary Report

Generated by [RRID](#) on Aug 2, 2026

## ChIPSeq Peak Finder

RRID:SCR\_002081

Type: Tool

### Proper Citation

ChIPSeq Peak Finder (RRID:SCR\_002081)

### Resource Information

**URL:** <http://cmb.gis.a-star.edu.sg/ChIPSeq/paperChIPSeq.htm>

**Proper Citation:** ChIPSeq Peak Finder (RRID:SCR\_002081)

**Description:** THIS RESOURCE IS NO LONGER IN SERVICE, documented on April 12, 2017. A software tool to find peaks from ChIPSeq data generated from the Solexa/Illumina platform., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

**Synonyms:** ChIPSeq Toolbox

**Resource Type:** data processing software, data analysis software, software application, software resource

**Defining Citation:** [PMID:27863463](#)

**Keywords:** chipseq, sequencing, dna, analysis, solexa, illumina

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

**Resource Name:** ChIPSeq Peak Finder

**Resource ID:** SCR\_002081

**Alternate IDs:** OMICS\_00434

**Alternate URLs:** <https://sources.debian.org/src/chip-seq/>

**Record Creation Time:** 20220129T080211+0000

**Record Last Update:** 20260801T190207+0000

### Ratings and Alerts

No rating or validation information has been found for ChIPSeq Peak Finder.

No alerts have been found for ChIPSeq Peak Finder.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [RRID](#) .

Hintermair C, et al. (2012) Threonine-4 of mammalian RNA polymerase II CTD is targeted by Polo-like kinase 3 and required for transcriptional elongation. The EMBO journal, 31(12), 2784.