

Resource Summary Report

Generated by [kravitz2](#) on Jul 31, 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, software application, data analysis software, 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: 20260731T042630+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.

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 [kravitz2](#) .

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.