Resource Summary Report

Generated by SPARC SAWG on Aug 9, 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: software resource, software application, data analysis software, data processing software

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: 20260808T185737+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 SPARC SAWG .

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.