

Resource Summary Report

Generated by [RRID](#) on Apr 17, 2025

FindPeaks

RRID:SCR_010857

Type: Tool

Proper Citation

FindPeaks (RRID:SCR_010857)

Resource Information

URL: <http://sourceforge.net/apps/mediawiki/vancouvershortr/index.php?title=FindPeaks>

Proper Citation: FindPeaks (RRID:SCR_010857)

Description: Software application that can be used for converting Eland, Maq (.map), BED or other files into WIG files and identifying areas of enrichment (ChIP-Seq analysis).

Abbreviations: FindPeaks

Resource Type: software resource

Keywords: chip-seq

Funding: BC Cancer Agency ;
Michael Smith Foundation for Health Research

Resource Name: FindPeaks

Resource ID: SCR_010857

Alternate IDs: OMICS_00440

Record Creation Time: 20220129T080301+0000

Record Last Update: 20250410T070025+0000

Ratings and Alerts

No rating or validation information has been found for FindPeaks.

No alerts have been found for FindPeaks.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 297 mentions in open access literature.

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

Zhou Z, et al. (2025) GnRH pulse generator activity in mouse models of polycystic ovary syndrome. *eLife*, 13.

Boubou Y, et al. (2025) Rural-urban transformation shapes oasis agriculture in Morocco's High Atlas Mountains. *Scientific reports*, 15(1), 3433.

Zhao M, et al. (2025) Genetic variation in IL-4 activated tissue resident macrophages determines strain-specific synergistic responses to LPS epigenetically. *Nature communications*, 16(1), 1030.

Han SY, et al. (2025) Multi-dimensional oscillatory activity of mouse GnRH neurons in vivo. *eLife*, 13.

Mestre-Fos S, et al. (2025) eIF3 engages with 3'-UTR termini of highly translated mRNAs. *eLife*, 13.

Pindwarawala M, et al. (2025) Defective glycosylation and ELFN1 binding of mGluR6 congenital stationary night blindness mutants. *Life science alliance*, 8(3).

Chan JD, et al. (2024) FOXO1 enhances CAR T cell stemness, metabolic fitness and efficacy. *Nature*, 629(8010), 201.

Chandrasekaran V, et al. (2024) Bivalent chromatin accommodates survivin and BRG1/SWI complex to activate DNA damage response in CD4+ cells. *Cell communication and signaling : CCS*, 22(1), 440.

Chapeau EA, et al. (2024) Direct and selective pharmacological disruption of the YAP-TEAD interface by IAG933 inhibits Hippo-dependent and RAS-MAPK-altered cancers. *Nature cancer*, 5(7), 1102.

Vitarelli MdO, et al. (2024) Integrating high-throughput analysis to create an atlas of replication origins in *Trypanosoma cruzi* in the context of genome structure and variability. *mBio*, 15(4), e0031924.

Liu B, et al. (2024) Nuclear respiratory factor 1 regulates super enhancer-controlled SPIDR

to protect hepatocellular carcinoma cells from oxidative stress. *BMC gastroenterology*, 24(1), 97.

Cipriano A, et al. (2024) Transcriptional and epigenetic characterization of a new in vitro platform to model the formation of human pharyngeal endoderm. *Genome biology*, 25(1), 211.

Nguyen CDK, et al. (2024) PRMT1 promotes epigenetic reprogramming associated with acquired chemoresistance in pancreatic cancer. *Cell reports*, 43(5), 114176.

Lee HK, et al. (2024) STAT5B leukemic mutations, altering SH2 tyrosine 665, have opposing impacts on immune gene programs. *bioRxiv : the preprint server for biology*.

Parolia A, et al. (2024) NSD2 is a requisite subunit of the AR/FOXA1 neo-enhanceosome in promoting prostate tumorigenesis. *bioRxiv : the preprint server for biology*.

Helminen L, et al. (2024) Chromatin accessibility and pioneer factor FOXA1 restrict glucocorticoid receptor action in prostate cancer. *Nucleic acids research*, 52(2), 625.

Cheng S, et al. (2024) Review and Evaluate the Bioinformatics Analysis Strategies of ATAC-seq and CUT&Tag Data. *Genomics, proteomics & bioinformatics*, 22(3).

Povoleri GAM, et al. (2024) Identification of a transcription factor network regulating anti-TNF mediated IL10 expression in human CD4+ T cells. *Discovery immunology*, 3(1), kyae013.

Zhang Y, et al. (2024) MSC-derived mitochondria promote axonal regeneration via Atf3 gene up-regulation by ROS induced DNA double strand breaks at transcription initiation region. *Cell communication and signaling : CCS*, 22(1), 240.

Tremblay BJM, et al. (2024) Interplay between coding and non-coding regulation drives the Arabidopsis seed-to-seedling transition. *Nature communications*, 15(1), 1724.