

Resource Summary Report

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FindPeaks

RRID:SCR_010857

Type: Tool

Proper Citation

FindPeaks (RRID:SCR_010857)

Resource Information

URL: <http://sourceforge.net/apps/mediawiki/vancouvershorttr/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: 20260801T190417+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

Usage and Citation Metrics

We found 328 mentions in open access literature.

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

Kama Y, et al. (2026) Notch interaction with RUNX factors regulates initiation of the T-lineage program. *The Journal of experimental medicine*, 223(2).

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

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

Ozturk H, et al. (2025) High PRMT5 levels, maintained by KEAP1 inhibition, drive chemoresistance in high-grade serous ovarian cancer. *The Journal of clinical investigation*, 135(6).

Wang G, et al. (2025) Machine learning-based prediction reveals kinase MAP4K4 regulates neutrophil differentiation through phosphorylating apoptosis-related proteins. *PLoS computational biology*, 21(3), e1012877.

Sek K, et al. (2025) Tumor site-directed A1R expression enhances CAR T cell function and improves efficacy against solid tumors. *Nature communications*, 16(1), 6123.

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.

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

Kaarijärvi R, et al. (2025) Matrix stiffness modulates androgen response genes and chromatin state in prostate cancer. *NAR cancer*, 7(1), zcaf010.

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

Lee HK, et al. (2025) STAT5B leukemic mutations, altering SH2 tyrosine 665, have opposing impacts on immune gene programs. *Life science alliance*, 8(7).

Chang X, et al. (2025) ZMYND11 functions in bimodal regulation of latent genes and brain-like splicing to safeguard corticogenesis. *Nature communications*, 16(1), 9010.

Yang HJ, et al. (2025) ACTL6A depletion induces KLF4-mediated anti-tumorigenic effects in colorectal cancer. *Cell death & disease*, 16(1), 653.

Bonilla B, et al. (2025) esBAF and INO80C fine-tune subcompartments and differentially regulate enhancer-promoter interactions. *bioRxiv : the preprint server for biology*.

Lee EC, et al. (2025) 3D enhancer architecture coordinated by CTCF determines immune-related gene expression patterns via RNA polymerase II pause-release in CD4+ T cells. *Nucleic acids research*, 53(22).

Stüve P, et al. (2025) ACC1 is a dual metabolic-epigenetic regulator of Treg stability and immune tolerance. *Molecular metabolism*, 94, 102111.

Van de Velde S, et al. (2025) Med14 phosphorylation shapes genomic response to GLP-1 Agonist. *bioRxiv : the preprint server for biology*.

Schneider Y, et al. (2025) Vimentin network dysregulation mediates neurite deficits in SNCA duplication Parkinson's patient-derived midbrain neurons. *Science advances*, 11(23), eadq2742.

Suryavanshi P, et al. (2025) Dendritic beading during early brain development impairs signal transmission and synaptic plasticity. *Acta neuropathologica communications*, 13(1), 212.

Ogikubo K, et al. (2025) OCT-2 Is Associated With Pro-Metastatic Epigenomic Properties of Triple-Negative Breast Cancer Cells. *Cancer science*, 116(8), 2150.