

Resource Summary Report

Generated by [ASWG](#) on Aug 4, 2026

Pscan-ChIP

RRID:SCR_010885

Type: Tool

Proper Citation

Pscan-ChIP (RRID:SCR_010885)

Resource Information

URL: http://159.149.160.51/pscan_chip_dev/

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Description: Web server that, starting from a collection of genomic regions derived from a ChIP-Seq experiment, scans them using motif descriptors like JASPAR or TRANSFAC position-specific frequency matrices, or descriptors uploaded by users, and it evaluates both motif enrichment and positional bias within the regions according to different measures and criteria.

Abbreviations: Pscan-ChIP

Resource Type: data analysis service, service resource, analysis service resource, production service resource

Defining Citation: [PMID:23748563](#), [DOI:10.1093/nar/gkt448](#)

Keywords: bio.tools

Resource Name: Pscan-ChIP

Resource ID: SCR_010885

Alternate IDs: biotools:pscanchip, OMICS_00490

Alternate URLs: <https://bio.tools/pscanchip>, <https://sources.debian.org/src/pscan-chip/>

Record Creation Time: 20220129T080301+0000

Record Last Update: 20260804T043435+0000

Ratings and Alerts

No rating or validation information has been found for Pscan-ChIP.

No alerts have been found for Pscan-ChIP.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

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

Pan M, et al. (2025) Bone morphogenetic protein (BMP) signaling determines neuroblastoma cell fate and sensitivity to retinoic acid. Nature communications, 16(1), 2036.

Chavan A, et al. (2025) Epigenetic regulation of MED12: a key contributor to the leukemic chromatin landscape and transcriptional dysregulation. Epigenetics & chromatin, 18(1), 44.

Hui-Yuen J, et al. (2023) B lymphocytes in treatment-naive paediatric patients with lupus are epigenetically distinct from healthy children. Lupus science & medicine, 10(1).

Tarallo R, et al. (2017) The nuclear receptor ER α engages AGO2 in regulation of gene transcription, RNA splicing and RISC loading. Genome biology, 18(1), 189.

Nault R, et al. (2017) Loss of liver-specific and sexually dimorphic gene expression by aryl hydrocarbon receptor activation in C57BL/6 mice. PloS one, 12(9), e0184842.