

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

DBChIP

RRID:SCR_010872

Type: Tool

Proper Citation

DBChIP (RRID:SCR_010872)

Resource Information

URL: <http://pages.cs.wisc.edu/~kliang/DBChIP/>

Proper Citation: DBChIP (RRID:SCR_010872)

Description: Detects differential binding of transcription factors with ChIP-seq.

Abbreviations: DBChIP

Resource Type: software resource

Resource Name: DBChIP

Resource ID: SCR_010872

Alternate IDs: OMICS_00470

Record Creation Time: 20220129T080301+0000

Record Last Update: 20260725T190706+0000

Ratings and Alerts

No rating or validation information has been found for DBChIP.

No alerts have been found for DBChIP.

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 [nidm-terms](#) .

Hu S, et al. (2020) ncHMR detector: a computational framework to systematically reveal non-classical functions of histone modification regulators. *Genome biology*, 21(1), 48.

Lun AT, et al. (2016) csaw: a Bioconductor package for differential binding analysis of ChIP-seq data using sliding windows. *Nucleic acids research*, 44(5), e45.

Lopez-Anido C, et al. (2015) Differential Sox10 genomic occupancy in myelinating glia. *Glia*, 63(11), 1897.

Myers KS, et al. (2015) Defining bacterial regulons using ChIP-seq. *Methods (San Diego, Calif.)*, 86, 80.

Beauchene NA, et al. (2015) Impact of Anaerobiosis on Expression of the Iron-Responsive Fur and RyhB Regulons. *mBio*, 6(6), e01947.