Resource Summary Report

Generated by kravitz2 on Jul 31, 2026

diChIPMunk

RRID:SCR_010879
Type: Tool

Proper Citation

diChIPMunk (RRID:SCR_010879)

Resource Information

URL: http://autosome.ru/dichipmunk/

Proper Citation: diChIPMunk (RRID:SCR_010879)

Description: Software for motif discovery using dinucleotide position weight matrices (PWMs).

Abbreviations: diChIPMunk

Synonyms: diChIPMunk - motif discovery using dinucleotide PWMs

Resource Type: software resource

Defining Citation: PMID:23427986

Keywords: positional weight matrix, motif

Funding: Dynasty Foundation ;
Russian Foundation for Basic Research ;
Program Cell and Molecular Biology of the Presidium of Russian Academy of Sciences ;
Russian Federation Government

Resource Name: diChIPMunk

Resource ID: SCR_010879

Alternate IDs: OMICS_00481

Record Creation Time: 20220129T080301+0000

Record Last Update: 20260725T190706+0000

Ratings and Alerts

No rating or validation information has been found for diChIPMunk.

No alerts have been found for diChIPMunk.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 3 mentions in open access literature.

Listed below are recent publications. The full list is available at kravitz2 .

Seo SW, et al. (2017) Revealing genome-scale transcriptional regulatory landscape of OmpR highlights its expanded regulatory roles under osmotic stress in Escherichia coli K-12 MG1655. Scientific reports, 7(1), 2181.

Kulakovskiy IV, et al. (2016) HOCOMOCO: expansion and enhancement of the collection of transcription factor binding sites models. Nucleic acids research, 44(D1), D116.

Zolotarev N, et al. (2016) Architectural proteins Pita, Zw5,and ZIPIC contain homodimerization domain and support specific long-range interactions in Drosophila. Nucleic acids research, 44(15), 7228.