

Resource Summary Report

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MAnorm

RRID:SCR_010869

Type: Tool

Proper Citation

MAnorm (RRID:SCR_010869)

Resource Information

URL: <http://bcb.dfci.harvard.edu/~gcyuan/MAnorm/MAnorm.htm>

Proper Citation: MAnorm (RRID:SCR_010869)

Description: A robust software package for quantitative comparison of ChIP-Seq data sets.

Abbreviations: MAnorm

Synonyms: Manorm - a robust model for quantitative comparison of ChIP-Seq data sets

Resource Type: software resource

Defining Citation: [PMID:22424423](#)

Keywords: bio.tools

Funding:

Resource Name: MAnorm

Resource ID: SCR_010869

Alternate IDs: OMICS_00467, biotools:manorm

Alternate URLs: <https://bio.tools/manorm>

Record Creation Time: 20220129T080301+0000

Record Last Update: 20250410T070026+0000

Ratings and Alerts

No rating or validation information has been found for MAnorm.

No alerts have been found for MAnorm.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 94 mentions in open access literature.

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

Maezawa S, et al. (2025) Site-specific DNA demethylation during spermatogenesis presets the sites of nucleosome retention in mouse sperm. *bioRxiv : the preprint server for biology*.

Zhou L, et al. (2024) Mina53 demethylates histone H4 arginine 3 asymmetric dimethylation to regulate neural stem/progenitor cell identity. *Nature communications*, 15(1), 10227.

Poza-Viejo L, et al. (2024) Brassica rapa CURLY LEAF is a major H3K27 methyltransferase regulating flowering time. *Planta*, 260(1), 27.

Lee BK, et al. (2024) Dynamic and distinct histone modifications facilitate human trophoblast lineage differentiation. *Scientific reports*, 14(1), 4505.

Wen C, et al. (2024) MCM8 interacts with DDX5 to promote R-loop resolution. *The EMBO journal*, 43(14), 3044.

John E, et al. (2024) Regulatory insight for a Zn2Cys6 transcription factor controlling effector-mediated virulence in a fungal pathogen of wheat. *PLoS pathogens*, 20(9), e1012536.

Feng Y, et al. (2024) The roles of DNA methylation on pH dependent i-motif (iM) formation in rice. *Nucleic acids research*, 52(3), 1243.

Kim M, et al. (2024) The transcriptional regulatory network modulating human trophoblast stem cells to extravillous trophoblast differentiation. *Nature communications*, 15(1), 1285.

Zhu Y, et al. (2024) ZBTB7B is a permissive regulator of hepatocellular carcinoma initiation by repressing c-Jun expression and function. *Cell death & disease*, 15(1), 55.

Xiang G, et al. (2024) JMnorm: a novel joint multi-feature normalization method for integrative and comparative epigenomics. *Nucleic acids research*, 52(2), e11.

Du Y, et al. (2024) ASH1L in Hepatoma Cells and Hepatic Stellate Cells Promotes Fibrosis-Associated Hepatocellular Carcinoma by Modulating Tumor-Associated Macrophages. *Advanced science* (Weinheim, Baden-Wurttemberg, Germany), 11(45), e2404756.

Jiang Z, et al. (2024) Ferroptosis in Osteocytes as a Target for Protection Against Postmenopausal Osteoporosis. *Advanced science* (Weinheim, Baden-Wurttemberg, Germany), 11(12), e2307388.

Puerto M, et al. (2024) The zinc-finger protein Z4 cooperates with condensin II to regulate somatic chromosome pairing and 3D chromatin organization. *Nucleic acids research*, 52(10), 5596.

Veerappa AM, et al. (2024) CloudATAC: a cloud-based framework for ATAC-Seq data analysis. *Briefings in bioinformatics*, 25(Supplement_1).

Templeton CW, et al. (2024) HPV induced R-loop formation represses innate immune gene expression while activating DNA damage repair pathways. *PLoS pathogens*, 20(8), e1012454.

Du K, et al. (2024) The chromatin remodeling factor OsINO80 promotes H3K27me3 and H3K9me2 deposition and maintains TE silencing in rice. *Nature communications*, 15(1), 10919.

Li J, et al. (2024) ZmELP1, an Elongator complex subunit, is required for the maintenance of histone acetylation and RNA Pol II phosphorylation in maize kernels. *Plant biotechnology journal*, 22(5), 1251.

Yang Z, et al. (2024) Tumor-Associated Monocytes Reprogram CD8+ T Cells into Central Memory-Like Cells with Potent Antitumor Effects. *Advanced science* (Weinheim, Baden-Wurttemberg, Germany), 11(16), e2304501.

Uneme Y, et al. (2024) Morc1 reestablishes H3K9me3 heterochromatin on piRNA-targeted transposons in gonocytes. *Proceedings of the National Academy of Sciences of the United States of America*, 121(13), e2317095121.

Smith AL, et al. (2024) BET inhibition reforms the immune microenvironment and alleviates T cell dysfunction in chronic lymphocytic leukemia. *JCI insight*, 9(10).