

Resource Summary Report

Generated by [RRID](#) on Sep 10, 2026

NOrMAL

RRID:SCR_010889

Type: Tool

Proper Citation

NOrMAL (RRID:SCR_010889)

Resource Information

URL: <http://www.cs.ucr.edu/~polishka/>

Proper Citation: NOrMAL (RRID:SCR_010889)

Description: A command line software tool for accurate placing of the nucleosomes using a Modified Gaussian Mixture Model. It was designed to resolve overlapping nucleosomes and extract extra information (fuzziness, probability, etc.) of nucleosome placement. To achieve this goal the tool clusters the input tags according to Nucleosome Model (see the paper for detailed description) using EM learning process. The tool is written in C++. There are no special requirements except for g++ compiler and *nix environment to compile and use the tool. It was checked to compile using g++ compiler under Ubuntu 11.04 and Mac OS X 10.6

Abbreviations: NOrMAL

Synonyms: NOrMAL: Accurate Nucleosome Positioning using a Modified Gaussian Mixture Model

Resource Type: software resource

Keywords: bio.tools

Availability: Free for academic use

Resource Name: NOrMAL

Resource ID: SCR_010889

Alternate IDs: OMICS_00504, biotools:normal

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

Record Creation Time: 20220129T080301+0000

Record Last Update: 20260905T132649+0000

Ratings and Alerts

No rating or validation information has been found for NOrMAL.

No alerts have been found for NOrMAL.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 84 mentions in open access literature.

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

Pourbabaki R, et al. (2026) Genetic polymorphisms of GSTM1 and GSTT1 genes: effects on susceptibility to formaldehyde-induced hematotoxicity. Molecular biology research communications, 15(1), 71.

Zeng X, et al. (2025) Stochastic-based learning for image classification in chest X-ray diagnosis. Digital health, 11, 20552076251361745.

Chun J, et al. (2025) Genome-wide association and selection studies for pod dehiscence resistance in the USDA soybean germplasm collection. PloS one, 20(3), e0318815.

Ak?n T, et al. (2025) Phlegmonous appendicitis as a variant of uncomplicated appendicitis. Scientific reports, 15(1), 4387.

Nir I, et al. (2025) The Biology, Microclimate, and Geology of a Distinctive Ecosystem Within the Sandstone of Hyper-Arid Timna Valley, Israel. Environmental microbiology reports, 17(5), e70188.

Huang X, et al. (2025) The mechanism by which MALAT1/CREG1 regulates premature rupture of fetal membrane through autophagy mediated differentiation of amniotic fibroblasts. Non-coding RNA research, 13, 29.

Alves LKS, et al. (2025) Performance, carcass, and pork traits in barrows and gilts slaughtered over 130 kg: insights from a Brazilian perspective. Translational animal science, 9, txaf023.

Moiz S, et al. (2025) Differential Expression of miRNAs Between Young-Onset and Late-Onset Indian Colorectal Carcinoma Patients. Non-coding RNA, 11(1).

Bakhshalinejad R, et al. (2025) Comparative efficacy of hydroxychloride and organic sources of zinc, copper, and manganese on hatching eggs, embryo and hatchlings attributes. Poultry science, 104(2), 104728.

Gencer G, et al. (2025) Advanced retinal disease detection from OCT images using a hybrid squeeze and excitation enhanced model. PloS one, 20(2), e0318657.

Siqueiros J, et al. (2025) Influence of loud auditory noise on postural stability in autistic children: an exploratory study. *Scientific reports*, 15(1), 19882.

Takahashi S, et al. (2025) Flexible school attendance strategies for epidemic outbreaks: simulation-based decision support. *BMC public health*, 25(1), 4205.

Jannat FE, et al. (2025) OCT-SelfNet: a self-supervised framework with multi-source datasets for generalized retinal disease detection. *Frontiers in big data*, 8, 1609124.

Powroznik P, et al. (2025) Residual self-attention vision transformer for detecting acquired vitelliform lesions and age-related macular drusen. *Scientific reports*, 15(1), 17107.

Srinivasan A, et al. (2025) Encoding of speech modes and loudness in ventral precentral gyrus. *bioRxiv : the preprint server for biology*.

Deyhle RT, et al. (2025) Development of a method for evaluation of absorbed doses in preclinical in vivo γ CT. *Radiation protection dosimetry*, 201(13-14), 868.

Bissacco J, et al. (2025) CSF phospho-tau levels at Parkinson's disease onset predict the risk for development of motor complications. *Journal of neurology*, 272(9), 573.

Pakarinen S, et al. (2025) Workload management measures for supporting nuclear industry main control room operators and emergency response organization personnel during crises-a scoping review. *Industrial health*, 63(3), 214.

Qin H, et al. (2024) A novel LGALS1-depended and immune-associated fatty acid metabolism risk model in acute myeloid leukemia stem cells. *Cell death & disease*, 15(7), 482.

Asegu LM, et al. (2024) The economic burden of nosocomial infections for hospitals: evidence from Germany. *BMC infectious diseases*, 24(1), 1294.