

Resource Summary Report

Generated by ASWG on Aug 2, 2026

ROSE

RRID:SCR_017390

Type: Tool

Proper Citation

ROSE (RRID:SCR_017390)

Resource Information

URL: http://younglab.wi.mit.edu/super_enhancer_code.html

Proper Citation: ROSE (RRID:SCR_017390)

Description: To create stitched enhancers, and to separate super enhancers from typical enhancers using sequencing data given file of previously identified constituent enhancers .

Abbreviations: ROSE

Synonyms: Rank Ordering of Super Enhancers, RANK ORDERING OF SUPER-ENHANCERS

Resource Type: software resource, software application, data processing software

Keywords: Stitched, enhancer, separate, super, sequencing, data, file, identified, previously, constituent

Availability: Free, Freely available

Resource Name: ROSE

Resource ID: SCR_017390

License: MIT X11 License

Record Creation Time: 20220129T080335+0000

Record Last Update: 20260801T190836+0000

Ratings and Alerts

No rating or validation information has been found for ROSE.

No alerts have been found for ROSE.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 56 mentions in open access literature.

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

Renatino Canevarolo R, et al. (2025) Epigenetic Plasticity Drives Carcinogenesis and Multi-Therapy Resistance in Multiple Myeloma. Research square.

Lang JD, et al. (2025) Super-enhancers and efficacy of triptolide in small cell carcinoma of the ovary hypercalcemic type. iScience, 28(2), 111770.

Cigrang M, et al. (2025) Pan-inhibition of super-enhancer-driven oncogenic transcription by next-generation synthetic ecteinascidins yields potent anti-cancer activity. Nature communications, 16(1), 512.

Zamuner FT, et al. (2025) Epigenetic profiling reveals key super-enhancer networks driving oncogenesis in HPV-positive HNSCC. iScience, 28(11), 113911.

Liksitwatanakul P, et al. (2025) Chemical Perturbations Impacting Histone Acetylation Govern Colorectal Cancer Differentiation. Gastroenterology.

Kaufman ME, et al. (2024) Characterizing Relationships between T-cell Inflammation and Outcomes in Patients with High-Risk Neuroblastoma According to Mesenchymal and Adrenergic Signatures. Cancer research communications, 4(8), 2255.

Prutsch N, et al. (2024) STAT3 couples activated tyrosine kinase signaling to the oncogenic core transcriptional regulatory circuitry of anaplastic large cell lymphoma. Cell reports. Medicine, 5(3), 101472.

Abeywardana T, et al. (2024) Regulation of Enhancers by SUMOylation Through TFAP2C Binding and Recruitment of HDAC Complex to the Chromatin. Research square.

Shan L, et al. (2024) SP1 undergoes phase separation and activates RGS20 expression through super-enhancers to promote lung adenocarcinoma progression. Proceedings of the National Academy of Sciences of the United States of America, 121(29), e2401834121.

Wei X, et al. (2024) Super-enhancer-driven ZFP36L1 promotes PD-L1 expression in infiltrative gastric cancer. eLife, 13.

Weng Y, et al. (2024) Zfp260 choreographs the early stage osteo-lineage commitment of skeletal stem cells. Nature communications, 15(1), 10186.

Obinata D, et al. (2024) Patient-derived castration-resistant prostate cancer model revealed CTBP2 upregulation mediated by OCT1 and androgen receptor. BMC cancer, 24(1), 554.

Likasitwatanakul P, et al. (2024) Chemical perturbations impacting histone acetylation govern colorectal cancer differentiation. *bioRxiv : the preprint server for biology*.

Kim M, et al. (2023) Super-enhancer-associated transcription factors collaboratively regulate trophoblast-active gene expression programs in human trophoblast stem cells. *Nucleic acids research*, 51(8), 3806.

Hamamoto R, et al. (2023) Analysis of super-enhancer using machine learning and its application to medical biology. *Briefings in bioinformatics*, 24(3).

Tanaka M, et al. (2023) HEY1-NCOA2 expression modulates chondrogenic differentiation and induces mesenchymal chondrosarcoma in mice. *JCI insight*, 8(10).

Youngblood MW, et al. (2023) Super-enhancer hijacking drives ectopic expression of hedgehog pathway ligands in meningiomas. *Nature communications*, 14(1), 6279.

Tanaka M, et al. (2023) ASPSCR1::TFE3 orchestrates the angiogenic program of alveolar soft part sarcoma. *Nature communications*, 14(1), 1957.

Mohan DR, et al. (2023) β -Catenin-Driven Differentiation Is a Tissue-Specific Epigenetic Vulnerability in Adrenal Cancer. *Cancer research*, 83(13), 2123.

Singh DK, et al. (2023) 5-Azacytidine- and retinoic-acid-induced reprogramming of DCCs into dormancy suppresses metastasis via restored TGF- β -SMAD4 signaling. *Cell reports*, 42(6), 112560.