

Resource Summary Report

Generated by [kravitz2](#) on Sep 4, 2026

iceLogo

RRID:SCR_012137

Type: Tool

Proper Citation

iceLogo (RRID:SCR_012137)

Resource Information

URL: <https://code.google.com/p/icelogo/>

Proper Citation: iceLogo (RRID:SCR_012137)

Description: Software that builds on probability theory to visualize significant conserved sequence patterns in multiple peptide sequence alignments against background (reference) sequence sets that can be tailored to the studied system and the used protocol.

Resource Type: software resource

Defining Citation: [PMID:19876014](#)

Keywords: standalone software, web app, bio.tools

Availability: Apache License

Resource Name: iceLogo

Resource ID: SCR_012137

Alternate IDs: biotools:icelogo, OMICS_05885

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

Record Creation Time: 20220129T080308+0000

Record Last Update: 20260903T235142+0000

Ratings and Alerts

No rating or validation information has been found for iceLogo.

No alerts have been found for iceLogo.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 185 mentions in open access literature.

Listed below are recent publications. The full list is available at [kravitz2](#) .

Zhang H, et al. (2026) Identification of protein lysine lactylation and potential targets in esophageal squamous cell carcinoma. *Translational cancer research*, 15(5), 404.

Zabala-Letona A, et al. (2026) Polyamine-dependent metabolic shielding regulates alternative splicing. *Nature*, 651(8106), 819.

Cho H, et al. (2026) Glucocorticoid-Induced Proteome and Phosphoproteome Changes in Breast Cancer Cell Lines. *Journal of proteome research*, 25(4), 2110.

Fu Q, et al. (2026) Identification of the regulatory elements and protein substrates of lysine acetoacetylation. *eLife*, 14.

Cho H, et al. (2026) Quantitative proteomics and phosphoproteomics reveal glucocorticoid stimulation of TLR and Rho GTPase signaling in neutrophil-like cells. *Genome biology*, 27(1).

Boswell EL, et al. (2026) An AI-Ready Phosphorylation Meta-Analysis for *Saccharomyces cerevisiae*. *Journal of proteome research*, 25(6), 2915.

Yee DH, et al. (2026) Pepsin Digestion for Proteomic Studies of the Human Hair Shaft. *Rapid communications in mass spectrometry : RCM*, 40(6), e70018.

Zhang J, et al. (2026) d-lactate drives lysine d-lactylation to regulate metabolism in *Escherichia coli*. *The Journal of biological chemistry*, 302(5), 111374.

Wang J, et al. (2026) EBV infection and HLA-DR15 jointly drive multiple sclerosis by myelin peptide presentation. *Cell*.

Barbosa da Silva E, et al. (2025) The impact of pH on proteolytic activity in wound fluid: Implications for acid therapy. *The Journal of biological chemistry*, 301(10), 110723.

Silva EB, et al. (2025) Enhancing schistosomiasis drug discovery approaches with optimized proteasome substrates. *Protein science : a publication of the Protein Society*, 34(6), e70180.

Yan M, et al. (2025) Lactylated Proteomic Analysis Reveals Functional Implications of Lysine Lactylation In Asthenozoospermia. *Molecular & cellular proteomics : MCP*, 24(12), 101439.

Li L, et al. (2025) PRMT5 Maintains Homeostasis of the Intestinal Epithelium by Modulating Cell Proliferation and Survival. *Advanced science (Weinheim, Baden-*

Wurttemberg, Germany), 12(12), e2415559.

Roumbo L, et al. (2025) The MAST kinase KIN-4 carries out mitotic entry functions of Greatwall in *C. elegans*. *The EMBO journal*, 44(7), 1943.

Steinberger S, et al. (2025) Proteasome caspase-like activity regulates stress granules and proteasome condensates. *Frontiers in cell and developmental biology*, 13, 1570499.

Liu G, et al. (2025) Deciphering the Metabolic Impact and Clinical Relevance of N-Glycosylation in Colorectal Cancer through Comprehensive Glycoproteomic Profiling. *Advanced science* (Weinheim, Baden-Wurttemberg, Germany), 12(21), e2415645.

de Groot R, et al. (2025) Cleavage of Cartilage Oligomeric Matrix Protein (COMP) by ADAMTS4 generates a neoepitope associated with osteoarthritis and other forms of degenerative joint disease. *Matrix biology : journal of the International Society for Matrix Biology*, 135, 106.

Tyl MD, et al. (2025) Infection-induced lysine lactylation enables herpesvirus immune evasion. *Science advances*, 11(2), eads6215.

Chambwe N, et al. (2025) Cellular heterogeneity and therapeutic response profiling of human IDH + glioma stem cell cultures. *Scientific reports*, 16(1), 3177.

Kauko O, et al. (2025) Diverse oncogenes use common mechanisms to drive growth of major forms of human cancer. *Science advances*, 11(34), eadt1798.