Resource Summary Report

Generated by dkNET on May 17, 2025

<u>lumi</u>

RRID:SCR_012781

Type: Tool

Proper Citation

Iumi (RRID:SCR_012781)

Resource Information

URL: http://bioconductor.org/packages/release/bioc/html/lumi.html

Proper Citation: lumi (RRID:SCR_012781)

Description: Software that provides an integrated solution for the Illumina microarray data

analysis.

Abbreviations: lumi

Resource Type: software resource

Keywords: bio.tools

Funding:

Resource Name: lumi

Resource ID: SCR_012781

Alternate IDs: biotools:lumi, OMICS_00770

Alternate URLs: https://bio.tools/lumi

Record Creation Time: 20220129T080312+0000

Record Last Update: 20250420T014619+0000

Ratings and Alerts

No rating or validation information has been found for lumi.

No alerts have been found for lumi.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 311 mentions in open access literature.

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

Kim SJ, et al. (2025) Disruption of bioenergetics enhances the radio-sensitivity of patient-derived glioblastoma tumorspheres. Translational oncology, 51, 102197.

Andersen GT, et al. (2024) Multi-bioinformatics revealed potential biomarkers and repurposed drugs for gastric adenocarcinoma-related gastric intestinal metaplasia. NPJ systems biology and applications, 10(1), 127.

Wlaz?owski G, et al. (2024) Fermionic quantum turbulence: Pushing the limits of high-performance computing. PNAS nexus, 3(5), pgae160.

Liu R, et al. (2024) In vitro assessment of thyroid peroxidase inhibition by chemical exposure: comparison of cell models and detection methods. Archives of toxicology, 98(8), 2631.

Graça M, et al. (2024) Distributed transformer for high order epistasis detection in large-scale datasets. Scientific reports, 14(1), 14579.

Park J, et al. (2024) Comparison of Glioblastoma Cell Culture Platforms Based on Transcriptional Similarity with Paired Tissue. Pharmaceuticals (Basel, Switzerland), 17(4).

Nicze M, et al. (2024) The Current and Promising Oral Delivery Methods for Protein- and Peptide-Based Drugs. International journal of molecular sciences, 25(2).

Chen XF, et al. (2024) Integrative high-throughput enhancer surveying and functional verification divulges a YY2-condensed regulatory axis conferring risk for osteoporosis. Cell genomics, 4(3), 100501.

Wu Y, et al. (2024) Flu-CED: A comparative transcriptomics database of influenza virus-infected human and animal models. Animal models and experimental medicine, 7(6), 881.

Irani Shemirani M, et al. (2024) Transcriptional markers classifying Escherichia coli and Staphylococcus aureus induced sepsis in adults: A data-driven approach. PloS one, 19(7), e0305920.

Jang J, et al. (2024) Endocardial HDAC3 is required for myocardial trabeculation. Nature communications, 15(1), 4166.

Dor H, et al. (2024) Schizophrenia Biomarkers: Blood Transcriptome Suggests Two Molecular Subtypes. Neuromolecular medicine, 26(1), 50.

Habgood-Coote D, et al. (2023) Diagnosis of childhood febrile illness using a multi-class blood RNA molecular signature. Med (New York, N.Y.), 4(9), 635.

Li Z, et al. (2023) The EstroGene Database Reveals Diverse Temporal, Context-Dependent, and Bidirectional Estrogen Receptor Regulomes in Breast Cancer. Cancer research, 83(16), 2656.

Meligova AK, et al. (2023) ER?1 Sensitizes and ER?2 Desensitizes ER?-Positive Breast Cancer Cells to the Inhibitory Effects of Tamoxifen, Fulvestrant and Their Combination with All-Trans Retinoic Acid. International journal of molecular sciences, 24(4).

Zeng R, et al. (2023) Investigating Causality and Shared Genetic Architecture between Neurodegenerative Disorders and Inflammatory Bowel Disease. Aging and disease, 14(4), 1349.

Seddon AR, et al. (2023) Site-specific decreases in DNA methylation in replicating cells following exposure to oxidative stress. Human molecular genetics, 32(4), 632.

Nomoto H, et al. (2023) Potential usage of anterior nasal sampling in clinical practice with three rapid antigen tests for SARS-CoV-2. Journal of infection and chemotherapy: official journal of the Japan Society of Chemotherapy, 29(1), 15.

Krum-Hansen S, et al. (2023) Associations of breast cancer related exposures and gene expression profiles in normal breast tissue-The Norwegian Women and Cancer normal breast tissue study. Cancer reports (Hoboken, N.J.), 6(4), e1777.

Jay F, et al. (2023) A contemporary reassessment of the enhanced transient expression system based on the tombusviral silencing suppressor protein P19. The Plant journal: for cell and molecular biology, 113(1), 186.