

# Resource Summary Report

Generated by SPARC SAWG on Sep 20, 2026

## Kyoto Encyclopedia of Genes and Genomes Expression Database

RRID:SCR\_001120

Type: Tool

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### Proper Citation

Kyoto Encyclopedia of Genes and Genomes Expression Database (RRID:SCR\_001120)

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### Resource Information

**URL:** <http://www.genome.jp/kegg/expression/>

**Proper Citation:** Kyoto Encyclopedia of Genes and Genomes Expression Database (RRID:SCR\_001120)

**Description:** Database for mapping gene expression profiles to pathways and genomes. Repository of microarray gene expression profile data for *Synechocystis* PCC6803 (syn), *Bacillus subtilis* (bsu), *Escherichia coli* W3110 (ecj), *Anabaena* PCC7120 (ana), and other species contributed by the Japanese research community.

**Abbreviations:** KEGG Expression Database

**Synonyms:** Kyoto Encyclopedia of Genes and Genomes Expression Database

**Resource Type:** data or information resource, data repository, database, service resource, storage service resource

**Defining Citation:** [PMID:9847135](#), [PMID:10592173](#)

**Keywords:** encyclopedia, endogenous, environment, enzyme, *escherichia coli*, exogenous, expression, family, functional, gene, genetic, *anabaena*, *bacillus subtilis*, biological system, biology, building block, cell, cellular, chemical, community, complex, genome, genomic, hierarchy, interaction, japanese, mapping, metabolic, metabolic pathway databases, microarray, molecular, molecular wiring, nomenclature, order, organism, ortholog, pathway, process, protein, reaction, research, sequence, specie, substance, *synechocystis*, FASEB list

**Availability:** Free, Available for download, Freely available

**Resource Name:** Kyoto Encyclopedia of Genes and Genomes Expression Database

**Resource ID:** SCR\_001120

**Alternate IDs:** r3d100011570, nif-0000-21234

**Alternate URLs:** <https://doi.org/10.17616/R3792T>

**Record Creation Time:** 20220129T080205+0000

**Record Last Update:** 20260919T194934+0000

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## Ratings and Alerts

No rating or validation information has been found for Kyoto Encyclopedia of Genes and Genomes Expression Database.

No alerts have been found for Kyoto Encyclopedia of Genes and Genomes Expression Database.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 1279 mentions in open access literature.

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

Yan X, et al. (2026) Activation of T Cell-Intrinsic p53 by Acetylation Elicits Antitumor Immunity to Boost Cancer Immunotherapy. *Cancer discovery*, 16(1), 155.

Li G, et al. (2026) Spatial Multiomics Analyses Reveal That Diabetes Promotes Pancreatic Cancer Progression by Stimulating Cholesterol-Induced Neutrophil Extracellular Trap Formation. *Cancer research*, 86(10), 2327.

Jiang YY, et al. (2026) Ze-qi Decoction Inhibits Neutrophil Extracellular Trap Formation to Suppress the Progression and Metastasis of Non-Small Cell Lung Cancer. *Integrative cancer therapies*, 25, 15347354261420750.

Kim DH, et al. (2026) Tumor Exosomal L1 Cell Adhesion Molecule Promotes Brain Metastasis of Lung Cancer. *Research (Washington, D.C.)*, 9, 1126.

Zink A, et al. (2026) Pluripotent stem-cell-based screening uncovers sildenafil as a mitochondrial disease therapy. *Cell*, 189(6), 1656.

Haberberger RV, et al. (2025) Transcriptomic and Histological Characterization of Telocytes in the Human Dorsal Root Ganglion. *The Journal of comparative neurology*, 533(3), e70044.

Menozzi E, et al. (2025) Exploring the relationship between GBA1 host genotype and gut microbiome in the GBA1 L444P/WT mouse model: implications for Parkinson's disease pathogenesis. *Frontiers in neuroscience*, 19, 1546203.

Hurst C, et al. (2025) Social Isolation Induces Sex-Specific Differences in Behavior and Gut Microbiota Composition in Stress-Sensitive Rats. *Brain and behavior*, 15(6), e70621.

Zhang T, et al. (2025) Intratumoral *Fusobacterium nucleatum* Recruits Tumor-Associated Neutrophils to Promote Gastric Cancer Progression and Immune Evasion. *Cancer research*, 85(10), 1819.

Pacheco-Fernandez T, et al. (2025) Protocol for identifying immune mediators during *Leishmania* infection in mice using metabolomic analysis. *STAR protocols*, 6(3), 103817.

Mahat DB, et al. (2025) Mutant p53 exploits enhancers to elevate immunosuppressive chemokine expression and impair immune checkpoint inhibitors in pancreatic cancer. *Immunity*, 58(7), 1688.

Huang J, et al. (2024) BBOX1 mediates metabolic reprogramming driven by hypoxia and participates in the malignant progress of high-grade serous ovarian cancer. *Biochimica et biophysica acta. Molecular cell research*, 1871(8), 119830.

Liang L, et al. (2024) Correlation between intestinal flora and GLP-1 receptor agonist dulaglutide in type 2 diabetes mellitus treatment-A preliminary longitudinal study. *iScience*, 27(5), 109784.

Wan S, et al. (2024) SPARC Stabilizes ApoE to Induce Cholesterol-Dependent Invasion and Sorafenib Resistance in Hepatocellular Carcinoma. *Cancer research*, 84(11), 1872.

Roy TA, et al. (2024) Discovery and validation of genes driving drug-intake and related behavioral traits in mice. *Genes, brain, and behavior*, 23(1), e12875.

Tabachnick-Cherny S, et al. (2024) Characterization of Immunosuppressive Myeloid Cells in Merkel Cell Carcinoma: Correlation with Resistance to PD-1 Pathway Blockade. *Clinical cancer research : an official journal of the American Association for Cancer Research*, 30(6), 1189.

Oljuskin T, et al. (2023) *Leishmania* major centrin knock-out parasites reprogram tryptophan metabolism to induce a pro-inflammatory response. *iScience*, 26(9), 107593.

Liu Y, et al. (2023) Immunogenetic Determinants of Susceptibility to Head and Neck Cancer in the Million Veteran Program Cohort. *Cancer research*, 83(3), 386.

Roy TA, et al. (2023) DISCOVERY AND VALIDATION OF GENES DRIVING DRUG-INTAKE AND RELATED BEHAVIORAL TRAITS IN MICE. *bioRxiv : the preprint server for biology*.

Iqbal J, et al. (2023) Differential regulation of hippocampal transcriptome by circulating estrogen. *Functional & integrative genomics*, 23(4), 309.