

Resource Summary Report

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GENCODE

RRID:SCR_014966

Type: Tool

Proper Citation

GENCODE (RRID:SCR_014966)

Resource Information

URL: <https://www.encodegenes.org>

Proper Citation: GENCODE (RRID:SCR_014966)

Description: Human and mouse genome annotation project which aims to identify all gene features in the human genome using computational analysis, manual annotation, and experimental validation.

Synonyms: ENCODE

Resource Type: data or information resource, dataset, portal, project portal

Defining Citation: [PMID:22955987](#)

Keywords: human, mouse, genome, annotation, sequence, gene features, bio.tools

Funding: NHGRI 5U54HG004555;
Wellcome Trust WT098051

Availability: Free

Resource Name: GENCODE

Resource ID: SCR_014966

Alternate IDs: biotools:GENCODE

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

Record Creation Time: 20220129T080323+0000

Record Last Update: 20260829T182844+0000

Ratings and Alerts

- Used for genome assembly by the Human Islet Research Network community.
Contact(s): [Golnaz Vahedi](#), [Klaus Kaestner](#) - Human Islets Research Network
<https://hirnetwork.org/>

No alerts have been found for GENCODE.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 8811 mentions in open access literature.

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

Horii M, et al. (2026) Clinical and Molecular Differences of Hypertensive Disorders During Pregnancy. Arteriosclerosis, thrombosis, and vascular biology, 46(3), e323457.

Balamurugan K, et al. (2026) 3D "Emboli" Culture Models Epithelial Breast Cancer Cell Oxidative Mitochondrial Metabolism with Relevance for Lung Metastasis. Cancer research communications, 6(3), 600.

Gadad SS, et al. (2026) X-linked cancer-associated polypeptide (XCP) from lncRNA1456 modulates PHF8 histone demethylase activity to regulate the epigenome, gene expression, and cellular pathways in breast cancer. Oncogene, 45(17), 1557.

Ye Y, et al. (2026) DCPS modulates TDP-43-linked neurodegeneration through P-body-mediated RNA decay. Neuron, 114(11), 1970.

Alvaro-Espinosa L, et al. (2026) MIF-Induced CD74+ Microglia and Macrophages Promote Progression of Brain Metastasis and Are Clinically Relevant across Central Nervous System Disorders. Cancer research, 86(13), 3249.

Sutcliffe MD, et al. (2026) Transcriptomic profiling of mouse mammary tumors enables prognostic and predictive biomarker discovery for human breast cancers. bioRxiv : the preprint server for biology.

Alsina M, et al. (2026) Anti-B-cell Maturation Antigen Chimeric Antigen Receptor T-cell Therapy bb21217 for Relapsed and Refractory Multiple Myeloma: Results from the Phase I CRB-402 Study. Cancer immunology research, 14(4), 528.

Sakrepatna Nagaraj A, et al. (2026) Ex Vivo Immuno-Oncology Platform Reveals Spatial T-cell Infiltration Patterns Linked to ATR Inhibition Responses in High-Grade Serous Ovarian Cancer. Cancer immunology research, 14(4), 625.

Zhao Y, et al. (2026) IDH1 Reprograms Nucleotide Metabolism by Inducing Chromatin Remodeling and DHODH Transcription to Drive Chemoresistance in Nasopharyngeal Carcinoma. Cancer research, 86(8), 2004.

Smolen C, et al. (2026) Protocol for identifying genetic modifiers of phenotypes in individuals with disease-associated variants. STAR protocols, 7(2), 104546.

Fahmy AM, et al. (2026) STING dampens the unfolded protein response to enable the presentation of self-antigens on MHC-I during inflammation. Cell reports, 45(6), 117497.

Swift SZ, et al. (2026) Integration of Short- and Long-Read RNA Sequencing Enables the Discovery of Circular RNAs. Cancer research, 86(5), 1300.

Centonze A, et al. (2026) A Plastic EMP1+ to LGR5+ Cell State Conversion as a Bypass to KRASG12D Pharmacologic Inhibition in Metastatic Colorectal Cancer. Cancer discovery, 16(2), 320.

Hasegawa K, et al. (2026) Age-related decline in nuclear envelope LINC complex drives neuronal aging via axon initial segment dysfunction. EMBO reports, 27(13), 3788.

Kuo HH, et al. (2026) Patient-derived organoids across cancers reveal conserved tumor heterogeneity and actionable therapeutic vulnerabilities. Science advances, 12(26), ead3351.

Suresh V, et al. (2026) Modelling EWS::FLI1 protein fluctuations reveal determinants of tumor plasticity in Ewing sarcoma. EMBO molecular medicine, 18(2), 646.

Eriksson J, et al. (2026) Single-Cell Lineage Tracing Uncovers Resistance Signatures and Sensitizing Strategies to FLT3 Inhibitors in Acute Myeloid Leukemia. Cancer research, 86(7), 1724.

Garralda E, et al. (2026) Broad Utility of Ultrasensitive Analysis of ctDNA Dynamics across Solid Tumors Treated with Immunotherapy. Clinical cancer research : an official journal of the American Association for Cancer Research, 32(2), 333.

Bernard MJ, et al. (2026) OGDHL Promotes Prostate Cancer Progression and Regulates Neuroendocrine Marker Expression and Nucleotide Abundance. Molecular cancer research : MCR, 24(5), 356.

Li C, et al. (2026) New Insights Into the Role of Polo-Like Kinase 3 in Lung Tumorigenesis. Journal of cellular biochemistry, 127(2), e70084.