Resource Summary Report

Generated by Kravitz on Sep 10, 2026

ZENODO

RRID:SCR_004129
Type: Tool

Proper Citation

ZENODO (RRID:SCR_004129)

Resource Information

URL: https://zenodo.org/

Proper Citation: ZENODO (RRID:SCR_004129)

Description: Repository for all research outputs from across all fields of science in any file format as well as both positive and negative results. They assign all publicly available uploads a Digital Object Identifier (DOI) to make the upload easily and uniquely citeable. They further support harvesting of all content via the OAI-PMH protocol. They promote peer-reviewed openly accessible research, and curate uploads. ZENODO allows users to create their own collection and accept or reject all uploads to it. They allow for uploading under a multitude of different licenses and access levels.

Abbreviations: ZENODO

Synonyms: Zenodo

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

Keywords: data set, software resource, video resource, audio track, image, poster, presentation, publication, digital preservation, digital archive, persistent identifier, digital object identifier, openaire orphan record repository

Funding: European Union FP7 OpenAIREplus 283595

Resource Name: ZENODO

Resource ID: SCR_004129

Alternate IDs: DOI:10.5281, nlx_158614, DOI:10.17616/R3QP53, DOI:10.25504/FAIRsharing.wy4egf, r3d100011858

Alternate URLs: https://doi.org/10.17616/R3QP53, https://doi.org/10.17616/r3QP53, https://doi.org/10.5281/, https://dx.doi.org/10.5281/,

https://fairsharing.org/10.25504/FAIRsharing.wy4egf, https://doi.org/10.17616/R35W56

Record Creation Time: 20220129T080222+0000

Record Last Update: 20260905T132515+0000

Ratings and Alerts

No rating or validation information has been found for ZENODO.

No alerts have been found for ZENODO.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 6074 mentions in open access literature.

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

Maeda M, et al. (2026) CRISPR screen of human pancreatic cancer xenografts identifies a KLF5 proliferation vulnerability through epigenetic modifiers NCAPD2 and MTHFD1. Molecular cancer, 25(1).

de Blas A, et al. (2026) Enhancing Proteasome Activity in T Cells Alleviates Exhaustion and Improves Antitumor Immunity. Cancer research, 86(12), 2896.

Hofvander J, et al. (2026) Transcriptomic Subgroups in Soft Tissue Tumors Correlate with Morphologic Subtype, Genomic Features, and Outcome. Clinical cancer research : an official journal of the American Association for Cancer Research, 32(9), 1825.

Min J, et al. (2026) AI-Powered Deep Visual Proteomics Reveals Critical Molecular Transitions in Pancreatic Cancer Precursors. Cancer discovery, 16(7), 1323.

Francette AM, et al. (2026) Evolutionary analysis of transcription elongation factors reveals conserved and lineage-specific regulatory domains. PLoS biology, 24(6), e3003855.

Hofer I, et al. (2026) An Extracellular Matrix-Producing Subset of Cancer-Associated Fibroblasts Drives Chemoresistance in Breast Cancer via SRC Activation and G0S2 Upregulation. Cancer research, 86(4), 1054.

Rutz A, et al. (2026) MultiMS2: A curated multi-modal, multi-energy spectral library for metabolomics. GigaScience, 15.

Mariancini A, et al. (2026) IL1R2 Deficiency Unleashes Neutrophil-Mediated Antitumor Potential in Sarcoma. Cancer immunology research, 14(5), 751.

Belanger K, et al. (2026) Cancer-Associated Fibroblasts Promote Epithelial-Mesenchymal Transition and Classical to Basal Subtype Shift in Pancreatic Cancer. Cancer research, 86(14), 3604.

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

Zhu Z, et al. (2026) A neurovascular survival axis for adult sympathetic neurons. Neuron.

Chen CX, et al. (2026) Continuous flash suppression of neural responses and population orientation coding in macaque V1. eLife, 14.

Koladiya A, et al. (2026) CellFuse Enables Multimodal Integration of Single-Cell and Spatial Proteomics Data for Systems-Level Analysis in Cancer. Cancer research, 86(14), 3379.

Balin S, et al. (2026) Single-cell analysis reveals corticosteroid-associated impairment of tumor-infiltrating NK cells in glioblastoma patients. Scientific reports.

Aggrey-Fynn JE, et al. (2026) Epigenetic context defines the transcriptional activity of canonical and noncanonical NF-??B signaling in pancreatic cancer. Cell death discovery, 12(1).

Williams W, et al. (2026) Chromosome segregation synchrony in S. pombe is noise limited and arises without positive feedback. The Journal of cell biology, 225(7).

Kern S, et al. (2026) Challenges in replay detection by TDLM in post-encoding resting state. eLife, 14.

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.

Izzo LT, et al. (2026) KLF4 Promotes a KRT13+ Hillock-Like State in Lung Squamous Cell Carcinoma. Cancer research.

Smith JW, et al. (2026) Direct optical activation of human IRE1 identifies unique patterns of transcriptional and post-transcriptional mRNA regulation in the unfolded protein response. The Journal of biological chemistry, 302(2), 111067.