

Resource Summary Report

Generated by [dkNET](#) on Aug 24, 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: 20260821T193723+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 [dkNET](#).

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- κ 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.