

Resource Summary Report

Generated by PRECISE-TBI on Jul 31, 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: software repository, service resource, data repository, software resource, catalog, data or information resource, database, 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

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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: 20260731T042701+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 4900 mentions in open access literature.

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

Rotem A, et al. (2026) Differentiation between regulated and disrupted growth arrests allows tailoring of effective treatments for antibiotic persistence. *Science advances*, 12(1), eadt6577.

Daniel MM, et al. (2026) Integrative Bioinformatics Reveals Novel Molecular Mechanisms and Therapeutic Targets in Acute Myeloid Leukaemia. *Journal of cellular and molecular medicine*, 30(1), e71007.

Bettinelli A, et al. (2026) Assessing the agreement of radiomic tools for dosiomics: A multi-software comparative study. *Medical physics*, 53(1), e70203.

Kot E, et al. (2026) Crystal structure of fatty acid thioesterase A bound by 129 fragments provides diverse development opportunities. *Pest management science*, 82(1), 151.

Martin-Gonzalez E, et al. (2026) Multi-Ancestry Epigenome-Wide Association Study of Asthma Exacerbations. *Allergy*, 81(1), 263.

Berselli A, et al. (2026) The effect of ionic strength on PETase enzymes: An experimental and computational study. *Protein science : a publication of the Protein Society*, 35(1), e70386.

Li X, et al. (2026) Talin-tensin3 interactions regulate fibrillar adhesion formation and tensin3 phase separation. *The Journal of cell biology*, 225(1).

Saviluoto A, et al. (2026) Association of Cerebral Oxygenation During Prehospital Anaesthesia and Functional Outcome: A Prospective, Observational Multi-Centre Cohort Study of 1014 Patients. *Acta anaesthesiologica Scandinavica*, 70(1), e70161.

- Zanini A, et al. (2026) Mapping functional homologies between human and marmoset brain networks using movie-driven ultra-high field fMRI. *Communications biology*, 9(1), 21.
- Andonie CR, et al. (2026) Comparative Analysis of Glucagon Receptor Agonists vs. Resmetirom in MASLD and MASH: Network Meta-Analysis of Clinical Trials. *Endocrinology, diabetes & metabolism*, 9(1), e70157.
- Yilmaz V, et al. (2026) Comparative Transcriptomic Profiling Identifies IL-1 β as a Non-Canonical Driver of Human Dendritic Cell Maturation. *Iranian journal of biotechnology*, 24(1), e4185.
- Zhao Z, et al. (2026) Multi-omics analysis unveils tumor heterogeneity and immunotherapy predictive model in breast cancer for precision medicine and early detection. *Neoplasia (New York, N.Y.)*, 71, 101260.
- Yan Y, et al. (2026) A novel mouse cell line model reveals the tumor intrinsic and immune characteristics of EGFR-mutant lung cancer. *Theranostics*, 16(2), 1006.
- Zappon E, et al. (2026) An efficient end-to-end computational framework for the generation of ECG calibrated volumetric models of human atrial electrophysiology. *Medical image analysis*, 107(Pt B), 103822.
- Freudenthal J, et al. (2026) A Novel Protistan Trait Database Reveals Functional Redundancy and Complementarity in Terrestrial Protists (Amoebozoa and Rhizaria). *Molecular ecology resources*, 26(1), e70064.
- Verna G, et al. (2026) A missense mutation in Muc2 promotes gut microbiome and metabolome-dependent colitis-associated tumorigenesis. *The Journal of clinical investigation*, 136(1).
- Widomski C, et al. (2026) Beyond the emm-type: fine-tuning Group A Streptococcus typing with Enn and Mrp. *Microbiology spectrum*, 14(1), e0204725.
- Michel M, et al. (2025) Noninvasive Multicancer Detection Using DNA Hypomethylation of LINE-1 Retrotransposons. *Clinical cancer research : an official journal of the American Association for Cancer Research*, 31(7), 1275.
- Kondi β T, et al. (2025) Shinyscreen: mass spectrometry data inspection and quality checking utility. *Journal of cheminformatics*, 17(1), 98.
- Roca MS, et al. (2025) Valproic acid improves the efficacy of oxaliplatin/fluoropyrimidine-based chemotherapy by targeting cancer stem cell via β -Catenin modulation in colorectal cancer. *Cell death & disease*, 16(1), 583.