

Resource Summary Report

Generated by [NIF](#) on Apr 9, 2025

SourceForge

RRID:SCR_004365

Type: Tool

Proper Citation

SourceForge (RRID:SCR_004365)

Resource Information

URL: <http://sourceforge.net/>

Proper Citation: SourceForge (RRID:SCR_004365)

Description: Web based service that offers software developers centralized online location to control and manage free and open source software projects. Open source software tool and business public software platform.

Resource Type: web service, software resource, data access protocol

Keywords: software, platform, tool, resource, catalog, community, project, , FASEB list

Funding:

Availability: Free, Freely available

Resource Name: SourceForge

Resource ID: SCR_004365

Alternate IDs: nlx_38463

License URLs: <http://slashdotmedia.com/terms-of-use/>

Record Creation Time: 20220129T080224+0000

Record Last Update: 20250409T060353+0000

Ratings and Alerts

No rating or validation information has been found for SourceForge.

No alerts have been found for SourceForge.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 885 mentions in open access literature.

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

Du Q, et al. (2025) Acylglycerol kinase inhibits macrophage anti-tumor activity via limiting mtDNA release and cGAS-STING-type I IFN response. *Theranostics*, 15(4), 1304.

Park JY, et al. (2025) Biological augmentation of anterior cruciate ligament reconstruction with bone marrow aspirate concentrate: a systematic review and meta-analysis of randomised controlled trials. *International orthopaedics*, 49(1), 35.

Bulduk BK, et al. (2025) High frequency of mitochondrial DNA rearrangements in the peripheral blood of adults with intellectual disability. *Journal of intellectual disability research : JIDR*, 69(2), 137.

Uesaka K, et al. (2024) Deciphering the genomes of motility-deficient mutants of *Vibrio alginolyticus* 138-2. *PeerJ*, 12, e17126.

Wu H, et al. (2024) Characterization of novel PHEX variants in X-linked hypophosphatemic rickets and genotype-PHEX activity correlation. *The Journal of clinical endocrinology and metabolism*.

Yi B, et al. (2024) Prognostic and clinical significance of tumor-associated macrophages in esophageal squamous cell carcinoma after surgery: do biomarkers and distributions matter? *Bioscience reports*, 44(4).

Foss CA, et al. (2024) Imaging tumor and ascites-associated macrophages in a mouse model of metastatic ovarian cancer. *EJNMMI research*, 14(1), 121.

Hu NQ, et al. (2024) Hypokalemic Periodic Paralysis Type 2 Due to SCN4A Val1105Met Mutation: A Case Study. *Cureus*, 16(1), e52063.

Warden CD, et al. (2024) Critical Differential Expression Assessment for Individual Bulk RNA-Seq Projects. *bioRxiv : the preprint server for biology*.

Li S, et al. (2024) A humanized monoclonal antibody targeting an ectonucleotidase rescues cardiac metabolism and heart function after myocardial infarction. *Cell reports. Medicine*,

5(11), 101795.

Wang X, et al. (2024) Clinicopathological and prognostic features of colorectal mucinous adenocarcinomas: a systematic review and meta-analysis. *BMC cancer*, 24(1), 1161.

Debernardi S, et al. (2024) Molecular characteristics of early-onset pancreatic ductal adenocarcinoma. *Molecular oncology*, 18(3), 677.

Schopfer LM, et al. (2024) Mass Spectrometry of Putrescine, Spermidine, and Spermine Covalently Attached to *Francisella tularensis* Universal Stress Protein and Bovine Albumin. *Biochemistry research international*, 2024, 7120208.

Selvaraju D, et al. (2024) Experimentally evolving *Drosophila erecta* populations may fail to establish an effective piRNA-based host defense against invading P-elements. *Genome research*, 34(3), 410.

Fu D, et al. (2024) Establishing a model composed of immune-related gene-modules to predict tumor immunotherapy response. *Scientific reports*, 14(1), 16630.

Martín Del Pico E, et al. (2024) FAIRsoft-a practical implementation of FAIR principles for research software. *Bioinformatics (Oxford, England)*, 40(8).

Yang M, et al. (2024) Antibiotic-Induced Gut Microbiota Dysbiosis Modulates Host Transcriptome and m6A Epitranscriptome via Bile Acid Metabolism. *Advanced science (Weinheim, Baden-Wurtemberg, Germany)*, 11(28), e2307981.

Galbusera F, et al. (2024) Image annotation and curation in radiology: an overview for machine learning practitioners. *European radiology experimental*, 8(1), 11.

Lo YL, et al. (2024) Modulating versatile pathways using a cleavable PEG shell and EGFR-targeted nanoparticles to deliver CRISPR-Cas9 and docetaxel for triple-negative breast cancer inhibition. *Archives of pharmacal research*, 47(10-11), 829.

Rong X, et al. (2024) Biocrusts Mediate the Niche Distribution and Diversity of Ammonia-Oxidizing Microorganisms in the Gurbantunggut Desert, Northwestern China. *Microbial ecology*, 87(1), 148.