

Resource Summary Report

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GitLab

RRID:SCR_013983

Type: Tool

Proper Citation

GitLab (RRID:SCR_013983)

Resource Information

URL: <https://about.gitlab.com>

Proper Citation: GitLab (RRID:SCR_013983)

Description: A software application which provides Git repository management with fine grained access controls, code reviews, issue tracking, activity feeds, wikis, and continuous integration. Four different versions are available, each with different packaging pricing: Community edition, Enterprise edition, Continuous integration, and GitLab.com.

Resource Type: software application, software resource, data management software

Keywords: software application, Git repository management, Git repository, data management

Availability: Fee (for certain editions and packages)

Resource Name: GitLab

Resource ID: SCR_013983

Alternate URLs: <https://about.gitlab.com/gitlab-com/>

License URLs: <https://about.gitlab.com/terms/>

Record Creation Time: 20220129T080318+0000

Record Last Update: 20260803T040623+0000

Ratings and Alerts

No rating or validation information has been found for GitLab.

No alerts have been found for GitLab.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 63 mentions in open access literature.

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

Sanz-Robinson J, et al. (2025) Open-source platforms to investigate analytical flexibility in neuroimaging. *Imaging neuroscience* (Cambridge, Mass.), 3.

Jannasch A, et al. (2025) Setting up an institutional OMERO environment for bioimage data: Perspectives from both facility staff and users. *Journal of microscopy*, 297(1), 105.

Miller D, et al. (2025) Pooled PPIseq: Screening the SARS-CoV-2 and human interface with a scalable multiplexed protein-protein interaction assay platform. *PloS one*, 20(1), e0299440.

Ko MC, et al. (2025) From silence to song: Testosterone triggers extensive transcriptional changes in the female canary HVC. *Journal of neuroendocrinology*, 37(6), e13476.

Goldmann U, et al. (2025) Data- and knowledge-derived functional landscape of human solute carriers. *Molecular systems biology*, 21(6), 599.

Gustafsson OJR, et al. (2025) WorkflowHub: a registry for computational workflows. *Scientific data*, 12(1), 837.

Pelot NA, et al. (2025) The human vagal complex: from gross anatomy to single neurons, from brainstem to abdomen. *Research square*.

Fraga-González G, et al. (2025) Affording reusable data: recommendations for researchers from a data-intensive project. *Scientific data*, 12(1), 258.

Abboush M, et al. (2025) Advancing real-time validation of automotive software systems via continuous integration and intelligent failure analysis. *Scientific reports*, 15(1), 32936.

Vancoillie L, et al. (2025) Relevance and application of digital project management in academic research: A case study in a Medical Physics lab and practice. *Journal of applied clinical medical physics*, 26(9), e70242.

Ferenc K, et al. (2024) Improving bioinformatics software quality through teamwork. *Bioinformatics* (Oxford, England), 40(11).

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

Massey A, et al. (2024) Real-time forecasting of COVID-19-related hospital strain in France using a non-Markovian mechanistic model. *PLoS computational biology*, 20(5), e1012124.

Karimian Sichani E, et al. (2024) Creating High-Quality Synthetic Health Data: Framework for Model Development and Validation. JMIR formative research, 8, e53241.

Spannenkrebs JB, et al. (2024) Closing the loop: establishing an autonomous test-learn cycle to optimize induction of bacterial systems using a robotic platform. Frontiers in bioengineering and biotechnology, 12, 1528224.

Vilstrup AP, et al. (2024) A germline PAF1 paralog complex ensures cell type-specific gene expression. Genes & development, 38(17-20), 866.

Afiaz A, et al. (2024) Best practices to evaluate the impact of biomedical research software-metric collection beyond citations. Bioinformatics (Oxford, England), 40(8).

Hing B, et al. (2024) Transcriptomic Evaluation of a Stress Vulnerability Network Using Single-Cell RNA Sequencing in Mouse Prefrontal Cortex. Biological psychiatry, 96(11), 886.

Vuillaume T, et al. (2023) The ESCAPE Open-source Software and Service Repository. Open research Europe, 3, 46.

Luo S, et al. (2023) Examining asymmetric pairwise pre-reaction and transition states in enzymatic catalysis by molecular dynamics simulation and quantum mechanics/molecular mechanics calculation. STAR protocols, 4(2), 102263.