Resource Summary Report

Generated by dkNET on May 16, 2025

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 resource, software application, data management software

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

Funding:

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: 20250513T061517+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 52 mentions in open access literature.

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

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.

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

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.

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

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

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

Poline JB, et al. (2023) Data and Tools Integration in the Canadian Open Neuroscience Platform. Scientific data, 10(1), 189.

Schulze A, et al. (2023) Ensuring privacy protection in the era of big laparoscopic video data: development and validation of an inside outside discrimination algorithm (IODA). Surgical endoscopy, 37(8), 6153.

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.

Morton C, et al. (2022) Software development skills for health data researchers. BMJ health & care informatics, 29(1).

Du X, et al. (2022) A Checklist for Reproducible Computational Analysis in Clinical Metabolomics Research. Metabolites, 12(1).

Talavera Andújar B, et al. (2022) Studying the Parkinson's disease metabolome and exposome in biological samples through different analytical and cheminformatics approaches: a pilot study. Analytical and bioanalytical chemistry, 414(25), 7399.

Sica MP, et al. (2022) Protocol to study the oligomeric organization of single-span transmembrane peptides using molecular dynamics simulations. STAR protocols, 3(3), 101636.

Djeddi S, et al. (2021) Multi-omics comparisons of different forms of centronuclear myopathies and the effects of several therapeutic strategies. Molecular therapy: the journal of the American Society of Gene Therapy, 29(8), 2514.

Mbareche H, et al. (2021) In Silico Study Suggesting the Bias of Primers Choice in the Molecular Identification of Fungal Aerosols. Journal of fungi (Basel, Switzerland), 7(2).

Cresswell K, et al. (2021) Understanding Public Perceptions of COVID-19 Contact Tracing Apps: Artificial Intelligence-Enabled Social Media Analysis. Journal of medical Internet research, 23(5), e26618.

Le Rutte EA, et al. (2021) Modelling the impact of COVID-19-related programme interruptions on visceral leishmaniasis in India. Transactions of the Royal Society of Tropical Medicine and Hygiene, 115(3), 229.

Allain F, et al. (2021) Geniac: Automatic Configuration GENerator and Installer for nextflow pipelines. Open research Europe, 1, 76.