

Resource Summary Report

Generated by RRID on Aug 8, 2026

PyPI

RRID:SCR_023145

Type: Tool

Proper Citation

PyPI (RRID:SCR_023145)

Resource Information

URL: <https://pypi.org/>

Proper Citation: PyPI (RRID:SCR_023145)

Description: Software repository for Python programming language.Used to find and install software developed and shared by Python community.

Abbreviations: PyPI

Synonyms: Python Package Index

Resource Type: software repository, software resource

Keywords: Python, find and install Python software, Python community,

Availability: Free, Available for download, Freely available

Resource Name: PyPI

Resource ID: SCR_023145

Record Creation Time: 20230118T050205+0000

Record Last Update: 20260808T190536+0000

Ratings and Alerts

No rating or validation information has been found for PyPI.

No alerts have been found for PyPI.

Data and Source Information

Usage and Citation Metrics

We found 78 mentions in open access literature.

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

Rodrigues GVP, et al. (2026) ViralQuest: a user-friendly interactive pipeline for viral-sequences analysis and curation. BMC bioinformatics, 27(1).

Becerra B, et al. (2026) Nucleotide-resolution mapping of regulatory elements via allelic readout of tiled base editing. Nature communications, 17(1).

Klim L, et al. (2026) Does Previous Anaphylaxis Determine Differences Between Patients Undergoing Oral Food Challenges to Cow's Milk and Hen's Egg? Nutrients, 18(2).

Salek Ardestani S, et al. (2026) GENOVIS: a Python package for the visualization of population genetic analyses. BMC genomics, 27(1), 190.

McArthur RN, et al. (2026) piqtree: a python package for seamless phylogenetic inference with IQ-TREE. Molecular biology and evolution, 43(3).

Rraku E, et al. (2026) Developing Del2Phen: A Novel Phenotype Description Tool for Chromosome Deletions. Human mutation, 2026, 5239482.

Vidal E, et al. (2026) ABaCo: addressing heterogeneity challenges in metagenomic data integration with adversarial generative models. Nucleic acids research, 54(5).

Feng S, et al. (2026) China's lake expansion amplified rapid CO2 emissions. Science advances, 12(17), eaea4657.

Daste S, et al. (2026) Projection-specific routing of odor information in the olfactory cortex. Current biology : CB, 36(11), 2890.

Lalli G, et al. (2025) JINet: easy and secure private data analysis for everyone. BMC bioinformatics, 26(1), 248.

Bueckle A, et al. (2025) Construction, Deployment, and Usage of the Human Reference Atlas Knowledge Graph for Linked Open Data. bioRxiv : the preprint server for biology.

Roth J, et al. (2025) Ten principles for reliable, efficient, and adaptable coding in psychology and cognitive neuroscience. Communications psychology, 3(1), 62.

Peng G, et al. (2025) Formal validation of a deep learning-based automated interpretation system for cardiac structure and function in adult echocardiography. Quantitative imaging in medicine and surgery, 15(4), 3093.

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

Greene D, et al. (2025) Biallelic variants in RNU2-2 cause the most prevalent known recessive neurodevelopmental disorder. medRxiv : the preprint server for health sciences.

Robinson JT, et al. (2025) igv-reports: Embedding interactive genomic visualizations in HTML reports to aid variant review. bioRxiv : the preprint server for biology.

Reig-Viader R, et al. (2025) Synaptic proteome diversity is shaped by the levels of glutamate receptors and their regulatory proteins. Nature communications, 16(1), 10487.

Liu SQ, et al. (2025) Decoding the Primacy of Transportation Emissions of Formaldehyde Pollution in an Urban Atmosphere. Toxics, 13(8).

De T, et al. (2025) PyPlaque is an open-source python package for phenotypic analysis of virus plaque assays. Scientific reports, 15(1), 36187.

Larracy R, et al. (2025) A dataset of high-resolution plantar pressures for gait analysis across varying footwear and walking speeds. Scientific data, 12(1), 1415.