

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).

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.

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

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

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

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.

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

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.

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

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

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.

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.

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

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.

Huang S, et al. (2025) GDBIG: A Pioneering Birth Cohort Genomic Platform Facilitating Intergenerational Genetic Research. Genomics, proteomics & bioinformatics, 23(5).

Wanichananan P, et al. (2025) Morphological characteristics and genome-wide association analysis among local *Andrographis paniculata* from Thailand under controlled environment in plant factory. PloS one, 20(4), e0320667.