

Resource Summary Report

Generated by [SPARC SAWG](#) on Jul 28, 2026

[infercnvpy](#)

RRID:SCR_025804

Type: Tool

Proper Citation

infercnvpy (RRID:SCR_025804)

Resource Information

URL: <https://holonet-tutorial.readthedocs.io/en/latest/index.html>

Proper Citation: infercnvpy (RRID:SCR_025804)

Description: Software application to infer copy number variation from single-cell transcriptomics data. Scalable Python library to infer CNV events from single cell transcriptomics data.

Resource Type: software library, software resource, software toolkit

Keywords: infer copy number variation, copy number variation, Copy Number Variation, CNV, single-cell transcriptomics data,

Availability: Free, Freely available,

Resource Name: infercnvpy

Resource ID: SCR_025804

Record Creation Time: 20240927T053302+0000

Record Last Update: 20260728T043729+0000

Ratings and Alerts

No rating or validation information has been found for infercnvpy.

No alerts have been found for infercnvpy.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Lin GT, et al. (2025) Combining Apatinib and Oxaliplatin Remodels the Immunosuppressive Tumor Microenvironment and Sensitizes Desert-Type Gastric Cancer to Immunotherapy. Cancer research, 85(11), 2117.

Liu Y, et al. (2025) SpatialSNV: A novel method for identifying and analyzing spatially resolved SNVs in tumor microenvironments. GigaScience, 14.