

Resource Summary Report

Generated by [Kravitz](#) on Sep 18, 2026

[svaNUMT](#)

RRID:SCR_021381

Type: Tool

Proper Citation

svaNUMT (RRID:SCR_021381)

Resource Information

URL: <https://github.com/PapenfussLab/svaNUMT>

Proper Citation: svaNUMT (RRID:SCR_021381)

Description: Software R package for Nuclear Mitochondrial integration events NUMT detection using structural variant calls.

Resource Type: data analysis software, data processing software, software application, software resource

Keywords: Nuclear mitochondrial integration events, NUMT, NUMT detection, structural variant calls.

Availability: Free, Available for download, Freely available

Resource Name: svaNUMT

Resource ID: SCR_021381

Alternate URLs: <https://bioconductor.org/packages/svaNUMT/>

License: GNU General Public License v3.0

Record Creation Time: 20220129T080355+0000

Record Last Update: 20260912T195955+0000

Ratings and Alerts

No rating or validation information has been found for svaNUMT.

No alerts have been found for svaNUMT.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Dong R, et al. (2022) svaRetro and svaNUMT: modular packages for annotating retrotransposed transcripts and nuclear integration of mitochondrial DNA in genome sequencing data. GigaByte (Hong Kong, China), 2022, gigabyte70.