

# Resource Summary Report

Generated by [kravitz2](#) on Sep 18, 2026

## PyroHMMvar

RRID:SCR\_002073

Type: Tool

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### Proper Citation

PyroHMMvar (RRID:SCR\_002073)

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### Resource Information

**URL:** <https://github.com/homopolymer/PyroTools/>

**Proper Citation:** PyroHMMvar (RRID:SCR\_002073)

**Description:** A software program to call short indels and SNPs for Ion Torrent and 454 data.

**Abbreviations:** PyroHMMvar

**Resource Type:** software resource

**Defining Citation:** [PMID:23995392](#)

**Availability:** Free, Available for download, Freely available

**Resource Name:** PyroHMMvar

**Resource ID:** SCR\_002073

**Alternate IDs:** OMICS\_00069

**Old URLs:** <https://code.google.com/p/pyrohmmvar/>

**Record Creation Time:** 20220129T080211+0000

**Record Last Update:** 20260912T195536+0000

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### Ratings and Alerts

No rating or validation information has been found for PyroHMMvar.

No alerts have been found for PyroHMMvar.

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### Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [kravitz2](#) .

Thangam M, et al. (2015) CRCDA--Comprehensive resources for cancer NGS data analysis. Database : the journal of biological databases and curation, 2015.