

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

TAPS

RRID:SCR_000356

Type: Tool

Proper Citation

TAPS (RRID:SCR_000356)

Resource Information

URL: <https://code.google.com/p/taps/>

Proper Citation: TAPS (RRID:SCR_000356)

Description: A bioinformatic tool for the identification of allele-specific copy numbers in tumor samples using data from Affymetrix SNP arrays.

Synonyms: Tumor Aberration Prediction Suite

Resource Type: software resource

Defining Citation: [PMID:22023820](#)

Keywords: standalone software, affymetrix

Resource Name: TAPS

Resource ID: SCR_000356

Alternate IDs: OMICS_02571

License: GNU General Public License

Record Creation Time: 20220129T080201+0000

Record Last Update: 20260905T132419+0000

Ratings and Alerts

No rating or validation information has been found for TAPS.

No alerts have been found for TAPS.

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 [Kravitz](#) .

Hofvander J, et al. (2026) Transcriptomic Subgroups in Soft Tissue Tumors Correlate with Morphologic Subtype, Genomic Features, and Outcome. Clinical cancer research : an official journal of the American Association for Cancer Research, 32(9), 1825.

Andersson N, et al. (2025) Tracing metastatic spread in pediatric solid tumors using copy number and targeted deep sequencing. The Journal of pathology.