

Resource Summary Report

Generated by SPARC SAWG on Aug 29, 2026

CNACS

RRID:SCR_022988

Type: Tool

Proper Citation

CNACS (RRID:SCR_022988)

Resource Information

URL: http://plaza.umin.ac.jp/kyoto_tumorpatho/soft.html

Proper Citation: CNACS (RRID:SCR_022988)

Description: Software tool for copy number analysis.

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

Keywords: copy number analysis

Availability: Free, Available for download, Freely available

Resource Name: CNACS

Resource ID: SCR_022988

Alternate URLs: <https://github.com/OgawaLabTumPath/CNACS>,
<https://github.com/OgawaLabTumPath>

License: MIT license

Record Creation Time: 20221123T050214+0000

Record Last Update: 20260821T194243+0000

Ratings and Alerts

No rating or validation information has been found for CNACS.

No alerts have been found for CNACS.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Yoshizato T, et al. (2025) Stable clonal contribution of lineage-restricted stem cells to human hematopoiesis. Nature genetics, 57(12), 3088.

Ushijima Y, et al. (2024) Initiating-clone analysis in patients with acute myeloid leukemia secondary to essential thrombocythemia. Scientific reports, 14(1), 15906.

Tabata M, et al. (2023) Inter- and intra-tumor heterogeneity of genetic and immune profiles in inherited renal cell carcinoma. Cell reports, 42(7), 112736.