# **Resource Summary Report**

Generated by RRID on May 16, 2025

# AbsCN-seq

RRID:SCR\_006409

Type: Tool

## **Proper Citation**

AbsCN-seq (RRID:SCR\_006409)

### **Resource Information**

#### **URL:**

http://bioinformatics.oxfordjournals.org/content/early/2014/01/02/bioinformatics.btt759.abstract?sid=e6226dc-428b-ba24-99e92a277d77

**Proper Citation:** AbsCN-seq (RRID:SCR\_006409)

**Description:** Statistical software to estimate tumor purity, ploidy and absolute copy numbers from next generation sequencing data.

**Abbreviations:** AbsCN-seq

**Resource Type:** software resource

**Defining Citation: PMID:24389661** 

**Keywords:** r, statistics, purity, ploidy, absolute copy number, next-generation sequencing

Related Condition: Tumor, Cancer

**Funding:** 

Availability: Free, Public

Resource Name: AbsCN-seq

Resource ID: SCR\_006409

Alternate IDs: OMICS\_02202

**Record Creation Time:** 20220129T080236+0000

Record Last Update: 20250420T014326+0000

## **Ratings and Alerts**

No rating or validation information has been found for AbsCN-seq.

No alerts have been found for AbsCN-seq.

### Data and Source Information

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 6 mentions in open access literature.

**Listed below are recent publications.** The full list is available at RRID.

Sandmann S, et al. (2022) clevRvis: visualization techniques for clonal evolution. GigaScience, 12.

Zhu G, et al. (2021) Tissue-specific cell-free DNA degradation quantifies circulating tumor DNA burden. Nature communications, 12(1), 2229.

Cai W, et al. (2018) MHC class II restricted neoantigen peptides predicted by clonal mutation analysis in lung adenocarcinoma patients: implications on prognostic immunological biomarker and vaccine design. BMC genomics, 19(1), 582.

Bao L, et al. (2015) Mutational Profiling Can Establish Clonal or Independent Origin in Synchronous Bilateral Breast and Other Tumors. PloS one, 10(11), e0142487.

Favero F, et al. (2015) Sequenza: allele-specific copy number and mutation profiles from tumor sequencing data. Annals of oncology: official journal of the European Society for Medical Oncology, 26(1), 64.

Moon HG, et al. (2015) The Clinical Significance and Molecular Features of the Spatial Tumor Shapes in Breast Cancers. PloS one, 10(12), e0143811.