

Resource Summary Report

Generated by [RRID](#) on Aug 2, 2026

ContEst

RRID:SCR_000595

Type: Tool

Proper Citation

ContEst (RRID:SCR_000595)

Resource Information

URL: <http://www.broadinstitute.org/cancer/cga/contest>

Proper Citation: ContEst (RRID:SCR_000595)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on August 8, 2025. A software tool (and method) for estimating the amount of cross-sample contamination in next generation sequencing data.

Abbreviations: ContEst

Resource Type: software resource

Defining Citation: [PMID:21803805](#)

Keywords: java, contamination, next-generation sequencing, bio.tools

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: ContEst

Resource ID: SCR_000595

Alternate IDs: OMICS_01038, biotools:contest

Alternate URLs: <https://bio.tools/contest>

License: BSD-style License, Account required

Record Creation Time: 20220129T080202+0000

Record Last Update: 20260801T190125+0000

Ratings and Alerts

No rating or validation information has been found for ContEst.

No alerts have been found for ContEst.

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](#) .

Chaudhary R, et al. (2025) Long-term Survival and Molecular Biomarker Evaluation of a Phase II Cetuximab and Nivolumab Clinical Trial in Recurrent/Metastatic Head and Neck Cancer. *Clinical cancer research : an official journal of the American Association for Cancer Research*.

Torres MJ, et al. (2023) *Stenotrophomonas goyi* sp. nov., a novel bacterium associated with the alga *Chlamydomonas reinhardtii*. *F1000Research*, 12, 1373.

Tsuji J, et al. (2022) Clinical Efficacy and Whole-Exome Sequencing of Liquid Biopsies in a Phase IB/II Study of Bazedoxifene and Palbociclib in Advanced Hormone Receptor-Positive Breast Cancer. *Clinical cancer research : an official journal of the American Association for Cancer Research*, 28(23), 5066.

Hurst CD, et al. (2021) Stage-stratified molecular profiling of non-muscle-invasive bladder cancer enhances biological, clinical, and therapeutic insight. *Cell reports. Medicine*, 2(12), 100472.

Sade-Feldman M, et al. (2018) Defining T Cell States Associated with Response to Checkpoint Immunotherapy in Melanoma. *Cell*, 175(4), 998.

, et al. (2017) Comprehensive and Integrated Genomic Characterization of Adult Soft Tissue Sarcomas. *Cell*, 171(4), 950.