

# Resource Summary Report

Generated by T1D on Sep 20, 2026

## ContEst

RRID:SCR\_000595

Type: Tool

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### Proper Citation

ContEst (RRID:SCR\_000595)

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### 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:** 20260919T194926+0000

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### Ratings and Alerts

No rating or validation information has been found for ContEst.

No alerts have been found for ContEst.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 6 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [T1D](#) .

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.