Resource Summary Report

Generated by dkNET on Sep 5, 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: 20260903T234351+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 dkNET .

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.