Resource Summary Report

Generated by NIF on Apr 25, 2025

CEQer

RRID:SCR_010813
Type: Tool

Proper Citation

CEQer (RRID:SCR_010813)

Resource Information

URL: http://www.ngsbicocca.org/html/ceqer.html

Proper Citation: CEQer (RRID:SCR_010813)

Description: A graphical, event-driven tool for CNA/AI-coupled analysis of exome sequencing reads.

Abbreviations: CEQer

Synonyms: Comparative Exome Quantification analyzer

Resource Type: software resource

Defining Citation: PMID:24124457

Keywords: bio.tools

Funding:

Availability: Commercial license, Free

Resource Name: CEQer

Resource ID: SCR_010813

Alternate IDs: biotools:ceqer, OMICS_00329

Alternate URLs: https://bio.tools/ceqer

Record Creation Time: 20220129T080300+0000

Record Last Update: 20250420T014510+0000

Ratings and Alerts

No rating or validation information has been found for CEQer.

No alerts have been found for CEQer.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 7 mentions in open access literature.

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

Fontana D, et al. (2024) Late relapse of chronic myeloid leukemia after allogeneic bone marrow transplantation points to KANSARL (KANSL1::ARL17A) alteration: a case report with insights on the molecular landscape. Annals of hematology, 103(5), 1561.

Franceschi S, et al. (2018) Cancer astrocytes have a more conserved molecular status in long recurrence free survival (RFS) IDH1 wild-type glioblastoma patients: new emerging cancer players. Oncotarget, 9(35), 24014.

Piazza R, et al. (2017) OncoScore: a novel, Internet-based tool to assess the oncogenic potential of genes. Scientific reports, 7, 46290.

Miyazaki J, et al. (2015) Intragenic duplication in the PKHD1 gene in autosomal recessive polycystic kidney disease. BMC medical genetics, 16, 98.

Yin S, et al. (2014) Exome sequencing identifies frequent mutation of MLL2 in non-small cell lung carcinoma from Chinese patients. Scientific reports, 4, 6036.

Reimann E, et al. (2014) Whole exome sequencing of a single osteosarcoma case--integrative analysis with whole transcriptome RNA-seq data. Human genomics, 8(1), 20.

Piazza R, et al. (2013) CEQer: a graphical tool for copy number and allelic imbalance detection from whole-exome sequencing data. PloS one, 8(10), e74825.