Resource Summary Report

Generated by RRID on Apr 28, 2025

CopySeq

RRID:SCR_010758
Type: Tool

Proper Citation

CopySeq (RRID:SCR_010758)

Resource Information

URL: http://www.embl.de/~korbel/CopySeq/

Proper Citation: CopySeq (RRID:SCR_010758)

Description: A computational tool that analyzes the depth-of-coverage of high-throughput DNA sequencing reads, and can integrate paired-end and breakpoint junction analysis based CNV-analysis approaches, to infer locus copy-number genotypes.

Abbreviations: CopySeq

Resource Type: software resource

Defining Citation: PMID:21085617

Keywords: java, bio.tools

Funding:

Resource Name: CopySeq

Resource ID: SCR_010758

Alternate IDs: biotools:copyseq, OMICS_00055

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

Record Creation Time: 20220129T080300+0000

Record Last Update: 20250420T014508+0000

Ratings and Alerts

No rating or validation information has been found for CopySeq.

No alerts have been found for CopySeq.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Zichner T, et al. (2013) Impact of genomic structural variation in Drosophila melanogaster based on population-scale sequencing. Genome research, 23(3), 568.