

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 8, 2026

DupRecover

RRID:SCR_006410

Type: Tool

Proper Citation

DupRecover (RRID:SCR_006410)

Resource Information

URL: <https://bitbucket.org/wandering/duprecover/overview>

Proper Citation: DupRecover (RRID:SCR_006410)

Description: Software that facilitates accurate estimation for sampling-induced read duplication in deep sequencing experiments.

Abbreviations: DupRecover

Resource Type: software resource

Defining Citation: [PMID:24389657](#)

Keywords: python, overcorrection, variant, allele fraction, copy number variation

Funding: MD Anderson Odyssey recruitment fellowship ;
The MD Anderson Cancer Center Sheikh Khalifa Ben Zayed Al Nahyan Institute of
Personalized Cancer Therapy ;
NCI R01CA172652-01;
NCI P30CA016672

Availability: Free, Public

Resource Name: DupRecover

Resource ID: SCR_006410

Alternate IDs: OMICS_02201

Record Creation Time: 20220129T080236+0000

Record Last Update: 20260905T132553+0000

Ratings and Alerts

No rating or validation information has been found for DupRecover.

No alerts have been found for DupRecover.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.